

Sequence	Accession	Sequence	Sequence	Fold Change	P-value	Ratio	Intensity1	Intensity2	
A_23_P21	NM_00196	Homo sapi	EGR1,TIS	364.7508		0	364.7508	973.7065	360481.4
A_32_P87	NM_00058	interleukin	IL8,K60,NA	275.0465		0	275.0465	639.3284	178368.3
A_23_P46	NM_00039	Homo sapi	EGR2,CM1	230.6523		0	230.6523	380.7201	89860.08
A_23_P34	NM_00402	Homo sapi	ATF3	224.9476		0	224.9476	1184.506	278668.1
A_24_P18	NM_00209	GRO3 onc	CXCL3,GR	155.1512		0	155.1512	794.812	124349.3
A_23_P10	NM_00525	v-fos FBJ r	FOS,c-fos	144.5045		0	144.5045	813.9425	118480.7
A_23_P20	NM_00222	Homo sapi	JUN,AP1	108.2045		0	108.2045	5188.445	567405.8
A_32_P74	BC052560	Hypothetic	LOC38776	91.09306		0	91.09306	509.6938	47857.56
A_23_P31	NM_00208	Homo sapi	CXCL2,GR	83.82861		0	83.82861	5648.745	478005.6
A_23_P38	NM_00196	Homo sapi	EGR4,NGF	80.83742		0	80.83742	1561.651	130449
A_24_P77	NM_00096	prostaglan	PTGS2,CC	79.10538		0	79.10538	451.059	36212.66
A_24_P33	NM_00402	activating t	ATF3	77.01963		0	77.01963	49.93195	3948.699
A_24_P25	NM_00096	prostaglan	PTGS2,CC	76.65108		0	76.65108	665.0615	52571.88
A_23_P12	NM_00213	Homo sapi	NR4A1,HV	73.1127		0	73.1127	6323.902	470426.3
A_23_P21	NM_00408	Homo sapi	DDIT3,CH	58.93116		0	58.93116	1840.167	110907.1
A_23_P72	NM_00057	Homo sapi	IL1A,IL1,IL	58.62542		0	58.62542	180.4437	10725.47
A_23_P21	NM_03141	Homo sapi	NFKBIZ,IK	48.98331		0	48.98331	303.3748	15391.12
A_23_P46	NM_01894	Gene 33V	ERRF1,MI	47.71919		0	47.71919	1988.694	96520.39
A_23_P17	NM_00459	small induc	CCL20,CK	42.67418	3.24E-36	42.67418	293.8745	13530.45	
A_23_P92	NM_02504	Homo sapi	ARF7,FLJ2	42.4731		0	42.4731	618.0479	26863.43
A_23_P12	NM_00285	Homo sapi	PTX3,TSG	42.36779		0	42.36779	2431.42	105479.3
A_23_P11	NM_00441	Homo sapi	DUSP1,HV	39.99293		0	39.99293	2823.082	113259.5
A_23_P39	NM_00340	Homo sapi	ZFP36,TF	38.61203		0	38.61203	3777.193	147900.3
A_23_P25	NM_00165	Homo sapi	AREG,AR,	38.2891		0	38.2891	871.5422	33612.21
A_23_P15	NM_00490	Homo sapi	IER2,ETR1	35.62196		0	35.62196	15858.6	566215.1
A_23_P63	NM_00130	Homo sapi	KLF6,GBF	32.58106		0	32.58106	1343.563	44456.11
A_24_P93	BC015987	Homo sapi	BC015987	32.53113		0	32.53113	172.4678	5748.764
A_23_P54	NM_01468	Homo sapi	HERPUD1	30.28991		0	30.28991	18585	562104.5
A_23_P15	NM_00441	dual specif	DUSP5,DL	29.12084		0	29.12084	1749.302	51424.36
A_24_P25	NM_00208	GRO2 onc	CXCL2,GR	28.59851		0	28.59851	4556.324	132706
A_23_P39	NM_17320	Homo sapi	NR4A3,CH	27.75576		0	27.75576	834.923	23637.61
A_24_P23	NM_01567	growth arre	GADD45B,	27.60541		0	27.60541	634.1123	17492.71
A_23_P42	NM_00389	Homo sapi	IER3,DIF2,	27.22347		0	27.22347	7761.453	212525.3
A_23_P13	NM_00297	Homo sapi	SAT,DC21	26.22969		0	26.22969	3473.385	91888.64
A_23_P13	NM_00618	nuclear rec	NR4A2,NC	25.55886		0	25.55886	408.6301	10382.05
A_23_P43	BC015544	Homo sapi	histone ger	24.60132		0	24.60132	325.5655	8035.791
A_23_P71	NM_00060	Homo sapi	IL6,HGF,H	24.54554		0	24.54554	990.0678	25250.94
A_23_P98	NM_00116	Homo sapi	BIRC3,AIP	23.97868		0	23.97868	174.4978	4165.446
A_23_P10	NM_02052	Homo sapi	NFKBIA,IK	23.88633		0	23.88633	11955.21	284790.7
A_24_P37	NM_00441	dual specif	DUSP2,PA	23.75824		0	23.75824	1252.757	29894.29
A_32_P15	AK092260	CDNA FLJ	DUSP5P,F	22.88293		0	22.88293	1983.302	46773.09
A_23_P25	NM_02013	chromoson	C8orf4,TC	22.85955	2.47E-40	22.85955	63.68555	1490.205	
A_23_P37	NM_00297	spermidine	SAT,DC21	21.13361		0	21.13361	1889.734	40771.37
A_32_P36	NM_00490	immediate	IER2,ETR1	20.18139		0	20.18139	27779.3	565821
A_23_P71	NM_00151	Homo sapi	CXCL1,GR	20.1136		0	20.1136	12630.2	254842.4
A_23_P13	NM_00194	Homo sapi	DUSP6,M	19.42457		0	19.42457	694.07	13690.04
A_32_P20	1976	Unknown	A_32_P20	19.21936	2.43E-29	19.21936	97.02917	1913.014	
A_32_P21	13948	BC003099	THC18819	17.67172		0	17.67172	471.8857	8407.649
A_24_P15	NM_00629	tumor necr	TNFAIP3,A	17.42476		0	17.42476	1483.655	26348.39
A_24_P11	NM_01894	Gene 33V	ERRF1,MI	17.41031		0	17.41031	50.7515	892.6777
A_24_P18	NM_00720	dual specif	DUSP10,M	17.39115		0	17.39115	118.5049	2069.749

A_23_P25	NM_00442	Homo sapi	EFNA1,B6	17.04084	0	17.04084	5055.609	86537.84
A_32_P97	BC024020	Transmem	VMP1	17.03171	0	17.03171	1028.898	17677.72
A_32_P91773		Unknown	A_32_P91	15.5054	0	15.5054	439.1943	6788.117
A_23_P76	BC018929	Pleckstrin	PHLDA1	15.28906	0	15.28906	5251.608	80069.48
A_24_P31	NM_02196	myeloid ce	MCL1,TM,I	15.06818	0	15.06818	573.4915	8705.725
A_24_P25	NM_00209	GRO3 onc	CXCL3,GR	14.80162	0	14.80162	9108.271	138607.1
A_23_P41	NM_00143	Homo sapi	EREG,ER	14.69961	0	14.69961	337.9906	4981.022
A_23_P16	NM_00486	Homo sapi	GDF15,PD	14.65333	0	14.65333	26417.96	386516.8
A_23_P32	NM_00538	Homo sapi	NFIL3,E4B	14.63351	0	14.63351	2146.466	31559.82
A_23_P35	NM_00354	Homo sapi	HIST1H4F	14.4706	0	14.4706	1318.646	19499.22
A_24_P53	AK057151	CAMP res	CREB5	14.36642	0	14.36642	461.1774	6747.169
A_23_P70	NM_00423	Homo sapi	CD83,BL1	14.12439	0	14.12439	1040.343	15073.05
A_32_P100365		Unknown	A_32_P10	14.03749	0	14.03749	271.6455	3830.605
A_23_P10	NM_00441	Homo sapi	DUSP2,PA	13.38213	0	13.38213	10682.76	143581.3
A_23_P25	NM_00526	Homo sapi	GEM,KIR,M	13.24889	1.15E-41	13.24889	398.0292	5467.242
A_32_P19	AK027019	Hypothetic	MGC45731	13.18919	0	13.18919	145.7916	1952.238
A_23_P21	NM_01337	CDK4-bind	SERTAD1	12.84502	0	12.84502	2325.226	29882.53
A_23_P36	NM_17851	hypothetic	LOC28348	12.84384	0	12.84384	661.5734	8708.133
A_23_P12	NM_02519	Homo sapi	TRIB1,C8F	12.57111	3.34E-23	12.57111	43.74789	549.2775
A_23_P20	NM_00395	Homo sapi	SOCS3,CI	12.54404	0	12.54404	7347.088	91740.66
A_23_P43	NM_00354	Homo sapi	HIST2H4,F	12.37577	0	12.37577	3088.948	39020.8
A_23_P48	NM_00222	Homo sapi	JUNB	12.32662	8.31E-43	12.32662	10512.42	133118.7
A_23_P88	NM_00416	Homo sapi	RRAD,RAI	12.01993	0	12.01993	1263.162	15493.34
A_23_P16	NM_02053	Homo sapi	OSM,MGC	12.00629	0	12.00629	1310.552	16064.59
A_23_P11	NM_01608	Homo sapi	RASD1,AG	11.74184	3.50E-34	11.74184	2784.976	33900.93
A_23_P13	NM_02030	cyclin L	aniCCNL1,BV	11.6135	0	11.6135	1757.241	21115.57
A_23_P11	NM_01627	Homo sapi	KLF2,LKLF	11.48539	0	11.48539	1378.169	16077.79
A_24_P16	NM_00629	tumor necr	TNFAIP3,A	11.42864	2.45E-17	11.42864	69.48774	811.5823
A_23_P21	NM_00353	Homo sapi	HIST1H3D	11.14795	0	11.14795	289.6968	3276.898
A_24_P29	AK124426	CDNA clon	AK124426	11.06368	0	11.06368	235.1308	2629.223
A_24_P36	AK092260	CDNA FLJ	DUSP5P,F	10.99182	1.57E-28	10.99182	2029.542	23528.88
A_23_P36	NM_01519	activity-reg	ARC,KIAA	10.65329	0	10.65329	1006.246	10822.17
A_23_P42	NM_00673	Homo sapi	FOSB,G0S	10.62813	4.08E-34	10.62813	853.3803	9406.883
A_32_P99804		Unknown	THC18224	10.52986	0	10.52986	1048.075	11172.82
A_24_P89	AK090468	TNF recept	TRAF1	10.52581	2.98E-37	10.52581	142.7888	1542.501
A_23_P23	NM_00192	Homo sapi	GADD45A,	10.52102	0	10.52102	3964.243	42276.14
A_24_P69	NM_00130	core promc	KLF6,GBF	10.47921	0	10.47921	915.0872	9730.409
A_23_P21	NM_00043	Homo sapi	PCSK1,PC	10.24813	0	10.24813	792.088	8154.718
A_23_P32	NM_00354	Homo sapi	HIST1H4H	10.16306	0	10.16306	1340.114	13900.64
A_23_P13	BC000297	3-hydroxy-	HMGCS1	10.06308	0	10.06308	564.5565	5704.263
A_23_P30	NM_00455	Homo sapi	NFKBIE,IK	10.05967	0	10.05967	796.6807	8052.825
A_24_P17	BC066345	AT rich inte	ARID5B	10.00671	0	10.00671	503.7797	5092.476
A_24_P18	NM_00534	heat shock	HSPA5,BIF	9.76762	0	9.76762	7115.133	69357.29
A_24_P916853		Unknown	A_24_P91	9.50619	0	9.50619	1102.331	10605.45
A_23_P21	NM_00443	Homo sapi	EGR3,PILC	9.26944	3.14E-11	9.26944	98.58073	965.897
A_23_P51	NM_00720	dual specif	DUSP10,M	9.26679	2.76E-33	9.26679	289.6567	2768.937
A_24_P33	NM_02030	cyclin L	aniCCNL1,BV	9.24454	0	9.24454	1491.106	13919.56
A_24_P33	NM_02196	myeloid ce	MCL1,TM,I	9.182	0	9.182	29298.77	270355.4
A_24_P27	NM_02080	Homo sapi	ARRDC3,K	9.1527	0	9.1527	848.7037	7893.808
A_24_P95	AK129833	Tissue fact	TFPI2	9.14452	0	9.14452	478.5648	4415.433
A_23_P35	AL137506	ELOVL fan	ELOVL7	9.1024	0	9.1024	91.79646	834.4548
A_24_P93	XM_37947	hypothetic	LOC40131	9.09061	6.75E-36	9.09061	292.161	2706.564

A_24_P26:	NM_00416	Ras-related	RRAD,RAI	9.08477	3.39E-22	9.08477	268.2578	2519.069
A_24_P401090	Unknown	ENST0000		9.06036	0	9.06036	1968.863	17902.69
A_24_P932388	Unknown	THC18617		8.98113	2.42E-43	8.98113	114.7027	1043.856
A_24_P33:	NM_02196	myeloid ce	MCL1,TM,I	8.95564	0	8.95564	1549.511	13949.87
A_32_P11:	CD104030	Transcribe	CD104030	8.75698	0	8.75698	513.3524	4518.074
A_32_P103776	Unknown	A_32_P10:		8.75282	0	8.75282	3412.469	30143.78
A_32_P30898	Unknown	A_32_P30:		8.66616	1.01E-29	8.66616	441.3976	3957.137
A_24_P20:	NM_00349	H4 histone	HIST1H4I,I	8.37203	0	8.37203	1298.178	10998.42
A_24_P10:	NM_02281	microtubule	MAP1LC3E	8.1772	0	8.1772	1229.959	10106.73
A_23_P10:	NM_01905	Homo sapi	DDIT4,Dig:	8.02532	0	8.02532	48283.03	387596.3
A_23_P37:	NM_02012	Homo sapi	TUFT1	7.98171	1.82E-20	7.98171	1425.449	12102.02
A_23_P93:	NM_00344	Homo sapi	ZNF165,LC	7.93511	0	7.93511	496.0256	3988.212
A_32_P177685	Unknown	A_32_P17:		7.88466	0	7.88466	647.361	5237.598
A_24_P710681	Unknown	THC19808		7.85215	2.51E-28	7.85215	252.5378	2033.511
A_23_P90:	NM_00614	Homo sapi	DNAJB1,H	7.84845	0	7.84845	3721.935	29399.45
A_23_P32:	NM_00292	Homo sapi	RGS16,RC	7.76509	0	7.76509	4494.509	35704.32
A_24_P88:	NM_00346	secretogran	SCG2,CHC	7.75208	0	7.75208	149.5961	1169.891
A_23_P13:	NM_00063	Homo sapi	SOD2,IPO	7.67419	0	7.67419	9935.857	76222.88
A_24_P668572	ALU5_HUM	THC19090		7.58417	7.30E-42	7.58417	433.2757	3390.013
A_23_P16:	NM_00565	Homo sapi	KLF10,EGI	7.55558	0	7.55558	1960.479	14826.47
A_24_P15:	NM_00329	tumor rejec	TRA1,ECC	7.5162	6.17E-44	7.5162	10951.62	84661.73
A_24_P92:	BC020539	KIAA1961	KIAA1961	7.36432	0	7.36432	471.1518	3473.829
A_32_P21:	XM_37947	hypothetic	LOC40131	7.33357	3.47E-30	7.33357	2872.19	21654.6
A_24_P34:	NM_01608	ras-related	RASD1,AG	7.31701	2.01E-13	7.31701	67.96687	502.9086
A_23_P55:	NM_00650	Homo sapi	RELB,I-RE	7.25622	0	7.25622	1698.809	12399.76
A_23_P21:	NM_00086	Homo sapi	HRH1,H1-I	7.25036	9.31E-38	7.25036	319.1467	2375.808
A_24_P30:	NM_00154	Homo sapi	IFIT2,G10F	7.22314	5.05E-25	7.22314	140.6856	1045.387
A_24_P32:	NM_00442	dual specif	DUSP8,HE	7.20746	6.87E-40	7.20746	580.2201	4270.663
A_23_P52:	NM_03095	likely ortho	NUAK2,SN	7.15102	1.86E-26	7.15102	1596.294	11947.59
A_24_P75948	Unknown	A_24_P75:		7.04823	0	7.04823	5986.042	42112.65
A_23_P74:	AF277181	PNAS-123	LOC85028	7.01689	0	7.01689	7636.487	54115.66
A_23_P35:	NM_01959	hypothetic	KIAA1434,	6.99367	1.49E-38	6.99367	5500.518	39435.73
A_23_P35:	NM_02509	KIAA1453	USP36,DU	6.94828	0	6.94828	802.5862	5566.891
A_24_P50:	AK056455	Poliovirus	rPVRL1	6.92402	2.21E-42	6.92402	5101.891	35466.82
A_24_P12:	NM_18178	Homo sapi	NCOA7,ES	6.91904	0	6.91904	5964.12	41334.79
A_23_P32:	NM_13839	hypothetic	LOC93082	6.87572	4.39E-14	6.87572	349.2139	2531.994
A_23_P21:	NM_00292	Homo sapi	RGS16,RC	6.8701	0	6.8701	1216.573	8394.928
A_32_P215304	Unknown	THC19098		6.85673	7.84E-20	6.85673	82.64811	564.9962
A_32_P232327	Unknown	A_32_P23:		6.83129	0	6.83129	375.3496	2565.059
A_24_P10:	NM_20339	Homo sapi	GK,GKD	6.79319	1.28E-34	6.79319	100.2065	681.6688
A_32_P10403	ALU1_HUM	THC18991		6.73678	0	6.73678	579.8876	3911.659
A_23_P12:	NM_03302	AXIN1 up-r	AXUD1,UR	6.6619	0	6.6619	363.4485	2435.429
A_23_P20:	NM_02112	Homo sapi	PMAIP1,AI	6.61844	0	6.61844	1784.784	11934.89
A_23_P12:	NM_02499	Homo sapi	ATF7IP2,M	6.4945	5.70E-23	6.4945	305.1532	2059.744
A_32_P10:	A1744472	BTG family	BTG2	6.49281	0	6.49281	152.9497	992.9325
A_23_P57836	Unknown	A_23_P57:		6.46164	0	6.46164	3953.084	25954.41
A_23_P17:	NM_00552	Homo sapi	HES1,HHL	6.45042	0	6.45042	2547.429	16467.46
A_23_P26:	NM_00329	Homo sapi	TRA1,ECC	6.4335	0	6.4335	86638.2	555466
A_23_P89:	NM_00261	Homo sapi	PER1,PER	6.395	0	6.395	1351.429	8659.314
A_24_P98:	NM_00534	heat shock	HSPA5,BIF	6.37136	0	6.37136	1707.919	10976.78
A_23_P47:	NM_00331	Homo sapi	PHLDA2,IF	6.35576	0	6.35576	17165.62	109629.7
A_23_P25:	NM_00523	Homo sapi	ETS2	6.33044	0	6.33044	13156.41	82977.28

A_24_P11	NM_00052	low density LDLR,FH,F	6.23955	0	6.23955	6606.489	41963.29
A_23_P51	NM_00407	Homo sapi PLK3,CNK	6.14899	0	6.14899	1456.043	8993.385
A_23_P21	NM_00118	Homo sapi BACH1	6.11384	0	6.11384	683.2474	4203.987
A_24_P87	AF001540	Metastasis MALAT1	6.113	0	6.113	1094.375	6610.297
A_23_P32	NM_02507	MCP-1 trez FLJ23231	6.10763	0	6.10763	2974.727	18190.5
A_24_P12	NM_18178	Homo sapi NCOA7,ES	6.10587	0	6.10587	1549.901	9612.936
A_23_P10	NM_00299	Homo sapi SDC4,SYN	6.04726	1.22E-29	6.04726	3561.837	21690.34
A_23_P74	NM_01571	putative lyr G0S2,RP1	6.04433	0	6.04433	14664.24	88646.19
A_24_P93	AL050388	Superoxide SOD2	6.02094	8.18E-19	6.02094	265.9128	1647.271
A_23_P16	BC017379	Homo sapi HIST1H2A	5.97736	0	5.97736	1171.101	7095.825
A_23_P33	NM_00735	pleckstrin t PHLDA1,P	5.96094	0	5.96094	667.1821	3984.186
A_24_P17	NR_00145	Homo sapi LOC85391	5.93793	4.53E-37	5.93793	132.1201	787.194
A_24_P56	AK027191	Trophoblas TncRNA	5.92904	0	5.92904	4321.317	25611.68
A_23_P29	NM_00531	Homo sapi H1F0,H10,	5.90942	0	5.90942	840.5389	5043.831
A_23_P13	NM_00731	Homo sapi ABL2,ARG	5.90192	0	5.90192	601.0023	3592.605
A_32_P67680	ENPL_HUI	THC18153	5.80411	8.92E-29	5.80411	983.9991	5757.962
A_23_P32	NM_02283	hypothetic FLJ22833,	5.79989	0	5.79989	2484.671	14355.68
A_23_P34	NM_00525	Homo sapi FOSL2,FR.	5.7321	0	5.7321	259.7281	1500.344
A_23_P13	NM_00112	Homo sapi ADFP,ADF	5.7093	0	5.7093	668.8935	3834.371
A_24_P38	NM_18259	hypothetic FLJ39575,	5.70006	7.70E-40	5.70006	571.6335	3321.73
A_24_P63	AK095492	3-hydroxy-; HMGCS1	5.68884	0	5.68884	2115.537	12041.34
A_24_P37	BC036402	MAX dimer MAD	5.68718	0	5.68718	553.0721	3187.447
A_23_P21	NM_00525	FOS-like al FOSL2,FR.	5.68201	6.93E-31	5.68201	1317.6	7598.929
A_32_P174121	Unknown	A_32_P17	5.63759	2.83E-13	5.63759	1457.974	8558.426
A_23_P62	NM_00676	Homo sapi BTG2,PC3	5.63315	1.03E-19	5.63315	4734.039	27855.27
A_24_P29	AF187554	Homo sapi sperm anti	5.54473	2.08E-27	5.54473	9484.39	52825.43
A_23_P90	NM_01433	Homo sapi PPP1R15A	5.51875	8.25E-34	5.51875	1258.027	6948.093
A_23_P14	NM_00681	Homo sapi PNR1,B4	5.46493	9.74E-40	5.46493	2835.305	15770.05
A_24_P82	BX538238	Metastasis MALAT1	5.44355	0	5.44355	51401.86	280876.6
A_23_P43	NM_03250	T-cell activ; KBTBD8,T.	5.44174	2.18E-23	5.44174	76.70815	415.4224
A_32_P173177	AF537132	THC19115	5.42431	0	5.42431	204.2273	1107.204
A_24_P26	NM_02411	hypothetic MGC5306	5.41943	0	5.41943	4080.681	22430.68
A_24_P73	AL832183	Hypothetic; LOC28445	5.41727	0	5.41727	2009.811	10962.86
A_24_P41	NM_02080	Homo sapi ARRDC3,K	5.35585	1.12E-14	5.35585	53.00171	282.2153
A_23_P73096	Unknown	A_23_P73	5.35493	9.92E-28	5.35493	629.0762	3433.147
A_24_P29	NM_01897	amino acid SLC38A2,	5.2955	0	5.2955	8652.986	46145.55
A_24_P95	NM_01504	KIAA0625 ALS4,AOA	5.2821	0	5.2821	3077.746	16441.84
A_23_P11	NM_00292	Homo sapi RGS2,G0S	5.26368	0	5.26368	8531.412	44939.88
A_23_P14	BC047030	Uridine phc UPP1	5.22162	0	5.22162	3534.489	18611.19
A_23_P32	NM_00439	Homo sapi DDX10,HR	5.22067	0	5.22067	12933.34	68854.23
A_24_P683829	Unknown	THC18075	5.2165	0	5.2165	392.4777	2051.942
A_24_P94	AF220656	Pleckstrin t PHLDA1	5.2148	2.78E-26	5.2148	733.2963	3903.984
A_24_P14	NM_01647	hypothetic C3orf19,FL	5.20455	0	5.20455	680.8366	3596.021
A_24_P74	AK022379	Beta-2-mic B2M	5.18394	5.64E-11	5.18394	226.0785	1227.376
A_24_P41	NM_03307	Homo sapi SYNE1,8B	5.08749	1.29E-20	5.08749	123.5898	631.0134
A_23_P30	NM_00574	Homo sapi PBEF1,PB	5.03001	0	5.03001	10368.33	52393.93
A_23_P251002	Unknown	A_23_P25	4.99801	1.76E-23	4.99801	329.8145	1702.157
A_23_P25	NM_01492	Homo sapi FNDC3A,F	4.95049	1.36E-39	4.95049	1805.813	9131.848
A_23_P71	NM_18159	Homo sapi UPP1,UP,	4.94929	2.03E-21	4.94929	375.6776	1911.486
A_23_P79	NM_00057	Homo sapi IL1B,IL-1,	4.92804	1.50E-27	4.92804	1448.056	7267.618
A_23_P12	NM_00508	Homo sapi XBP1,XBP	4.88683	0	4.88683	12806.81	63279.4
A_24_P91	NM_01504	KIAA0625 ALS4,AOA	4.82705	1.28E-16	4.82705	112.9251	544.9136

A_23_P14:	NM_00516 Homo sapi	RND3,ARF	4.80891	0	4.80891	5224.832	25291.51
A_23_P20:	NM_00052 low density LDLR,	FH,F	4.8082	0	4.8082	368.3842	1775.174
A_23_P32:	NM_00543 Homo sapi	FOSL1,FR	4.80291	1.39E-29	4.80291	117.4895	564.2455
A_24_P67:	AF307332 Meningiom	MGEA5	4.7982	0	4.7982	1849.436	8914.285
A_23_P30:	NM_00424 Jumonji do	JMJD1C	4.78904	1.12E-14	4.78904	417.3115	2144.621
A_24_P22:	NM_02283 hypothetic	FLJ22833,I	4.75824	5.86E-42	4.75824	287.4417	1384.667
A_23_P16:	AK128645 Syndecan	SDCBP	4.7533	1.18E-29	4.7533	278.0299	1342.46
A_23_P41:	NM_02017 Homo sapi	FEM1C,FE	4.71397	7.92E-06	4.71397	41.00797	189.9298
A_23_P14:	NM_00329 tropomyosi	TPM4	4.71355	0	4.71355	3449.11	16297.71
A_23_P24:	NM_00154 Homo sapi	IFIT2,G10F	4.69758	2.34E-27	4.69758	1792.773	8666.872
A_23_P20:	NM_01539 Homo sapi	ZNF10,KO	4.62189	1.81E-25	4.62189	330.3763	1567.344
A_24_P93:	NM_00552 hairy (Dros	HES1,HHL	4.60366	7.90E-30	4.60366	2802.169	13169.24
A_32_P154079	AB028926 THC18921		4.59116	5.47E-21	4.59116	2261.164	10431.71
A_24_P35:	NM_19826 similar to s	FLJ11021	4.59093	3.51E-42	4.59093	1375.5	6399.279
A_23_P11:	NM_00299 Homo sapi	CXCL5,SC	4.58824	3.54E-31	4.58824	3652.033	17272.23
A_23_P11:	NM_00510 A kinase (F	AKAP12,Al	4.57061	2.82E-07	4.57061	50.18372	230.549
A_23_P93:	NM_00234 Homo sapi	LTB,p33,TI	4.56466	7.01E-45	4.56466	701.2149	3210.444
A_23_P65:	NM_00552 hairy (Dros	HES1,HHL	4.55866	5.58E-06	4.55866	46.53979	206.5902
A_23_P34:	BX647822 FOS-like al	FOSL2	4.54974	0	4.54974	1778.288	8128.134
A_23_P77:	NM_02281 Homo sapi	MAP1LC3E	4.54491	3.00E-28	4.54491	1498.608	6919.541
A_23_P61:	BC017083 Pim-3 onc	PIM3	4.53697	0	4.53697	17929.38	81255.58
A_32_P78:	BM994509 Superoxide	SOD2	4.53241	0	4.53241	479.4221	2174.788
A_24_P31:	AB037855 Hypothetic	KIAA1434	4.52826	1.09E-08	4.52826	101.536	461.7121
A_32_P139894	ALU7_HUM	THC19992	4.52797	0	4.52797	1323.639	6063.703
A_23_P20:	NM_02476 hypothetic	WDR78,FL	4.51014	0	4.51014	910.1536	4135.608
A_23_P43:	NM_01826 hypothetic	H2AFJ,MG	4.48789	1.99E-16	4.48789	161.0198	731.3973
A_23_P16:	NM_01328 Homo sapi	FLRT3	4.48058	5.51E-10	4.48058	181.8868	839.657
A_23_P21:	NM_01897 Homo sapi	SLC38A2,/	4.47625	3.20E-32	4.47625	827.4671	3800.883
A_24_P25:	BC036529 Enhancer c	EPC1	4.47166	4.76E-38	4.47166	519.475	2372.992
A_23_P12:	NM_00219 Homo sapi	INHBA,ED	4.45358	0	4.45358	453.8213	2021.733
A_32_P16:	NM_03326 winged hel	FOXQ1,HF	4.45027	7.38E-28	4.45027	1475.784	6737.691
A_23_P14:	NM_01550 TCDD-indu	TIPARP	4.4044	4.15E-22	4.4044	2950.747	13445.29
A_23_P53:	NM_02301 similar to s	FLJ11021	4.39064	0	4.39064	8417.396	37088.82
A_32_P57:	CD365380 V-ets eryth	ETS2	4.38866	3.25E-17	4.38866	1320.472	5870.148
A_23_P13:	NM_17335 Homo sapi	SNF1LK,M	4.38389	0	4.38389	985.3085	4333.63
A_23_P34:	AK091203 Zinc finger	ZNF484	4.35601	9.37E-20	4.35601	119.2236	522.4338
A_23_P13:	NM_17335 Homo sapi	SNF1LK,M	4.34403	0	4.34403	6787	29850.71
A_24_P94:	AB014525 Amyotroph	ALS4	4.33745	8.76E-31	4.33745	1569.208	7001.927
A_23_P15:	NM_18178 Homo sapi	NCOA7,ES	4.3203	5.20E-29	4.3203	2160.026	9509.705
A_24_P26:	AL831836 Hypothetic	LOC28368	4.28882	0	4.28882	1462.265	6301.401
A_23_P31:	NM_01326 myosin reg	MYLIP,MIF	4.2874	2.16E-21	4.2874	183.7804	795.3765
A_32_P38:	BC039555 CDNA clon	BC039555	4.2819	0	4.2819	27076.98	115355.3
A_24_P25:	NM_02519 phosphopr	TRIB1,C8F	4.24362	2.07E-43	4.24362	4753.263	20454.05
A_23_P37:	NM_17881 Homo sapi	ARL8	4.23225	3.45E-23	4.23225	160.7245	682.8499
A_23_P39:	NM_00442 dual specif	DUSP8,HE	4.22976	3.55E-24	4.22976	1876.857	8090.063
A_32_P171488	Unknown	A_32_P17	4.20979	2.19E-23	4.20979	247.9327	1052.732
A_24_P17:	NM_00043 proprotein	PCSK1,PC	4.17779	1.57E-30	4.17779	793.5391	3378.3
A_24_P91:	NM_01644 hypothetic	LOC51233	4.16931	0	4.16931	1842.818	7677.845
A_23_P59:	NM_02105 Homo sapi	HIST1H2A	4.14298	0	4.14298	3183.281	13254.13
A_24_P31:	NM_00523 v-ets avian	ETS2	4.14105	1.86E-36	4.14105	865.5968	3654.124
A_32_P80295	Unknown	THC18796	4.13948	0	4.13948	1715.727	7084.372
A_23_P86:	NM_00732 Homo sapi	DMBT1,GF	4.13316	6.73E-31	4.13316	1910.071	8005.242

A_23_P13	NM_00483	Homo sapi	EIF2AK3,P	4.12631	0	4.12631	323.5236	1339.791
A_23_P10	NM_00351	Homo sapi	HIST2H2A	4.1073	0	4.1073	10949.46	45111.99
A_32_P21	NM_00329	tropomyosi	TPM4	4.09994	0	4.09994	3786.834	15730.22
A_24_P26	NM_00367	basic helix-	BHLHB2,D	4.09673	1.24E-31	4.09673	2185.143	9124.707
A_23_P18	NM_03204	Homo sapi	B3GNT5,B	4.0824	0	4.0824	1284.721	5245.513
A_23_P21	NM_02115	Homo sapi	TRIB3,NIP	4.08126	0	4.08126	48399.81	196790.3
A_32_P23	AL110141	Growth arr	GAS5	4.06864	0	4.06864	21580.12	87566.05
A_24_P73	NM_00469	solute carri	SLC16A6,I	4.05906	2.61E-35	4.05906	343.2677	1406.676
A_23_P103951		Unknown	A_23_P10	4.05813	0	4.05813	11538.02	47035.59
A_23_P35	AB051487	Dual specif	DUSP16	4.05723	0	4.05723	2294.969	9424.347
A_23_P12	NM_00112	Homo sapi	ADM,AM	4.05502	0	4.05502	4847.948	19800.46
A_24_P545200		Unknown	A_24_P54	4.04516	0	4.04516	22318.9	90394.33
A_23_P16	NM_00384	Homo sapi	TNFRSF1C	4.03251	1.59E-14	4.03251	511.1829	2130.368
A_24_P516728		Unknown	A_24_P51	4.02861	0	4.02861	14221.75	57480.7
A_24_P33	AK024229	Hypothetic	FLJ14167	4.02488	0	4.02488	1125.237	4532.425
A_23_P13	NM_00373	Homo sapi	OASL,TRIF	4.01854	2.01E-27	4.01854	3825.389	15624.81
A_23_P16	NM_00711	Homo sapi	TNFAIP6,T	4.01213	1.22E-23	4.01213	138.0156	554.1433
A_23_P52	NM_00154	Homo sapi	IFIT1,G10F	3.99987	0	3.99987	1355.498	5494.233
A_23_P20	NM_18301	Homo sapi	CREM,ICE	3.99113	4.92E-39	3.99113	2870.084	11619.28
A_24_P27	NM_00086	histamine r	HRH1,H1-I	3.97971	3.56E-21	3.97971	1035.658	4217.397
A_32_P12	BC066906	Similar to	LOC44009	3.97935	0	3.97935	73094.8	290058.8
A_24_P707530		Unknown	THC18182	3.96686	0	3.96686	732.9843	2915.321
A_24_P94	NM_17881	Homo sapi	ARL8	3.95813	9.94E-21	3.95813	299.342	1207.329
A_24_P27	NM_00299	small induc	CXCL5,SC	3.94542	0	3.94542	29404.23	115803.3
A_23_P36	NM_00535	Homo sapi	JUND	3.93183	0	3.93183	19465.71	76902.82
A_24_P38	AK022915	ADP-ribosy	ARL8	3.92384	9.44E-35	3.92384	918.7028	3679.271
A_24_P10	NM_00508	X-box bind	XBP1,XBP	3.91945	0	3.91945	448.3186	1764.158
A_24_P11	NM_01223	RING1 and	RYBP,AAF	3.90613	2.16E-33	3.90613	1414.781	5602.746
A_24_P21	NM_00384	tumor necr	TNFRSF1C	3.8847	0	3.8847	3814.416	14771.58
A_23_P15	NM_01847	Homo sapi	BEX1,BEX	3.87077	3.52E-08	3.87077	119.8327	464.5019
A_23_P41	NM_01208	Homo sapi	ELL2	3.86589	1.40E-45	3.86589	4402.375	17264.17
A_23_P87	NM_00173	B-cell trans	BTG1	3.85937	1.20E-34	3.85937	2837.5	11069.18
A_23_P42	NM_00424	Jumonji do	JMJD1C	3.85115	3.89E-36	3.85115	589.9503	2310.784
A_24_P82	NM_00329	tropomyosi	TPM4	3.84992	6.06E-36	3.84992	2651.698	10336.51
A_23_P86	NM_15278	hypothetic	MGC4040	3.84684	9.38E-41	3.84684	1776.423	6957.978
A_32_P208654	A47582	B-	THC19033	3.82716	0.00047	3.82716	63.48295	244.0279
A_23_P96	NM_00602	H1 histone	H1FX,H1X	3.80933	0	3.80933	1273.966	4858.438
A_23_P16	NM_00541	Homo sapi	SLC20A1,F	3.80174	0	3.80174	24640.39	93445.2
A_23_P39	NM_00652	Homo sapi	TFPI2,PP5	3.80063	0	3.80063	15317.62	58072.14
A_32_P22	NM_02281	microtubule	MAP1LC3E	3.79985	0	3.79985	3635.903	13802.91
A_24_P592591	PBEF_HUI	THC18675		3.79077	0	3.79077	15837.12	60006.96
A_23_P51	NM_00404	Homo sapi	RHOB,ARL	3.78909	2.10E-31	3.78909	864.5391	3281.594
A_23_P41	NM_00318	Homo sapi	TAF9,CIP,	3.78843	0	3.78843	18484.88	69934.22
A_24_P11	NM_00122	caspase 9,	CASP9,MC	3.77602	0	3.77602	4586.408	17272.79
A_32_P79	NM_00574	pre-B-cell c	PBEF1,PB	3.7758	0	3.7758	21167.05	79823.5
A_23_P96	NM_00016	Homo sapi	GK,GKD	3.76847	0	3.76847	936.6318	3548.7
A_32_P73143	ALU7_HUI	THC19625		3.75515	8.87E-27	3.75515	203.2431	765.952
A_23_P21	BX538238	Metastasis	MALAT1	3.75267	1.32E-30	3.75267	1170.254	4400.719
A_23_P97	NM_00122	Homo sapi	CASP9,MC	3.74185	0	3.74185	3004.547	11255.19
A_32_P90	NM_00318	TATA box	ITAF9,CIP,	3.73778	0	3.73778	9864.988	36886.49
A_23_P10	NM_00443	Homo sapi	ELF3,ERT,	3.7328	0	3.7328	343.3076	1287.187
A_24_P37	NM_00299	syndecan	SDC4,SYN	3.71607	5.18E-44	3.71607	161.8956	601.5899

A_23_P42	NM_00353	Homo sapi	HIST1H3G	3.71435	1.50E-39	3.71435	4424.042	16696.36
A_24_P67552		Unknown	A_24_P67	3.71099	0	3.71099	1814.982	6799.074
A_23_P83	NM_20340	Homo sapi	C9orf150,b	3.70307	7.60E-23	3.70307	624.3042	2343.417
A_32_P39	AK095791	Hypothetic	LOC39995	3.69641	1.08E-40	3.69641	643.0397	2401.254
A_32_P167176		Unknown	THC18837	3.68993	2.66E-39	3.68993	1046.219	3922.533
A_24_P82	NM_00532	H3 histone	H3F3B,H3.	3.68551	0	3.68551	42694.56	157405.5
A_23_P39	NM_17366	hypothetic	FLJ33718,l	3.677	1.79E-17	3.677	215.4091	801.0072
A_32_P181548		C40201 art	THC19339	3.67597	2.48E-06	3.67597	59.27254	213.7735
A_23_P16	NM_18342	TSC22 dor	TGFB1I4	3.67382	6.17E-44	3.67382	1375.794	5082.733
A_23_P11	NM_00442	Homo sapi	EFNA1,B6	3.66893	9.97E-34	3.66893	2475.784	9114.779
A_24_P93	NM_00353	H3 histone	HIST1H3I,l	3.66374	3.71E-39	3.66374	4692.84	17442.95
A_23_P15	NM_00532	Homo sapi	H3F3B,H3.	3.6447	0	3.6447	75219.84	273668
A_23_P53	NM_00173	Homo sapi	KLF5,CKLI	3.63502	4.88E-39	3.63502	5269.127	19491.54
A_23_P21	NM_00162	Homo sapi	AHR	3.62526	0	3.62526	1370.576	5004.608
A_24_P23	NM_18172	Ankyrin re	Lrp2bp	3.60968	0	3.60968	624.9395	2261.819
A_23_P57	NM_00616	Homo sapi	NFE2L2,NI	3.60779	0	3.60779	5559.723	20047.66
A_24_P26	NM_00731	v-abl Abels	ABL2,ARG	3.60083	1.30E-11	3.60083	209.2259	762.7982
A_23_P12	NM_00531	H1 histone	HIST1H1C	3.6	0	3.6	11218.5	40283.18
A_23_P40	NM_00235	Homo sapi	MXD1,MAI	3.58706	7.63E-11	3.58706	239.6121	880.38
A_23_P74	NM_03201	Homo sapi	STK40,SHI	3.58649	0	3.58649	1884.973	6779.598
A_32_P222571		Unknown	THC18595	3.57752	7.03E-25	3.57752	5096.476	18446.68
A_24_P40	NM_18279	Homo sapi	PBEF1,PB	3.57731	0	3.57731	7745.355	27642.11
A_24_P36	NM_00397	sperm assc	SPAG9,HS	3.57623	0	3.57623	2042.175	7278.51
A_23_P26	NM_13426	Homo sapi	RORA,ROI	3.56762	2.15E-38	3.56762	255.8599	910.5092
A_23_P12	NM_00190	Homo sapi	CTH,MGC	3.55443	1.40E-45	3.55443	4883.928	17567.64
A_23_P20	NM_00395	Homo sapi	MAP3K14,	3.55415	0	3.55415	909.579	3236.243
A_23_P91	AB037855	Hypothetic	KIAA1434	3.5386	7.33E-22	3.5386	2404.727	8619.352
A_24_P17	NM_00353	H3 histone	HIST1H3B	3.53845	0	3.53845	14874.35	52806.74
A_24_P935782		Z121_HUM	ENST0000	3.5248	7.61E-06	3.5248	66.58802	238.4745
A_23_P38	NM_01441	Homo sapi	BBC3,JFY	3.51355	0	3.51355	2926.18	10262.74
A_24_P917810		Unknown	A_24_P91	3.51169	0.00004	3.51169	53.47765	183.7692
A_23_P37	NM_00490	Wilms' tum	WTAP,MG	3.51071	0	3.51071	3235.705	11409.92
A_23_P13	NM_00353	Homo sapi	HIST1H3C	3.51049	0	3.51049	30140.89	105981.4
A_24_P35	NM_05290	Homo sapi	SLC25A25	3.49959	0	3.49959	2127.84	7495.306
A_23_P98	NM_01816	hypothetic	FLJ10652,l	3.48625	6.72E-33	3.48625	1087.944	3866.627
A_23_P15	NM_00469	Homo sapi	SLC16A6,l	3.48346	5.83E-41	3.48346	479.0795	1680.741
A_23_P11	NM_01907	Homo sapi	ING3,ING2	3.46979	0	3.46979	1891.881	6603.94
A_23_P49	NM_01663	Homo sapi	TNFRSF12	3.46209	0	3.46209	6390.884	22051.45
A_24_P17	NM_18301	Homo sapi	CREM,ICE	3.45745	0	3.45745	4354.738	15216.36
A_23_P30	AB051487	Dual specif	DUSP16	3.45719	7.96E-42	3.45719	1330.845	4637.808
A_23_P25	NM_03062	Homo sapi	CPEB4,KI	3.45678	7.12E-41	3.45678	1729.22	6063.26
A_23_P88	NM_00077	cytochrom	CYP3A5,C	3.45677	4.94E-07	3.45677	70.47058	244.4454
A_24_P25	AL390214	Triple funct	TRIO	3.44742	2.11E-25	3.44742	610.9446	2128.645
A_24_P68	NM_17506	Homo sapi	HIST2H2A	3.44618	3.31E-39	3.44618	5823.209	20320.92
A_24_P917703		Unknown	A_24_P91	3.43948	1.91E-06	3.43948	39.42934	135.6229
A_32_P20	AF085847	Full length	AF085847	3.43031	2.02E-12	3.43031	85.83305	295.2969
A_24_P58	XM_37327	similar to r	LOC39228	3.41104	3.63E-28	3.41104	977.1371	3408.243
A_24_P802562		Unknown	A_24_P80	3.39165	0	3.39165	1157.285	3939.647
A_23_P70	NM_00337	vascular er	VEGF,VPF	3.37491	9.39E-44	3.37491	809.0848	2755.379
A_23_P56	NM_00290	Homo sapi	REL,C-Rel	3.37051	2.08E-18	3.37051	506.9486	1751.378
A_23_P32	NM_02100	Homo sapi	UBC,HMG	3.36851	0	3.36851	166425.6	562470.5
A_24_P110213		BC036372	ENST0000	3.36226	1.34E-07	3.36226	139.8546	465.9047

A_23_P25	NM_01232 Homo sapi	DNAJB9,M	3.36045	8.80E-14	3.36045	2609.766	8948.541
A_23_P22	NM_00554 Homo sapi	INSIG1,CL	3.35978	0	3.35978	2100.591	7118.537
A_24_P29	NM_00600 zinc finger	ZA20D2,ZI	3.35648	1.25E-35	3.35648	3624.809	12333.7
A_23_P35	NM_03145 sestrin 2	SESN2,HIS	3.35594	1.54E-21	3.35594	1332.414	4507.088
A_24_P27	AK000703 Hypothetic	FLJ10652	3.355	0.00001	3.355	54.03096	179.7895
A_24_P26	NM_03096 sprouty (Dr	SPRY4	3.34564	1.58E-19	3.34564	328.556	1110.026
A_23_P93	NM_00353 Homo sapi	HIST1H3B	3.34313	4.38E-22	3.34313	1930.407	6564.047
A_24_P849245	Unknown	THC18695	3.33939	2.53E-36	3.33939	1865.692	6285.785
A_23_P35	BC009198 RRN3 RN	RRN3	3.33881	2.56E-43	3.33881	406.5239	1364.774
A_23_P37	NM_00059 Homo sapi	TNF,DIF,TI	3.33237	9.07E-10	3.33237	424.3825	1441.88
A_24_P25	AF132202 Homo sapi	PRO1859	3.32788	0	3.32788	658.9485	2190.606
A_23_P12	NM_01336 zinc finger	ZNF222	3.32414	1.56E-10	3.32414	140.6509	467.1927
A_23_P42	NM_02106 Homo sapi	HIST1H2A	3.31807	0	3.31807	9032.278	30063.87
A_23_P14	NM_00120 bone morp	BMP2,BMF	3.31424	3.60E-08	3.31424	67.87014	224.5747
A_24_P24	NM_15278 hypothetic	MGC40405	3.31386	2.44E-31	3.31386	985.6773	3327.855
A_23_P42	NM_00073 Homo sapi	CGA,HCG,	3.31347	2.57E-16	3.31347	421.8192	1421.708
A_32_P450879	Unknown	THC19005	3.2965	5.42E-30	3.2965	849.7274	2789.832
A_23_P12	NM_00117 Homo sapi	ARG2	3.27244	0	3.27244	476.4318	1565.659
A_32_P192842	Unknown	A_32_P19	3.27216	3.11E-16	3.27216	581.4349	1945.576
A_23_P41	NM_00427 Homo sapi	HOMER1,S	3.27113	0	3.27113	2868.521	9437.52
A_24_P36	NM_00595 metallothio	MT2A,MT2	3.2711	0	3.2711	27484.39	90322.16
A_32_P9931	ROU_HUM	THC18914	3.26603	0	3.26603	326.7173	1069.52
A_24_P39	NM_19821 Homo sapi	ING1,p33,p	3.25827	3.11E-28	3.25827	812.7235	2670.587
A_32_P16	AL832253 MRNA;	cDIAL832253	3.24791	1.26E-34	3.24791	250.8072	816.3108
A_24_P24	NM_19889 Homo sapi	ZNF160,F1	3.24789	1.52E-17	3.24789	473.1383	1563.183
A_24_P90	AK021804 Hypothetic	LOC39995	3.24768	4.47E-12	3.24768	225.6256	740.1037
A_23_P42	NM_02172 nuclear rec	NR1D1,EA	3.24129	5.66E-16	3.24129	2353.405	7776.253
A_23_P13	NM_01400 Homo sapi	ZNF297B,Z	3.23791	3.26E-31	3.23791	1014.253	3308.593
A_23_P90	NM_02434 hypothetic	ZNF557,M	3.23716	3.73E-37	3.23716	314.3685	1022.086
A_24_P16	BC032480 Similar to	FHSUP1	3.2323	0	3.2323	54742.23	177844.3
A_32_P71212	TR2A_HUM	THC19236	3.22959	2.28E-13	3.22959	122.2576	393.3342
A_24_P324465	H3 histone	HIST2H3C	3.22641	0	3.22641	26491.79	85883.56
A_32_P69	XM_37348 hypothetic	LOC38774	3.22593	0.00012	3.22593	97.18069	310.3104
A_24_P93	AK096192 SPOC dom	SPOCD1	3.22526	5.47E-11	3.22526	179.0804	581.2543
A_23_P33	AB075862 KIAA1982	KIAA1982	3.22465	7.15E-35	3.22465	1309.564	4259.237
A_24_P358305	Unknown	A_24_P35	3.22159	9.24E-24	3.22159	1296.956	4256.206
A_24_P40	AK000959 Brain expr	BEXL1	3.21291	0.00001	3.21291	45.50821	146.4598
A_24_P91	BC002724 Homo sapi	SNX5	3.2123	0	3.2123	6285.39	20200.89
A_23_P31	AK128423 Cold induci	CIRBP	3.20995	2.80E-45	3.20995	1008.979	3254.747
A_23_P40	BC021238 Hypothetic	MGC19764	3.20632	1.47E-28	3.20632	1002.478	3258.354
A_23_P36	NM_02123 selenoprot	SELK,HSP	3.20494	0	3.20494	9090.346	29270.62
A_23_P15	NM_13931 Homo sapi	ANGPTL4,	3.20311	1.35E-26	3.20311	151.8651	486.9249
A_23_P16	NM_00543 FOS-like al	FOSL1,FR	3.19936	6.70E-12	3.19936	819.1472	2680.207
A_23_P14	NM_00167 Homo sapi	ASNS,TS1	3.1991	3.98E-32	3.1991	50838.16	161794.2
A_24_P12	NM_00595 metallothio	MT1X,MT1	3.19713	0	3.19713	5570.877	17926.84
A_32_P49	XM_37204 similar to a	LOC38965	3.19627	2.11E-41	3.19627	21942.32	69733.28
A_24_P307104	Unknown	A_24_P30	3.19326	1.73E-24	3.19326	556.5137	1807.85
A_23_P30	NM_00605 Homo sapi	TNIP1,VAN	3.19235	0	3.19235	3447.592	11017.67
A_23_P13	BX640843 Hypothetic	BX640843	3.18975	0	3.18975	6366.918	20285.76
A_24_P92	AB052759 Transform	TRA2A	3.18749	1.06E-15	3.18749	169.9563	544.4796
A_23_P11	NM_02105 Homo sapi	HIST2H3C	3.18681	0	3.18681	16332.05	52136.66
A_23_P30	NM_01223 Homo sapi	RYBP,AAF	3.17714	2.19E-31	3.17714	1564.548	5087.742

A_24_P21:	NM_01211	CCR4-like	CCRN4L,N	3.1729	6.86E-13	3.1729	657.3008	2156.748
A_23_P170713		Unknown	A_23_P170	3.17194	6.56E-39	3.17194	6929.847	22296.65
A_32_P38821		Unknown	A_32_P388	3.17168	1.00E-35	3.17168	180.6952	573.2959
A_23_P30:	NM_00595	Homo sapi	MT1X,MT1	3.1711	0	3.1711	5969.264	19056.11
A_23_P21:	NM_00607	Homo sapi	CITED2,MI	3.16505	0	3.16505	826.2153	2632.137
A_23_P147404		Unknown	A_23_P147	3.16087	0	3.16087	4194.605	13247.48
A_24_P15:	NM_15278	hypothetic	MGC4040	3.1553	1.82E-15	3.1553	3587.344	11828.32
A_23_P31:	NM_00519	Homo sapi	CEBPD,CE	3.14451	0	3.14451	8961.221	28092.43
A_23_P35:	NM_00180	Homo sapi	CEBPG,GF	3.14279	0	3.14279	4525.747	14255.97
A_24_P27:	NM_00600	zinc finger	ZA20D2,ZI	3.13983	7.18E-27	3.13983	768.171	2453.144
A_23_P25:	BC007034	Metallothio	MT2A	3.13872	0	3.13872	27038.82	84978.95
A_23_P11:	NM_00392	Homo sapi	BCL10,CL	3.13336	1.45E-20	3.13336	1007.478	3248.81
A_24_P83:	XM_37247	Homo sapi	XM_37247	3.12957	2.27E-21	3.12957	1525.228	4886.017
A_32_P10:	BF084890	Trophoblas	TncRNA	3.11873	8.38E-13	3.11873	645.545	2034.292
A_32_P44628		Unknown	A_32_P446	3.11449	4.51E-12	3.11449	1546.017	4847.725
A_24_P91:	NM_00532	H3 histone	H3F3B,H3	3.11188	0	3.11188	14683.89	45548.29
A_23_P41:	NM_17562	Homo sapi	MT1J,MT1	3.11164	4.32E-43	3.11164	6643.203	20892.48
A_23_P12:	NM_02520	PH domain	pp9099,PP	3.11064	1.37E-26	3.11064	535.5676	1678.275
A_23_P42:	X97261	Metallothio	MT1L	3.10059	3.95E-31	3.10059	5915.768	18632.52
A_32_P28:	NM_00329	tropomyosi	TPM4	3.09496	1.84E-22	3.09496	9195.065	28816.46
A_24_P26:	NM_02100	Homo sapi	UBC,HMG;	3.08872	0	3.08872	182161.7	561928.5
A_23_P37:	NM_00057	Homo sapi	DAF,CR,T	3.08736	4.59E-26	3.08736	4193.88	13148.32
A_23_P41:	NM_00519	Homo sapi	CEBPB,LA	3.08271	0	3.08271	38501.78	118820
A_23_P32:	NM_00220	Homo sapi	ISG20,HE	3.08247	5.46E-35	3.08247	3996.884	12439.08
A_32_P10:	NM_00600	zinc finger	ZA20D2,ZI	3.07877	3.54E-16	3.07877	951.3778	3015.826
A_23_P10:	NM_00595	Homo sapi	MT2A,MT2	3.07862	0	3.07862	38197.54	117690.3
A_23_P39:	NM_15375	Homo sapi	C21orf81	3.06956	1.29E-17	3.06956	268.0909	829.8317
A_23_P12:	NM_00196	Homo sapi	EIF4A2,DC	3.06753	0	3.06753	15077.71	46364.76
A_24_P24:	NM_00222	jun B proto	JUNB	3.06702	0	3.06702	1170.667	3580.155
A_24_P21:	NM_00143	epiregulin	EREG,ER	3.06044	0.00001	3.06044	211.0109	663.9509
A_23_P46:	NM_00517	Homo sapi	BCL3,BCL	3.04159	3.66E-27	3.04159	5097.768	15418.63
A_24_P27:	NM_18301	Homo sapi	CREM,ICE	3.02257	2.95E-35	3.02257	1318.694	4029.814
A_24_P10:	NM_20339	Homo sapi	GK,GKD	3.0225	1.59E-22	3.0225	1461.638	4495.037
A_24_P68:	BX640843	Hypothetic	BX640843	3.022	0	3.022	9933.264	30113.87
A_23_P38:	NM_15334	Homo sapi	C6orf141,Λ	3.00694	1.95E-20	3.00694	1120.362	3411.705
A_24_P28:	NM_00264	phosphatid	PIGA,GPI3	2.9965	0	2.9965	1041.911	3139.115
A_23_P52:	NM_02513	hypothetic	HKDC1,FL	2.99307	1.01E-35	2.99307	1109.134	3343.339
A_23_P11:	NM_00371	stanniocalc	STC2,STC	2.99239	0	2.99239	1676.442	5010.997
A_24_P185018		Unknown	THC18203	2.99188	1.76E-13	2.99188	235.0013	708.1547
A_24_P33:	AK057830	Homo sapi	AK057830	2.9878	6.10E-11	2.9878	102.4884	305.765
A_23_P21:	NM_00246	Homo sapi	MYC,c-My	2.98474	0	2.98474	29802.97	88786.91
A_23_P20:	NM_02485	hypothetic	FLJ22028	2.98205	0	2.98205	472.4476	1411.768
A_24_P91:	BC037430	CDNA clon	BC037430	2.98185	0	2.98185	1928.547	5790.51
A_24_P27:	NM_00600	zinc finger	ZA20D2,ZI	2.97945	2.04E-26	2.97945	5163.316	15608.63
A_23_P15:	NM_00490	Homo sapi	CREB5,CF	2.97886	2.39E-16	2.97886	733.0953	2216.078
A_23_P10:	NM_00239	mouse dou	MDM4,MD	2.96861	2.20E-17	2.96861	692.5753	2097.007
A_23_P41:	AB046797	Hypothetic	MGC4040	2.95544	1.72E-13	2.95544	2290.601	7001.858
A_23_P10:	NM_15247	Homo sapi	ZNF565,FL	2.95504	0	2.95504	1271.46	3752.262
A_23_P21:	NM_00021	Homo sapi	JAG1,AGS	2.95154	3.07E-12	2.95154	206.9471	612.7159
A_24_P29:	U60873	Trophoblas	TncRNA	2.95026	3.29E-26	2.95026	450.566	1337.49
A_23_P43:	NM_02518	Homo sapi	KIAA1539,	2.94894	1.08E-27	2.94894	4560.317	13385.8
A_23_P30:	NM_00399	Homo sapi	NFKB1,KB	2.94843	2.69E-15	2.94843	3843.796	11577.28

A_32_P17	BX324301	Zinc finger ZNF297B	2.94363	2.78E-21	2.94363	284.1459	834.9197
A_23_P33	NM_00448	Homo sapi GOLGB1,C	2.91848	6.22E-35	2.91848	2874.806	8465.139
A_24_P87	NM_00351	Homo sapi HIST2H2A	2.89976	0	2.89976	8456.257	24634.66
A_32_P20	AF116620	Homo sapi PRO1068	2.89499	3.77E-17	2.89499	197.8346	574.1888
A_23_P42	NM_00491	protein dist PDIA4,ERF	2.88826	0	2.88826	7746.08	22366.94
A_24_P30	NM_00510	bromodom BRD2,NAT	2.88821	0	2.88821	21780.16	62959.68
A_24_P64	NM_03225	hypothetic ANKRD20,	2.86852	0.00005	2.86852	68.05563	192.5624
A_24_P24	NM_00257	serine (or c SERPINB2	2.86703	5.68E-09	2.86703	98.45478	281.6091
A_24_P16	NM_01336	zinc finger ZNF222	2.85784	4.97E-20	2.85784	503.4435	1459.679
A_23_P22	NM_03262	Homo sapi BEX2,BEX	2.85524	2.70E-23	2.85524	2596.977	7480.576
A_32_P33	AF390550	Zinc finger ZNF434	2.85108	1.55E-37	2.85108	432.9654	1240.938
A_23_P50	NM_03322	tripartite m TRIM15,R	2.84504	0	2.84504	793.2034	2268.579
A_23_P43	AK055561	Endoplasm ERN1	2.83617	4.01E-10	2.83617	729.8668	2114.178
A_24_P12	BC065522	Vascular ei VEGF	2.83454	8.65E-23	2.83454	702.5976	2016.219
A_32_P41	267	Unknown THC18482	2.83002	1.75E-16	2.83002	347.2493	990.8351
A_23_P12	3645	Unknown THC19529	2.82933	2.03E-13	2.82933	427.3824	1240.776
A_24_P25	3588	E40201 art THC18422	2.82452	1.95E-10	2.82452	580.0252	1639.284
A_24_P75	3760	153353 tro THC18313	2.82316	1.59E-18	2.82316	3009.96	8585.351
A_23_P71	NM_00600	Homo sapi ZA20D2,ZI	2.82222	0	2.82222	10468.07	29537.89
A_23_P46	NM_02520	Homo sapi EPC1,DKF	2.81783	7.79E-06	2.81783	520.2167	1524.347
A_23_P15	NM_00020	Homo sapi ICAM1,BB	2.81731	0.00032	2.81731	374.3325	1063.774
A_32_P15	BC029608	Likely orthc AVO3	2.81063	0.00018	2.81063	97.77858	276.1479
A_32_P20	BC035518	Homo sapi BC035518	2.80636	0	2.80636	620.5259	1742.919
A_23_P37	AB011145	Thioredoxi TXNDC4	2.79527	2.99E-16	2.79527	190.6101	532.2305
A_23_P35	NM_00541	Homo sapi SKIL,SNO,	2.79479	1.87E-11	2.79479	149.1518	414.9476
A_23_P29	NM_00180	Homo sapi CEBPE,CF	2.79164	1.49E-40	2.79164	889.1646	2491.446
A_23_P99	NM_00492	Homo sapi ZFP36L1,E	2.78923	8.84E-25	2.78923	4052.903	11446.78
A_23_P24	NM_00265	Homo sapi PLAU,ATF	2.78494	1.85E-15	2.78494	3106.807	8772.003
A_32_P17	8800	ALU7_HU THC19573	2.78475	0.00005	2.78475	85.74364	236.8856
A_23_P30	NM_02196	Homo sapi HIST1H4J,	2.78295	1.77E-42	2.78295	3107.763	8698.788
A_24_P63	5355	Unknown A_24_P63	2.78114	1.25E-13	2.78114	1236.737	3506.786
A_23_P42	NM_00629	Homo sapi TNFAIP2,E	2.77477	8.67E-28	2.77477	40001.86	110590.1
A_24_P38	NM_01660	chromoson FAM53C,C	2.76821	1.56E-32	2.76821	2332.508	6505.521
A_23_P14	NM_03344	Homo sapi HIST3H2A	2.75911	3.35E-39	2.75911	5964.997	16577.08
A_24_P50	AK096179	Hypothetic LOC28595	2.75589	0	2.75589	6748.202	18626.41
A_24_P94	NM_01765	Abelson he AHI1	2.73631	1.13E-10	2.73631	185.3597	506.2846
A_23_P58	NM_05300	TIGA1 TIGA1,MG	2.73407	0	2.73407	24776.96	67595.55
A_23_P79	NM_00532	Homo sapi HIST1H1E	2.73137	8.50E-23	2.73137	3497.27	9751.332
A_24_P41	NM_20330	Homo sapi FBXO33,FI	2.72922	7.04E-24	2.72922	1000.055	2779.087
A_23_P81	NM_00346	Homo sapi PTP4A1,HI	2.72591	2.57E-42	2.72591	49078.76	133349
A_23_P35	NM_00154	interferon-i IFIT3,IRG2	2.72567	7.34E-29	2.72567	391.4327	1072.499
A_24_P89	BC016022	Similar to a BC016022	2.7164	2.80E-32	2.7164	259.6699	705.681
A_23_P11	NM_00492	aquaporin : AQP3	2.71481	0	2.71481	4692.604	12748.82
A_23_P35	AF242519	Homo sapi zinc finger	2.71296	9.24E-11	2.71296	120.2826	325.2282
A_23_P39	NM_01563	Homo sapi FGFR1OP	2.70685	0	2.70685	556.1519	1507.959
A_23_P20	NM_17561	Homo sapi MT1E,MT1	2.70498	1.57E-33	2.70498	3142.959	8601.749
A_23_P16	NM_01660	Homo sapi FAM53C,C	2.70441	1.49E-24	2.70441	1078.436	2949.764
A_23_P35	NM_18159	Homo sapi UPP1,UP,I	2.70083	3.47E-29	2.70083	2062.416	5577.459
A_23_P88	NM_00506	Homo sapi SEL1L,IBD	2.69321	4.61E-37	2.69321	1253.315	3397.085
A_23_P25	NM_00510	Homo sapi BRD2,NAT	2.69193	0	2.69193	441.4958	1188.986
A_24_P25	NM_00407	CDC-like k CLK1,CLK	2.69137	3.22E-25	2.69137	265.986	716.783
A_32_P20	8076	AF512556 THC18470	2.689	4.84E-22	2.689	1035.879	2817.316

A_23_P12:	NM_01838	hypothetic	FLJ11292	2.68681	1.01E-25	2.68681	1780.466	4854.893
A_24_P45:	NM_02105	Homo sapi	HIST2H3C	2.68625	7.04E-37	2.68625	7747.499	20951.11
A_24_P78:	AK095849	RNA, U22	RNU22	2.67878	2.16E-33	2.67878	8735.309	23656.73
A_32_P18:	BX641020	AT rich inte	ARID5B	2.6783	9.81E-14	2.6783	2968.558	8046.231
A_23_P53:	NM_00557	Homo sapi	LMO2,TTG	2.67448	2.83E-18	2.67448	252.4663	673.4807
A_23_P38:	NM_02487	Homo sapi	TNIP3,LINI	2.6655	2.18E-10	2.6655	166.1753	441.2802
A_24_P33:	NM_01763	hypothetic	FLJ20035,	2.66129	3.49E-21	2.66129	255.0101	678.9081
A_32_P123743	U12390	be	THC19144	2.65692	1.44E-43	2.65692	4182.894	11173.21
A_23_P21:	NM_00673	Human imr	HIVEP2	2.65367	0	2.65367	505.0625	1341.734
A_24_P60:	AK095242	Catenin (c	CTNNB1	2.64938	6.43E-29	2.64938	207.9453	550.8426
A_23_P50:	NM_15262	Homo sapi	ZNF92,TF1	2.64559	0	2.64559	1282.964	3391.534
A_24_P732106	Unknown	THC18621		2.64082	3.91E-27	2.64082	409.8102	1088.209
A_32_P17:	NM_00016	glycerol kin	GK,GKD	2.6408	6.92E-08	2.6408	982.6899	2598.254
A_24_P29:	BC032478	Mitogen-ac	MAP2K3	2.63679	0	2.63679	9533.283	25173.75
A_24_P10:	NM_00541	solute carri	SLC20A1,F	2.63635	0	2.63635	899.8374	2372.352
A_23_P21:	NM_01583	Adenosine	ADARB1	2.63538	0	2.63538	2674.604	7055.265
A_24_P67:	AK090611	Hypothetic	LOC15909	2.62936	4.98E-11	2.62936	477.6999	1284.173
A_24_P63:	BC008632	Chromosor	C6orf176	2.6285	0	2.6285	5188.954	13690.68
A_23_P81:	NM_00510	interferon- ϵ	G1P2,UCR	2.61881	0	2.61881	8089.264	21163.71
A_32_P89277	Unknown	A_32_P89:		2.61335	6.15E-11	2.61335	254.734	669.5485
A_23_P60:	NM_00595	Homo sapi	MT1G,MT1	2.6102	1.80E-32	2.6102	8077.077	21256.36
A_24_P10:	NM_00450	isopenteny	IDI1,IPP1,I	2.60524	2.67E-27	2.60524	5349.363	14031.34
A_23_P12:	NM_00584	Homo sapi	SPRY2,hS	2.60166	2.42E-09	2.60166	526.8026	1403.395
A_23_P72448	SKIL_HUM	THC18076		2.59511	0.00036	2.59511	117.411	303.7761
A_32_P44:	AK091220	Homo sapi	HIST3H2B	2.594	1.06E-06	2.594	211.0578	553.3166
A_23_P40:	NM_00218	Homo sapi	IL7R,CD12	2.59106	7.79E-09	2.59106	1125.716	2994.262
A_23_P15:	NM_03204	BRCA1-int	BRIP1,OF,	2.5908	0.00026	2.5908	90.3495	231.774
A_24_P87:	BC043357	Discs, larg	DLGAP1	2.5903	4.64E-12	2.5903	353.4638	928.8679
A_24_P25:	AK097130	Ariadne ho	ARIH1	2.58821	5.94E-06	2.58821	123.6694	318.6552
A_32_P16:	AL833005	MRNA; cDI	AL833005	2.58765	4.41E-21	2.58765	985.558	2572.831
A_23_P20:	NM_13432	Homo sapi	SLC26A9	2.58727	8.41E-10	2.58727	422.3492	1101.456
A_23_P14:	NM_01829	Zinc finger	ZNF654	2.58355	0.00008	2.58355	120.3599	307.7697
A_24_P24:	NM_00448	golgi autoa	GOLGB1,C	2.57301	2.15E-30	2.57301	1036.568	2689.435
A_23_P15:	NM_00422	Interleukin	NK4	2.57276	4.76E-15	2.57276	2215.764	5768.252
A_23_P12:	NM_03093	Homo sapi	TMEM49,v	2.57093	1.59E-31	2.57093	5544.194	14347.7
A_32_P124728	Unknown	THC18931		2.56932	8.68E-18	2.56932	353.654	915.4767
A_24_P84:	NM_19827	Homo sapi	CXorf23	2.56923	5.22E-08	2.56923	146.9598	373.1853
A_23_P71:	NM_00182	Homo sapi	CKS2,CKS	2.56853	1.59E-30	2.56853	55675.45	144332.3
A_32_P22:	NM_13344	Homo sapi	CTGLF1,M	2.56691	5.70E-39	2.56691	1533.356	3962.918
A_23_P31:	NM_00110	Homo sapi	ACTB	2.56653	1.40E-45	2.56653	70640.14	181074.7
A_32_P85:	BC008632	Chromosor	C6orf176	2.56537	0	2.56537	10284.8	26397.21
A_24_P11:	NM_00366	basic leucit	BLZF1,JEM	2.56184	0.00011	2.56184	96.93286	244.4533
A_23_P30:	NM_00351	Homo sapi	HIST2H2A	2.55948	1.00E-19	2.55948	1989.085	5175.403
A_32_P105940	Unknown	A_32_P10:		2.55765	1.68E-08	2.55765	455.2558	1187.241
A_23_P11:	(AB002301	Microtubul	MAST4	2.55222	0	2.55222	1178.875	3014.661
A_23_P32:	AK002200	SMC4 struc	SMC4L1	2.54695	7.85E-18	2.54695	274.3065	700.2861
A_23_P50:	NM_00250	Homo sapi	NFYA,HAF	2.54198	9.72E-07	2.54198	269.5816	695.0298
A_32_P87:	NM_20330	Homo sapi	FBXO33,FI	2.53945	8.13E-44	2.53945	2281.334	5829.832
A_24_P25:	NM_00346	protein tyrc	PTP4A1,HI	2.53689	1.43E-39	2.53689	2031.415	5181.821
A_23_P14:	NM_00095	Homo sapi	PTGER4,E	2.53511	0	2.53511	640.7662	1621.292
A_24_P86:	NM_00351	H2A histon	HIST1H2A	2.53484	0	2.53484	6811.805	17296.04
A_23_P12:	NM_01694	Chromosor	C6orf48	2.5334	0	2.5334	38474.09	97284.14

A_23_P40	NM_15360	Homo sapi	LOC15322	2.53072	1.77E-11	2.53072	355.2847	909.8673
A_24_P33	NM_00696	zinc finger	ZNF24,KO	2.52936	1.70E-16	2.52936	188.5464	477.7987
A_23_P37	AK056767	V-maf mus	MAFK	2.52798	2.80E-43	2.52798	3572.282	9036.285
A_24_P29	NM_01766	hypothetic	ANKRD10,	2.52596	0	2.52596	1114.362	2808.557
A_23_P14	NM_01612	putative rin	RNF12,RL	2.51157	1.17E-21	2.51157	1261.943	3215.842
A_23_P16	NM_03283	protein phc	PPP1R15E	2.50985	1.67E-16	2.50985	643.036	1631.978
A_23_P30	NM_00713	Homo sapi	ZNF79,pT7	2.5075	3.89E-33	2.5075	307.3164	770.4336
A_23_P16	NM_00407	Homo sapi	CLK1,CLK	2.50586	1.76E-21	2.50586	978.5382	2480.212
A_32_P14	AF086205	Tuftelin 1	TUFT1	2.50254	4.81E-32	2.50254	361.0618	903.6362
A_23_P21	NM_00652	Homo sapi	ZNF217,Z/	2.5015	2.55E-12	2.5015	2228.989	5740.878
A_32_P23	NM_13344	Homo sapi	CTGLF1,M	2.5004	1.54E-30	2.5004	1405.943	3547.46
A_24_P153234		Unknown	ENST0000	2.49726	1.91E-21	2.49726	232.1252	579.5698
A_23_P10	NM_00145	Homo sapi	FHL2,DRA	2.49588	2.70E-34	2.49588	4190.125	10501.31
A_24_P26	NM_00532	H1 histone	HIST1H1D	2.49185	0	2.49185	7614.271	18995.04
A_23_P16	NM_00351	H2B histon	HIST1H2B	2.4878	0	2.4878	1332.004	3305.645
A_23_P123234		Unknown	A_23_P12	2.48699	7.98E-31	2.48699	498.3937	1241.575
A_24_P14	NM_19883	Homo sapi	SERPINB8	2.48679	2.20E-12	2.48679	484.6729	1211.03
A_32_P15	AK091904	CDNA FLJ	AK091904	2.48151	1.48E-15	2.48151	247.5096	614.8256
A_24_P73	NM_15231	hypothetic	FLJ38968,	2.48098	1.12E-38	2.48098	731.5074	1821.498
A_23_P10	NM_02103	Homo sapi	ZNF14,KO	2.47821	9.09E-39	2.47821	432.2653	1070.644
A_24_P264416		Unknown	A_24_P26	2.47674	0	2.47674	9794.945	24223.32
A_24_P13	AB040932	Nuclear prc	NPL4	2.47611	5.79E-08	2.47611	247.4253	614.2351
A_23_P34	NM_01989	Homo sapi	ERO1LB	2.47299	4.30E-12	2.47299	262.4708	651.2923
A_32_P14	AL122093	MRNA; cDIAL	AL122093	2.47001	5.64E-25	2.47001	6649.871	16402.75
A_24_P20	NM_01660	chromoson	C5orf5,N61	2.46792	3.92E-09	2.46792	206.7808	512.9249
A_24_P931955		Unknown	THC18191	2.46732	4.25E-14	2.46732	636.8849	1601.204
A_23_P78	NM_02403	hypothetic	MIS12,hMi	2.46364	9.88E-31	2.46364	2166.912	5388.295
A_23_P11	NM_00492	Homo sapi	AQP3	2.45539	1.53E-33	2.45539	1292.003	3170.111
A_23_P21	NM_00195	Homo sapi	EDN1,ET1	2.45534	1.60E-13	2.45534	633.2557	1569.716
A_23_P37	NM_00594	Homo sapi	MT1B,MT1	2.4523	7.43E-25	2.4523	9440.436	23414.99
A_32_P16	AL833005	MRNA; cDIAL	AL833005	2.44663	5.78E-26	2.44663	1212.012	2977.985
A_23_P86	NM_02480	hypothetic	ZNF669,FL	2.44602	1.91E-12	2.44602	136.123	332.7491
A_23_P25	NM_02461	Homo sapi	HSPBAP1,	2.44455	4.64E-06	2.44455	823.8439	2039.767
A_32_P86517		Unknown	A_32_P86	2.43776	7.69E-14	2.43776	242.518	592.2212
A_24_P16	AB002333	Human mR	KIAA0335	2.43704	4.17E-10	2.43704	295.2661	725.5648
A_23_P38	AB051513	KIAA1726	KIAA1726	2.43459	1.53E-12	2.43459	1271.504	3115.323
A_24_P20	NM_00413	potassium	KCNMB1,S	2.43412	1.68E-20	2.43412	8434.295	20553.29
A_32_P15	BC001103	GULP, eng	GULP1	2.43349	4.77E-06	2.43349	138.3328	336.0025
A_23_P16	NM_01837	Receptor-ir	RIF1	2.43195	1.10E-21	2.43195	637.2687	1565.496
A_24_P12	NM_20703	Homo sapi	C1orf63,NF	2.4293	2.80E-14	2.4293	203.9888	495.1985
A_23_P87	NM_02392	Homo sapi	C1QDC1,E	2.42763	5.54E-16	2.42763	740.4303	1825.374
A_23_P10	NM_00675	Homo sapi	SSFA2,CS	2.4272	1.01E-29	2.4272	3159.578	7743.918
A_23_P93	NM_00635	Homo sapi	TRIM38,RI	2.42637	0.00005	2.42637	103.0651	248.3358
A_24_P21	NM_02471	multiple C2	MCTP1,FL	2.42595	3.66E-08	2.42595	100.6009	243.6323
A_32_P88905		Unknown	A_32_P88	2.42569	7.18E-20	2.42569	1914.343	4664.361
A_24_P39	NM_02106	H2A histon	HIST1H2A	2.4249	1.44E-14	2.4249	2198.922	5416.183
A_24_P16361		Unknown	A_24_P16	2.42269	1.26E-32	2.42269	350.3634	849.6997
A_23_P10	NM_01622	Chromosor	C1orf9	2.41983	2.78E-14	2.41983	3349.79	8267.47
A_24_P578445		Unknown	A_24_P57	2.41738	0	2.41738	6621.773	15976.37
A_23_P67	NM_00064	Homo sapi	IL11,AGIF	2.41661	1.28E-11	2.41661	413.3421	1000.256
A_24_P49800		Unknown	A_24_P49	2.41604	1.24E-32	2.41604	20779.41	49989.52
A_23_P83	NM_00476	Homo sapi	PSCD1,B2	2.4115	1.01E-19	2.4115	3600.313	8777.69

A_24_P30	NM_02115	protein kin	TRIB3,NIP	2.41138	2.80E-45	2.41138	1050.755	2537.754
A_32_P68350		Unknown	THC19611	2.40691	0.00062	2.40691	87.92558	206.8259
A_23_P20	AK023077	Ubiquitin s	USP36	2.405	0	2.405	1326.932	3184.789
A_24_P72	AL133620	Likely hom	KIDINS22C	2.40444	8.11E-07	2.40444	183.865	439.762
A_24_P38	NM_18249	hypothetic	LRP5L,DKI	2.39981	4.80E-14	2.39981	407.1343	987.079
A_23_P25	NM_01763	collaborate	CARF,FLJ;	2.3998	4.93E-26	2.3998	1902.804	4591.301
A_32_P82	BX538288	Hypothetic	DKFZp779	2.39939	1.24E-23	2.39939	341.1295	820.3436
A_32_P22	BF977562	Transcribe	BF977562	2.39878	1.91E-11	2.39878	254.6581	611.7004
A_23_P71	NM_20334	MSFL2541	UNQ2541	2.39735	2.12E-25	2.39735	485.2498	1163.938
A_23_P58	NM_00598	Homo sapi	S100P,MIC	2.39669	0	2.39669	106779.6	256689.8
A_23_P12	NM_01345	Homo sapi	BAZ2B,DK	2.39616	1.62E-31	2.39616	268.5186	643.4138
A_23_P81	BC024725	KIAA1223	KIAA1223	2.3959	5.95E-10	2.3959	471.798	1152.953
A_24_P24	NM_00181	centromere	CENPC1,C	2.3902	2.29E-28	2.3902	525.3062	1259.846
A_24_P58	AY358924	KIAA1201	KIAA1201	2.38937	1.47E-18	2.38937	1564.569	3788.499
A_23_P17	NM_05295	Homo sapi	ACRC,NA/	2.38851	3.60E-32	2.38851	498.6926	1195.231
A_23_P12	NM_00168	Homo sapi	ATP2B1,PI	2.383	8.29E-18	2.383	993.9053	2411.998
A_24_P32	NM_01424	zinc finger	ZMYM5,M'	2.3828	1.85E-16	2.3828	245.7361	585.6101
A_32_P17	NM_02020	Homo sapi	C9orf102,S	2.37783	2.99E-15	2.37783	393.2223	942.4117
A_24_P630916		ALU6_HUM	THC19031	2.37528	0	2.37528	820.4533	1948.931
A_23_P66	NM_00341	Homo sapi	ZNF267,H	2.37478	2.80E-45	2.37478	663.2995	1576.907
A_32_P19	AK129547	Homeobox	LOC36003	2.37465	1.22E-40	2.37465	397.0997	943.4599
A_23_P31	NM_00335	UDP-gluc	UGCG,GC	2.37322	3.25E-16	2.37322	700.553	1692.962
A_23_P11	NM_14511	Homo sapi	MAP2K3,M	2.37116	0	2.37116	646.7366	1533.604
A_24_P697437		ALU8_HUM	THC18822	2.36934	4.02E-13	2.36934	207.9619	492.2752
A_23_P80	NM_00108	Homo sapi	AADAC,DA	2.36893	1.30E-27	2.36893	1301.956	3107.801
A_23_P40	NM_00352	Homo sapi	HIST1H2B	2.36836	1.81E-42	2.36836	3136.143	7471.35
A_23_P34	AB067502	KIAA1915	KIAA1915	2.36697	6.07E-17	2.36697	256.6518	607.522
A_24_P15	NM_17346	hypothetic	DKFZp434	2.36431	0.00006	2.36431	91.61853	214.3717
A_24_P92	NM_00675	Homo sapi	SSFA2,CS	2.36159	0.00013	2.36159	199.8925	473.5423
A_23_P14	NM_01867	Homo sapi	THSD1,TM	2.36085	6.38E-15	2.36085	478.4719	1143.534
A_23_P78	NM_02163	Homo sapi	ZNF350,ZF	2.3571	4.94E-12	2.3571	725.0504	1740.977
A_23_P16	NM_00181	Homo sapi	CENPC1,C	2.35562	1.95E-15	2.35562	417.3243	992.2074
A_24_P22	NM_02276	hypothetic	FNDC3B,F	2.35502	1.50E-36	2.35502	1564.558	3705.442
A_23_P70	NM_00354	Homo sapi	HIST1H4L,	2.35296	4.66E-35	2.35296	50601.95	119942.9
A_23_P55	NM_14590	Homo sapi	RIOK3,SUI	2.35159	2.78E-23	2.35159	572.5343	1355.511
A_23_P46673		Unknown	ENST0000	2.35074	2.94E-39	2.35074	1732.919	4091.643
A_24_P16	NM_00354	H4 histone	HIST1H4B	2.3431	3.99E-33	2.3431	55217.18	130396.8
A_32_P10	NM_14506	Homo sapi	C6orf113,d	2.33515	2.83E-27	2.33515	1054.016	2486.285
A_24_P88	XM_38010	hypothetic	LOC40250	2.33411	1.03E-11	2.33411	1301.785	3097.624
A_23_P14	NM_05816	Homo sapi	MOGAT1,M	2.33313	0.00095	2.33313	65.88981	153.548
A_23_P87	NM_00576	Homo sapi	AASS,LKR	2.33076	6.56E-08	2.33076	241.4301	565.1577
A_24_P14	NM_02106	H2B histon	HIST1H2B	2.32976	0	2.32976	1350.233	3148.698
A_23_P37	NM_17321	Homo sapi	NFAT5,NF	2.3291	1.40E-10	2.3291	257.9332	601.9346
A_23_P49	NM_02430	Homo sapi	FA2H,FAA	2.32601	1.87E-18	2.32601	1027.844	2423.882
A_23_P13	NM_17072	Homo sapi	KCNJ14,IR	2.32528	1.17E-14	2.32528	771.4744	1825.625
A_23_P19	NM_00605	Nef-associ	TNIP1,VAN	2.32186	3.34E-11	2.32186	217.9049	505.7519
A_32_P206308		Unknown	THC19453	2.32171	1.61E-07	2.32171	1057.835	2514.231
A_23_P21	NM_02208	Chromosor	C1orf24	2.31988	1.94E-16	2.31988	246.232	571.3063
A_32_P19	NM_19828	hypothetic	LOC34911	2.3191	1.76E-07	2.3191	647.8763	1548.437
A_23_P32	NM_03075	Oxysterol t	OSBP2	2.31597	0.00041	2.31597	337.9874	798.3864
A_24_P91	NM_01326	myosin reg	MYLIP,MIF	2.31386	1.58E-19	2.31386	358.673	832.5121
A_24_P91	AY129015	Amylase, a	AMY2B	2.31025	0	2.31025	2685.101	6198.005

A_32_P26144	Unknown	THC18395	2.30646	0.00074	2.30646	103.7226	236.6449
A_24_P33	NM_01909	hypothetic C1orf183,F	2.30354	9.26E-13	2.30354	10257.18	23864.45
A_23_P93	NM_00352	Homo sapi HIST1H2B	2.3026	1.67E-29	2.3026	1353.281	3151.125
A_24_P13	AB037771	Ubiquitin s USP53	2.30118	0.00043	2.30118	712.9099	1645.214
A_23_P98	NM_01223	Homo sapi SIRT1,SIR	2.30028	6.00E-14	2.30028	1465.471	3437.497
A_24_P54	BC068044	Histone 2, HIST2H2A	2.30013	9.71E-28	2.30013	3100.199	7205.91
A_24_P24	NM_14575	Homo sapi PHLDB2,LI	2.29939	2.12E-17	2.29939	2312.197	5371.855
A_24_P76210	TVMVRR	THC19532	2.29832	6.98E-31	2.29832	6672.347	15295.77
A_23_P23	NM_01496	Homo sapi WDR47,KI	2.29777	0.00007	2.29777	129.8173	295.6436
A_24_P59	BC063888	Homo sapi BC063888	2.29492	2.47E-21	2.29492	5284.596	12131.04
A_24_P255786	Unknown	A_24_P25	2.2901	9.29E-27	2.2901	17390.78	39802.38
A_23_P50	NM_07847	Homo sapi BTN2A1,B	2.28974	3.77E-38	2.28974	551.2116	1264.81
A_23_P28397	Unknown	A_23_P28	2.28752	5.02E-35	2.28752	1318.01	3027.697
A_23_P21	NM_00354	H4 histone HIST1H4C	2.28692	4.68E-22	2.28692	182297.6	421384.6
A_24_P49	AL050210	Metastasis MALAT1	2.28666	2.47E-14	2.28666	1308.589	3020.819
A_23_P32	NM_01535	Homo sapi MORC3,N	2.28489	9.06E-10	2.28489	1375.992	3203.305
A_24_P93	AY082339	Homo sapi unknown	2.2835	5.77E-06	2.2835	97.66119	222.2672
A_23_P34	NM_03339	Homo sapi KIAA1754,	2.28018	3.80E-08	2.28018	4303.913	9936.298
A_23_P21	NM_01771	Homo sapi SNRK,HSN	2.27832	5.48E-10	2.27832	232.3261	529.0494
A_23_P20	NM_00696	zinc finger ZNF24,KO	2.27588	0	2.27588	15875.75	36241.88
A_23_P57	NM_02110	claudin 1 CLDN1,CL	2.27539	1.33E-14	2.27539	636.4771	1470.782
A_23_P31	NM_17379	Homo sapi PAPD4,FL	2.27516	4.82E-10	2.27516	368.657	847.7979
A_23_P86	NM_00022	Homo sapi LAMB3,LA	2.27393	1.07E-38	2.27393	16067.88	36419.71
A_24_P15	NM_00374	insulin rece IRS2	2.27235	7.44E-24	2.27235	5699.801	13107.78
A_24_P28	NM_00315	serine/thre CDKL5,ST	2.27222	7.27E-09	2.27222	170.8426	386.9252
A_23_P16	NM_03141	Homo sapi GABARAP	2.27189	1.38E-24	2.27189	876.1502	1991.24
A_23_P39	NM_00353	H4 histone HIST1H4D	2.27104	3.32E-20	2.27104	15077.77	34672.43
A_24_P297302	Unknown	A_24_P29	2.26776	1.85E-06	2.26776	163.6165	376.9058
A_23_P33	AB007870	Homo sapi KIAA0410	2.26558	6.40E-41	2.26558	867.5499	1971.456
A_24_P25	BC001892	NPD014 pr NPD014	2.26421	4.20E-24	2.26421	993.3457	2273.188
A_23_P11	NM_00016	Homo sapi GIPR,MGC	2.26222	1.25E-33	2.26222	3563.954	8032.497
A_32_P27	BC010117	Brevican BCAN	2.26042	8.61E-17	2.26042	9193.217	20629.1
A_24_P18	NM_00057	decay acce DAF,CR,T	2.25818	1.42E-31	2.25818	7215.53	16392.71
A_23_P15	NM_01532	Homo sapi FEM1B,FI	2.25802	3.30E-22	2.25802	411.8882	933.1338
A_23_P16	NM_00242	Homo sapi MMP1,CLC	2.25741	8.97E-09	2.25741	224.9648	507.9626
A_23_P39	NM_01527	Homo sapi KIAA1033	2.25671	3.37E-40	2.25671	455.8391	1028.244
A_24_P850172	Unknown	A_24_P85	2.2563	1.24E-30	2.2563	27148.63	61130.3
A_24_P84	AB007962	KIAA0493 KIAA0493	2.25441	0.00003	2.25441	237.5967	539.5592
A_23_P13	NM_00601	Homo sapi ARMET,AF	2.25324	1.76E-32	2.25324	96190.62	216296.7
A_23_P11	NM_00352	H2B histon HIST1H2B	2.25223	4.60E-23	2.25223	4352.798	9925.146
A_24_P16	NM_02110	claudin 1 CLDN1,CL	2.24545	0.00007	2.24545	187.7298	419.106
A_24_P92	NM_00675	Homo sapi SSFA2,CS	2.24542	0.00005	2.24542	92.19785	208.1438
A_23_P16	NM_00595	Homo sapi MT1H,MT1	2.24364	6.37E-17	2.24364	10359.84	23622.29
A_23_P92	NM_01477	Homo sapi TSC22D2,	2.24023	2.25E-10	2.24023	551.5019	1251.165
A_32_P23	NM_15278	hypothetic MGC4040	2.23863	2.40E-18	2.23863	735.8832	1658.407
A_24_P38	NM_00523	v-ets avian ETS2	2.23645	6.71E-33	2.23645	1290.893	2882.601
A_23_P65	AK098698	Thioredoxin TXNRD1	2.23641	3.77E-35	2.23641	1297.695	2914.347
A_23_P16	BC005130	Hypothetic MGC1108	2.23573	1.16E-09	2.23573	183.2192	407.9964
A_23_P94	NM_00070	Homo sapi ANXA1,AN	2.23476	0	2.23476	85284	190155.5
A_23_P10	NM_01232	Homo sapi MAFF,U-M	2.23331	9.49E-07	2.23331	612.0698	1363.382
A_24_P264413	Unknown	A_24_P26	2.22708	5.00E-21	2.22708	486.8766	1082.289
A_24_P37	NM_00352	H2B histon HIST1H2B	2.22466	1.40E-17	2.22466	4089.383	9259.465

A_32_P142363	Unknown	THC19324	2.21375	2.76E-08	2.21375	232.5699	513.5953
A_23_P29	AB029551	RING1 and RYBP	2.21324	8.39E-37	2.21324	494.6155	1095.862
A_23_P20	NM_00595	metallothio MT1G,MT1	2.2112	0.00003	2.2112	3956.327	8842.035
A_23_P35	NM_01563	DKFZP566 EPC2,DKF	2.2092	1.15E-29	2.2092	1075.687	2392.583
A_23_P21	NM_01744	Homo sapi C21orf91,Y	2.20832	2.10E-07	2.20832	344.9337	771.4896
A_23_P30	NM_00351	Homo sapi HIST2H2A	2.20777	6.69E-11	2.20777	1145.421	2544.294
A_24_P12	NM_00515	actin, alpha ACTC	2.20745	0	2.20745	7043.409	15547.74
A_24_P914102	Unknown	A_24_P91	2.20549	6.21E-22	2.20549	261.417	576.5096
A_23_P76	NM_01658	Homo sapi IL23A,P19,	2.2033	2.61E-08	2.2033	298.5809	660.8878
A_23_P42	NM_01445	Homo sapi TNFRSF21	2.20063	3.44E-06	2.20063	112.3422	247.5664
A_24_P161393	Unknown	A_24_P16	2.19987	6.98E-27	2.19987	22539.14	49395.13
A_24_P34	NM_00358	dual-specific DYRK3,RE	2.19954	9.70E-35	2.19954	350.4449	770.8522
A_24_P24	NM_00220	integrin, alpha ITGA2,BR,	2.19879	4.74E-09	2.19879	211.1223	463.6426
A_23_P11	NM_02479	Homo sapi ABHD9,FL	2.19381	5.70E-24	2.19381	30976.16	68118.86
A_24_P91	NM_00071	bradykinin BDKRB1,B	2.19339	1.75E-06	2.19339	294.6257	652.0642
A_23_P36	NM_02291	vasculin GPBP1,GF	2.1916	1.23E-31	2.1916	5527.979	12198.49
A_24_P475115	Unknown	A_24_P47	2.19039	0	2.19039	5941.249	12999.87
A_32_P79719	Unknown	A_32_P79	2.19029	0	2.19029	9700.165	21210.32
A_23_P34	NM_01804	Homo sapi GPATC2,F	2.18844	2.60E-12	2.18844	830.7869	1842.514
A_32_P43	NM_00481	Homo sapi C9orf61,X1	2.18782	0.00026	2.18782	100.0839	216.9224
A_32_P13	AB051464	Kelch-like KLHL15	2.18736	0	2.18736	1187.484	2595.336
A_32_P88	AK090515	Hypothetic LOC28366	2.18411	6.79E-08	2.18411	169.8648	370.2606
A_32_P11	AK128746	Hypothetic LOC39974	2.18192	1.28E-31	2.18192	943.5535	2067.431
A_23_P13	NM_00598	Homo sapi SNAI1,SN	2.18113	1.12E-44	2.18113	2015.901	4405.136
A_23_P86	NM_01654	Homo sapi IER5,SBBI	2.17786	3.07E-31	2.17786	21806.56	47310.2
A_23_P32	NM_00712	ubiquitous UTY,UTY1	2.1725	1.19E-06	2.1725	174.583	376.8597
A_23_P16	NM_02106	H2A histon HIST1H2A	2.16995	3.27E-33	2.16995	5131.91	11194.36
A_23_P20	NM_00229	lymphocyte LCP1,CP6	2.16759	0	2.16759	6770.962	14692.14
A_32_P178945	Unknown	THC18077	2.16541	2.77E-13	2.16541	2649.369	5849.61
A_23_P10	NM_00585	Homo sapi CDK2AP2,	2.16477	1.33E-37	2.16477	1074.54	2328.355
A_23_P39	NM_17358	Homo sapi PGM2L1,B	2.16367	9.05E-10	2.16367	289.107	627.2326
A_23_P10	NM_17583	Homo sapi SMOX,PA	2.15983	1.10E-19	2.15983	18780.61	40928.41
A_23_P33	NM_17505	Homo sapi HIST3H2B	2.15839	5.82E-17	2.15839	2862.95	6268.549
A_32_P225870	Unknown	A_32_P22	2.15656	3.83E-25	2.15656	727.7316	1563.12
A_24_P187626	Unknown	A_24_P18	2.15537	1.84E-18	2.15537	32994.86	70903.69
A_23_P31	AK024335	Intracellular PLA2(GAM	2.154	7.38E-13	2.154	2078.196	4534.618
A_23_P32	AL832398	Hypothetic MGC26717	2.15399	7.66E-27	2.15399	527.3909	1140.123
A_23_P40	NM_01744	Homo sapi H2BFS	2.14911	7.57E-44	2.14911	1004.057	2163.026
A_24_P37	NM_00403	interferon gamma IRF7,IRF7,	2.14873	2.07E-18	2.14873	1301.085	2815.803
A_24_P71	BC033793	KIAA1238 KIAA1238	2.14757	1.95E-08	2.14757	180.5543	387.6097
A_32_P15	AY014272	Similar to FFKSG30	2.14683	2.16E-12	2.14683	55398.74	118884.3
A_24_P12	NM_00230	leukemia inducer LIF,CDF,H	2.14485	0	2.14485	2002.21	4294.996
A_32_P15	NM_00161	actin, gamma ACTG1,AC	2.1416	0	2.1416	107955.3	230807.9
A_23_P44	NM_00161	Homo sapi ACTG1,AC	2.14002	0	2.14002	124599.7	266325.9
A_23_P55	BC006340	Zinc finger, ZCCHC2	2.1393	0.00003	2.1393	171.851	365.6005
A_23_P15	NM_02478	Homo sapi TMC5,FLJ	2.13853	0.00004	2.13853	95.28458	204.1029
A_24_P35	NM_00409	ephrin-B2 EFNB2,HT	2.1366	1.47E-10	2.1366	2673.903	5734.859
A_23_P15	NM_00562	syndecan 1 SDCBP,ST	2.13359	2.23E-19	2.13359	5359.517	11570.66
A_23_P24	AK075342	Hypothetic MGC1094	2.13111	5.95E-13	2.13111	566.6217	1221.166
A_23_P34	AB051464	Kelch-like KLHL15	2.13106	2.14E-24	2.13106	1005.857	2160.356
A_32_P90	BM726576	UI-E-EJ0-a BM726576	2.13085	1.06E-08	2.13085	262.963	561.2302
A_24_P19	NM_02195	guanine nucleotide GNGT1,GN	2.13042	0.0008	2.13042	86.71893	186.3189

A_24_P15	NM_00352	H2B histon	HIST2H2B	2.12779	1.26E-44	2.12779	2720.836	5799.152
A_24_P55	NM_02105	H2B histon	HIST1H2B	2.12741	1.25E-17	2.12741	1223.905	2635.432
A_24_P29	D50917	SERTA do	SERTAD2	2.12403	1.24E-16	2.12403	843.8306	1818.943
A_23_P10	BC050568	Meteorin, g	METRNL	2.12209	2.55E-40	2.12209	4958.763	10540.26
A_24_P30	NM_00535	LIM domain	LMO7, LOM	2.12186	9.09E-18	2.12186	522.7336	1116.495
A_24_P45	AK123446	CDNA FLJ	AK123446	2.11174	2.48E-20	2.11174	2015.828	4267.68
A_32_P134451	Unknown	A_32_P13		2.11152	7.68E-13	2.11152	580.0494	1225.404
A_23_P64	NM_15232	Homo sapi	ZNF641, FL	2.11023	3.68E-07	2.11023	216.7188	459.3959
A_23_P27	NM_01527	ubiquitin-pr	NEDD4L, R	2.10807	3.19E-26	2.10807	902.7368	1917.769
A_24_P52	BC031320	LOC44030	BC031320	2.10624	0	2.10624	6614.901	13944.22
A_24_P91	NM_15260	hypothetic	ZNF709, FL	2.10405	2.02E-25	2.10405	817.6272	1735.876
A_23_P40	NM_14466	hypothetic	FAM76B, M	2.10399	3.31E-09	2.10399	351.406	744.4456
A_32_P107797	Unknown	A_32_P10		2.10253	2.21E-15	2.10253	1021.414	2175.812
A_23_P34	NM_00508	DNA segm	DXYS155E	2.1024	9.26E-43	2.1024	12022.47	25336.33
A_24_P212875	Unknown	A_24_P21		2.10229	7.11E-08	2.10229	160.3657	335.6579
A_23_P36	NM_15228	Homo sapi	SCYL1BP1	2.10153	2.00E-27	2.10153	983.6519	2074.942
A_24_P306614	Unknown	A_24_P30		2.09892	8.30E-32	2.09892	4056.7	8527.21
A_23_P14	NM_01634	PDZ doma	RAPGEF6,	2.09889	3.86E-10	2.09889	387.4167	819.0833
A_32_P49035	Unknown	A_32_P49		2.09458	1.68E-23	2.09458	2517.822	5321.758
A_23_P15	NM_00714	Homo sapi	ZNF184	2.0904	1.73E-12	2.0904	580.5469	1219.177
A_24_P92	AK096990	Homo sapi	CHCHD5, C	2.08964	1.01E-09	2.08964	7526.935	15763.79
A_23_P43	NM_00492	calbindin 1	CALB1, CA	2.08691	0.00002	2.08691	312.0068	656.0155
A_23_P21	NM_00348	Homo sapi	NRIP1, RIP	2.08372	2.95E-21	2.08372	2054.039	4310.447
A_23_P45	NM_01622	Homo sapi	ZNF588, ZF	2.08356	7.51E-17	2.08356	274.5458	571.8985
A_23_P20	NM_01791	Hsp90-ass	CDC37L1, I	2.08098	4.20E-45	2.08098	648.923	1350.691
A_24_P23	BC063888	Homo sapi	BC063888	2.08083	4.98E-15	2.08083	38932.29	80723.34
A_23_P30	NM_00352	Homo sapi	HIST1H2B	2.08066	0	2.08066	5869.821	12212.98
A_24_P90	BQ188387	Hypothetic	MGC2747	2.07828	8.91E-12	2.07828	667.1538	1402.4
A_23_P15	NM_17199	Homo sapi	RAB39B	2.07738	2.04E-07	2.07738	423.2269	889.8165
A_23_P30	NM_00512	Homo sapi	NR1D2, RV	2.07681	3.71E-11	2.07681	832.4982	1743.753
A_23_P52	NM_01234	Homo sapi	BAMBI, NM	2.07673	3.22E-18	2.07673	1894.471	3962.834
A_32_P108567	Unknown	A_32_P10		2.07654	3.55E-12	2.07654	693.2669	1442.047
A_23_P16	NM_00574	Homo sapi	TOB1, TOB	2.07645	6.12E-33	2.07645	6353.943	13251.62
A_23_P64	NM_01500	Homo sapi	STK38L, NI	2.07604	1.06E-22	2.07604	581.5581	1211.068
A_23_P42	NM_00352	Homo sapi	HIST1H2B	2.07493	1.53E-37	2.07493	8353.654	17282.52
A_24_P23	AB062480	SH3 multip	SH3MD2	2.07037	1.47E-08	2.07037	124.6247	257.8898
A_32_P60687	Unknown	A_32_P60		2.06755	7.74E-07	2.06755	2306.346	4875.816
A_24_P39	NM_00703	DnaJ (Hsp	DNAJB4, D	2.06755	6.26E-17	2.06755	1243.518	2592.913
A_24_P21	NM_00351	H2A histon	HIST1H2A	2.06736	0	2.06736	9409.798	19469.97
A_23_P10	NM_00030	Homo sapi	PMP22, DS	2.06653	3.42E-36	2.06653	4009.026	8307.38
A_23_P38	NM_01774	hypothetic	FLJ20291, I	2.06372	1.19E-16	2.06372	415.5278	860.1361
A_23_P12	NM_00358	Homo sapi	DYRK3, RE	2.06182	5.42E-19	2.06182	698.7147	1452.587
A_23_P35	NM_13871	Homo sapi	NFAT5, NF	2.06151	8.84E-07	2.06151	943.7672	1992.882
A_23_P10	NM_00014	Homo sapi	FUT1, H, H	2.06145	1.37E-07	2.06145	338.2642	699.6088
A_24_P92	AF085846	KIAA1961	KIAA1961	2.05985	5.80E-30	2.05985	492.1307	1012.837
A_23_P14	BC028291	FERM dom	FRMD4B	2.05949	5.79E-07	2.05949	556.6456	1154.897
A_23_P31	NM_01766	Homo sapi	SUHW3, ZN	2.0575	1.24E-14	2.0575	1276.225	2652.805
A_23_P13	NM_01826	hypothetic	ZNF701, FL	2.05743	1.95E-08	2.05743	242.613	500.5132
A_23_P17	NM_00645	Homo sapi	PDLIM5, L	2.05611	0.00002	2.05611	1436.064	2969.918
A_32_P1434	ALU6_HUM	THC18775		2.05317	6.20E-06	2.05317	199.6741	409.359
A_24_P41	XM_37295	Homo sapi	ENST0000	2.05065	1.47E-11	2.05065	40651.71	82999.77
A_24_P383901	Unknown	A_24_P38		2.04795	6.45E-23	2.04795	28704.12	58592.11

A_23_P30	NM_13836	hypothetic	LOC90799	2.04625	6.96E-32	2.04625	5030.531	10336.39
A_23_P38	NM_01527	ubiquitin-pr	NEDD4L,R	2.04612	1.25E-07	2.04612	755.731	1570.891
A_23_P65	NM_01424	Homo sapi	ZMYM5,M	2.04565	1.79E-14	2.04565	534.6992	1100.53
A_23_P156609	Unknown	A_23_P15		2.04493	0.0001	2.04493	560.2736	1155.957
A_23_P45	NM_00264	Homo sapi	SERPINB8	2.04234	4.43E-08	2.04234	6848.859	13924.33
A_24_P247175	Unknown	ENST0000		2.04149	4.87E-28	2.04149	59777.48	121752.2
A_23_P20	NM_01472	Homo sapi	SLK,STK2,	2.04011	1.72E-34	2.04011	1902.724	3894.685
A_23_P10	NM_01865	golgin-like	GOLGA6,C	2.03785	3.27E-14	2.03785	516.9908	1054.386
A_23_P20	NM_01431	Homo sapi	DDX58,RIC	2.03746	3.65E-36	2.03746	605.1359	1233.801
A_24_P32	NM_13836	hypothetic	LOC90799	2.03454	4.12E-12	2.03454	840.8709	1735.033
A_23_P67	NM_15325	gonadotrop	GIOT-1,GI	2.02772	8.16E-07	2.02772	234.868	475.9133
A_23_P394562	X-linked re	RPGR		2.0261	0.00003	2.0261	121.4555	244.7406
A_32_P113533	Unknown	A_32_P11		2.0251	1.14E-27	2.0251	8321.632	16819.48
A_32_P60632	Unknown	A_32_P60		2.02496	2.85E-11	2.02496	1467.279	2954.832
A_23_P15	NM_01635	epithelial p	EPLIN,SRE	2.02491	1.67E-16	2.02491	4581.374	9422.16
A_24_P88	NM_02100	Homo sapi	UBC,HMG	2.0245	1.33E-18	2.0245	277959.7	560889
A_23_P25	NM_00341	Homo sapi	ZNF7,KOX	2.02351	5.38E-40	2.02351	763.434	1546.06
A_23_P70	NM_00220	Homo sapi	ITGA2,BR,	2.02189	1.44E-06	2.02189	201.4324	403.8861
A_24_P17	NM_01520	KIAA0874	ANKRD12,	2.02054	7.65E-10	2.02054	431.9592	881.1141
A_23_P13	U09410	Zinc finger	ZNF131	2.0193	5.00E-21	2.0193	3022.429	6157.759
A_23_P21	AF218008	LOC44050	unknown	2.01843	1.23E-10	2.01843	38212.16	76802.35
A_23_P37	AB046824	KIAA1604	KIAA1604	2.01833	4.47E-12	2.01833	590.9015	1214.988
A_23_P13	NM_15371	Homo sapi	TTL,MGC4	2.01447	2.26E-06	2.01447	240.1216	483.6455
A_24_P15	NM_02248	zinc finger	ZNF336,G	2.01362	2.37E-06	2.01362	197.6503	397.2903
A_32_P187975	U39779	be THC19107		2.01164	4.52E-11	2.01164	1686.824	3417.152
A_23_P77	NM_02471	hypothetic	C15orf29,F	2.01134	0.00004	2.01134	189.205	377.1272
A_23_P80	NM_00351	Homo sapi	HIST1H2B	2.01108	0	2.01108	10663.45	21436.57
A_24_P11	NM_02407	hypothetic	SECISBP2	2.011	3.40E-15	2.011	4255.769	8694.088
A_23_P55	NM_02023	putative pr	NBLA0383	2.01052	0.0001	2.01052	813.9639	1616.266
A_32_P14	NM_01765	hypothetic	ZNF562,FL	2.00938	8.18E-18	2.00938	371.0181	746.2604
A_23_P23	NM_14523	Homo sapi	ZNF31,KO	2.0091	5.53E-13	2.0091	399.7789	805.1424
A_32_P23	NM_18107	88-kDa gol	GOLGA8A	2.00815	2.62E-10	2.00815	461.0069	930.1319
A_23_P14	NM_01567	Homo sapi	GADD45B,	2.00709	4.51E-15	2.00709	2736.788	5519.608
A_23_P31	NM_01248	zinc finger	ZNF281,ZE	2.00692	3.77E-21	2.00692	1300.086	2631.707
A_32_P50	BQ188770	Transcribe	BQ188770	2.00536	0.00063	2.00536	236.6468	478.2938
A_24_P15	XM_37300	similar to H	LOC39156	2.00392	1.16E-17	2.00392	488.7856	983.0941
A_23_P85	NM_03229	hypothetic	GON4,GOI	2.00338	5.08E-15	2.00338	1480.047	2993.5
A_24_P22	NM_00110	actin, beta	ACTB	2.00271	5.04E-20	2.00271	72887.89	144815.2
A_24_P13	NM_00155	interferon-r	IFRD1,PC	2.002	1.40E-45	2.002	6929.858	13892.02
A_24_P84	XM_37328	Homo sapi	XM_37328	2.00157	0.00007	2.00157	477.713	982.4038
A_24_P10	AK130825	Isolate A8	AK130825	2.00103	3.84E-08	2.00103	582.4628	1170.651
A_32_P135450	AL163227	THC19683		2.00082	4.51E-07	2.00082	202.8693	403.5605
A_32_P17	BC035034	Transmem	TMEM41B	1.999	2.11E-15	1.999	1380.323	2784.685
A_24_P84	AK055809	Hypothetic	LOC14870	1.99769	0	1.99769	6665.501	13315.99
A_32_P11	NM_00196	eukaryotic	EIF4A2,DC	1.99745	8.99E-11	1.99745	2010.344	4043.471
A_23_P41	NM_02017	Homo sapi	FEM1C,FE	1.99424	2.87E-12	1.99424	779.5378	1561.119
A_23_P10	NM_02282	Homo sapi	MARCH7,/	1.99392	7.41E-15	1.99392	642.8432	1289.237
A_24_P27	AK023542	Homo sapi	FLJ13480	1.9932	1.34E-10	1.9932	368.392	735.9269
A_23_P14	NM_00667	Homo sapi	ARID5A,M	1.9929	1.47E-06	1.9929	286.9729	574.0554
A_23_P15	NM_03216	Homo sapi	ZNF394,FL	1.99288	3.13E-23	1.99288	1578.888	3156.3
A_23_P41	NM_15240	Homo sapi	GRPEL2,F	1.99271	2.11E-15	1.99271	3367.405	6741.56
A_23_P21	NM_14449	Homo sapi	AKAP12,A	1.99226	0.00001	1.99226	659.9252	1306.954

A_24_P13	NM_00688	CD86 antigen CD86,B70,	1.99187	7.72E-07	1.99187	693.7468	1426.819
A_23_P56	NM_00333	ubiquitin A-UBA52,CE	1.99157	2.19E-15	1.99157	40695.75	80615.92
A_32_P46191	Unknown	A_32_P46	1.98912	2.61E-22	1.98912	4101.464	8207.423
A_23_P14	NM_00334	ubiquitin-cc UBE2H,UE	1.98881	1.67E-27	1.98881	1804.125	3595.911
A_23_P35	NM_00113	Homo sapi AFG3L1,Al	1.98836	1.19E-22	1.98836	4355.354	8705.507
A_24_P153207	zinc finger	ZNF565	1.98776	1.44E-07	1.98776	489.9846	984.6052
A_23_P10	NM_03219	homolog of INO80	1.98752	6.01E-37	1.98752	6262.22	12466.87
A_24_P126731	Unknown	ENST0000	1.98618	1.24E-25	1.98618	3297.19	6555.077
A_32_P12703	Unknown	THC18723	1.98613	2.69E-20	1.98613	661.6346	1324.476
A_23_P22	NM_00116	baculoviral BIRC4,API	1.98193	3.26E-08	1.98193	828.2522	1682.597
A_23_P89	NM_02404	hypothetic MGC3020	1.9794	2.67E-12	1.9794	424.8116	841.0431
A_23_P51	NM_00703	Homo sapi DNAJB4,D	1.97807	4.02E-22	1.97807	710.4805	1411.084
A_24_P35	AB037791	Hypothetic FLJ10980	1.9774	4.95E-10	1.9774	673.084	1341.421
A_23_P12	NM_02111	Homo sapi CRY2,HCF	1.97739	3.08E-13	1.97739	761.7329	1512.661
A_23_P12	NM_01242	Homo sapi RLF,Zn-15	1.97517	1.24E-09	1.97517	2252.647	4478.698
A_23_P10	NM_02514	hypothetic FLJ13448	1.97509	1.42E-06	1.97509	204.5324	402.6626
A_23_P20	NM_01759	transmemb VEZATIN,I	1.97402	3.71E-20	1.97402	3007.044	5992.767
A_23_P39	AL833005	MRNA; cDI AL833005	1.97392	5.46E-25	1.97392	5078.688	10025.15
A_24_P19	NM_00342	zinc finger ZNF85,HP	1.97162	5.19E-28	1.97162	2234.891	4425.224
A_32_P128773	Unknown	A_32_P12	1.96993	1.43E-14	1.96993	679.8842	1341.148
A_23_P20	NM_05824	Homo sapi CCNT2	1.96932	0.00004	1.96932	139.2574	276.299
A_32_P68148	Unknown	A_32_P68	1.96829	7.22E-16	1.96829	1980.904	3936.166
A_23_P52	NM_17074	Homo sapi UNC5B,UN	1.96676	2.09E-22	1.96676	1088.092	2148.167
A_24_P38	AK074675	Protein kin PRKAG2	1.96665	0.00053	1.96665	139.386	274.4764
A_24_P62	AY358241	Hypothetic MGC40368	1.96551	0.00022	1.96551	102.2357	201.4805
A_23_P25	NM_01959	Homo sapi LRRC8A,L	1.96446	4.27E-34	1.96446	6791.851	13368.47
A_23_P30	NM_01586	PR domain PRDM2,RI	1.96398	1.93E-07	1.96398	448.9625	891.5665
A_24_P18	NM_03064	Homo sapi DUSP16,M	1.95985	2.07E-16	1.95985	486.0822	954.6267
A_32_P51	BC053362	Zinc finger ZC3HDC5	1.95925	6.72E-16	1.95925	1967.034	3890.468
A_23_P94	NM_01339	Homo sapi TMEM2	1.95873	2.09E-31	1.95873	1837.336	3604.235
A_32_P10	BQ013066	Vestigial lik VGLL4	1.95786	0.00005	1.95786	318.7122	628.4039
A_23_P40	NM_19935	Hypothetic LOC37575	1.95701	1.83E-08	1.95701	492.4428	953.7123
A_32_P19	NM_01899	hypothetic KCTD5,FL	1.95557	6.40E-09	1.95557	2982.503	5858.088
A_23_P25	NM_00155	Homo sapi IFRD1,PC4	1.95436	5.88E-34	1.95436	6855.621	13438.53
A_23_P35	NM_17591	Homo sapi ZNF493,FL	1.95428	9.44E-22	1.95428	2065.132	4056.746
A_24_P63	AF009267	Homo sapi AF009267	1.95327	3.42E-11	1.95327	2318.156	4494.34
A_32_P162716	Unknown	A_32_P16	1.95261	0.00003	1.95261	201.9636	393.0814
A_32_P161140	Unknown	A_32_P16	1.9519	2.88E-12	1.9519	562.2553	1097.821
A_32_P214611	Unknown	THC19045	1.94859	9.51E-25	1.94859	25380.39	49580.18
A_24_P21	NM_00173	Kruppel-lik KLF5,CKLI	1.94794	5.02E-12	1.94794	1238.056	2423.314
A_32_P52	AK026328	Hypothetic FLJ22675	1.94623	1.12E-44	1.94623	5972.626	11616.59
A_24_P13	NM_03089	hypothetic ZNF323,ZN	1.94512	1.21E-22	1.94512	938.0228	1833.419
A_23_P38	NM_01656	Homo sapi TBX3,UMS	1.94511	1.32E-25	1.94511	370.582	720.7815
A_32_P18	BX103476	Transcribe BX103476	1.94464	8.32E-10	1.94464	564.1166	1099.379
A_32_P199217	Unknown	THC19521	1.94444	0.00004	1.94444	124.7654	241.6296
A_32_P18	NM_17855	Homo sapi ZNF680,FL	1.94286	8.70E-22	1.94286	2500.7	4881.166
A_24_P48	NM_19827	Homo sapi CXorf23	1.94238	1.79E-06	1.94238	274.2987	532.8352
A_23_P34	NM_00151	Homo sapi GTF2B,TF	1.9409	2.13E-14	1.9409	2390.531	4707.028
A_23_P42	NM_15243	Homo sapi PIWIL4,HIL	1.94041	0.00004	1.94041	204.1431	397.5714
A_23_P36	NM_00352	Homo sapi HIST1H2B	1.93944	0	1.93944	8663.23	16794.46
A_23_P43	NM_14456	Homo sapi SPOCD1,F	1.93929	7.87E-09	1.93929	2503.196	4822.171
A_23_P15	NM_00171	Homo sapi BF,GBG,C	1.93719	1.42E-15	1.93719	295.0428	571.6429

A_24_P27	NM_00354	H4 histone	HIST1H4G	1.9354	0.00005	1.9354	200.0088	386.004
A_24_P25	NM_03218	hypothetic	ZNF644,Ze	1.93455	2.29E-25	1.93455	514.0606	993.4541
A_32_P86	BX537972	Hypothetic	LOC11606	1.93417	2.87E-21	1.93417	1856.569	3603.913
A_23_P59	NM_00352	Homo sapi	HIST1H2B	1.93356	0	1.93356	7139.488	13809.48
A_23_P92	NM_13363	DNA helica	HEL308,M	1.93308	1.36E-07	1.93308	254.9382	492.2997
A_24_P20	NM_00547	lymphocyte	LNK	1.93296	7.01E-11	1.93296	562.1202	1099.159
A_24_P14	AB007874	Homo sapi	KIAA0414	1.93253	2.40E-08	1.93253	509.0685	985.0574
A_24_P33	NM_00458	Rab gerany	RABGGTB	1.93098	1.50E-24	1.93098	6830.799	13274.02
A_23_P35	NM_00666	phosphodi	PDE10A,H	1.93088	1.79E-06	1.93088	238.4223	459.1351
A_23_P12	NM_02455	Homo sapi	C14orf138,	1.93068	4.45E-09	1.93068	605.8608	1186.692
A_24_P17	NM_00499	methionine	MARS,ME	1.93008	1.18E-11	1.93008	6749.045	12944.68
A_32_P61	975	Unknown	A_32_P61	1.92855	6.84E-08	1.92855	266.6034	514.7545
A_23_P81	NM_08059	Homo sapi	HIST1H2A	1.92644	1.13E-20	1.92644	10019.94	19411.72
A_32_P19	2804	Unknown	A_32_P19	1.92524	5.19E-09	1.92524	188.7393	363.1134
A_23_P61	NM_01745	Homo sapi	BAIAP2,BA	1.92487	8.24E-11	1.92487	1596.862	3098.25
A_32_P21	NM_13344	Homo sapi	CTGLF1,M	1.9247	3.33E-31	1.9247	774.0707	1491.479
A_23_P25	NM_00502	protein ger	PGGT1B,E	1.92423	7.63E-06	1.92423	159.6297	305.4617
A_24_P15	NM_20328	Homo sapi	ZNF539	1.92151	9.06E-13	1.92151	1648.811	3205.362
A_32_P12	7501	Unknown	A_32_P12	1.91968	6.10E-31	1.91968	654.6903	1256.261
A_23_P25	NM_01450	putative R	RABGEF1,	1.91702	1.03E-19	1.91702	1322.03	2512.003
A_23_P11	NM_02106	Homo sapi	HIST1H2B	1.91678	1.95E-25	1.91678	2671.951	5139.251
A_23_P16	NM_01758	Homo sapi	SLC2A6,G	1.91626	8.95E-09	1.91626	207.1681	397.635
A_23_P13	NM_00650	Homo sapi	PTPRE,PT	1.91611	2.51E-06	1.91611	405.5815	783.8004
A_24_P94	NM_01801	hypothetic	C10orf118,	1.91555	1.63E-06	1.91555	276.1538	529.0757
A_23_P21	NM_00194	Homo sapi	HBEGF,DT	1.91495	9.89E-07	1.91495	659.6122	1257.577
A_23_P43	NM_13426	Homo sapi	WSB1,SW	1.9148	1.62E-10	1.9148	4431.521	8613.162
A_24_P17	NM_00538	Natural kill	NKTR	1.91215	1.57E-09	1.91215	343.5505	658.3043
A_24_P39	U16307	GLI pathog	GLIPR1	1.91192	3.29E-26	1.91192	1305.36	2506.772
A_23_P99	AB037791	Hypothetic	FLJ10980	1.91106	1.33E-08	1.91106	895.8281	1742.162
A_23_P30	NM_00062	Homo sapi	BDKRB2,B	1.91048	6.01E-10	1.91048	1136.101	2182.95
A_24_P41	NM_01543	double ring	RNF19,DO	1.9103	0.00003	1.9103	412.8994	800.3206
A_24_P75	3638	Unknown	A_24_P75	1.90938	0.00039	1.90938	7908.056	15207.39
A_24_P41	NM_00620	Phosphodi	PDE4D	1.90759	0.0008	1.90759	174.9571	331.3442
A_32_P21	BC021677	Homo sapi	BC021677	1.90548	0.00002	1.90548	290.3347	555.523
A_24_P46	XM_29281	Homo sapi	XM_29281	1.90428	0	1.90428	5820.56	11083.63
A_23_P20	BC007524	Sperm ass	SPAG9	1.90284	8.34E-09	1.90284	560.6188	1070.954
A_23_P51	NM_00156	Homo sapi	TNFRSF9,	1.90243	9.98E-08	1.90243	1191.886	2270.637
A_24_P64	895	ALU6_HUM	THC18914	1.90232	1.63E-08	1.90232	175.4034	333.9717
A_23_P93	NM_00508	U2 small nt	U2AF1RS1	1.90229	0.00004	1.90229	250.7772	477.9221
A_23_P16	NM_03225	hypothetic	TBC1D3,P	1.89916	3.18E-08	1.89916	1272.553	2460.303
A_23_P88	NM_00646	Homo sapi	ARID3B,BI	1.89854	2.62E-07	1.89854	3394.2	6387.599
A_23_P85	NM_00458	Homo sapi	RABGGTB	1.89807	1.07E-37	1.89807	27353.18	52022.48
A_23_P31	NM_00433	Homo sapi	BNIP2,NIP	1.89763	2.98E-18	1.89763	3783.729	7222.036
A_23_P98	AK002085	Hypothetic	LOC14443	1.89683	2.66E-32	1.89683	1649.689	3135.136
A_23_P21	NM_02114	Homo sapi	UTX,bA38	1.89607	1.32E-09	1.89607	1967.494	3767.423
A_23_P56	NM_00393	Homo sapi	KYNU	1.89566	1.40E-45	1.89566	65158.69	123583.3
A_32_P12	NM_19828	hypothetic	LOC34911	1.89423	4.50E-23	1.89423	8852.566	16797.75
A_24_P83	2831	AF247142	THC19660	1.89401	0.00004	1.89401	177.5263	337.8528
A_24_P86	AK022745	Hypothetic	RNU71B,U	1.89377	6.59E-10	1.89377	13577.38	26162.66
A_23_P20	NM_01798	Homo sapi	SCYL2,CV	1.89327	5.88E-10	1.89327	5554.346	10547.5
A_23_P15	NM_01632	Homo sapi	RHCG,RH	1.8924	4.79E-09	1.8924	1006.82	1909.389
A_24_P16	NM_01214	deleted in	DDX26,HD	1.89183	2.36E-12	1.89183	1547.86	2974.67

A_24_P695669	Unknown	A_24_P695669	1.89112	2.60E-10	1.89112	334.8829	634.1044
A_23_P84	AF257098 Homo sapi	OVN9-3	1.88664	1.16E-06	1.88664	1313.504	2561.629
A_23_P14	(NM_18184 amphoterin	AMIGO2,A	1.88606	1.48E-11	1.88606	2848.246	5477.122
A_23_P97	NM_00585 Homo sapi	IRX5,IRX-2	1.88549	3.88E-06	1.88549	1032.417	1992.02
A_23_P16	NM_00265 plasminoge	PLAUR,CC	1.88521	1.90E-15	1.88521	684.6727	1290.14
A_24_P41	(NM_19807 golgi autoa	FLJ40113,I	1.88453	2.32E-16	1.88453	3400.535	6461.903
A_23_P21	(NM_01336 zinc finger	ZNF222	1.88277	8.50E-14	1.88277	818.917	1543.315
A_23_P10	(NM_02246 Homo sapi	AZI2,AZ2,N	1.88268	2.88E-16	1.88268	4251.837	8087.913
A_23_P32	(NM_00274 Homo sapi	MAPK6,EF	1.88252	2.28E-15	1.88252	4649.068	8812.992
A_23_P29	(NM_00553 interferon	IFNGR2,AI	1.88244	4.85E-11	1.88244	634.1393	1206.334
A_23_P55	(NM_14497 Homo sapi	ZNF564,Mi	1.87583	1.44E-08	1.87583	224.4551	420.6307
A_23_P216071	Unknown	A_23_P216071	1.87442	2.24E-21	1.87442	495.1492	929.2542
A_23_P21	(NM_03220 Homo sapi	PHF20L1,C	1.87399	1.25E-09	1.87399	501.1366	945.915
A_32_P18034	Unknown	THC19642	1.87336	0.00001	1.87336	337.8083	636.2163
A_23_P31	(AK026275 RAB11 fan	RAB11FIP	1.8733	1.81E-15	1.8733	2254.643	4246.531
A_23_P10	(NM_00303 Homo sapi	SIAH1,Sial	1.87031	9.47E-13	1.87031	904.8347	1709.155
A_24_P11	(NM_00116 baculoviral	BIRC2,API	1.87015	1.22E-25	1.87015	1111.603	2086.007
A_23_P43	AF115510 Leucine ric	LRRFIP1	1.86989	1.77E-06	1.86989	232.0228	432.1189
A_23_P40	NM_00113 Homo sapi	AFG3L1,AI	1.86783	3.35E-18	1.86783	1888.617	3547.131
A_32_P141257	BC036372	ENST0000	1.86396	1.29E-08	1.86396	349.9945	653.2106
A_23_P88	NM_01549 DKFZP564	DKFZP564	1.86313	3.24E-28	1.86313	8443.639	15690.24
A_24_P25	(NM_02213 secreted m	SMOC1	1.86243	0.00001	1.86243	4313.059	8026.609
A_24_P11	(NM_00696 zinc finger	ZNF24,KO	1.86145	1.65E-33	1.86145	3767.917	7023.13
A_23_P13	(NM_01324 Homo sapi	CLCF1,CL	1.86095	9.19E-26	1.86095	755.1229	1404.34
A_23_P36	(NM_01395 neuregulin	NRG1,GGI	1.86025	7.83E-09	1.86025	614.2235	1150.946
A_24_P38	NM_01842 RNA polymr	RRN3,TIFI	1.86015	4.30E-15	1.86015	1952.571	3667.231
A_24_P16	XM_29317 similar to	PLOC34371	1.85808	1.18E-07	1.85808	586.74	1102.383
A_23_P12	AK023471 Zinc finger,	ZCCHC6	1.85801	3.94E-11	1.85801	593.3474	1105.159
A_24_P76	AK097080 Hypothetic	AK097080	1.85761	2.95E-09	1.85761	505.9954	947.3848
A_23_P27	NM_00227 Homo sapi	KRT15,K15	1.85686	2.05E-25	1.85686	4911.203	9104.037
A_24_P17	(NM_19845 Homo sapi	ZNF600,KF	1.85685	6.25E-07	1.85685	224.8857	419.3109
A_24_P55	NM_15282 Homo sapi	TES,TESS	1.85673	2.15E-18	1.85673	2758.705	5147.324
A_24_P22	(NM_00351 H2A histon	HIST1H2A	1.85595	8.58E-27	1.85595	1610.126	3001.129
A_23_P36	(NM_01835 hypothetic	C14orf106,	1.8524	9.35E-19	1.8524	1279.727	2381.58
A_24_P14	(NM_00194 diphtheria	tHBEGF,DT	1.85161	1.67E-09	1.85161	704.673	1302.538
A_23_P15	(NM_02485 hypothetic	FLJ22028	1.85126	1.72E-08	1.85126	385.8149	716.7667
A_23_P10	(NM_03249 Homo sapi	ZC3H8,Fliz	1.84964	7.21E-32	1.84964	2528.822	4667.327
A_23_P14	(NM_00443 Homo sapi	EPHA2,EC	1.84898	1.34E-10	1.84898	185.5237	343.0923
A_23_P73	(NM_01955 hypothetic	MOSPD1,I	1.84882	3.88E-12	1.84882	564.7297	1048.969
A_24_P38	(NM_00699 zinc finger	ZNF197,P1	1.84881	6.37E-16	1.84881	390.7145	722.9435
A_23_P32	(NM_00423 Homo sapi	KLF4,EZF,	1.84681	1.59E-16	1.84681	1042.557	1926.061
A_32_P146890	ALU1_HUI	THC18615	1.84636	4.25E-06	1.84636	376.7673	697.2532
A_23_P81	(AF022375 Vascular ei	VEGF	1.84603	2.87E-10	1.84603	2299.363	4224.494
A_23_P97	(BX641020 AT rich inte	ARID5B	1.84527	5.43E-07	1.84527	924.6466	1701.876
A_32_P85230	Unknown	A_32_P85230	1.84481	8.96E-17	1.84481	970.4265	1796.636
A_24_P10	(NM_19853 Homo sapi	ZNF568,FL	1.84468	8.36E-08	1.84468	243.5399	448.3615
A_23_P30	(NM_00354 Homo sapi	HIST1H4K	1.84117	5.86E-07	1.84117	3356.349	6179.595
A_32_P14	(NM_20328 Homo sapi	ZNF539	1.8407	5.74E-18	1.8407	3771.044	6986.199
A_24_P32	(NM_18289 Homo sapi	ARL2L1,Mi	1.8403	1.96E-08	1.8403	3221.242	6027.287
A_23_P56	(NM_00566 Homo sapi	RNF103,KI	1.83974	1.93E-11	1.83974	1190.207	2224.577
A_23_P34	(NM_01989 endoplasm	ERO1LB	1.83955	0.00027	1.83955	156.0387	286.7758
A_24_P475753	ARH5_HUI	THC19878	1.83706	0.00007	1.83706	365.9495	677.7151

A_23_P81: NM_00390	sequestosc	SQSTM1,p	1.83669	3.77E-11	1.83669	81606.73	148771.7
A_24_P256073	Unknown	A_24_P256073	1.83632	3.79E-06	1.83632	148.4265	272.725
A_24_P80: BC041359	Hypothetic	MGC2477	1.83538	6.37E-32	1.83538	1746.996	3209.082
A_32_P118013	ALU1_HUM	THC18274	1.83524	7.92E-06	1.83524	321.305	591.6556
A_24_P24: BC017902	PDZ and L	PDLIM5	1.83513	2.01E-16	1.83513	514.8264	942.3394
A_32_P211026	Unknown	THC18491	1.8342	1.93E-15	1.8342	790.0322	1456.472
A_32_P23731	Unknown	A_32_P23731	1.83354	1.94E-06	1.83354	6300.588	11678.15
A_24_P37: NM_00073	glycoprotei	CGA,HCG,	1.83307	3.88E-09	1.83307	1505.555	2769.22
A_23_P13: NM_02031	Homo sapi	CMKOR1,F	1.83271	1.75E-07	1.83271	536.7412	991.7299
A_24_P27: AL713659	Zinc finger	FLJ31295	1.83171	1.96E-09	1.83171	207.6505	380.0125
A_23_P22: NM_02481	Homo sapi	CXorf45,FL	1.83164	1.47E-19	1.83164	1823.854	3359.325
A_23_P35: AL834280	Homo sapi	DKFZp547	1.83064	0.00048	1.83064	181.7708	333.6885
A_24_P814872	Unknown	A_24_P814872	1.82896	3.66E-09	1.82896	1060.126	1973.112
A_24_P31: AK094969	Hypothetic	FLJ10808	1.8286	4.61E-06	1.8286	215.1644	392.4322
A_24_P22: NM_01637	AT rich inte	ARID4B	1.82832	1.66E-14	1.82832	922.0787	1693.047
A_23_P62: NM_02070	Homo sapi	SFRS15,S	1.82705	0.00022	1.82705	178.1854	323.0576
A_23_P25: NM_01394	Homo sapi	CLIC4,H1,I	1.82661	4.72E-19	1.82661	1448.607	2660.948
A_32_P134090	AB046048	THC19465	1.82654	0.00002	1.82654	195.3411	357.6551
A_24_P897062	Unknown	A_24_P897062	1.82516	0.00009	1.82516	372.871	683.908
A_32_P165823	Unknown	A_32_P165823	1.82396	3.83E-06	1.82396	1076.183	1985.644
A_24_P11: NM_01895	ubiquitin B	UBB,MGC	1.82358	1.06E-15	1.82358	308416.1	560698.4
A_23_P25: NM_15238	hypothetic	FLJ31438	1.82353	2.50E-07	1.82353	292.1605	533.1971
A_23_P42: AL834393	Hypothetic	FLJ39441	1.82078	1.47E-09	1.82078	697.3421	1288.177
A_23_P93: NM_00385	Homo sapi	TRIM24,P1	1.82037	4.33E-17	1.82037	9406.584	17036
A_23_P12: NM_00629	Homo sapi	ZNF193,Pf	1.81979	2.14E-17	1.81979	742.9803	1355.489
A_23_P30: NM_01445	Homo sapi	TNFRSF21	1.81823	9.53E-09	1.81823	1935.25	3529.885
A_23_P13: AK091330	Multiple C2	MCTP1	1.8172	2.64E-08	1.8172	368.887	671.9635
A_23_P61: NM_00659	Homo sapi	SLC12A7,I	1.81667	1.02E-24	1.81667	14126.96	25677.3
A_24_P92: AF113674	Homo sapi	PRO0398	1.8156	1.01E-07	1.8156	175.9182	319.0887
A_24_P78: NM_17237	Homo sapi	ELF1	1.81538	7.04E-27	1.81538	474.2329	860.9509
A_32_P171253	U12390 be	THC18375	1.81487	1.13E-07	1.81487	1601.618	2940.264
A_24_P35: NM_01538	Nipped-B h	NIPBL	1.81473	2.56E-13	1.81473	656.8345	1200.108
A_23_P18: NM_00577	Homo sapi	RBM5,G15	1.81236	2.65E-30	1.81236	6089.633	11058.01
A_23_P37: NM_00380	Homo sapi	RIPK1,RIP	1.81179	0.0002	1.81179	391.2653	715.5717
A_32_P13: BC042539	Homo sapi	LOC44242	1.81067	3.97E-09	1.81067	7058.896	12761.49
A_32_P12: NM_01416	HSPC128	HSPC128,I	1.81029	8.54E-10	1.81029	4094.044	7535.172
A_24_P367326	Unknown	ENST0000	1.81001	1.89E-09	1.81001	675.9241	1226.62
A_24_P15: BX537779	ATPase, C	ATP11B	1.80921	5.61E-13	1.80921	534.176	970.9285
A_24_P35: NM_00603	histone de	HDAC4,HC	1.80889	1.22E-16	1.80889	411.4816	744.8551
A_32_P15: BF210146	Transcribe	BF210146	1.80864	4.74E-15	1.80864	1064.534	1929.018
A_23_P85: NM_00412	Homo sapi	GBP2	1.80747	2.91E-06	1.80747	477.5869	867.4954
A_24_P18: NM_01415	HSPC056	ARMC8,S8	1.80746	4.51E-09	1.80746	757.1381	1384.844
A_23_P21: NM_13869	premature	LOC91431	1.80613	0.00006	1.80613	320.4463	580.7466
A_24_P23: NM_01450	putative R	RABGEF1,	1.8059	1.07E-16	1.8059	1326.211	2409.214
A_24_P799680	Unknown	A_24_P799680	1.80551	0.0003	1.80551	163.3113	294.3188
A_23_P19: AK021777	UDP-N-ac	GALNT10	1.80447	6.48E-35	1.80447	920.9253	1661.771
A_23_P25: NM_01648	Homo sapi	C9orf156,N	1.80428	5.05E-18	1.80428	1505.017	2726.457
A_32_P20: NM_00505	Homo sapi	RANBP2L1	1.80394	2.56E-10	1.80394	933.0891	1697.696
A_24_P19: AB040906	Zinc finger	ZNF492	1.80315	1.92E-34	1.80315	960.5263	1732.555
A_24_P204639	Unknown	ENST0000	1.80305	1.33E-14	1.80305	1523.514	2764.485
A_24_P932253	Unknown	A_24_P932253	1.8029	2.86E-16	1.8029	8715.574	15677.96
A_23_P91: NM_01810	hypothetic	ZNF334	1.80258	3.41E-18	1.80258	533.1317	960.4053

A_23_P22	NM_00362	Homo sapi	PIK3R3,p5	1.80151	1.98E-08	1.80151	619.8718	1129.056
A_24_P11	NM_00393	kynurenina	KYNU	1.80112	7.44E-36	1.80112	8491.77	15312.24
A_24_P92	NM_00538	Natural kill	NKTR	1.80064	4.27E-15	1.80064	3157.774	5712.64
A_24_P30	NM_02106	H2A histon	HIST1H2A	1.80027	3.20E-31	1.80027	7727.347	13937.77
A_23_P21	1468	Unknown	A_23_P21	1.79947	3.48E-15	1.79947	307.8266	553.8835
A_24_P13	NM_00361	oral-facial-c	OFD1,71-7	1.79839	5.17E-20	1.79839	1503.9	2716.402
A_23_P75	NM_00239	solute carri	SLC3A2,4F	1.79837	4.26E-25	1.79837	3563.304	6402.047
A_23_P43	BC042539	Homo sapi	LOC44242	1.79774	8.57E-10	1.79774	1264.325	2265.098
A_23_P32	NM_02092	Homo sapi	WNK3,PR	1.79722	5.65E-06	1.79722	274.4605	493.13
A_24_P84	AK123296	Zinc finger	ZNF585A	1.79652	5.99E-10	1.79652	244.0261	438.7943
A_23_P91	NM_00036	Homo sapi	THBD,TM,	1.79646	9.24E-19	1.79646	3951.847	7143.372
A_24_P81	4096	AE003449	THC18772	1.7947	3.26E-12	1.7947	5875.774	10606.24
A_32_P43	320	Unknown	THC19590	1.79441	7.16E-06	1.79441	147.1328	263.7858
A_32_P19	AW939148	Leiomodin	LMOD1	1.79415	0.00001	1.79415	432.4533	784.2173
A_23_P15	NM_00344	Homo sapi	ZNF140,p	1.79374	1.85E-12	1.79374	382.9724	687.1921
A_23_P15	NM_00298	small induc	CCL5,SISc	1.79257	0.00002	1.79257	563.7584	1010.541
A_32_P88	NM_13344	Homo sapi	CTGLF1,M	1.79246	4.10E-12	1.79246	407.8133	731.7156
A_24_P37	NM_01828	Homo sapi	GBP3,FLJ	1.78961	0.00041	1.78961	203.7359	364.6549
A_24_P17	AY008274	F-box and	'FBXW7	1.78917	3.08E-19	1.78917	1477.03	2657.862
A_23_P34	NM_02031	Homo sapi	C1orf63,NF	1.78711	3.60E-07	1.78711	815.4117	1467.878
A_23_P40	NM_19856	FLJ32363	FLJ32363,	1.78674	8.62E-06	1.78674	407.0109	732.6105
A_23_P88	NM_00664	Homo sapi	HSPH1,HS	1.78654	7.49E-35	1.78654	28534.82	50922.76
A_24_P21	NM_18174	Homo sapi	LINS1,WIN	1.78546	6.38E-09	1.78546	182.1313	325.118
A_23_P86	NM_02114	cyclin D bir	DMTF1,DM	1.78244	9.22E-25	1.78244	922.36	1647.676
A_23_P50	NM_00221	Homo sapi	ITGAV,CD	1.78177	1.02E-39	1.78177	2136.198	3806.511
A_24_P62	NM_00584	signal trans	STAM2,Hb	1.77918	0.00051	1.77918	236.4456	417.8056
A_23_P64	NM_00356	early endo	EEA1,ZFY	1.7783	0.00007	1.7783	171.6086	303.5908
A_24_P18	NM_00207	golgi autoa	GOLGA4,C	1.77742	3.13E-14	1.77742	3895.348	6980.218
A_23_P41	NM_00510	Homo sapi	OXS1,OS	1.77637	1.21E-10	1.77637	5326.098	9557.186
A_32_P23	NM_17587	Homo sapi	EXOC8,EX	1.77555	2.14E-14	1.77555	1867.385	3351.231
A_23_P77	AK128671	Hypothetic	LOC12441	1.77419	3.23E-17	1.77419	760.0913	1352.357
A_24_P20	NM_15238	hypothetic	FLJ37953,	1.77385	7.13E-07	1.77385	486.5712	870.8782
A_23_P27	NM_03214	hypothetic	DKFZP434	1.77368	1.40E-12	1.77368	2695.443	4823.922
A_24_P47	NM_01400	vinculin	VCL	1.77253	3.27E-19	1.77253	3300.373	5871.731
A_23_P42	NM_00442	early devel	PHC2,PH2	1.77246	1.13E-10	1.77246	7055.452	12536.21
A_24_P17	NM_03091	protein kin	CDADC1,M	1.7718	2.12E-15	1.7718	861.8076	1533.418
A_24_P24	NM_00188	activating t	ATF2,HB1	1.77168	3.80E-13	1.77168	542.892	964.0604
A_23_P50	AK056006	Hypothetic	LOC14820	1.7706	2.29E-13	1.7706	1682.589	2991.841
A_24_P19	NM_00713	Homo sapi	ZNF92,TF1	1.76985	8.53E-36	1.76985	2670.565	4728.868
A_23_P77	NM_20341	similar to	RLOC92162	1.76864	3.79E-06	1.76864	1671.85	2916.617
A_24_P30	BX161510	Trafficking	TRAPPC6	1.76852	3.38E-12	1.76852	478.0432	848.4684
A_24_P94	AF213987	ALL1 fusec	AF5Q31	1.76832	0.00081	1.76832	123.6858	218.4323
A_32_P57	453	Unknown	THC18074	1.76559	0.00004	1.76559	203.8923	359.4346
A_23_P11	NM_00548	Homo sapi	TOM1L1,S	1.76553	2.78E-16	1.76553	1803.171	3205.584
A_23_P74	NM_00595	Homo sapi	MTF1,MTF	1.76497	6.26E-09	1.76497	4233.274	7550.122
A_24_P46	NM_00599	serine/thre	STK10,LOI	1.76453	2.72E-13	1.76453	1251.713	2220.733
A_23_P13	NM_00216	Homo sapi	ID3,HEIR-1	1.76425	2.00E-28	1.76425	8334.961	14742.49
A_23_P34	NM_00525	Homo sapi	FOSL2,FR	1.76412	4.98E-10	1.76412	4267.281	7425.868
A_23_P21	NM_00466	Homo sapi	DDX3Y,DE	1.76002	4.22E-09	1.76002	5414.154	9713.695
A_23_P15	NM_00201	Homo sapi	FOXO1A,F	1.75851	2.90E-28	1.75851	1376.391	2423.969
A_24_P34	NM_00449	hepatocyte	FOXA1,HN	1.75848	1.01E-11	1.75848	4728.18	8445.172
A_23_P56	NM_02518	Homo sapi	ZNF430,FL	1.75825	7.53E-12	1.75825	2659.838	4706.277

A_32_P13	(BQ441611 Transcriber	BQ441611	1.75656	3.62E-08	1.75656	205.5165	360.7156
A_24_P94	NM_00620 Phosphodi	PDE4D	1.75574	1.01E-13	1.75574	1422.171	2506.237
A_32_P12	AF085848 Ubiquitin s	USP53	1.75536	0.00023	1.75536	2383.001	4149.402
A_23_P39	NM_01530 Homo sapi	FNBP4,FB	1.75407	1.14E-15	1.75407	2589.257	4578.834
A_24_P21	NM_01959 zinc finger	ZNF26,KO	1.75318	1.12E-08	1.75318	774.221	1363.566
A_23_P21	AL050270 Homo sapi	DKFZP564	1.75289	4.86E-11	1.75289	611.3372	1072.831
A_32_P30966	Unknown	THC19873	1.75224	1.17E-13	1.75224	552.4711	970.4838
A_23_P13	AB018326 PHD finger	PHF14	1.75222	6.40E-06	1.75222	144.6273	253.2583
A_23_P45	NM_00446 Homo sapi	FIGF,VEGI	1.75162	3.53E-12	1.75162	1184.954	2079.436
A_32_P51787	IAP-like pr	ENST0000	1.75078	0.00015	1.75078	172.0591	300.5972
A_24_P89	NM_19828 hypothetic	LOC34911	1.74989	6.69E-10	1.74989	3899.749	6833.415
A_24_P30	AK097947 Similar to j	KIAA1971	1.74981	1.30E-16	1.74981	1926.447	3393.509
A_24_P40	NM_05292 Homo sapi	LENG8,pp	1.74946	1.26E-12	1.74946	552.5604	969.9201
A_32_P148824	Unknown	A_32_P14	1.74928	2.52E-21	1.74928	3272.076	5749.473
A_23_P37	AB020721 Family with	FAM13A1	1.74911	6.59E-08	1.74911	2074.128	3616.902
A_23_P21	NM_00416 RAB5A, m	RAB5A,RA	1.74869	8.99E-38	1.74869	3011.969	5267.978
A_24_P23	(NM_02094 Mesoderm	MI-ER1	1.74783	1.08E-08	1.74783	1237.198	2195.304
A_32_P17	AK128192 Hypothetic	AK128192	1.74777	1.96E-09	1.74777	22130.82	38773.98
A_23_P20	NM_01400 vinculin	VCL	1.74639	1.86E-13	1.74639	1026.259	1796.791
A_32_P18	NM_17494 Homo sapi	GAS2L3	1.74617	0.00011	1.74617	199.8975	347.6718
A_23_P81	NM_00564 Homo sapi	TAF7,TAF	1.7459	6.33E-11	1.7459	20076.25	35526.11
A_24_P358868	Zinc finger	ENST0000	1.74536	0.00086	1.74536	241.5342	424.076
A_24_P27	BC036731 Zinc finger	ZBTB24	1.74529	0.00027	1.74529	313.4067	548.7536
A_24_P81	NM_00207 glutamic-o	GOT1	1.74499	1.17E-31	1.74499	3479.879	6078.82
A_32_P172453	Unknown	THC19832	1.74465	9.66E-08	1.74465	510.8695	892.1773
A_23_P43	NM_02095 Homo sapi	ZNF529,KI	1.74413	6.48E-10	1.74413	520.4691	909.984
A_23_P16	NM_00306 Homo sapi	SNAI2,SLL	1.74296	9.91E-10	1.74296	3064.782	5321.081
A_32_P22	AK022044 Ribosomal	RPL10A	1.74276	1.78E-20	1.74276	439.0743	765.243
A_24_P39	AK026328 Hypothetic	FLJ22675	1.7406	9.24E-25	1.7406	1612.685	2805.424
A_23_P40	NM_15240 Homo sapi	GRPEL2,F	1.74053	6.82E-07	1.74053	2559.44	4506.525
A_24_P12	NM_00713 Homo sapi	ZNF79,pT7	1.73999	1.09E-11	1.73999	744.5101	1294.927
A_24_P717305	Unknown	A_24_P71	1.73991	1.85E-06	1.73991	375.5013	655.3267
A_32_P193908	Unknown	A_32_P19	1.73764	0.00089	1.73764	1330.56	2322.665
A_24_P94	(AB051476 B double p	BDP1	1.73752	1.24E-13	1.73752	952.0839	1667.666
A_23_P87	NM_00239 solute carri	SLC3A2,4F	1.73618	1.16E-12	1.73618	25098.08	43763.66
A_23_P84016	Unknown	A_23_P84	1.73584	1.32E-13	1.73584	5632.362	9862.661
A_24_P33	NM_02492 hypothetic	DNAJB14,I	1.73495	7.46E-15	1.73495	1071.874	1874.489
A_32_P84	(AK098044 CDNA FLJ	AK098044	1.73464	5.55E-15	1.73464	465.4827	808.4301
A_32_P75	BG400725 60246411C	RNU2,U2	1.73463	1.68E-23	1.73463	1940.267	3366.514
A_23_P30	NM_02094 Mesoderm	MI-ER1	1.73408	7.49E-09	1.73408	825.3588	1445.38
A_24_P32	NM_00577 RNA bindir	RBM5,G15	1.73405	1.12E-13	1.73405	2445.382	4270.636
A_24_P34	NM_01765 hypothetic	ZNF562,FL	1.73291	8.91E-13	1.73291	1115.225	1948.073
A_23_P14	NM_01794 Homo sapi	USP47,TR	1.73103	1.16E-17	1.73103	1599.242	2783.631
A_24_P213296	Unknown	ENST0000	1.73001	0.00098	1.73001	162.9686	280.4783
A_24_P94	AB032974 Putative an	DKFZP564	1.72982	5.34E-14	1.72982	721.1738	1254.839
A_24_P20	NM_02519 Homo sapi	ITPKC	1.72973	2.42E-08	1.72973	862.5678	1496.333
A_23_P25	NM_01865 Homo sapi	ZNF167,ZF	1.72881	1.39E-11	1.72881	301.8397	521.9781
A_24_P14	U87836 Splicing fac	SFRS10	1.72805	3.84E-06	1.72805	3714.263	6588.02
A_32_P10	AK129982 Homo sapi	AK129982	1.72682	6.60E-07	1.72682	501.702	871.0173
A_23_P8812	Unknown	A_23_P88	1.72663	3.10E-15	1.72663	2182.411	3793.928
A_23_P42	NM_01765 aftiphilin pr	AFTIPHILII	1.72561	9.52E-10	1.72561	1433.377	2492.424
A_23_P90	NM_18291 TIR domair	TICAM1,TF	1.72512	4.06E-34	1.72512	3153.91	5442.619

A_23_P37	NM_00068	Homo sapi	ADRA2B,A	1.72476	0.00005	1.72476	213.1114	368.4887
A_24_P38	NM_03214	hypothetic	DKFZP434	1.72438	1.36E-26	1.72438	2454.995	4240.078
A_23_P25	NM_00644	Homo sapi	USP16,UB	1.72418	4.11E-21	1.72418	3592.402	6218.253
A_23_P78	NM_02472	Homo sapi	MYH14,DF	1.72405	7.10E-08	1.72405	46158.86	79864.33
A_23_P42	NM_00007	Homo sapi	CDKN1C,E	1.72403	2.47E-27	1.72403	914.7286	1576.136
A_24_P93	BX538183	Hypothetic	LOC90321	1.72351	0.00006	1.72351	268.5983	462.1099
A_24_P10	AY117438	Zinc finger	ZNF468	1.72274	7.05E-11	1.72274	313.7042	540.3729
A_23_P16	NM_00542	Homo sapi	VEGFC,VF	1.7216	7.57E-13	1.7216	3294.048	5695.317
A_23_P10	NM_00714	zinc finger	ZNF146,O	1.72084	4.01E-18	1.72084	926.4408	1588.324
A_24_P52	NM_00321	thymine-D	TDG	1.72048	1.01E-08	1.72048	6279.443	10993.48
A_24_P66	BC062355	Hypothetic	BC062355	1.71953	9.84E-12	1.71953	552.9211	952.6202
A_23_P97	NM_00356	Homo sapi	BCAR3,NS	1.71872	4.94E-15	1.71872	5405.049	9324.774
A_32_P224638	T47874	sc	THC19589	1.7164	5.14E-08	1.7164	725.4404	1257.549
A_32_P181271	Unknown		THC19491	1.71603	0.00023	1.71603	322.9449	554.7698
A_24_P94	AK129652	Sperm ass	SPAG9	1.71575	1.21E-13	1.71575	1053.256	1821.809
A_32_P184937	Unknown		A_32_P18	1.71494	5.40E-22	1.71494	5356.206	9198.663
A_23_P15	AB037770	Zinc finger	ZNF624	1.71478	0.001	1.71478	104.3381	179.0215
A_24_P39	NM_01442	ALL1 fusec	AFF4,MCE	1.71418	7.82E-10	1.71418	370.1154	634.8784
A_23_P10	NM_00342	Homo sapi	ZNF84,HP	1.71307	3.61E-11	1.71307	1439.695	2470.636
A_23_P55	NM_14590	Homo sapi	RIOK3,SUI	1.71266	4.97E-16	1.71266	535.4398	918.3915
A_23_P76	NM_01397	Homo sapi	BNIP1,NIP	1.71192	1.98E-06	1.71192	939.6653	1632.33
A_23_P26	NM_00547	Homo sapi	HDAC5,HC	1.71119	6.03E-08	1.71119	460.9882	790.1507
A_24_P823768	Unknown		A_24_P82	1.71114	4.81E-11	1.71114	2422.651	4141.07
A_24_P27	BC008810	Hypothetic	LOC19626	1.71083	4.41E-07	1.71083	621.4379	1067.224
A_24_P24	NM_00261	pyruvate d	PK4	1.71063	7.65E-12	1.71063	4654.343	7942.772
A_24_P11	AB075851	Similar to j	KIAA1971	1.7095	0.00002	1.7095	1726.247	3030.144
A_24_P79	AK021552	Hypothetic	LOC39995	1.70783	4.65E-08	1.70783	261.7116	447.079
A_23_P16	NM_00486	Homo sapi	TBPL1,TLF	1.7075	3.00E-10	1.7075	3351.179	5759.356
A_24_P84	NM_17234	Homo sapi	SPAG9,HS	1.70674	1.06E-07	1.70674	644.4305	1093.783
A_23_P12	NM_00587	Homo sapi	SMNDC1,S	1.70578	8.80E-14	1.70578	6028.682	10367.4
A_24_P27	NM_01487	KIAA0063	KIAA0063,	1.7045	6.95E-11	1.7045	3692.043	6310.597
A_23_P41	NM_01763	hypothetic	FLJ20035,	1.70436	1.62E-12	1.70436	736.0179	1257.287
A_32_P14	BC051298	Similar to A	FLJ00312	1.70393	0.00067	1.70393	186.601	319.7176
A_23_P15	NM_03080	Homo sapi	C12orf22,C	1.7029	5.87E-10	1.7029	1061.081	1815.3
A_23_P20	NM_00321	Homo sapi	TDG	1.70216	9.41E-30	1.70216	14707.37	25066.15
A_24_P252739	U51869	Bc	THC19236	1.70123	7.18E-07	1.70123	235.2341	400.3562
A_24_P24	AL833358	DnaJ (Hsp	DNAJB12	1.69989	1.65E-14	1.69989	1397.659	2388.711
A_32_P22	NM_20750	FLJ45248	FLJ45248	1.69739	1.24E-09	1.69739	504.1508	857.882
A_23_P39	NM_01843	putative zir	JMJD1A,T	1.69675	5.18E-07	1.69675	369.2345	627.5544
A_23_P40	NM_02494	Homo sapi	PHC3,EDF	1.69583	0.00012	1.69583	465.5416	796.9346
A_23_P87	NM_00481	Homo sapi	LPXN,LDP	1.69551	0.00007	1.69551	286.8344	487.6079
A_24_P36	NM_00239	mouse dou	MDM4,MD	1.69539	1.57E-07	1.69539	395.6409	672.0574
A_32_P129621	AD17_HU		THC18575	1.69523	1.26E-17	1.69523	4061.588	6865.813
A_23_P20	NM_20126	Homo sapi	ZNF644,Ze	1.69501	2.32E-06	1.69501	1171.222	2025.687
A_23_P11	NM_01553	Homo sapi	ZZZ3,DKF	1.69477	0	1.69477	909.9276	1540.325
A_24_P92	AK096270	Ras associ	RAPH1	1.69378	0.00071	1.69378	201.1603	338.6139
A_24_P13	NM_00694	stress 70 p	STCH	1.69313	1.00E-29	1.69313	1871.636	3167.3
A_23_P108394	BC041692	THC18680		1.6928	9.02E-15	1.6928	4384.156	7486.565
A_32_P46	BC018064	Activating t	ATF6	1.69161	6.78E-08	1.69161	415.4565	702.7101
A_23_P20	NM_00270	Homo sapi	PPM1B,PF	1.69144	3.40E-09	1.69144	2754.531	4655.88
A_32_P21	NM_00321	thymine-D	TDG	1.69135	4.40E-31	1.69135	10936.96	18484.75
A_24_P12	NM_01744	chromoson	C21orf91,Y	1.69133	0.00002	1.69133	434.957	739.9349

A_23_P30	NM_00328	Homo sapi	TOP1, TOP	1.69092	7.65E-16	1.69092	2965.592	5040.429
A_23_P79	NM_01788	hypothetic	FLJ20558	1.69	0.00002	1.69	362.895	614.9724
A_23_P32	NM_01441	nasophary	NAG8, NSC	1.68926	0.00001	1.68926	393.6333	667.3661
A_23_P75	NM_01766	Homo sapi	ZCCHC10,	1.68755	6.93E-20	1.68755	2310.438	3907.487
A_23_P32	NM_01468	Homo sapi	KIAA0355	1.68717	5.20E-08	1.68717	726.6403	1238.558
A_24_P24	NM_00445	fatty-acid-C	ACSL3, AC	1.68666	1.63E-18	1.68666	1973.56	3344.971
A_23_P17	NM_02209	Homo sapi	ZNF335, NI	1.68642	1.38E-14	1.68642	496.3488	837.928
A_24_P37	NR_00128	Homo sapi	TOP1P2	1.68623	5.10E-11	1.68623	4134.879	7001.507
A_23_P385427		DNA-bindir	NP111954	1.68572	7.53E-15	1.68572	929.0831	1570.042
A_24_P94	BC015929	Nuclear rec	NR1D2	1.68556	6.92E-14	1.68556	573.9099	965.3539
A_23_P82	BC011254	Glucocortic	GLCCI1	1.68524	1.33E-15	1.68524	4000.328	6779.749
A_23_P27	Y10936	F-box prote	FBXO46	1.68306	1.23E-11	1.68306	1513.406	2559.698
A_32_P10	AK096221	Zinc finger	ZNF626	1.683	0.00015	1.683	283.191	475.29
A_23_P31	NM_02114	Homo sapi	ZNF273, H	1.68291	0.00001	1.68291	330.2219	556.5512
A_23_P16	NM_00253	Homo sapi	ODC1	1.68289	9.64E-13	1.68289	39404.43	65986.93
A_24_P16	XM_37295	similar to P	LOC39144	1.68243	0.00036	1.68243	259.2105	436.8942
A_23_P42	NM_17362	Hypothetic	FLJ36674	1.68207	1.79E-06	1.68207	314.283	528.9953
A_24_P21	AB062434	Procollage	P4HB	1.68128	1.04E-21	1.68128	48842.2	82309.48
A_23_P58	NM_15240	hypothetic	FLJ39370	1.68117	1.76E-11	1.68117	1841.929	3077.368
A_23_P19	NM_17504	Homo sapi	PILRB, FDF	1.68104	2.26E-34	1.68104	8382.907	14091.58
A_32_P49116		Unknown	A_32_P49	1.68021	3.14E-09	1.68021	384.3153	645.6629
A_24_P14	NM_17071	Homo sapi	RASSF1, 1:	1.67995	1.80E-06	1.67995	190.9984	320.8439
A_23_P10	NM_00091	Procollage	P4HB	1.67942	4.77E-15	1.67942	8798.11	14731.3
A_24_P93	BC038247	LOC44044	TRIM25, EF	1.67926	6.52E-13	1.67926	1295.746	2179.807
A_24_P35	NM_05286	TRAF-inter	TIFA, T2BP	1.67856	5.56E-13	1.67856	2001.233	3376.765
A_23_P25	NM_18264	Homo sapi	DLC1, HP, /	1.67836	1.39E-09	1.67836	1911.053	3201.627
A_23_P62	NM_01233	nasophary	CCDC19, N	1.67829	0.0003	1.67829	256.4052	432.5325
A_24_P281801		Zinc finger	ZNF27	1.67742	1.21E-09	1.67742	1234.57	2091.905
A_23_P14	NM_00573	ADP-ribosy	ARL4, ARL	1.67666	1.41E-09	1.67666	1465.597	2488.477
A_32_P11	BC063309	Golgi auto	LOC39053	1.67666	0.00037	1.67666	515.7032	877.7905
A_23_P21	NM_00321	Homo sapi	TEC, PSCT	1.67631	0.00067	1.67631	132.01	221.6278
A_23_P28743		Unknown	A_23_P28	1.67601	2.47E-09	1.67601	3256.607	5488.159
A_32_P85	NM_17361	hypothetic	FLJ35785	1.67569	0.00017	1.67569	494.4761	839.3469
A_23_P84	NM_01464	Homo sapi	IQCB1, NPI	1.67544	1.85E-08	1.67544	3013.397	5102.708
A_24_P28	NM_18404	Homo sapi	VPS13B, C	1.67493	0.00066	1.67493	318.7897	534.9384
A_23_P32	NM_13916	Homo sapi	STARD4	1.67136	1.87E-10	1.67136	3983.679	6669.585
A_23_P16	NM_01840	Decapping	DCP1A	1.67108	1.45E-21	1.67108	8508.577	14250.75
A_23_P37	NM_00308	small nucle	SNAPC1, S	1.67056	2.83E-07	1.67056	1566.732	2676.051
A_23_P36	AF332224	Chromosor	CYorf15A	1.66865	1.05E-15	1.66865	427.1038	712.8389
A_23_P52	NM_15264	Homo sapi	FAM24B, M	1.66792	2.66E-07	1.66792	1530.497	2556.271
A_24_P23	NM_20126	Homo sapi	ZNF644, Ze	1.66781	8.68E-11	1.66781	789.328	1323.134
A_23_P12	NM_01830	hypothetic	FLJ11046, l	1.6675	8.92E-07	1.6675	672.2728	1137.224
A_23_P79	NM_02513	Homo sapi	FBXO11, V	1.66619	1.62E-16	1.66619	4635.016	7745.703
A_23_P12	NM_02301	Homo sapi	C2orf26	1.66605	1.46E-17	1.66605	1672.063	2800.466
A_24_P33	NM_02097	Homo sapi	KIAA1641,	1.66595	0.00045	1.66595	218.978	362.9121
A_32_P39	NM_19845	Homo sapi	ZNF600, KF	1.66554	3.94E-09	1.66554	529.9437	885.3557
A_23_P60189		Unknown	A_23_P60	1.66532	5.87E-07	1.66532	1453.239	2412.309
A_23_P41	NM_02451	Homo sapi	C4orf15, IT	1.66509	1.22E-08	1.66509	1097.051	1851.351
A_23_P58	NM_02461	TPA-induct	TTMP	1.66489	0.00071	1.66489	525.5906	879.0441
A_32_P63562		U16359 nit	THC18591	1.66451	1.44E-27	1.66451	1971.02	3278.144
A_23_P58	AK127224	Nedd4 binc	N4BP3	1.66427	8.55E-20	1.66427	466.3551	776.1871
A_23_P43	NM_00166	Homo sapi	ARF4	1.66372	4.00E-10	1.66372	15973.94	26835.47

A_32_P14	AK092349	CDNA FLJ:AK092349	1.66323	0.00039	1.66323	129.9657	216.382
A_24_P40	NM_02392	hypothetical C1QDC1, E	1.6629	8.65E-08	1.6629	234.7507	390.225
A_23_P32	NM_02283	Homo sapi PLEKHG2,	1.66235	5.23E-14	1.66235	757.2549	1261.846
A_23_P50	NM_00657	UDP-GlcN/ B3GNT1, B	1.66185	0.00002	1.66185	372.502	620.7069
A_32_P44	NM_18250	Homo sapi C9orf85, M	1.66174	0.00001	1.66174	718.1605	1214.138
A_23_P12	NM_15302	Homo sapi ZNF75A, FL	1.65994	0.00002	1.65994	164.4079	272.688
A_23_P21	NM_03145	Homo sapi FBXW10, H	1.65987	0.00028	1.65987	227.8372	377.4163
A_23_P25	AB014563	KIAA0663 KIAA0663	1.65976	4.85E-07	1.65976	11790.57	19870.28
A_23_P16	NM_02469	hypothetical ZNF419, FL	1.65971	2.66E-17	1.65971	423.1213	702.2399
A_23_P21	NM_03095	Homo sapi TIGD6, DK	1.65944	1.04E-12	1.65944	950.3563	1578.48
A_23_P16	NM_00699	Homo sapi SLC19A2, T	1.65862	1.82E-25	1.65862	4550.167	7560.517
A_23_P16	NM_00460	syntaxin 1/ STX1A, ST	1.65854	4.54E-12	1.65854	2551.725	4240.398
A_24_P339869	Z295_HUM	THC18809	1.65839	1.01E-06	1.65839	901.2885	1501.679
A_23_P82	NM_00460	Homo sapi STX1A, ST	1.65784	7.00E-18	1.65784	2783.566	4616.209
A_23_P16	AK128435	Cofactor re CRSP7	1.65732	1.68E-14	1.65732	1694.535	2817.89
A_23_P22	NM_00226	Homo sapi KLRC2, NK	1.65627	9.43E-06	1.65627	192.0083	318.4365
A_23_P34	NM_17498	Homo sapi C19orf28, F	1.65588	1.12E-26	1.65588	2742.198	4540.148
A_23_P58	AB014563	KIAA0663 KIAA0663	1.65568	4.82E-08	1.65568	10026.91	16901.54
A_23_P37	NM_13828	Homo sapi ZNF681, FL	1.65524	7.80E-15	1.65524	2039.167	3391.325
A_32_P90	BC045657	MRNA; cD1C6orf204	1.65521	6.44E-10	1.65521	672.4041	1117.846
A_23_P15	AK054969	Ubiquitin p1 UBR1	1.65517	5.32E-12	1.65517	2762.991	4610.09
A_23_P11	NM_00633	Homo sapi HMG20B, S	1.65515	9.55E-30	1.65515	1547.48	2561.465
A_24_P19	NM_00596	NGFI-A bin NAB1	1.65382	6.30E-08	1.65382	1249.262	2082.044
A_23_P34	NM_17506	Homo sapi HIST2H2A	1.65365	3.31E-12	1.65365	382.8932	633.1651
A_24_P85	NM_03222	Homo sapi ZCCHC7, F	1.65362	0.00018	1.65362	728.4677	1226.837
A_24_P38	NM_00318	TATA box ITAF2, TAF	1.65289	2.31E-09	1.65289	832.9146	1389.175
A_23_P12	NM_01959	Homo sapi ZNF26, KO	1.65276	3.67E-22	1.65276	905.6368	1495.245
A_23_P30	AL833066	KIAA1040 KIAA1040	1.65269	1.52E-18	1.65269	909.8781	1505.077
A_23_P14	NM_01757	G protein-c MKNK2, M	1.65182	6.65E-22	1.65182	35863.74	59410.45
A_23_P35	AB046766	KIAA1546 KIAA1546	1.65156	2.91E-09	1.65156	313.8333	518.5712
A_32_P22	NM_00351	H2A histon HIST1H2A	1.65117	3.91E-08	1.65117	640.1273	1062.667
A_23_P58	AK001520	Serine PI K SPINK5L3	1.65102	6.62E-21	1.65102	1347.1	2228.303
A_23_P33	NM_13833	Homo sapi ZNF675, T	1.65076	4.24E-22	1.65076	1548.646	2551.026
A_23_P30	AF158555	Glutaminase GLS	1.65056	1.83E-08	1.65056	2498.903	4172.827
A_23_P31	NM_01519	Homo sapi COBL, KIA	1.64986	3.27E-08	1.64986	1094.837	1818.849
A_23_P11	NM_00337	Homo sapi VASP	1.64955	0.00005	1.64955	2798.005	4743.085
A_23_P25	NM_01214	Homo sapi DDX26, HD	1.64949	4.03E-10	1.64949	2778.53	4622.38
A_23_P46	NM_00735	Homo sapi MTF2, M96	1.64916	2.66E-07	1.64916	1127.28	1885.275
A_32_P14	BC015064	Transducer TOB1	1.64869	3.05E-17	1.64869	1622.19	2676.111
A_32_P21	AK023660	CDNA FLJ AK023660	1.64795	1.65E-10	1.64795	1283.385	2129.063
A_24_P48	NM_01774	hypothetical FLJ20291, I	1.64723	8.80E-07	1.64723	485.6064	800.6882
A_23_P21	NM_01848	Homo sapi DDEF1, PA	1.64684	0.00005	1.64684	498.8839	827.6026
A_32_P14	XM_29283	Homo sapi ZNF66, FLJ	1.64569	7.24E-15	1.64569	1675.409	2766.998
A_23_P21	NM_00505	Homo sapi RANBP2L1	1.64478	2.44E-06	1.64478	1610.818	2678.557
A_24_P24	AL833358	DnaJ (Hsp) DNAJB12	1.64264	2.27E-06	1.64264	589.6793	976.8629
A_24_P204290	Unknown	ENST0000	1.64209	0.00001	1.64209	278.7472	457.2694
A_32_P177097	Unknown	THC19530	1.64168	1.54E-10	1.64168	1619.494	2660.04
A_23_P61	NM_18254	Homo sapi NSUN6, NC	1.64139	1.09E-06	1.64139	1001.409	1662.927
A_23_P20	NM_00423	Homo sapi SOCS6, CI	1.6407	6.78E-06	1.6407	195.6749	320.6588
A_23_P136849	Unknown	THC19937	1.64008	3.42E-07	1.64008	215.2212	352.8487
A_23_P37	NM_00449	Homo sapi FOXA1, HN	1.6398	7.15E-18	1.6398	2231.367	3689.447
A_23_P37	NM_20745	FLJ35767 FLJ35767	1.63965	1.04E-09	1.63965	749.7626	1232.275

A_23_P14	(NM_14457 Homo sapi	C14orf32,M	1.63964	5.61E-10	1.63964	8194.49	13609.81
A_23_P94	(NM_05823 Homo sapi	ZNF354B,F	1.63951	0.00002	1.63951	316.3284	518.5392
A_23_P13	(NM_01442 Homo sapi	SNX5,FLJ1	1.63943	9.98E-29	1.63943	32961.55	54004.06
A_23_P40	(NM_15222 Homo sapi	CSNK1E,H	1.6378	4.05E-21	1.6378	14922.92	24487.64
A_23_P15	(NM_13350 Homo sapi	ZNF274,ZF	1.63753	3.01E-13	1.63753	1015.896	1671.221
A_24_P10	(AK092646 KIAA1040	KIAA1040	1.6373	8.36E-10	1.6373	327.8413	536.7004
A_32_P43	(AK097687 Zinc finger	ZNF21	1.63726	1.75E-22	1.63726	1037.922	1698.746
A_24_P25	(NM_02172 nuclear rec	NR1D1,EA	1.63721	2.46E-08	1.63721	1300.856	2131.168
A_24_P62	(NM_00335 urocortin	UCN,UI,UF	1.63701	1.44E-08	1.63701	345.5443	565.8815
A_24_P18	(NM_00553 inositol(my	IMPA1,IMF	1.63692	2.64E-08	1.63692	3718.306	6157.16
A_24_P10	(NM_17491 Homo sapi	UBR1	1.63676	0.00009	1.63676	673.1116	1123.223
A_24_P30	(NM_00128 chloride ch	CLCN6,CL	1.63673	1.97E-10	1.63673	815.9628	1342.924
A_24_P63	(AK023559 Excision re	PGBD3	1.63591	4.08E-12	1.63591	952.9418	1561.205
A_32_P54	(AK126842 Similar to	ZLOC16323	1.6359	1.07E-10	1.6359	1397.789	2296.857
A_24_P17	(NM_05824 Homo sapi	CCNT2	1.63573	6.68E-07	1.63573	475.2073	779.4268
A_23_P21	(NM_01225 Homo sapi	HBP1	1.63458	1.13E-08	1.63458	533.5714	874.0783
A_23_P30	(NM_00060 fibroblast g	FGFR1,H2	1.6345	1.55E-12	1.6345	648.1448	1062.636
A_23_P35	(NM_01647 Homo sapi	C3orf19,FL	1.63445	1.47E-09	1.63445	456.8238	747.0355
A_23_P35	(AK090500 M4-50 mR	IAK090500	1.63431	1.59E-13	1.63431	24112.27	39591.34
A_23_P13	(NM_00139 Homo sapi	DUSP4,TY	1.63419	2.22E-11	1.63419	7974.929	13065.12
A_24_P54	(NM_15240 hypothetic	FLJ39370	1.63329	2.32E-14	1.63329	3647.944	5993.233
A_23_P11	(NM_00289 Homo sapi	ARID4A,R	1.63321	0.00069	1.63321	238.9914	391.2237
A_23_P11644	Unknown A_23_P11		1.63303	0.00003	1.63303	625.186	1017.209
A_24_P94	(AB007928 KIAA0459	KIAA0459	1.63298	1.10E-07	1.63298	447.6372	732.2957
A_23_P95	(NM_01765 hypothetic	ZNF562,FL	1.6319	1.66E-08	1.6319	1413.136	2325.224
A_32_P22	(NM_01900 protein ass	ZA20D3,A	1.63151	3.63E-27	1.63151	8844.668	14442.49
A_24_P38	(AK057241 Hypothetic	FLJ32679	1.63066	0.00018	1.63066	373.5049	610.857
A_23_P35	(NM_14580 Homo sapi	ABTB2,DK	1.62957	1.03E-10	1.62957	2522.403	4102.444
A_24_P34	(U79263 RAP1 inter	RIF1	1.62843	3.08E-08	1.62843	362.5637	590.6277
A_24_P16	(NM_00487 golgi SNAF	GOSR1,P2	1.62825	9.04E-08	1.62825	1573.365	2597.945
A_23_P19	(NM_00190 Homo sapi	CTGF,CCN	1.62703	1.82E-08	1.62703	266.9627	434.1664
A_23_P21	(NM_00565 Homo sapi	TRAF1,EB	1.62595	3.09E-07	1.62595	2066.301	3326.638
A_23_P79	(NM_01647 Homo sapi	C20orf111,	1.62586	7.66E-26	1.62586	4709.405	7664.687
A_24_P56	(AF086032 Protein phc	PPP1R12A	1.62555	2.68E-18	1.62555	2836.895	4629.539
A_23_P41	(NM_00219 Homo sapi	IRF1,MAR,	1.6255	3.71E-14	1.6255	2069.404	3349.068
A_23_P51	(NM_01828 Homo sapi	GBP3,FLJ	1.62527	0.00004	1.62527	375.2741	611.1949
A_23_P43	(BC022822 Immunoglo	IGLC2	1.62448	8.73E-30	1.62448	3039.743	4938.118
A_23_P62	(NM_00325 tissue inhib	TIMP1,EP	1.62432	2.48E-12	1.62432	16345.6	26759.21
A_23_P35	(NM_14889 Homo sapi	FAM44A,F	1.62426	0.00005	1.62426	431.0911	703.9534
A_32_P80	(NM_00573 ADP-ribosy	ARL4,ARL	1.62333	6.79E-22	1.62333	1089.837	1771.117
A_32_P41	(AF086011 Potassium	KCNQ1OT	1.62259	0.00015	1.62259	259.7091	421.4757
A_23_P152570	Unknown THC18494		1.62241	8.01E-07	1.62241	1381.217	2278.246
A_23_P20	(NM_00737 Homo sapi	SHOC2,SC	1.62239	6.08E-29	1.62239	1842.086	2988.397
A_32_P118686	AF273048 THC18261		1.62158	1.55E-17	1.62158	2951.188	4774.842
A_24_P10	(NM_01843 putative zir	JMJD1A,T	1.62103	3.45E-06	1.62103	666.7838	1085.631
A_23_P41	(NM_17071 Homo sapi	RASSF1,1	1.62085	6.24E-12	1.62085	3560.169	5766.969
A_32_P113472	Unknown A_32_P11		1.6208	0.00002	1.6208	378.4439	614.6043
A_23_P38	(NM_00360 Homo sapi	OGT,HRN	1.61971	1.85E-16	1.61971	4798.238	7800.507
A_23_P67	(NM_00342 Homo sapi	ZNF85,HP	1.61934	1.18E-10	1.61934	1528.003	2487.295
A_24_P60	(NM_13924 Homo sapi	C9orf97,RF	1.61912	2.13E-08	1.61912	683.4592	1112.942
A_24_P12	(NM_01765 aftiphilin pr	AFTIPHILII	1.61901	2.33E-11	1.61901	1484.579	2418.214
A_24_P98948	ALU8_HUM THC18915		1.61858	7.45E-06	1.61858	1168.403	1902.953

A_23_P12: NM_00136	Homo sapi	DLG2,PSD	1.61828	0.00012	1.61828	498.5942	811.217
A_24_P17: BC013319	FYVE, Rho	FGD6	1.61793	1.01E-07	1.61793	262.8262	425.0508
A_24_P29: M28207	Major histo	HLA-C	1.61752	3.82E-20	1.61752	41819.84	67487.17
A_32_P124580	Unknown	A_32_P12:	1.61702	6.10E-07	1.61702	1943.281	3190.858
A_23_P33: NM_00353	Homo sapi	HIST1H3H	1.61664	6.96E-07	1.61664	1625.078	2665.877
A_23_P74: L12686	Phosphodi	PDE4B	1.61603	6.42E-12	1.61603	2569.03	4169.06
A_24_P48: NM_01524	cylindroma	CYLD,EAC	1.616	2.59E-11	1.616	500.2516	808.7652
A_23_P69: NM_00396	Homo sapi	CCRL2,HC	1.61476	0.00039	1.61476	406.6606	659.7512
A_23_P26: NM_03080	LIS1-intera	NDEL1,EO	1.61443	0.00003	1.61443	558.7118	901.863
A_32_P213543	Unknown	A_32_P21:	1.61436	1.69E-18	1.61436	1350.754	2179.148
A_23_P20: AL713775	Zinc finger	ZNF507	1.61416	0.00005	1.61416	380.9075	616.1588
A_23_P11: NM_01632	cyclin-E bir	HERC5,CE	1.61377	0.00081	1.61377	198.4011	318.3215
A_32_P141338	Unknown	A_32_P14:	1.61311	1.98E-09	1.61311	2378.388	3836.783
A_23_P17: NM_02430	hypothetic	CHCHD7,M	1.61281	1.05E-07	1.61281	1550.959	2516.713
A_23_P31: NM_05291	Establishm	ESCO1	1.6117	0.00002	1.6117	338.5455	546.2139
A_24_P94: AK092652	FLJ16231	FLJ16231	1.60964	0.00001	1.60964	293.5156	472.4232
A_24_P29: NM_01909	GTP bindin	GTPBP2,M	1.6096	0.00001	1.6096	189.7669	305.3897
A_23_P69: NM_00699	Homo sapi	ZNF197,P1	1.60905	0.00005	1.60905	210.5127	338.444
A_23_P90: NM_15265	Homo sapi	ZNF585A,F	1.609	5.76E-07	1.609	592.6815	959.0901
A_23_P38: NM_01509	Homo sapi	HIC2,HRG	1.60873	1.29E-10	1.60873	1637.808	2639.277
A_23_P14: NM_00382	Homo sapi	NAPG,GAM	1.60843	1.01E-14	1.60843	864.9821	1395.075
A_23_P77: NM_00703	protein tyrc	PTPN21,P	1.60797	2.18E-08	1.60797	258.8456	416.0739
A_23_P25: NM_01630	Homo sapi	CRBN,MR	1.60765	2.44E-10	1.60765	969.0012	1564.22
A_24_P34: NM_00313	Serum res	SRF	1.6072	1.96E-23	1.6072	7359.664	11833.43
A_24_P81: BC041882	Activating t	ATF7IP2	1.60689	0.00001	1.60689	515.509	832.7395
A_24_P21: NM_00353	H3 histone	HIST1H3D	1.60569	5.13E-21	1.60569	18574.85	29878.54
A_24_P18: NM_02040	protocadhe	PCDH9	1.60391	1.82E-14	1.60391	822.4121	1322.991
A_23_P21: NM_00139	Homo sapi	DYRK1A,M	1.60362	5.98E-17	1.60362	1267.073	2036.883
A_24_P81: AK097080	Hypothetic	AK097080	1.60328	2.35E-09	1.60328	1054.517	1698.231
A_23_P55: NM_00734	zinc finger	ZNF236,ZI	1.60236	1.03E-07	1.60236	350.1142	561.05
A_23_P20: NM_02276	hypothetic	ISG20L1,A	1.60205	9.49E-13	1.60205	968.8658	1556.595
A_24_P36: NM_00248	protein phc	PPP1R12A	1.60185	6.72E-09	1.60185	1842.591	2990.798
A_32_P57: NM_00631	ubiquitin s	USP15,UN	1.60103	9.73E-06	1.60103	2978.476	4881.851
A_23_P14: NM_03320	Homo sapi	ZNF101,H	1.60027	6.75E-11	1.60027	1902.951	3025.051
A_24_P94: AL832717	Multiple C2	MCTP2	1.60018	0.00009	1.60018	859.1051	1361.542
A_24_P91: BX647673	Golgin-67	GOLGIN-6	1.60001	0.00065	1.60001	424.595	682.1457
A_24_P20: NM_00416	RAB5A, m	RAB5A,RA	1.5999	0.00014	1.5999	360.1248	577.196
A_23_P15: NM_00345	Homo sapi	ZNF200,M	1.5987	6.70E-07	1.5987	1572.584	2539.819
A_24_P19: NM_13916	Homo sapi	SFRS12,S	1.59805	1.67E-10	1.59805	2145.005	3449.908
A_24_P6903	Unknown	ENST0000	1.59687	2.56E-13	1.59687	9797.256	15621.24
A_24_P18: NM_01656	apoptosis r	BFAR,BAR	1.59663	1.76E-08	1.59663	539.9322	865.6471
A_23_P33: AK093655	Glucocortic	GLCCI1	1.5966	5.63E-08	1.5966	1049.552	1696.314
A_24_P30: NM_01621	polymeras	POLK,DINI	1.59611	4.50E-12	1.59611	497.6313	794.9586
A_23_P25: NM_00055	Homo sapi	WRN,REC	1.59553	1.98E-12	1.59553	1612.294	2576.482
A_24_P20: NM_01557	CAZ-assoc	CAST1,CA	1.59531	1.60E-24	1.59531	1937.749	3090.049
A_23_P64: NM_01761	Homo sapi	ZCCHC8,C	1.59505	6.98E-16	1.59505	5986.411	9577.17
A_23_P20: NM_02513	hypothetic	HKDC1,FL	1.59497	4.03E-22	1.59497	5444.665	8697.603
A_23_P11: NM_00460	Homo sapi	TAF1,OF,E	1.59475	0.0009	1.59475	163.8096	261.9023
A_23_P10: NM_03081	hypothetic	MGC10471	1.59371	5.99E-18	1.59371	7756.495	12347.97
A_32_P12: NM_17585	Homo sapi	NM_17585	1.59245	0.0004	1.59245	541.5975	868.2524
A_24_P792130	AF071172	THC19993	1.5924	0.00026	1.5924	703.1321	1142.487
A_24_P53: AK026896	Serine/thre	STK24	1.59178	6.64E-06	1.59178	740.217	1178.875

A_24_P942798	AB093299	THC19806	1.59154	0.00008	1.59154	749.6983	1194.045
A_24_P91	AV647560	Transcriber AV647560	1.59144	5.12E-12	1.59144	6363.753	10139.41
A_24_P34	NM_02473	hypothetic ZNF665,ZF	1.59133	0.0002	1.59133	503.1153	804.4219
A_24_P621434	HS7C_HUI	THC18728	1.5907	8.13E-19	1.5907	32389.37	51374.42
A_23_P13	NM_00075	Homo sapi CSF2,GMC	1.59006	0.00029	1.59006	154.6795	245.8103
A_23_P15	NM_00479	Homo sapi PPIG,CYP	1.58925	8.74E-11	1.58925	5398.602	8604.918
A_23_P58	NM_01617	hypothetic LOC51149	1.58745	6.34E-16	1.58745	1270.292	2014.603
A_24_P51	NM_08092	Homo sapi DCBLD2,E	1.58686	0.00004	1.58686	541.2372	865.2576
A_24_P16	BC007197	Nuclear fac NFKBIB	1.5866	6.51E-15	1.5866	785.2994	1244.404
A_23_P42	BC063594	Zinc finger ZNF292	1.58643	5.07E-06	1.58643	639.502	1024.988
A_23_P32	NM_00579	Homo sapi DHRS2,HE	1.58635	2.52E-15	1.58635	857.9195	1360.259
A_32_P21	AK092942	Hypothetic AK092942	1.58602	6.28E-08	1.58602	598.5685	951.6091
A_24_P19	AB014563	KIAA0663 KIAA0663	1.58596	5.54E-21	1.58596	5806.326	9227.462
A_23_P31	NM_00720	Homo sapi AKAP10,PI	1.58545	3.59E-11	1.58545	492.3586	781.2473
A_23_P39	NM_17379	Homo sapi PAPD4,FL	1.5852	5.87E-09	1.5852	1231.787	1971.206
A_32_P19	AB082530	Likely orthc AVO3	1.58491	4.16E-12	1.58491	2098.056	3342.029
A_24_P14	NM_00352	H2B histon HIST2H2B	1.58448	2.23E-13	1.58448	963.5472	1535.378
A_24_P58	XM_06490	Similar to z ENST0000	1.58445	0.00008	1.58445	263.7651	417.6606
A_23_P25	NM_02276	Homo sapi FNDC3B,F	1.5842	5.05E-10	1.5842	3089.294	4933.226
A_23_P21	NM_03292	Homo sapi ZNF3,A8-5	1.5837	1.72E-12	1.5837	752.8689	1194.405
A_23_P12	NM_01771	Homo sapi C20orf13,F	1.58315	0.00014	1.58315	1138.653	1814.698
A_23_P15	NM_01831	chromoson SAPS3,SA	1.58268	2.22E-11	1.58268	842.9302	1339.223
A_23_P43	NM_19828	hypothetic LOC34911	1.58266	0.00004	1.58266	767.4386	1213.346
A_23_P25	BC000859	Zuotin relat ZRF1	1.58258	1.98E-20	1.58258	12203.7	19284.09
A_23_P10	NM_01416	HSPC128 HSPC128,I	1.58206	2.00E-10	1.58206	13881.79	22074.97
A_24_P91	NM_20347	selenoprot SELS,VIMI	1.58189	2.39E-06	1.58189	395.9759	627.2282
A_24_P15	NM_17353	Homo sapi ZNF100	1.58188	4.15E-20	1.58188	2511.031	3975.192
A_23_P25	NM_02037	Homo sapi C12orf4	1.58152	4.08E-07	1.58152	702.2606	1117.025
A_23_P10	NM_00663	Homo sapi ZNF266,H	1.58133	5.42E-14	1.58133	459.5803	726.8455
A_24_P27	NM_02470	Smad nucl SNIP1,FLJ	1.58119	5.98E-10	1.58119	370.0154	585.1024
A_23_P44	NM_00359	Homo sapi KLF11,FKL	1.58115	9.95E-09	1.58115	674.0685	1071.453
A_23_P38	NM_00427	phosphatid PIGL	1.58076	4.31E-06	1.58076	370.2	585.6681
A_23_P25	NM_00715	Homo sapi ZNF195,DI	1.57977	2.72E-14	1.57977	2055.074	3248.192
A_23_P95	NM_17855	Homo sapi ZNF680,FL	1.57942	2.45E-18	1.57942	2305.653	3646.003
A_23_P14	NM_08059	Homo sapi HIST1H2B	1.57934	4.05E-16	1.57934	13926.76	22037.45
A_23_P36	NM_00626	Homo sapi DNAJC3,P	1.57789	0.00031	1.57789	2088.595	3392.607
A_24_P38	D26067	Transmem TMEM41B	1.57667	5.06E-08	1.57667	2067.29	3278.036
A_23_P58	AK056503	KIAA1387 KIAA1387	1.57653	8.03E-08	1.57653	389.4158	613.8959
A_23_P48	NM_18290	Homo sapi DHRS2,HE	1.57648	8.06E-10	1.57648	1007.607	1586.213
A_23_P16	NM_03081	Homo sapi PDRG1,PC	1.5764	8.10E-09	1.5764	4505.133	7114.182
A_23_P348349	IAP-like prc	ENST0000	1.57606	8.79E-06	1.57606	473.9778	749.1797
A_32_P107441	Unknown	A_32_P10	1.57588	6.67E-08	1.57588	243.9202	384.4241
A_24_P33	NM_01664	HDCMA18 HDCMA18	1.57564	4.91E-14	1.57564	3657.869	5791.632
A_24_P56	AK093869	Homo sapi RNF3	1.57499	0.00007	1.57499	188.0251	296.3982
A_23_P74	NM_02042	Homo sapi SCYL3,PA	1.5746	3.34E-08	1.5746	970.3315	1538.734
A_24_P34	NM_01560	DKFZP564 HERC4,KI	1.57443	1.83E-12	1.57443	3457.474	5476.173
A_32_P31785	Unknown	A_32_P31	1.57351	4.49E-12	1.57351	972.3178	1524.558
A_23_P20	NM_01487	Homo sapi KIAA0020,	1.57346	3.31E-19	1.57346	9057.091	14232.9
A_24_P15	AB032978	Homo sapi KIAA1152	1.57275	4.09E-08	1.57275	576.9253	909.0248
A_24_P39	NM_00043	sialidase 1 NEU1,NEL	1.57271	2.56E-09	1.57271	1864.562	2956.435
A_23_P40	NM_00211	Major histo HLA-A	1.57239	7.53E-12	1.57239	10682.13	16712.79
A_24_P85	AF119913	Homo sapi PRO3077	1.5723	3.54E-10	1.5723	323.6691	508.8928

A_32_P93	BC036018	Protein tyrc	PTPRG	1.57169	0.00002	1.57169	250.3603	393.5552
A_23_P111	NM_00688	Homo sapi	ATBF1,ATI	1.57166	3.83E-21	1.57166	3060.662	4804.77
A_23_P67	NM_00434	Homo sapi	CALR,RO,	1.571	1.50E-08	1.571	18869.5	29344.1
A_23_P78	NM_01867	Homo sapi	ABCA5,AB	1.57073	3.87E-09	1.57073	679.1986	1070.624
A_23_P121	NM_00707	Huntingtin	HYPE,MG	1.56991	4.12E-08	1.56991	374.7417	588.4522
A_23_P28	NM_01784	Chromosor	C1orf27	1.56972	2.70E-12	1.56972	1839.841	2906.162
A_32_P62	NM_01457	schwannor	SCHIP1,SC	1.56903	3.09E-06	1.56903	1950.435	3062.643
A_23_P20	AL137667	Homo sapi	MAPK8,JN	1.56879	2.73E-09	1.56879	1337.756	2106.616
A_23_P25	NM_01843	putative zir	JMJD1A,T	1.56698	2.38E-11	1.56698	653.2262	1025.413
A_32_P12	NM_18258	Homo sapi	NM_18258	1.56667	0.00004	1.56667	968.1643	1537.524
A_24_P33	NM_00580	highly char	C13orf22,C	1.56625	5.78E-08	1.56625	864.5309	1363.621
A_23_P89	NM_14591	Homo sapi	ZNF23,KO	1.5659	2.97E-06	1.5659	307.4426	481.0513
A_24_P30	NM_01326	peroxisom	PPARGC1	1.56564	5.20E-09	1.56564	449.5672	703.3231
A_32_P13	NM_00659	heat shock	HSPA8,LA	1.56555	1.90E-08	1.56555	241934.5	377161.1
A_24_P28	NM_00659	heat shock	HSPA8,LA	1.56523	5.24E-10	1.56523	262685.7	409213.8
A_23_P38	NM_14468	Homo sapi	ZNF480,M	1.56507	1.09E-06	1.56507	1725.695	2738.057
A_23_P25	NM_15238	Homo sapi	BBS5	1.56465	2.68E-23	1.56465	12918.72	20224.79
A_24_P52	BI910970	Similar to	BI910970	1.56464	8.13E-09	1.56464	2453.892	3885.203
A_23_P51	NM_00472	Homo sapi	ARHGEF2	1.5644	4.56E-12	1.5644	1496.934	2343.728
A_23_P38	AB051515	TPR doma	TANC	1.56418	4.15E-08	1.56418	2323.733	3683.831
A_23_P15	NM_00418	Homo sapi	TANK,TRA	1.56404	1.80E-14	1.56404	3590.036	5618.883
A_23_P76	NM_01828	Homo sapi	PSPC1,PS	1.56379	1.63E-13	1.56379	899.0039	1408.417
A_23_P24	NM_00659	heat shock	HSPA8,LA	1.56321	2.31E-09	1.56321	258110.9	400622.5
A_24_P35	AF318360	MAX bindir	MNT	1.56267	1.36E-16	1.56267	3517.25	5484.241
A_23_P55	NM_00579	Homo sapi	MPHOSP	1.56204	7.34E-13	1.56204	6024.484	9456.441
A_23_P14	NM_00721	Homo sapi	RNF139,R	1.56189	1.75E-17	1.56189	2998.957	4697.974
A_23_P20	AB043587	Acyl-Coenz	ACBD3	1.56158	5.57E-17	1.56158	4312.103	6751.312
A_23_P37	NM_15239	Homo sapi	AHSA2,Hc	1.56069	4.82E-10	1.56069	3320.648	5216.976
A_23_P25	NM_02523	Homo sapi	ZNF435,Z	1.56036	0.00001	1.56036	243.0702	378.809
A_23_P37	NM_00404	beta-2-mic	B2M	1.55981	4.46E-22	1.55981	33963.18	52906.57
A_32_P20	NM_00399	CDC-like k	CLK3	1.55938	8.48E-09	1.55938	6646.415	10456.12
A_24_P22	NM_02485	hypothetic	FLJ22028	1.55907	0.00006	1.55907	296.8672	463.0808
A_24_P92	AK055915	CDC42 eff	CDC42EP	1.55892	2.01E-09	1.55892	459.4401	715.7018
A_23_P16	NM_00283	Homo sapi	PTPN4,PT	1.55827	2.82E-06	1.55827	721.4225	1134.627
A_24_P11	NM_01644	cysteine-ric	CRIM1,S5	1.55817	5.12E-11	1.55817	1047.617	1638.588
A_24_P86	AK092921	M4-50 mRI	AK092921	1.55798	2.69E-16	1.55798	17360.08	27122.08
A_23_P20	NM_02461	Homo sapi	PLEKHF2,	1.55774	4.05E-08	1.55774	938.5576	1471.907
A_23_P63	NM_00207	Homo sapi	GOT1	1.5577	4.86E-20	1.5577	10092.99	15728.18
A_24_P29	NM_03143	tethering fa	COG3,SEC	1.55745	1.72E-10	1.55745	1019.788	1598.672
A_23_P13	NM_13344	Homo sapi	CTGLF1,M	1.5574	2.20E-07	1.5574	400.8246	624.3322
A_24_P12	NM_01548	Homo sapi	KBTBD2,B	1.55688	2.21E-06	1.55688	654.5256	1022.419
A_23_P16	NM_00553	interferon,	IFI16,PYHI	1.5565	5.42E-19	1.5565	4521.231	7025.096
A_24_P85	NM_19427	Homo sapi	C14orf43,L	1.55615	9.72E-07	1.55615	4078.323	6323.132
A_23_P31	AB051518	KIAA1731	KIAA1731	1.55581	0.00099	1.55581	1564.527	2505.506
A_23_P11	AB029038	Homo sapi	KIAA1115	1.55562	0.00098	1.55562	267.0073	413.4122
A_32_P64	NM_15260	Chromosor	FLJ35382	1.5554	9.23E-20	1.5554	1072.443	1668.295
A_32_P169243		Unknown	A_32_P16	1.55532	1.77E-19	1.55532	849.0574	1320.675
A_23_P40	NM_01564	Homo sapi	ZBTB20,H	1.55451	2.87E-07	1.55451	4779.909	7427.652
A_23_P14	NM_00368	Homo sapi	PDLIM4,RI	1.5545	8.78E-22	1.5545	11046.36	17182.64
A_23_P24	NM_01847	Homo sapi	C10orf110,	1.55423	0.00006	1.55423	318.0661	495.0299
A_23_P13	NM_14529	Homo sapi	ZNF627,FL	1.55405	9.44E-06	1.55405	766.5386	1195.995
A_23_P13	NM_03148	Homo sapi	ITCH,AIF4,	1.55353	8.41E-07	1.55353	1201.938	1876.263

A_23_P170719	Unknown	A_23_P170719	1.55337	5.26E-10	1.55337	358.2748	556.5729
A_23_P201	NM_01584	methyl-CpG MBD1,RFT	1.55288	1.52E-12	1.55288	609.532	947.6564
A_32_P95	BX090181	Transcriber BX090181	1.55282	1.34E-06	1.55282	13287.77	20630.85
A_32_P181513	Unknown	A_32_P181513	1.55249	9.15E-08	1.55249	494.3019	769.6096
A_23_P23	U79751	Basic leucic BLZF1	1.55236	5.22E-14	1.55236	1949.233	3030.421
A_23_P11	M21963	Major histo HLA-C	1.55211	9.32E-15	1.55211	61302.63	94808.61
A_24_P15	NM_03130	DEAD (Asp) DKFZP564	1.55205	0.0004	1.55205	428.8297	669.0166
A_32_P75	AK128746	Hypothetic LOC39974	1.55165	2.45E-07	1.55165	1912.642	2958.496
A_23_P16	NM_00450	Homo sapi HRB,RAB,1	1.55132	2.52E-22	1.55132	7004.417	10857.5
A_23_P125109	Unknown	A_23_P125109	1.55045	1.90E-11	1.55045	16070.43	24835.43
A_23_P78	NM_01770	Homo sapi FBXL12,Ft	1.55034	1.89E-12	1.55034	1741.466	2697.339
A_23_P16	NM_18265	Homo sapi UTY,UTY1	1.54992	0.0008	1.54992	180.877	280.3763
A_24_P61	NM_01475	KIAA0187 BMS1L,KI/	1.54822	0.00023	1.54822	500.967	779.9513
A_24_P91	AK022299	Hypothetic LOC14818	1.54796	0.00016	1.54796	229.8544	354.793
A_23_P90	NM_02524	Homo sapi PBX4	1.54756	0.00003	1.54756	387.9561	600.877
A_23_P37	NM_01500	88-kDa gol GOLGA8A	1.54709	1.96E-06	1.54709	429.9352	666.0457
A_23_P11	AB029039	RNA bindir RBM16	1.54696	6.87E-08	1.54696	1963.763	3024.954
A_23_P41	NM_00634	Chromosor C13orf24	1.54655	0.00004	1.54655	671.5397	1042.92
A_23_P21	AK094968	Interferon, IFI16	1.54614	6.52E-23	1.54614	8420.968	13011.25
A_23_P25	X78926	Similar to ZHZF3	1.54567	3.21E-22	1.54567	3157.927	4884.132
A_32_P11	NM_19849	Homo sapi CTAGE4	1.54545	1.80E-06	1.54545	2189.127	3420.031
A_24_P38	NM_00316	ribosomal r RPS6KB1,	1.54517	3.46E-18	1.54517	1680.097	2599.349
A_24_P24	AF116642	Homo sapi PRO1618	1.54316	0.00077	1.54316	165.0383	254.2592
A_23_P10	NM_00284	Homo sapi PTPRH,SA	1.54256	0.00001	1.54256	419.4432	647.1043
A_23_P41	NM_01863	Homo sapi ETNK1,EK	1.54196	5.64E-06	1.54196	815.0502	1256.626
A_23_P10	AF454744	Zinc finger ZNF473	1.54153	1.27E-09	1.54153	406.2068	626.2634
A_24_P57	AK025344	Homo sapi AK025344	1.54103	7.51E-12	1.54103	3665.346	5674.625
A_32_P12372	ALU1_HU	THC19187	1.53997	0.00081	1.53997	1218.94	1860.677
A_24_P28	NM_01507	GTPase re ARHGAP2	1.53936	4.79E-16	1.53936	725.97	1116.956
A_23_P39	AL832556	Hypothetic LOC28533	1.5392	9.21E-06	1.5392	300.308	461.6477
A_23_P21	NM_00226	Homo sapi KPNA4,QII	1.53903	7.31E-08	1.53903	2172.978	3384.957
A_23_P57	AF288741	Oxysterol t OSBP2	1.53858	0.0009	1.53858	204.2804	315.5301
A_23_P90	NM_02217	TIA1 cytotc TIA1	1.5376	9.67E-06	1.5376	592.7667	917.2528
A_24_P913629	Unknown	A_24_P913629	1.53708	3.28E-06	1.53708	720.6307	1110.599
A_23_P20	NM_00535	Homo sapi LMO7,LON	1.53703	2.82E-10	1.53703	940.4226	1446.862
A_23_P21	NM_03322	Homo sapi TRIM15,RI	1.53697	4.78E-08	1.53697	315.3336	484.6726
A_23_P31	AK024388	Homo sapi FGFR1,H2	1.53571	0.00061	1.53571	201.4431	309.9479
A_23_P74	NM_01583	Homo sapi WARS2,Tr	1.53544	6.65E-18	1.53544	1604.605	2467.396
A_23_P12	NM_15333	Homo sapi C10orf89,N	1.53441	8.16E-09	1.53441	703.3065	1083.212
A_24_P36	AK023023	WD repeat WDR26	1.53401	2.20E-21	1.53401	1594.707	2445.164
A_23_P32	NM_00580	Homo sapi TOPORS,L	1.53308	2.47E-06	1.53308	659.681	1018.259
A_32_P41	NM_13344	Homo sapi CTGLF1,M	1.5326	8.00E-09	1.5326	386.6925	592.5968
A_24_P55	BC038432	Retinoic ac RARA	1.53226	0.00067	1.53226	849.8386	1296.748
A_32_P80	AK096255	CDNA FLJ: AK096255	1.53213	0.00002	1.53213	585.5454	901.083
A_32_P4882	Unknown	A_32_P4882	1.53187	5.67E-14	1.53187	9216.947	14189.17
A_23_P41	NM_14505	Homo sapi C18orf25,A	1.53179	0.00073	1.53179	595.3109	915.8627
A_24_P21	NM_01648	hypothetic C9orf156,N	1.53035	0.00001	1.53035	261.949	401.3854
A_24_P367421	MP10_HU	THC19471	1.52963	0.00001	1.52963	723.0588	1119.056
A_23_P79	NM_05294	Homo sapi NOSTRIN,	1.52961	3.51E-13	1.52961	1065.154	1628.756
A_24_P30	NM_02472	MICAL-like MICAL-L2	1.52931	2.40E-12	1.52931	3843.965	5893.679
A_24_P28	AF173886	ALL1 fusec AF5Q31	1.52874	1.23E-08	1.52874	1173.685	1802.499
A_23_P97	AB046826	Hypothetic FLJ20203	1.5281	1.29E-17	1.5281	2665.466	4068.42

A_24_P230416	Unknown	A_24_P230416	1.5278	1.12E-10	1.5278	8088.12	12441.46
A_32_P41	(AK092942 Hypothetic	AK092942	1.52769	0.00015	1.52769	750.11	1148.079
A_23_P45	NM_08063 Homo sapi	UPF3B,UP	1.52757	9.01E-06	1.52757	2780.551	4325.461
A_23_P21	NM_01777 hypothetic	ZNF673,FL	1.52738	0.00002	1.52738	1305.182	2008.333
A_23_P25	(NM_00207 Homo sapi	GOLGA4,C	1.52564	4.20E-08	1.52564	2308.866	3540.629
A_24_P40	(NM_00344 zinc finger	ZNF141,D	1.52507	2.54E-06	1.52507	585.2802	892.5503
A_23_P14	(NM_02117 Homo sapi	HES4	1.52471	1.19E-16	1.52471	11131.62	16983.88
A_23_P13	(NM_00179 Homo sapi	CDK7,CAK	1.52447	5.14E-23	1.52447	10023.54	15275.34
A_23_P78	(NM_01825 Homo sapi	STATIP1,E	1.52434	9.95E-11	1.52434	11807.14	17926.23
A_24_P43	(NM_01462 gene predi	MAD2L1BF	1.52397	1.80E-21	1.52397	3266.928	4980.384
A_24_P68	(NM_02100 Homo sapi	UBC,HMG	1.52342	5.30E-06	1.52342	367271.1	561075.8
A_23_P30	(NM_00212 HLA-G hist	HLA-G,MH	1.52246	0.00022	1.52246	371.8549	566.687
A_23_P14	(NM_02173 Homo sapi	AVP11,VIP	1.52191	5.88E-15	1.52191	16348.49	24830.12
A_24_P12	(NM_01826 hypothetic	TMEM39A	1.52132	3.29E-09	1.52132	841.5441	1282.616
A_24_P85	BC063702 SEC31-like	SEC31L1	1.5211	1.28E-06	1.5211	232.3975	353.46
A_23_P48	(NM_02455 Homo sapi	ADIPOR2,I	1.52041	8.78E-13	1.52041	5205.135	7916.651
A_23_P43	(NM_15254 Homo sapi	SRFBP1,P	1.51977	5.67E-06	1.51977	2388.707	3631.292
A_23_P21	(NM_00465 Homo sapi	USP9X,DF	1.51975	1.48E-11	1.51975	16320.42	24667.48
A_24_P44	(NM_01494 KIAA0872	MON1B,H	1.51927	1.23E-12	1.51927	1557.832	2376.478
A_23_P20	(NM_01831 Homo sapi	SAPS3,SA	1.51922	5.26E-08	1.51922	544.1015	828.4677
A_24_P20	(NM_20338 Homo sapi	ACSL5,AC	1.51922	3.99E-09	1.51922	1061.475	1622.112
A_23_P30	BC001864 Inositol hex	IHPK2	1.51856	1.02E-07	1.51856	11260.54	17084.19
A_23_P10	(NM_00423 Homo sapi	TRIP11,CE	1.51836	7.72E-08	1.51836	2349.135	3609.834
A_24_P68	(NM_00471 serological	SDCCAG1	1.51819	1.65E-09	1.51819	7154.721	10926.47
A_24_P932706	Unknown	A_24_P932706	1.51728	0.00085	1.51728	454.1641	693.3776
A_23_P21	(NM_14532 Homo sapi	OSBPL3,O	1.51705	0.00001	1.51705	902.9356	1382.804
A_23_P20	(NM_01481 centrosom	CAP350,gr	1.51703	1.98E-09	1.51703	1318.765	2013.464
A_23_P52	(NM_00679 Homo sapi	EHD1,PAS	1.51686	6.60E-09	1.51686	5192.477	7890.539
A_32_P10	(NM_00205 guanylate t	GBP1	1.51647	0.00005	1.51647	686.6461	1049.345
A_23_P43	(NM_14471 Homo sapi	SEPT10,FI	1.51637	6.64E-13	1.51637	8235.92	12546.45
A_23_P25	(NM_00706 Homo sapi	TBC1D8,A	1.51629	5.87E-10	1.51629	1732.364	2635.018
A_24_P27	(NM_05287 hypothetic	MGC4677,	1.51621	4.19E-11	1.51621	7998.434	12205.72
A_24_P24	(NM_02199 zinc finger	ZNF277,NI	1.51579	1.03E-17	1.51579	1944.154	2950.44
A_32_P21	(BM921567 Transcriber	BM921567	1.51514	0.00004	1.51514	273.9085	414.7975
A_23_P49	(NM_03289 hypothetic	MGC1437	1.51466	1.05E-09	1.51466	601.8012	912.4059
A_24_P83	(NM_19818 Homo sapi	ASTN2,KIA	1.51458	7.83E-09	1.51458	15470.14	23531.61
A_23_P25	(NM_03241 putative nu	ORF1-FL4	1.51456	3.08E-20	1.51456	12525.46	18956.16
A_23_P43	AB029040 KIAA1117	KIAA1117	1.5143	4.37E-08	1.5143	877.042	1332.632
A_23_P74	(NM_00568 Homo sapi	TAF1A,SL	1.51351	8.30E-09	1.51351	1758.741	2683.581
A_23_P42	(NM_03244 ligand-dep	MLR2,LCC	1.51248	7.92E-08	1.51248	1119.362	1709.874
A_24_P89	(NM_00345 zinc finger	ZNF189	1.51247	0.00001	1.51247	582.6848	886.8942
A_24_P28	(AB002303 Zinc finger,	ZFYVE16	1.51198	7.48E-08	1.51198	767.9166	1165.839
A_23_P35	(AB037838 Inhibitor of	IBTK	1.51169	2.20E-09	1.51169	6833.825	10407.7
A_24_P30	(NM_00718 WW domai	WBP4,FBF	1.51165	0.00004	1.51165	247.7787	374.1079
A_23_P14	BC036033 Ribosomal	RPS6KB1	1.51111	1.12E-11	1.51111	2209.465	3348.413
A_24_P40	(NM_01664 HDCMC28	C20orf6,AE	1.51087	0.00002	1.51087	1134.019	1735.887
A_23_P32	AK075506 Midnolin	MIDN	1.51053	2.72E-09	1.51053	1038.78	1563.436
A_32_P45	AB019569 Glycoprote	CGA	1.51042	0.00052	1.51042	153.6375	232.008
A_24_P20	(NM_01622 zinc finger	ZNF588,ZF	1.50994	8.30E-16	1.50994	665.1969	1004.341
A_32_P22	(AB020719 KIAA0912	Cep152	1.50992	0.00037	1.50992	555.0883	842.6696
A_23_P42	(NM_00477 Homo sapi	PPARBP,F	1.50917	2.61E-19	1.50917	7105.992	10733.89
A_32_P12	BC024745 Homo sapi	LOC39988	1.50866	1.86E-09	1.50866	611.4249	922.3872

A_23_P88	NM_17209	Homo sapi	CATSPER	1.50865	0.00032	1.50865	342.9828	517.4163
A_24_P30	NM_00504	reelin	RELN,RL	1.50805	0.00002	1.50805	588.3051	891.9547
A_24_P41	(NM_19933	Homo sapi	FLJ35171	1.50745	2.07E-10	1.50745	508.7554	767.212
A_23_P68	NM_01629	mammaliar	IHPK2,PiU	1.50703	0.00009	1.50703	275.285	415.6834
A_23_P19	NM_03245	serine/thre	STK19,G1	1.50681	2.33E-13	1.50681	676.6915	1019.693
A_23_P105732		Unknown	A_23_P10	1.50671	1.20E-08	1.50671	30960.64	46398.15
A_24_P77	(NM_00413	heat shock	HSPA9B,C	1.50632	1.85E-14	1.50632	18625.47	27983.04
A_23_P43	(NM_17587	Homo sapi	EXOC8,EX	1.50631	1.49E-09	1.50631	808.782	1222.373
A_23_P31	(NM_01895	major histo	HLA-F,HLA	1.50612	1.17E-09	1.50612	12574.19	18846.62
A_24_P94	(NM_00341	zinc finger	ZFY	1.50594	0.00001	1.50594	474.5698	715.1406
A_24_P738130		27486 MLN	THC19657	1.50594	0.00006	1.50594	677.3572	1023.867
A_23_P10	(AK000707	Hepatoma-	HDGF2	1.5054	9.26E-11	1.5054	8018.044	12102.71
A_24_P11	(NM_17854	Homo sapi	ZNF678,M	1.50492	0.0005	1.50492	371.1398	559.2559
A_24_P69	BC014776	Hypothetic	MGC4677	1.50485	5.04E-08	1.50485	7715.346	11731.86
A_23_P13	(NM_01394	chloride int	CLIC4,H1,I	1.50433	1.18E-17	1.50433	7854.102	11823.02
A_24_P34	(NM_03287	Homo sapi	C6orf111,I	1.50433	2.33E-06	1.50433	4634.142	7060.296
A_23_P13	(NM_02526	Homo sapi	THUMPD2	1.50417	1.53E-12	1.50417	1468.835	2216.333
A_24_P92	AJ584699	Ras associ	RAPH1	1.50401	0.00003	1.50401	2218.919	3376.21
A_23_P56	(NM_00577	Homo sapi	DHRS9,RC	1.50368	3.35E-06	1.50368	586.1893	883.3871
A_23_P11	AL136837	Hypothetic	DKFZp434	1.50354	5.65E-09	1.50354	3809.523	5734.561
A_32_P10067		Unknown	A_32_P10	1.50333	9.80E-09	1.50333	997.2343	1505.142
A_24_P17	(NM_00645	LIM protein	PDLIM5,L9	1.50291	0.00083	1.50291	449.9465	679.5093
A_23_P25	(NM_02163	apoptosis i	FKSG2	1.50271	2.56E-08	1.50271	21325.79	32344.38
A_24_P16	(NM_19849	Homo sapi	CTAGE4	1.50244	2.93E-10	1.50244	2913.644	4396.2
A_23_P74	(NM_03321	Zinc finger	MGC1246	1.50167	1.45E-09	1.50167	1201.08	1798.169
A_24_P91	AF287958	Major histo	HLA-A	1.50132	9.44E-08	1.50132	8581.838	12809.38
A_32_P32	(NM_01764	hypothetic	FAM29A,R	1.50106	5.84E-17	1.50106	1592.77	2388.637
A_23_P19	(NM_00337	Homo sapi	VIL2,CVL,C	1.50061	1.78E-12	1.50061	6101.257	9170.651
A_23_P21	(NM_00183	clusterin (c	CLU,CLI,A	1.50004	4.05E-21	1.50004	2912.262	4367.936
A_23_P13	(NM_00110	Homo sapi	ACTB	1.49989	1.69E-07	1.49989	92508.87	137507.8
A_24_P31	(NM_00212	HLA-G hist	HLA-G,MH	1.49975	5.15E-15	1.49975	37195.11	55669.41
A_23_P13	(NM_01519	Homo sapi	SNF1LK2,C	1.49956	3.19E-06	1.49956	473.9672	710.23
A_23_P53	(NM_01325	Homo sapi	ZNF215,B	1.49898	4.33E-07	1.49898	573.832	862.4192
A_32_P76	(NM_01657	TU12B1-T	TU12B1-T	1.49865	2.17E-08	1.49865	375.3974	562.5839
A_23_P35	(NM_15260	Homo sapi	ZNF567,M	1.49859	7.84E-06	1.49859	687.9818	1033.322
A_24_P10	(NM_20554	Homo sapi	ALS2CR16	1.4969	0.00019	1.4969	754.3051	1141.846
A_23_P40	(NM_00397	Homo sapi	BTAF1,MC	1.49661	7.35E-12	1.49661	622.9719	932.4554
A_24_P63	(NM_00624	protein phc	PPP1R2,IF	1.49637	1.39E-08	1.49637	1003.55	1511.101
A_23_P14	(NM_03212	implantatio	DKFZp564	1.49603	4.55E-16	1.49603	1093.301	1636.989
A_23_P16	(NM_01451	Homo sapi	CNOT2,NC	1.49581	3.30E-09	1.49581	6112.662	9214.27
A_23_P21	(AK025201	Nuclear tra	NFYA	1.49563	2.57E-08	1.49563	698.6526	1047.994
A_24_P36	(NM_19807	golgi autoa	FLJ40113,I	1.49545	1.18E-08	1.49545	1031.406	1546.821
A_23_P14	(NM_08065	similar to R	MGC17337	1.49521	1.85E-19	1.49521	2568.308	3838.202
A_32_P16	(NM_19849	Homo sapi	CTAGE4	1.49503	2.73E-09	1.49503	3657.775	5493.891
A_24_P22	BX537874	Hypothetic	DKFZp313	1.49473	0.00078	1.49473	612.1935	922.8817
A_32_P96	(NM_00329	tumor prote	TPT1,HRF	1.49462	2.73E-22	1.49462	121094.8	180983.8
A_32_P12	(NM_08065	similar to R	MGC17337	1.49442	2.20E-15	1.49442	2376.934	3545.802
A_24_P20	(NM_01664	HDCMC28	C20orf6,AE	1.49435	0.00057	1.49435	1280.54	1955.404
A_24_P65	BX648822	KIAA0912	Cep152	1.4943	0.00011	1.4943	327.9558	490.8158
A_23_P12	(NM_01243	SET domai	SETDB1,E	1.4943	1.95E-08	1.4943	4807.502	7211.328
A_23_P31	AY055760	Decay acc	DAF	1.49345	0.00019	1.49345	833.7439	1246.583
A_24_P35	AB051895	Epiplakin 1	EPPK1	1.49335	6.35E-06	1.49335	690.0231	1027.193

A_23_P35: NM_00638 Homo sapi	CDRT1,HR	1.49332	0.00012	1.49332	306.5038	457.5023
A_32_P213946	Unknown A_32_P21:	1.49278	7.81E-09	1.49278	549.4832	821.2231
A_24_P40: NM_00584 Homo sapi	MYCPBP,I	1.49166	2.65E-06	1.49166	439.6249	656.5051
A_23_P15: NM_00299 Homo sapi	CXCL6,GC	1.49166	0.00096	1.49166	177.1367	264.4823
A_32_P10: AB095935 Ankyrin re	FLJ35740	1.49105	0.00036	1.49105	836.2965	1266.651
A_23_P33: BC007197 Nuclear fac	NFKBIB	1.4909	0.00087	1.4909	310.0558	464.3127
A_23_P28: NM_17237 Homo sapi	ELF1	1.49085	5.91E-16	1.49085	1773.068	2644.029
A_24_P33: NM_00476 serine/thre	STK17A,DI	1.49078	1.17E-13	1.49078	4948.983	7402.049
A_32_P192615	Unknown THC18582	1.49063	5.90E-13	1.49063	4468.371	6680.865
A_24_P31: NM_01456 zinc finger	ZFP95,KIA	1.4906	0.00021	1.4906	313.5217	466.3171
A_32_P10: AY354203 RAN-bindir	LOC40096	1.48998	3.68E-08	1.48998	890.0779	1328.521
A_23_P20: NM_00625 Homo sapi	PRKAB1,A	1.48979	0.00004	1.48979	580.6814	869.2588
A_23_P25: NM_00495 Homo sapi	EBI2	1.48959	3.59E-09	1.48959	373.2026	555.9122
A_32_P33: XM_29282 similar to h	LOC34296	1.48958	4.44E-06	1.48958	229.8835	342.4524
A_32_P58: NM_14580 TL132 prot	LOC22059	1.48948	0.00001	1.48948	1199.981	1809.673
A_32_P22: NM_18248 Homo sapi	CPEB2,MC	1.48947	0.00007	1.48947	261.109	389.3533
A_23_P28: NM_14469 Homo sapi	ZNF558,FL	1.48853	1.90E-06	1.48853	698.6563	1042.069
A_23_P65: NM_01430 Homo sapi	TGDS,TDF	1.48768	0.00029	1.48768	2088.095	3165.599
A_23_P91: NM_14471 Homo sapi	SEPT10,FI	1.48757	0.0001	1.48757	393.5308	585.967
A_24_P94: U92014	Human clo U92014	1.48743	0.00048	1.48743	1546.786	2352.534
A_24_P41: BC020891 HLA-G hist	HLA-G	1.48668	6.00E-18	1.48668	16031.92	23864.88
A_24_P30: NM_00329 tumor prote	TPT1,HRF	1.48583	7.04E-09	1.48583	116819.5	172369.9
A_23_P117912	AY059636 THC20037	1.48558	8.26E-13	1.48558	2228.01	3311.226
A_23_P20: NM_02122 Homo sapi	NTN4,PRC	1.48427	2.63E-07	1.48427	468.4821	696.1898
A_24_P733308	Unknown A_24_P73:	1.48424	0.00008	1.48424	669.6811	994.0532
A_32_P104053	Unknown THC19019	1.48355	0.00009	1.48355	1923.218	2889.621
A_24_P16: NM_00318 TATA box	ITAF2,TAF2	1.48299	2.02E-06	1.48299	904.1952	1349.558
A_24_P64: NM_02165 SMA4	SMA4	1.48286	3.71E-06	1.48286	292.8613	434.2119
A_23_P15: NM_01587 Homo sapi	AZIN1,OA2	1.48235	2.33E-09	1.48235	5288.497	7898.751
A_32_P179646	Unknown A_32_P17:	1.48202	0.00002	1.48202	286.1702	424.1829
A_23_P14: AK057783 Zinc finger	ZNF18	1.48097	0.00093	1.48097	657.9434	977.6782
A_23_P20: NM_02431 Homo sapi	LENG1	1.48072	1.88E-06	1.48072	335.0318	496.3089
A_24_P17: NM_00329 tumor prote	TPT1,HRF	1.48068	5.99E-11	1.48068	207604.3	305774.3
A_32_P213002	Unknown A_32_P21:	1.48066	2.46E-06	1.48066	2570.928	3751.505
A_23_P11: NM_00667 MHC class	HCP5,P5-1	1.4806	5.02E-07	1.4806	529.8046	783.9779
A_32_P22: NM_05291 Establishm	ESCO1	1.47982	0.00027	1.47982	4268.537	6290.75
A_23_P88: NM_01610 Homo sapi	SCFD1,RA	1.4798	1.77E-16	1.4798	6742.563	9965.407
A_23_P12: NM_02429 Homo sapi	C20orf149,	1.47979	1.05E-14	1.47979	29864.89	44278.06
A_23_P49: NM_00125 Homo sapi	CDC6,CDC	1.47973	6.68E-18	1.47973	4948.337	7316.507
A_23_P10: NM_14499 Homo sapi	ARL2L1,M	1.47943	0.00008	1.47943	504.5033	750.101
A_23_P23: NM_02494 Homo sapi	C10orf97,C	1.4793	0.00003	1.4793	1024.64	1531.506
A_23_P31: NM_00653 nuclear rec	NCOA3,AC	1.47915	0.00008	1.47915	2598.658	3861.517
A_23_P13: NM_02080 Homo sapi	SIPA1L2,K	1.47914	0.00003	1.47914	251.0347	371.1529
A_24_P174824	ALU4_HUM THC19983	1.47863	6.82E-13	1.47863	2696.546	3999.515
A_24_P40: AK056071 Fibronectin	FNDC3A	1.47809	0.00008	1.47809	886.2955	1300.598
A_23_P11: NM_02018 Homo sapi	CNNM4,AC	1.47795	0.00002	1.47795	2269.636	3356.123
A_24_P32: NM_00723 SP140 nuc	SP140,LY9	1.47791	0.00028	1.47791	1442.397	2168.747
A_24_P50567	Unknown A_24_P50:	1.47782	1.07E-11	1.47782	119452.5	175685.3
A_24_P83: NM_17858 Homo sapi	WDFY3,AL	1.47774	0.00003	1.47774	700.413	1036.725
A_23_P13: NM_00658 Homo sapi	GMEB1,PI	1.47708	5.33E-07	1.47708	823.2145	1217.926
A_23_P21: AY358828 SLIT and N	SLITRK2	1.47701	8.85E-08	1.47701	44203.55	64877.7
A_24_P51: NM_02246 hypothetic	AZI2,AZ2,N	1.47692	3.14E-06	1.47692	883.9894	1314.516

A_23_P11(NM_01870 Homo sapi TRIM36,R	1.47655	4.18E-08	1.47655	289.6379	427.6678
A_23_P78975 Unknown A_23_P78	1.47586	8.28E-18	1.47586	34660.62	51103.95
A_24_P255252 Unknown ENST0000	1.47569	7.50E-14	1.47569	161262.4	237196.9
A_32_P20 BC007307 Hypothetic LOC91664	1.47531	2.68E-11	1.47531	1008.121	1488.858
A_24_P28 NM_00361 mitogen-ac MAP4K3,G	1.47499	0.00084	1.47499	197.2927	291.4759
A_32_P97 AF086126 Chromosor LOC14617	1.47492	0.00002	1.47492	1232.812	1836.589
A_23_P20 NM_03210 Homo sapi PPP1R12E	1.47371	0.00004	1.47371	1281.219	1900.69
A_32_P44 NM_00483 absent in n AIM2,PYHI	1.47351	0.00007	1.47351	293.4929	431.9042
A_24_P69 AK022339 Seryl-tRNA SARS	1.47243	3.82E-06	1.47243	904.4276	1339.876
A_24_P24 NM_01517 Homo sapi BAT2D1,X	1.47194	1.95E-10	1.47194	11459.89	16960.29
A_24_P86 NM_01450 putative Rε RABGEF1,	1.47151	8.89E-11	1.47151	546.4772	804.0888
A_23_P35 NM_00395 Homo sapi SOCS3,Cl	1.47141	3.33E-17	1.47141	1882.828	2767.86
A_23_P10 NM_01336 zinc finger ZNF223	1.47105	0.00015	1.47105	299.9697	441.4059
A_32_P200586 Unknown A_32_P20	1.47055	0.00003	1.47055	237.9234	349.9132
A_24_P37 NM_00642 step II splic SLU7,hSlu	1.46972	2.01E-14	1.46972	2100.429	3082.186
A_23_P42 NM_13363 Homo sapi RHOV,Chp	1.46892	1.38E-17	1.46892	2007.936	2949.669
A_24_P38 NM_00468 myotubular MTMR4,ZF	1.46806	1.70E-08	1.46806	3623.104	5356.48
A_24_P842962 Unknown THC19599	1.46693	0.00033	1.46693	488.3462	718.4134
A_24_P21 NM_00650 protein tyrc PTPRE,PT	1.46661	0.00037	1.46661	737.0237	1097.928
A_24_P40 NM_02476 hypotheticε WDR78,FL	1.46657	0.00005	1.46657	356.227	522.6146
A_23_P38 AL080138 Structural r KIAA0650	1.46618	6.86E-08	1.46618	1245.594	1833.792
A_24_P40 NM_17504 Homo sapi PILRB,FDF	1.4654	3.35E-12	1.4654	1835.885	2695.354
A_24_P89 NM_02125 ring finger j TRIM39,TF	1.46488	1.09E-06	1.46488	533.2124	782.676
A_23_P29 NM_01658 Homo sapi MBIP	1.4648	8.09E-06	1.4648	1525.766	2254.322
A_23_P10 NM_01765 aftiphilin pr AFTIPHILII	1.46476	0.00005	1.46476	1222.63	1819.236
A_24_P13 NM_00055 von Hippel VHL,RCA1	1.46421	8.55E-13	1.46421	899.8254	1316.044
A_24_P11 NM_02485 hypotheticε C17orf41,F	1.46421	0.00078	1.46421	454.994	668.6748
A_24_P92 BG429154 MRNA; cDI ZNF21	1.46419	0.00047	1.46419	223.1057	326.535
A_24_P33 NM_01457 nuclear aut STRN3,SG	1.46417	1.88E-06	1.46417	1523.687	2252.446
A_32_P703 Unknown A_32_P70	1.46406	0.0004	1.46406	284.9039	417.5745
A_32_P108203 Unknown A_32_P10	1.46376	5.26E-08	1.46376	773.0654	1131.326
A_24_P773958 Unknown A_24_P77	1.46321	0.00003	1.46321	267.4283	391.3896
A_24_P48 NM_03141 GABA(A) r GABARAP	1.46288	4.77E-09	1.46288	367.0149	536.9005
A_23_P30 NM_14717 A kinase (FAKAP9	1.46268	1.93E-06	1.46268	1967.179	2887.506
A_23_P12 NM_02403 hypotheticε MGC3130,	1.46188	2.78E-08	1.46188	358.2635	523.7629
A_23_P21 NM_15277 hypotheticε MGC3321	1.46107	2.05E-06	1.46107	848.0567	1247.638
A_23_P25 NM_00261 Homo sapi PEX13,ZW	1.46099	1.01E-09	1.46099	1921.128	2812.621
A_24_P85 NM_18145 Homo sapi GCC2,GC	1.46077	0.00002	1.46077	872.4155	1279.91
A_32_P47 BC060766 Solute carr SLC2A14	1.46076	0.00075	1.46076	770.6965	1119.517
A_32_P234570 Unknown THC19462	1.46059	4.27E-12	1.46059	29101.57	42441.88
A_23_P25 NM_00382 Homo sapi RIPK2,RIC	1.46055	7.16E-11	1.46055	2769.163	4054.395
A_23_P15 NM_18252 Homo sapi ZNF595,FL	1.45906	1.41E-08	1.45906	337.6735	492.6801
A_24_P58 NM_02114 Homo sapi ZNF273,Hi	1.459	0.0001	1.459	413.1715	603.1066
A_23_P30 NM_00719 Homo sapi POLI,RAD	1.45891	0.00033	1.45891	623.2454	916.2393
A_23_P12 NM_00199 Homo sapi F3,TF,TFA	1.45884	9.61E-06	1.45884	595.9379	871.3471
A_23_P50 AB027121 Chromosor C18orf12	1.45868	7.51E-08	1.45868	16509.98	24176.51
A_23_P62 NM_02464 Ischemia/rε FLJ23476	1.45856	1.99E-18	1.45856	5045.932	7361.698
A_24_P36 NM_00361 solute carri SLC4A7,NI	1.45765	0.00023	1.45765	883.8604	1304.451
A_23_P34 NM_02073 Homo sapi CCPG1,CF	1.45758	0.00002	1.45758	263.147	383.4154
A_24_P562242 Unknown A_24_P56	1.45751	0.00006	1.45751	550.8062	805.4159
A_24_P45 L40520 (clone DR3 L40520	1.45733	0.00035	1.45733	261.65	380.5688
A_23_P39 NM_01657 HBV pX as HBXAP,RS	1.45726	1.73E-11	1.45726	1174.56	1716.818

A_23_P11(NM_01635 Homo sapi IRX4	1.45676	0.00046	1.45676	730.2277	1072.529
A_24_P70(D55639 Kynureninase KYNU	1.45667	5.06E-06	1.45667	2035.122	2971.781
A_23_P87(NM_00283 Homo sapi PTPN12,PT	1.45594	9.36E-09	1.45594	3748.468	5499.048
A_23_P59(NM_00008 Homo sapi CLCN1,CL	1.45586	7.46E-13	1.45586	4007.136	5839.652
A_24_P68(BC043212 Hypothetical BC043212	1.45585	0.00001	1.45585	649.6814	950.9578
A_23_P10(NM_00214 Homo sapi HOXB2,K8	1.45561	3.28E-06	1.45561	429.1046	624.9387
A_23_P13(NM_00032 retinitis pig RPGR,CRI	1.45499	2.71E-07	1.45499	987.5922	1442.424
A_23_P15(NM_17069 Homo sapi TGIF,HPE4	1.45455	0.0001	1.45455	1726.4	2538.704
A_23_P20(NM_00584 Homo sapi STAM2,Hb	1.45416	3.10E-10	1.45416	556.3451	808.9818
A_23_P63(NM_01510 Homo sapi POGZ,SUF	1.45357	0.00005	1.45357	5912.255	8704.005
A_24_P17(NM_01325 muskelin 1 MKLN1,FL	1.45318	2.65E-07	1.45318	670.9058	978.0252
A_24_P32(XM_37056 Homo sapi ARL4P,AR	1.45299	0.00019	1.45299	536.8118	783.5273
A_24_P414376 Unknown THC20025	1.45296	0.00002	1.45296	925.0235	1357.261
A_23_P54(NM_03127 Testis expr TEX14	1.45296	0.00037	1.45296	199.8192	290.4633
A_24_P34(NM_14523 Homo sapi ZNF625	1.45276	0.00005	1.45276	364.5184	529.7319
A_32_P13(AB051515 TPR doma TANC	1.45257	1.55E-07	1.45257	1823.704	2659.921
A_23_P82(BC013179 TWIST nei TWISTNB	1.45187	0.00006	1.45187	6024.486	8902.406
A_24_P54(NM_01604 CGI-100 pr TMED5,CC	1.45185	0.00001	1.45185	4587.606	6751.427
A_23_P27(NM_01895 Homo sapi UBB,MGC4	1.45151	1.29E-09	1.45151	238999.8	345405.5
A_23_P39(NM_00488 CBF1 inter CIR	1.45079	7.67E-17	1.45079	1408.239	2042.714
A_23_P15(BC052980 CGG triplet CGGBP1	1.45075	3.03E-15	1.45075	5552.533	8068.344
A_24_P33(NM_15235 hypothetical MGC40157	1.45002	8.45E-14	1.45002	5699.176	8251.748
A_24_P33(AL133018 Hypothetical AL133018	1.44971	2.16E-13	1.44971	2563.552	3723.815
A_32_P32(NM_02206 hypothetical C10orf84,F	1.44902	9.38E-10	1.44902	3827.767	5576.523
A_24_P100695 Lipin 3 (Lip LPIN3	1.44893	0.00001	1.44893	770.5655	1108.265
A_24_P23(NM_03184 microtubule MAP2,MAF	1.44889	0.00031	1.44889	309.0507	447.211
A_23_P14(NM_00471 Homo sapi SDCCAG1	1.44858	1.16E-08	1.44858	3337.045	4862.627
A_23_P89(NM_00365 Homo sapi CBX4,PC2	1.44853	0.00002	1.44853	2201.776	3192.909
A_23_P11(NM_00576 Homo sapi CEBPZ,CE	1.44843	8.06E-17	1.44843	8088.959	11706.99
A_23_P11(NM_00642 step II splic SLU7,hSlu	1.44839	1.72E-18	1.44839	2962.407	4291.33
A_23_P93(NM_01657 Homo sapi GLULD1,Li	1.44814	8.65E-18	1.44814	2351.998	3407.16
A_23_P77(NM_01761 Homo sapi RNF111,Al	1.44799	0.0007	1.44799	1185.45	1711.584
A_23_P89(NM_01627 Homo sapi RNF138,H	1.44777	8.75E-09	1.44777	2236.24	3259.878
A_32_P52(NM_14569 Homo sapi LPIN1,KIA	1.44762	2.63E-08	1.44762	3851.161	5605.496
A_23_P75(NM_14580 Homo sapi TRAF6,RN	1.44718	0.00008	1.44718	883.4636	1275.116
A_23_P62(NM_00341 Homo sapi ZFY	1.44669	0.00032	1.44669	471.5776	683.6616
A_23_P29(NM_01547 transcriptio WWTR1,T	1.44632	1.27E-09	1.44632	2030.156	2939.265
A_23_P12(NM_00541 Homo sapi SEPP1,Se	1.44617	2.74E-10	1.44617	2152.404	3121.518
A_23_P41(NM_15240 hypothetical FLJ35779,l	1.44563	7.49E-08	1.44563	1228.745	1781.076
A_24_P24(AK098175 Zinc finger ZNF283	1.44469	5.86E-06	1.44469	449.6576	649.3402
A_32_P167883 ALU7_HUMAN THC19703	1.44448	0.00005	1.44448	277.705	401.0748
A_32_P78(NM_14457 Homo sapi C14orf32,M	1.44442	3.32E-09	1.44442	2057.679	2984.289
A_23_P10(NM_02504 hypothetical ZNF614,FL	1.44412	1.18E-10	1.44412	616.9642	891.2958
A_32_P146884 ALU1_HUMAN THC18615	1.44386	0.00052	1.44386	403.2046	582.3658
A_32_P89(NM_18258 Homo sapi NM_18258	1.44382	0.00009	1.44382	2004.379	2933.504
A_24_P40(NM_00407 cryptochrome CRY1,PHL	1.44374	0.00017	1.44374	347.1097	500.6177
A_32_P212764 AE011791 THC18475	1.44365	0.00027	1.44365	1666.362	2387.909
A_32_P20(AF075027 Full length AF075027	1.44351	9.55E-09	1.44351	5449.016	7876.658
A_24_P20(U28831 Protein imr LOC40098	1.44307	2.66E-06	1.44307	1636.927	2384.186
A_23_P20(NM_00662 cyclic AMP ARPP-19,A	1.4429	7.53E-07	1.4429	3616.057	5248.367
A_23_P37(AF426265 Chromosom C21orf87	1.44289	6.02E-08	1.44289	4229.193	6036.057
A_23_P16(NM_03310 hypothetical LOC85865	1.44229	3.27E-08	1.44229	2036.533	2946.064

A_23_P21	NM_03348	Homo sapi	CDC2L1,p	1.44164	5.70E-08	1.44164	10521.13	15257.38
A_23_P12	NM_00593	Homo sapi	CTAGE5,M	1.44142	3.86E-07	1.44142	4465.05	6470.026
A_23_P36	NM_00685	Homo sapi	GLIPR1,GI	1.44106	1.09E-09	1.44106	3676.483	5327.797
A_24_P31	NM_01231	antigenic d	KIN,BTCD,	1.44104	6.12E-06	1.44104	1553.407	2256.436
A_23_P25	NM_00346	zyxin	ZYX,ESP-2	1.44102	4.47E-06	1.44102	1171.362	1688.572
A_32_P16	NM_00213	heterogene	HNRPA1,F	1.44049	0.00002	1.44049	18076.31	25904.87
A_23_P74	NM_00407	Homo sapi	CSR1,CF	1.44047	4.97E-07	1.44047	6064.719	8815.317
A_23_P35	NM_15323	Homo sapi	ZNF550,M	1.44004	1.14E-06	1.44004	1748.934	2524.922
A_32_P18	NM_14523	Homo sapi	ZNF625	1.44	2.82E-06	1.44	1402.103	2021.188
A_23_P31	NM_01626	GIOT-3 for	ZNF12,KO	1.43942	8.88E-12	1.43942	1831.769	2643.378
A_24_P75	AL832683	Protein pre	PTAR1	1.4382	8.89E-11	1.4382	2928.937	4224.737
A_23_P33	NM_14564	Homo sapi	WDR67,Gr	1.4376	9.72E-07	1.4376	1505.643	2177.229
A_23_P19	NM_01809	G1 to S ph	GSPT2,GS	1.43741	0.00003	1.43741	2330.46	3392.878
A_24_P26	NM_01560	DKFZP547	C1orf77,pp	1.43699	2.70E-16	1.43699	8863.676	12730.82
A_24_P29	NM_15234	Homo sapi	SLC43A2,L	1.43684	9.36E-07	1.43684	974.8975	1403.765
A_24_P17	NM_05295	WD40- anc	WDFY2,WI	1.43644	0.00001	1.43644	925.5568	1336.859
A_23_P20	NM_00448	Homo sapi	GALNT3,M	1.43626	1.52E-07	1.43626	116878.5	166862.2
A_23_P31	NM_01629	receptor as	RAP80,X2I	1.43606	1.56E-07	1.43606	8570.945	12361.04
A_23_P10	NM_00265	Homo sapi	PIK4CA,pi	1.43589	1.98E-14	1.43589	996.8931	1431.209
A_24_P204484		Unknown	A_24_P20	1.43565	7.07E-06	1.43565	19872.02	28365.79
A_23_P20	NM_00076	cytochrom	CYP1A2,C	1.43503	2.27E-14	1.43503	9344.882	13437.12
A_23_P75	NM_03281	Homo sapi	ATAD1,AF	1.43451	8.15E-18	1.43451	3875.333	5560.124
A_32_P37	BX649059	Growth arr	GAS2L3	1.43409	9.03E-07	1.43409	421.1285	604.053
A_23_P39	NM_17348	hypothetic	LOC12629	1.43398	1.48E-08	1.43398	511.1719	733.0353
A_23_P80	BC007518	Homo sapi	HIST1H3H	1.43304	0.00002	1.43304	1081.795	1542.418
A_24_P705136		Unknown	A_24_P70	1.4324	0.00002	1.4324	546.4038	783.4177
A_23_P26	NM_00684	Homo sapi	SDS,SDH	1.43185	1.10E-10	1.43185	1717.923	2451.355
A_23_P87	NM_00471	Homo sapi	SFRS2IP,S	1.43011	6.07E-10	1.43011	2749.073	3948.423
A_24_P39	NM_17753	Homo sapi	CYP20A1,(1.42958	0.00064	1.42958	690.9592	986.7764
A_32_P18	NM_00213	heterogene	HNRPA1,F	1.42856	4.54E-06	1.42856	15457.45	22004.53
A_24_P20	NM_02109	myotubular	MTMR3,ZF	1.42823	4.92E-06	1.42823	825.9457	1187.502
A_32_P194740		Unknown	A_32_P19	1.42784	0.00007	1.42784	452.9357	647.9154
A_23_P10	NM_03258	Ubiquitin s	USP32	1.42756	1.97E-09	1.42756	823.6206	1173.362
A_23_P13	NM_01560	DKFZP547	C1orf77,pp	1.42725	1.92E-17	1.42725	4405.207	6286.715
A_23_P27	AK027693	Myeloid-as	MYADM	1.42678	0.00008	1.42678	2441.939	3468.478
A_24_P34	AL137546	KIAA0826	KIAA0826	1.42668	1.54E-07	1.42668	705.2116	1009.348
A_24_P17	NM_02273	golgi phos	ACBD3,PA	1.4266	0.00011	1.4266	942.9408	1364.345
A_32_P17	NM_00018	hexokinase	HK2,HKII,F	1.42657	8.78E-11	1.42657	6354.304	9089.446
A_23_P83	BC029735	KIAA1102	KIAA1102	1.42639	3.26E-08	1.42639	1854.728	2650.173
A_24_P25	NM_00213	heterogene	HNRPA1,F	1.42616	1.29E-07	1.42616	16083.13	22848.7
A_24_P10	NM_00149	UDP-Gal:b	B4GALT1,(1.42608	4.87E-16	1.42608	24544.25	34984.02
A_23_P21	NM_01538	Homo sapi	PREI3,2C4	1.42604	1.27E-07	1.42604	2933.451	4193.048
A_23_P53	NM_05410	H-rev107-li	HRLP5	1.42578	5.65E-07	1.42578	8986.448	12739.87
A_23_P10	NM_00600	Homo sapi	ZBTB16,PI	1.42561	9.82E-08	1.42561	416.2531	593.3762
A_32_P17	BC010934	Zinc finger	ZBTB20	1.42514	0.00039	1.42514	1837.448	2645.179
A_23_P21	NM_00189	Homo sapi	CSNK1A1,	1.42502	4.17E-07	1.42502	7118.959	10220.73
A_24_P18	AK024095	Homo sapi	HIF3A,IPA	1.42479	1.78E-06	1.42479	734.7654	1042.416
A_32_P20	NM_00399	CDC-like k	CLK3	1.42444	2.99E-09	1.42444	1828.861	2611.953
A_23_P12	NM_03286	Homo sapi	C6orf93,FL	1.42438	9.01E-18	1.42438	7515.425	10704.57
A_23_P17	NM_01757	Homo sapi	KIAA1212	1.42369	0.00002	1.42369	2984.137	4299.582
A_23_P21	NM_00538	Natural kill	NKTR	1.42274	0.00008	1.42274	1769.164	2550.161
A_23_P40	NM_01232	Homo sapi	MAPRE1,E	1.42261	2.26E-09	1.42261	2521.296	3597.451

A_23_P34	NM_00587	Homo sapi	BCAS2,DA	1.42261	2.62E-08	1.42261	5847.892	8294.041
A_24_P32	NM_01774	Homo sapi	ANKHD1,M	1.42202	0.00015	1.42202	5810.35	8379.903
A_24_P16	BC036263	Hypothetic	MGC9515	1.42182	1.31E-10	1.42182	44264.05	62775.53
A_23_P54	NM_01664	membrane	MIR16,363	1.42083	0.00028	1.42083	1163.484	1678.615
A_23_P97	NM_02489	Homo sapi	PAQR6,FL	1.4205	2.85E-11	1.4205	1575.446	2241.899
A_23_P12	NM_00339	Homo sapi	WEE1,WEI	1.42044	5.28E-06	1.42044	554.9547	790.1343
A_23_P38	U09848	Zinc finger	ZKSCAN1	1.42016	0.00005	1.42016	1097.373	1570.875
A_23_P21	NM_03217	Homo sapi	RNUXA,PT	1.41991	5.09E-16	1.41991	4934.945	7004.238
A_23_P30	NM_01331	Homo sapi	CNOT4,NC	1.41948	3.56E-07	1.41948	1618.234	2312.261
A_24_P18	NM_01469	KIAA0296	ZNF646,KI	1.41894	2.23E-07	1.41894	1767.659	2518.396
A_24_P366566		Unknown	ENST0000	1.4189	2.65E-06	1.4189	10972.31	15569.04
A_23_P11	AK023859	RNA bindir	RBM16	1.41884	1.47E-13	1.41884	842.9122	1196.031
A_23_P20	NM_00248	Homo sapi	PPP1R12A	1.41881	1.62E-06	1.41881	422.77	599.8208
A_23_P20	NM_00343	Homo sapi	ZNF137,pt	1.41793	5.68E-09	1.41793	782.5254	1109.038
A_23_P20	NM_00596	Homo sapi	NAB1	1.41781	1.57E-10	1.41781	1100.672	1562.588
A_24_P56	NM_00153	hypoxia-inc	HIF1A,MOI	1.41765	3.47E-10	1.41765	8769.674	12483.14
A_24_P214556		Unknown	THC19937	1.41756	2.25E-06	1.41756	523.0999	741.4498
A_24_P33	NM_17591	Homo sapi	ZNF493,FL	1.41702	1.59E-07	1.41702	995.7133	1418.208
A_23_P20	NM_00505	retinoblast	JARID1A,F	1.41612	0.00003	1.41612	850.5708	1211.962
A_23_P20	NM_01231	Homo sapi	KIN,BTCD,	1.41557	0.00004	1.41557	1014.286	1446.658
A_23_P70	NM_01515	Homo sapi	PHF3,KIAA	1.41528	1.52E-06	1.41528	1586.528	2253.874
A_23_P85	NM_00508	Homo sapi	U2AF1L2,L	1.41508	7.65E-08	1.41508	2652.636	3752.153
A_32_P12	AF316855	Hypothetic	FLJ22795	1.4145	2.18E-12	1.4145	6032.841	8551.468
A_23_P50195		Unknown	A_23_P50	1.4144	0.0003	1.4144	512.0986	725.4004
A_24_P29	NM_01631	REV1 (yea	REV1L,RE	1.41377	0.00012	1.41377	492.3836	697.5136
A_23_P20	NM_00250	Homo sapi	NFKB2,LY	1.41355	4.59E-06	1.41355	1934.871	2713.663
A_23_P66	NM_15333	hypothetic	FLJ90165,I	1.41338	2.86E-06	1.41338	163714.9	230069.8
A_24_P57	NM_03260	CREB/ATF	CREB3L3,I	1.41279	0.00003	1.41279	1697.242	2378.094
A_23_P37	BC035647	M4-50 mRI	BC035647	1.41224	6.60E-11	1.41224	7955.556	11223.88
A_24_P27	NM_01906	chromoson	EML4,C20	1.41223	2.65E-07	1.41223	8409.485	11942.86
A_23_P10	NM_01776	Homo sapi	PNRC2,FL	1.41172	7.06E-15	1.41172	4049.767	5714.919
A_23_P74	NM_02457	Homo sapi	C1orf54,FL	1.41168	0.00051	1.41168	510.3438	721.9799
A_23_P21	AK001750	Hypothetic	FLJ11046	1.41158	1.51E-13	1.41158	1047.187	1478.485
A_32_P47	NM_14504	ubiquitin-lik	UBLCP1,C	1.41146	2.35E-10	1.41146	868.6373	1227.18
A_32_P91	NM_17852	hypothetic	FLJ38792	1.41144	6.88E-09	1.41144	2034.838	2862.015
A_32_P153361		Unknown	A_32_P15	1.41137	9.05E-07	1.41137	1261.721	1785.906
A_23_P90	NM_15227	Homo sapi	ZNF585B,S	1.41094	1.51E-06	1.41094	415.1456	585.7483
A_23_P16	BC008889	Zinc finger	ZNF547	1.41059	0.00006	1.41059	2170.813	3094.925
A_23_P25	NM_00445	Homo sapi	EZH2,EZH	1.41055	3.84E-06	1.41055	10134.21	14405.44
A_23_P16	AB020670	KIAA0863	KIAA0863	1.41054	0.00061	1.41054	1030.558	1465.35
A_24_P68	AB002449	Hypothetic	AB002449	1.41053	0.00077	1.41053	787.1814	1118.852
A_32_P18	NM_17506	Homo sapi	WBSCR19	1.41016	0.00044	1.41016	310.9364	438.5703
A_23_P13	NM_00626	Homo sapi	RANBP2,T	1.40966	0.00016	1.40966	6578.062	9426.958
A_23_P18	NM_01326	peroxisom	PPARGC1	1.40963	0.00001	1.40963	802.0216	1135.277
A_23_P20	NM_00008	collagen, ty	COL1A1,O	1.40939	9.48E-06	1.40939	987.1015	1388.292
A_24_P713312		ALU8_HU	THC18707	1.40907	0.00057	1.40907	363.8173	512.8247
A_23_P20	NM_00371	Homo sapi	PPAP2B,V	1.40739	0.00002	1.40739	2648.183	3729.383
A_24_P12	NM_00254	oxidised lo	OLR1,LOX	1.40731	0.00034	1.40731	496.4467	698.7324
A_23_P14	NM_02187	Homo sapi	FGG	1.40713	0.00015	1.40713	271.1144	381.4043
A_23_P12	NM_00663	methylene	MTHFD2,N	1.4071	2.16E-06	1.4071	30098.21	42120.07
A_24_P23	NM_05296	Homo sapi	C1orf24,NI	1.40706	6.12E-06	1.40706	1497.315	2122.953
A_23_P53	NM_00187	Homo sapi	CPS1	1.407	3.85E-21	1.407	12724.5	17911.08

A_24_P17:	NM_02484	hypothetic	ZNF613,FL	1.4069	0.00011	1.4069	397.1838	558.9307
A_24_P67:	NM_01867	Homo sapi	ABCA5,AB	1.40665	0.0002	1.40665	440.9603	620.9204
A_23_P20:	NM_01531	Homo sapi	PPP1R13E	1.40651	0.00009	1.40651	325.8846	458.5595
A_23_P67:	NM_00381	Homo sapi	TNFSF9,4-	1.40641	1.35E-12	1.40641	1278.999	1799.644
A_23_P39:	NM_00145	Homo sapi	FOXC1,AR	1.40592	5.48E-16	1.40592	17513.29	24622.97
A_23_P13:	NM_01517	Homo sapi	FBXO28,FI	1.4059	3.47E-15	1.4059	2695.271	3789.398
A_24_P23:	NM_17792	Homo sapi	H2AFJ,MG	1.40551	6.26E-11	1.40551	39067.73	54986.39
A_23_P87:	NM_00579	Homo sapi	RFP2,CAR	1.40541	0.00002	1.40541	4840.228	6806.043
A_23_P16:	NM_02021	Homo sapi	PARP6	1.40536	1.04E-12	1.40536	1226.274	1724.395
A_32_P18:	NM_15263	Homo sapi	METT5D1,	1.40486	1.46E-10	1.40486	593.6514	834.0768
A_23_P4462	Unknown	A_23_P44		1.4045	2.80E-07	1.4045	776.5834	1093.876
A_32_P50463	Unknown	A_32_P50		1.40443	0.00004	1.40443	1806.105	2538.384
A_32_P16:	NM_14715	Homo sapi	TMEM23,M	1.40398	1.51E-09	1.40398	1349.17	1890.046
A_23_P91:	NM_00306	Homo sapi	SLPI,ALP,I	1.40373	5.06E-12	1.40373	44114.44	62049
A_32_P75867	ALU1_HUM	THC18592		1.40371	0.00017	1.40371	1095.338	1544.152
A_32_P220700	Unknown	THC19819		1.4035	3.51E-08	1.4035	989.913	1390.394
A_24_P32:	NM_00551	Major histo	HLA-E	1.40349	1.68E-10	1.40349	22670.02	31831.91
A_24_P11:	NM_00551	Major histo	HLA-B	1.40332	1.41E-11	1.40332	15558.46	21822.1
A_23_P12:	NM_03287	Homo sapi	C6orf111,I	1.40304	6.36E-08	1.40304	737.6051	1037.212
A_24_P58:	XM_06202	similar to	HLOC12036	1.40303	8.11E-10	1.40303	8371.898	11714.99
A_23_P45:	NM_00212	Homo sapi	HLA-DRB5	1.40271	1.04E-07	1.40271	1348.477	1881.79
A_23_P48637	Unknown	A_23_P48		1.40251	2.86E-22	1.40251	3219.464	4522.882
A_32_P11:	NM_02468	Coiled-coil	CCDC7	1.40249	0.00002	1.40249	552.6179	775.2434
A_24_P15:	NM_00204	glycyl-tRN	GARS,CM	1.4021	2.05E-10	1.4021	5502.799	7738.62
A_24_P37:	NM_14521	Homo sapi	TRIM11,BI	1.40153	3.53E-08	1.40153	10407.53	14591.87
A_23_P10:	NM_00031	Homo sapi	PRNP,CJD	1.40137	2.06E-10	1.40137	14502.15	20399.71
A_24_P14:	NM_00673	lymphoid b	AKAP13,BI	1.40092	0.00002	1.40092	2932.832	4112.155
A_23_P11:	NM_00244	Homo sapi	MSX1,HO	1.4006	0.00015	1.4006	463.454	650.066
A_23_P10:	NM_01432	origin reco	ORC6L,OF	1.4005	5.04E-12	1.4005	84419.63	117994.9
A_23_P20:	AK000796	Chromosor	C14orf129	1.40029	0.0001	1.40029	5664.474	8014.762
A_23_P41:	NM_03096	Homo sapi	SBF2,CMT	1.40021	0.00081	1.40021	335.8696	469.8428
A_23_P32:	AF116619	Homo sapi	PRO1051	1.40005	1.37E-07	1.40005	2061.939	2881.596
A_24_P94:	AB002331	Death assc	DATF1	1.39996	0.00042	1.39996	379.6958	531.6394
A_23_P18:	NM_15269	Homo sapi	SEN5,MC	1.39953	9.11E-13	1.39953	926.0748	1295.985
A_24_P33:	NM_00313	Serum res	SRF	1.3994	0.00026	1.3994	285.125	399.2463
A_32_P162095	Unknown	THC18384		1.39902	6.28E-08	1.39902	606.8649	849.933
A_23_P37:	NM_01909	GTP bindin	GTPBP2,IV	1.39836	2.20E-08	1.39836	6396.755	8919.783
A_23_P35:	AB051523	Hypothetic	FLJ20294	1.39832	4.90E-09	1.39832	1653.914	2306.534
A_23_P34:	NM_02031	Homo sapi	MNT,ROX,	1.39813	1.46E-06	1.39813	6115.407	8528.014
A_24_P19:	NM_00306	secretory k	SLPI,ALP,I	1.39803	1.71E-13	1.39803	7230.891	10123.27
A_32_P21:	NM_01435	TNF-induc	TNFAIP8,C	1.39802	4.27E-06	1.39802	1763.484	2471.212
A_23_P338113	DJ377H14	ENST0000		1.39793	6.80E-10	1.39793	6412.752	8960.027
A_23_P31:	NM_01533	Zinc finger,	ZDHHC17	1.39774	0.00064	1.39774	1533.96	2176.541
A_23_P45:	NM_00329	Homo sapi	TPR	1.39732	7.06E-08	1.39732	6844.853	9536.949
A_23_P31:	NM_02400	DEAD/H (A	DDX3X,DE	1.39727	7.42E-09	1.39727	3973.636	5578.327
A_24_P23:	NM_00463	RAB7, mer	RAB7,PSN	1.39726	0.00001	1.39726	3551.496	4966.782
A_23_P70:	NM_00127	Homo sapi	CHD1	1.39677	2.14E-06	1.39677	1457.856	2048.641
A_24_P29:	NM_00346	protein tyrc	PTP4A1,HI	1.39674	2.85E-07	1.39674	12505.81	17550.71
A_23_P30:	AL137514	Inositol hex	IHPK2	1.39653	2.76E-07	1.39653	5989.05	8329.723
A_32_P20:	C18665	C18665 H	C18665	1.39632	3.31E-10	1.39632	1226.903	1713.839
A_23_P56:	BC042674	Phospholip	PLB1	1.3962	0.00003	1.3962	760.8084	1057.37
A_23_P13:	NM_01481	Homo sapi	PJA2,RNF	1.39619	0.00014	1.39619	1657.852	2334.739

A_23_P20	NM_01637	AT rich inte	ARID4B	1.39584	0.00001	1.39584	1210.036	1687.415
A_32_P95	NM_00221	Homo sapi	ITGB1,CD3	1.39577	1.05E-09	1.39577	28422.05	39821.95
A_23_P38	NM_02275	Homo sapi	CCDC14,F	1.39569	0.00001	1.39569	4398.527	6145.955
A_24_P32	AK092090	Homo sapi	AK092090	1.39478	6.91E-10	1.39478	3991.561	5563.352
A_24_P25	NM_01219	rab6 GTPa	RABGAP1	1.39468	0.00005	1.39468	769.9884	1079.995
A_32_P23	NM_01422	Homo sapi	TM4SF1,LI	1.39449	5.99E-08	1.39449	49782.71	69125.48
A_23_P45	NM_00647	myosin reg	MRCL3,ML	1.39435	5.59E-06	1.39435	29999.28	42225.3
A_24_P72	NM_13344	Homo sapi	CTGLF1,M	1.39408	0.00002	1.39408	615.9034	860.3867
A_23_P14	NM_00718	WW domai	WBP4,FBF	1.39367	2.05E-06	1.39367	2097.892	2941.327
A_23_P21	NM_03256	esophagus	ECG2,ECF	1.39345	1.76E-09	1.39345	9033.841	12600.95
A_32_P16	NM_19900	Homo sapi	ZNF322B,Z	1.39324	0.00052	1.39324	464.8777	648.9227
A_24_P30	NM_01497	KIAA1049	KIAA1049,	1.39289	9.84E-08	1.39289	5725.343	8017.843
A_23_P12	NM_02508	hypothetic	FLJ22795	1.39286	5.88E-07	1.39286	6977.842	9758.893
A_24_P10	NM_01527	ubiquitin-pr	NEDD4L,R	1.39286	9.07E-13	1.39286	2724.875	3795.614
A_23_P12	NM_01868	Homo sapi	ZNF313,RI	1.39281	6.44E-06	1.39281	4989.522	6944.458
A_32_P25	NM_20747	FLJ42953	FLJ42953	1.3921	6.23E-08	1.3921	711.1166	988.9185
A_23_P63	NM_03075	nuclear rec	NRBF2,CC	1.39195	7.56E-08	1.39195	689.5692	961.0239
A_23_P20	NM_01601	CGI-62 pro	C8orf70,CC	1.39189	2.92E-08	1.39189	1751.34	2449.481
A_23_P70	NM_01883	Homo sapi	MATR3,MC	1.39134	7.75E-15	1.39134	6671.004	9285.085
A_23_P35	NM_19933	Homo sapi	FLJ35171	1.39089	0.0001	1.39089	411.3978	572.2192
A_23_P89	NM_01325	zinc finger	ZNF180,HI	1.39074	0.00012	1.39074	838.3694	1170.087
A_23_P25	NM_00615	Homo sapi	NCK1,NCK	1.39045	0.00002	1.39045	3143.143	4405.412
A_24_P15	AF218008	LOC44050	unknown	1.39027	1.48E-11	1.39027	1313.409	1824.56
A_24_P14	NM_01472	Ste20-relat	SLK,STK2,	1.39019	0.0001	1.39019	3059.91	4252.843
A_24_P15	NM_02093	Homo sapi	USP37,KIA	1.38995	0.00003	1.38995	709.7363	991.2847
A_23_P15	BC012609	Serine (or	SERPINB2	1.38983	6.04E-10	1.38983	2910.798	4069.084
A_23_P31	NM_02472	hypothetic	FLJ23518	1.38961	1.07E-07	1.38961	707.8658	985.1234
A_23_P40	NM_14525	LOC12440	LOC12440	1.38929	0.00028	1.38929	1371.595	1929.33
A_24_P92	AY358690	Eukaryotic	EEF1D	1.38791	0.00007	1.38791	526.5939	732.106
A_24_P135391	Unknown	A_24_P13		1.38748	9.60E-08	1.38748	9559.842	13224.23
A_23_P256487	Unknown	THC18188		1.38745	0.00002	1.38745	3102.661	4272.411
A_23_P57	NM_00577	Homo sapi	RBM6,3G2	1.3874	0.00016	1.3874	1286.134	1798.51
A_23_P22	NM_00704	HIV-1 rev t	HRB2,RIP-	1.38711	0.00027	1.38711	6451.26	9105.935
A_23_P41	NM_14529	similar to	RLOC19733	1.38699	9.19E-06	1.38699	4127.831	5752.094
A_23_P71	NM_02452	hypothetic	GCC1,GC	1.38678	2.06E-09	1.38678	1033.646	1431.9
A_32_P159939	Unknown	THC19593		1.38668	0.00044	1.38668	679.8972	942.8121
A_23_P20	NM_00644	Homo sapi	CDC42EP	1.3864	6.82E-11	1.3864	668.6175	927.0298
A_23_P11	BC050088	Similar to	CLOC13050	1.38624	9.37E-07	1.38624	3703.5	5154.573
A_23_P12	M26004	Compleme	CR2	1.38564	1.07E-07	1.38564	5357.079	7414.277
A_24_P35	BC047452	Hypothetic	LOC15838	1.38555	0.00053	1.38555	2002.676	2797.122
A_32_P46510	Unknown	A_32_P46		1.38547	0.00089	1.38547	495.7565	689.1424
A_23_P49	NM_00042	Homo sapi	KRT10,K1	1.38544	6.19E-11	1.38544	61135.75	84596.67
A_23_P20	NM_02289	Homo sapi	PAPOLG,F	1.38511	0.00058	1.38511	1628.972	2258.22
A_24_P26	NM_01865	myoneurin	MYNN,OS	1.38502	8.99E-08	1.38502	789.8453	1096.351
A_23_P16	NM_01792	hypothetic	PRPF39,FI	1.38485	0.00002	1.38485	935.458	1301.488
A_23_P36	NM_13837	hypothetic	MGC1604	1.38463	1.29E-07	1.38463	2018.532	2824.828
A_32_P21	NM_03115	heterogene	HNRPA1,F	1.38457	1.05E-11	1.38457	32676.17	45181.34
A_24_P92	BC039151	Chromosor	C20orf119	1.38429	0.00056	1.38429	622.6896	862.8593
A_23_P39	NM_13347	Homo sapi	ZNF431,KI	1.38419	0.00012	1.38419	619.5138	857.8638
A_23_P41	NM_17382	p18 splicin	P18SRP,FI	1.38416	8.92E-11	1.38416	5774.883	8015.102
A_24_P32	AK126454	Heterogen	LOC14498	1.38393	4.35E-11	1.38393	14316.53	19775.38
A_24_P28	AL050020	DKFZP564	DKFZP564	1.38374	1.48E-06	1.38374	10512.97	14429.19

A_23_P11(NM_02442 Homo sapi PLA2G4A,	1.38349	5.18E-10	1.38349	5716.121	7956.22
A_23_P30(NM_01814 centrosom: Cep72,FLJ	1.38341	6.66E-10	1.38341	2016.979	2789.03
A_24_P238186 Unknown A_24_P23	1.38339	5.16E-13	1.38339	3647.427	5048.844
A_23_P16(NM_15279 Homo sapi ZNF555,M	1.38332	0.00026	1.38332	1264.768	1739.769
A_32_P12(NM_19827 Homo sapi NHS,SCMI	1.38305	0.00002	1.38305	1430.744	1982.924
A_24_P38(BC010266 Similar to HNRPA1,H	1.38304	2.53E-14	1.38304	7640.855	10571.43
A_32_P22(NM_17490 Homo sapi PPP4R2	1.38297	0.00077	1.38297	636.4789	883.144
A_32_P169179 MSX2_HUI/THC19571	1.38248	1.60E-06	1.38248	1283.154	1778.207
A_23_P21(NM_02449 Homo sapi ZNF117,HI	1.38239	2.81E-08	1.38239	1493.759	2066.427
A_23_P56(NM_01847 likely ortho LEREPO4,	1.38217	3.57E-07	1.38217	15899.81	22033.49
A_24_P34(NM_03249 zinc finger ZC3H8,Fliz	1.38201	0.00013	1.38201	290.5709	401.3912
A_23_P14(NM_01859 chondroitin GALNACT	1.38175	0.00006	1.38175	1913.219	2622.608
A_24_P39(NM_00213 heterogene HNRPA2B	1.38142	1.40E-09	1.38142	23246.09	32033.21
A_24_P17(NM_00329 translocate TPR	1.38133	2.22E-07	1.38133	7096.911	9846.266
A_24_P10(NM_00673 DnaJ (Hsp DNAJB2,H	1.38091	1.04E-10	1.38091	865.8024	1195.056
A_23_P15(AF338192 Homo sapi FKSG63	1.38057	3.10E-07	1.38057	10565.01	14651.5
A_23_P42(NM_02525 Homo sapi C6orf27,G	1.38051	0.00005	1.38051	10918.36	14942.21
A_24_P332953 Unknown A_24_P33	1.38011	3.96E-07	1.38011	3710.801	5137.422
A_24_P61(NM_20691 Chromosom C6orf216	1.37972	5.33E-11	1.37972	2106.094	2903.393
A_23_P17(NM_01904 Homo sapi WDR44,RF	1.37971	6.36E-07	1.37971	3082.958	4261.67
A_24_P11(NM_00226 karyopherin KPNA4,QII	1.37931	7.56E-08	1.37931	7278.607	10073.61
A_23_P35(NM_03223 FLJ10378 LARP2,MC	1.37915	2.24E-10	1.37915	2338.917	3232.512
A_32_P45(BC002581 Similar to C DKFZP434	1.37903	1.09E-08	1.37903	15481.45	21306.79
A_32_P159787 AJ007829 THC19433	1.37844	0.00016	1.37844	4380.827	5987.162
A_23_P94(AK098175 Zinc finger ZNF283	1.37844	3.83E-07	1.37844	792.5899	1093.587
A_23_P45(NM_00425 Homo sapi RAB9A,RA	1.37824	2.10E-07	1.37824	6031.237	8363.217
A_23_P65(NM_00711 Homo sapi TEP1,TP1,	1.37818	0.00002	1.37818	854.188	1177.372
A_24_P27(NM_01238 p8 protein P8,COM1	1.37806	0.0001	1.37806	3723.541	5160.291
A_23_P14(NM_00432 Homo sapi BAG1	1.378	5.48E-09	1.378	29828.61	41239.23
A_23_P75(NM_00362 Homo sapi PPFIA1,LIF	1.37796	2.06E-06	1.37796	1846.422	2535.179
A_24_P10(AK001406 SUMO1/se SENP6	1.37753	0.00016	1.37753	306.7997	422.8604
A_24_P70(AK023907 Phosphata PHACTR4	1.37753	0.00005	1.37753	319.0272	439.5424
A_23_P99(NM_00329 Homo sapi TPP2	1.3774	9.19E-10	1.3774	2729.112	3741.207
A_23_P20(NM_00470 Homo sapi ATG12,AP	1.37717	6.00E-07	1.37717	4864.447	6742.937
A_23_P33(NM_18249 hypothetic ZFAND2A	1.3767	0.00052	1.3767	2361.287	3292.89
A_23_P13(NM_01212 cysteine ar CHORDC1	1.37606	6.38E-08	1.37606	5224.061	7213.946
A_24_P84(AK054826 Homo sapi AK054826	1.37599	2.45E-09	1.37599	3260.258	4499.374
A_23_P56(NM_02053 Homo sapi RTN4,ASY	1.37596	9.38E-13	1.37596	12838.74	17679.71
A_32_P96(BX537597 Clone TES BX537597	1.37557	7.90E-06	1.37557	1871.262	2581.727
A_23_P39(NM_00346 Homo sapi FZD5,HFZ	1.37548	2.43E-07	1.37548	1875.719	2582.702
A_23_P40(NM_13833 Homo sapi C9orf42,M	1.3752	6.98E-09	1.3752	1543.46	2121.704
A_23_P11(NM_01826 Sec3-like SEC3L1,SI	1.37498	0.00073	1.37498	4246.081	5910.427
A_23_P25(NM_00048 Homo sapi ARSA,MLC	1.37477	8.88E-10	1.37477	4789.313	6574.283
A_23_P59(D31887 Solute carr SLC39A14	1.37454	0.00033	1.37454	3897.212	5411.315
A_23_P52(NM_00428 BCL2-asso BAG3,BIS,	1.37437	2.51E-07	1.37437	9153.611	12633.21
A_32_P33802 Unknown A_32_P33	1.37405	1.80E-07	1.37405	944.8089	1295.001
A_23_P20(NM_01779 Homo sapi C14orf101,	1.37405	0.00003	1.37405	1047.203	1448.889
A_23_P11(NM_01447 Homo sapi PDLIM3,AL	1.37377	0.00014	1.37377	530.2836	730.2549
A_23_P25(NM_02511 Centrosom Cep290	1.37309	0.00099	1.37309	1373.844	1920.978
A_24_P122921 BIM_HUM/THC18776	1.37305	8.39E-09	1.37305	1833.662	2523.499
A_23_P25(NM_02454 Homo sapi C13orf7,FL	1.37292	0.00002	1.37292	4105.153	5651.232
A_24_P35(NM_00626 RAN bindir RANBP2,T	1.37205	0.0005	1.37205	1319.669	1828.979

A_23_P15	NM_00212	Homo sapi	HMGB2, HM	1.37198	1.08E-11	1.37198	13141.35	18013.25	
A_32_P39	NM_00678	SMA3	SMA3, b55	1.37195	3.87E-09	1.37195	551.5845	756.7777	
A_23_P16	NM_02182	hypothetic	PPCDC, MI	1.37163	1.74E-06	1.37163	1330.489	1834.768	
A_24_P20	NM_00333	ubiquitin-c	UBE2B, HR	1.37158	1.61E-09	1.37158	4181.83	5752.316	
A_23_P59	NM_00619	Homo sapi	PAX4	1.37072	8.37E-09	1.37072	7336.268	10093.29	
A_24_P94	X59739	Zinc finger	ZFX	1.37021	0.00001	1.37021	641.3295	881.2877	
A_24_P24	NM_18153	Homo sapi	BTN2A2, B	1.37012	0.00011	1.37012	2226.026	3030.436	
A_23_P21	NM_13343	Homo sapi	NIPBL, CDI	1.36976	0.0003	1.36976	1340.146	1845.913	
A_24_P94	AB002344	Jumonji do	JMJD3	1.36975	2.70E-06	1.36975	3075.085	4191.776	
A_23_P33	NM_01631	Homo sapi	GULP1, CE	1.36933	1.82E-08	1.36933	2415.584	3320.368	
A_24_P35	NM_03058	UDP-Gal:b	B4GALT2	1.36873	1.62E-07	1.36873	6912.531	9451.557	
A_23_P33	BC009539	Chromosor	C14orf46	1.36841	5.63E-08	1.36841	426.6201	583.7873	
A_24_P65	NM_15252	hypothetic	FLJ40432	1.36829	2.24E-06	1.36829	2269.508	3119.105	
A_24_P10	NM_01628	STE20-like	TAOK3, DP	1.36799	1.05E-06	1.36799	4007.497	5507.541	
A_24_P30	7075	Unknown	A_24_P30	1.36758	0.00002	1.36758	6656.408	9051.725	
A_23_P52	NM_01543	DKFZP434	C1orf73, D	1.36714	4.57E-09	1.36714	1390.572	1904.178	
A_23_P14	NM_00540	Homo sapi	PTPN14, P	1.36707	0.00023	1.36707	394.6888	539.6357	
A_32_P14	5790	DDRT heli	THC19773	1.36687	3.42E-09	1.36687	72343.57	98674.16	
A_23_P36	NM_15235	hypothetic	MGC40157	1.36603	0.00002	1.36603	96691.89	131176.8	
A_23_P76	NM_02117	Homo sapi	CCNB1IP1	1.36593	1.60E-09	1.36593	5752.209	7846.103	
A_32_P9	9715	AJ004940	THC18434	1.36494	0.00002	1.36494	2209.282	3012.42	
A_32_P6	2	AK023827	Homo sapi	BANP, SM	1.36477	0.00077	1.36477	307.2928	419.752
A_24_P9	4	NM_03308	EAF1 prote	EAF1	1.36454	5.41E-06	1.36454	1443.217	1974.74
A_23_P8	3	AB051895	Epiplakin 1	EPPK1	1.36454	0.00004	1.36454	6112.162	8259.104
A_23_P2	5	NM_01488	Homo sapi	ANAPC10,	1.36451	5.44E-07	1.36451	2276.954	3129.571
A_24_P5	1	NM_00213	heterogene	HNRPA1, F	1.36418	0.00014	1.36418	8575.653	11618.55
A_24_P9	1	X74861	H.sapiens	IHPX-2	1.36383	4.61E-06	1.36383	1107.915	1508.511
A_23_P1	4	NM_01764	Homo sapi	FAM29A, R	1.36362	1.16E-06	1.36362	916.5355	1253.775
A_24_P4	1	1890	Glucose 6-	ENST0000	1.36359	3.88E-06	1.36359	7370.195	10011.38
A_23_P2	0	NM_01462	Homo sapi	S100A6, 2A	1.36345	1.42E-08	1.36345	248393	340406.7
A_23_P1	0	NM_01467	Homo sapi	PUM1, PUM	1.36279	8.74E-07	1.36279	6497.901	8894.816
A_23_P1	8	NM_00470	Homo sapi	ATG12, AP	1.36263	2.14E-09	1.36263	2397.187	3261.359
A_24_P4	0	9521	Unknown	ENST0000	1.36261	2.51E-11	1.36261	6454.733	8789.164
A_24_P3	0	84239	Unknown	A_24_P3	1.36252	0.00006	1.36252	9452.672	13009.89
A_23_P9	7	NM_00469	U4/U6-ass	PRPF3, PR	1.36227	0.00004	1.36227	1099.054	1503.72
A_24_P4	0	NM_18258	hypothetic	DKFZp451	1.36219	3.35E-06	1.36219	381.8817	520.1793
A_32_P1	1	NM_00213	heterogene	HNRPA1, F	1.36215	0.00005	1.36215	8440.416	11429.29
A_32_P6	2	434	Unknown	A_32_P6	1.36167	0.00098	1.36167	1017.622	1377.05
A_23_P1	1	BX648200	DKFZP434	DKFZP434	1.36161	0.00011	1.36161	13422.57	18157.15
A_24_P4	9	3116	Unknown	A_24_P4	1.36158	1.56E-08	1.36158	6127.972	8370.508
A_23_P9	9	NM_00399	Homo sapi	CLK3	1.36158	5.10E-06	1.36158	584.181	795.0847
A_23_P8	3	NM_01661	Homo sapi	TTRAP, EA	1.36144	2.75E-11	1.36144	16229.67	22118.75
A_24_P9	1	BC005305	Collagen, t	COL4A6	1.36105	0.00007	1.36105	947.0764	1287.955
A_23_P2	5	NM_02456	hypothetic	NARG1L, M	1.36011	0.00002	1.36011	867.0472	1184.975
A_32_P1	0	NM_17347	hypothetic	DCUN1D3	1.35972	4.11E-06	1.35972	1776.895	2414.751
A_32_P7	6	060	Unknown	THC18807	1.35913	3.03E-07	1.35913	1818.888	2480.583
A_24_P7	1	AK025082	Staphyloco	SND1	1.3578	0.00087	1.3578	422.8655	574.1857
A_23_P1	1	NM_00215	Homo sapi	HSPA6	1.35779	0.00043	1.35779	395.4421	537.0017
A_24_P2	6	AK057480	Heterogene	HNRPDL	1.35779	0.00024	1.35779	772.4182	1054.537
A_23_P1	7	NM_01467	pumilio (Dr	PUM1, PUM	1.35763	7.08E-08	1.35763	5882.751	8013.9
A_32_P1	3	AB014563	KIAA0663	KIAA0663	1.35715	7.20E-10	1.35715	2999.4	4078.2
A_24_P3	4	9896	Unknown	A_24_P3	1.35711	1.84E-07	1.35711	6347.796	8589.571

A_24_P28	AB033038 KIAA1212 KIAA1212	1.35641	4.45E-08	1.35641	508.3896	689.6057
A_32_P46	X56841 Major histo HLA-E	1.35641	4.04E-08	1.35641	20390.47	27631.41
A_23_P12	NM_01787 interphase ICF45,FLJ	1.35625	3.37E-09	1.35625	2513.343	3416.823
A_23_P36	BC009220 LOC44116 C6orf160,M	1.35495	9.21E-14	1.35495	76482.11	103629.3
A_24_P605190	PGT1_HUI THC19577	1.35493	1.82E-06	1.35493	1082.615	1467.279
A_24_P12	NM_02075 Homo sapi SERINC1,	1.3544	5.97E-09	1.3544	2311.396	3140.467
A_23_P59	NM_01648 Homo sapi NT5C3,PN	1.35431	0.00001	1.35431	8468.088	11516.6
A_23_P71	(NM_02452 hypothetical GCC1,GCC	1.35408	6.20E-12	1.35408	3130.658	4236.613
A_24_P748389	Unknown THC19829	1.35389	0.00001	1.35389	323.2592	437.645
A_23_P57	NM_13893 Homo sapi BCL6,BCL	1.35375	0.00096	1.35375	1141.832	1559.879
A_23_P14	(NM_00366 Homo sapi INE1,MGC	1.35343	5.51E-13	1.35343	36198.26	48978.81
A_23_P14	(NM_02481 Homo sapi CBLL1,HA	1.35332	2.70E-08	1.35332	885.1266	1197.417
A_23_P12	NM_00620 Phosphodi PDE4D	1.35321	0.00002	1.35321	825.1209	1116.2
A_24_P25	(NM_01631 REV1 (yea REV1L,RE	1.35229	0.00033	1.35229	462.6395	626.2924
A_23_P13	BC006795 Hypothetic MGC5370	1.35201	0.00006	1.35201	743.463	1009.645
A_24_P93	AK125595 DEAD (As DDX56	1.35188	4.10E-08	1.35188	16434.1	22151.37
A_23_P37	NM_00692 Homo sapi SFRS5,HR	1.35132	0.00042	1.35132	16579.25	22226.18
A_23_P20	(NM_01221 integrin, al ITGA11,Hs	1.35122	6.02E-09	1.35122	1334.007	1804.097
A_23_P20	AF283776 Activating t ATF2	1.35081	1.51E-11	1.35081	1789.205	2415.54
A_24_P10	BC001663 Ribosomal RPL31	1.35026	2.34E-08	1.35026	1695.387	2296.59
A_24_P32	(NM_00059 insulin-like IGFBP3,IB	1.34959	2.20E-10	1.34959	1315.945	1774.507
A_23_P54	BC051317 Homo sapi RBBP6	1.34939	0.00003	1.34939	455.5273	614.5534
A_32_P87	U79293 Clone 239 U79293	1.34922	0.00002	1.34922	741.1616	998.3807
A_23_P97	AK001865 ERO1-like ERO1LB	1.34911	0.00002	1.34911	610.5301	822.8463
A_32_P96933	Unknown A_32_P96	1.34886	1.95E-10	1.34886	881.3414	1188.704
A_24_P92	AF321778 Toll interac TOLLIP	1.34881	1.13E-10	1.34881	5771.062	7774.777
A_23_P97	NM_01559 Homo sapi GPATC4	1.34859	7.09E-10	1.34859	13140.22	17746.21
A_23_P94	(NM_01620 Homo sapi LSM8,YJR	1.34848	0.00003	1.34848	15250.12	20762.23
A_23_P37	(NM_14529 Homo sapi ZNF626	1.34842	2.95E-07	1.34842	1864.713	2518.029
A_24_P29	(NM_15336 Homo sapi ZNF679,M	1.34826	1.31E-09	1.34826	3296.692	4446.418
A_24_P94	NM_00391 serine/thre PRPF4B,P	1.34815	0.00008	1.34815	1701.279	2310.167
A_24_P69	BX640887 Hypothetic RNU71B,U	1.34801	0.00039	1.34801	3444.969	4675.173
A_24_P94	BC017179 Hypothetic FLJ35867	1.34769	0.00009	1.34769	590.9092	798.3765
A_23_P15	(NM_03233 hypothetical C8orf53,M	1.3474	0.00001	1.3474	2747.485	3725.608
A_24_P70	(AL832246 Heterogen HNRPA3	1.34705	9.88E-06	1.34705	3208.459	4354.582
A_23_P83	(NM_03279 reserved SYAP1,PR	1.34619	6.99E-06	1.34619	11677.22	15810.72
A_23_P38	(NM_00194 Homo sapi E2F3,E2F-	1.3457	0.00013	1.3457	5979.243	8064.443
A_24_P63	(NM_01620 U6 snRNA LSM8,YJR	1.34568	3.14E-09	1.34568	14951.86	20017.36
A_23_P11	M15530 Human B-c BCGF1	1.34568	1.26E-12	1.34568	58927.23	79278.59
A_24_P161655	Unknown ENST0000	1.34546	3.03E-08	1.34546	12234.34	16493.54
A_24_P12	(NM_01241 rab3 GTPa RAB3GAP	1.34506	1.08E-07	1.34506	6595.008	8909.866
A_24_P8049	AF180473 THC19780	1.34481	0.00007	1.34481	1684.884	2287.326
A_23_P21	(NM_01228 Homo sapi CENTB2,A	1.34477	0.00039	1.34477	1469.339	1982.328
A_32_P55	BX537816 MRNA; cDIBX537816	1.34445	2.99E-06	1.34445	14269.69	19080.82
A_23_P94	(NM_00674 Homo sapi SLC20A2,(1.34329	0.00026	1.34329	1061.099	1434.655
A_23_P15	AB040906 Zinc finger ZNF492	1.3429	0.00011	1.3429	351.8152	472.3554
A_24_P24	(NM_17069 Homo sapi dJ222E13.	1.34278	0.00007	1.34278	12175.44	16363.7
A_32_P180199	Unknown A_32_P18	1.34262	0.00068	1.34262	718.343	968.833
A_23_P12	(NM_00515 ubiquitin s USP8,UBP	1.34218	8.77E-06	1.34218	1199.293	1615.192
A_24_P17	(NM_00450 ubiquitin s USP6,HRF	1.34202	0.00006	1.34202	399.0892	535.5604
A_23_P33	(NM_17072 Homo sapi PGBD2	1.34201	4.62E-08	1.34201	1172.031	1572.757
A_23_P20	(NM_00438 Homo sapi CTBS,CTB	1.3415	3.72E-06	1.3415	740.3729	995.1857

A_23_P21	NM_00291	Homo sapi	REV3L,PO	1.34145	8.85E-08	1.34145	1448.767	1946.292
A_23_P15	NM_15345	Homo sapi	MED19,LC	1.3414	0.00006	1.3414	2311.685	3117.264
A_24_P14	NM_01522	chromosom	POFUT2,F	1.34131	0.00001	1.34131	3408.244	4563.249
A_23_P44	NM_03202	eukaryotic	eIF2A,CDA	1.3411	0.00023	1.3411	3043.939	4117.961
A_24_P39	1960	Unknown	ENST0000	1.3411	8.92E-11	1.3411	41079.42	55021.59
A_23_P21	NM_01901	Homo sapi	PLEKHA5,	1.341	4.65E-07	1.341	924.46	1241.888
A_23_P12	NM_01802	Homo sapi	TMEM51,C	1.34087	0.00003	1.34087	2049.488	2766.505
A_24_P22	NM_03214	hypothetic	ZRANB3,M	1.34071	0.00002	1.34071	428.4023	574.3481
A_23_P12	NM_00551	Major histo	HLA-B	1.34065	1.50E-10	1.34065	6631.061	8884.207
A_23_P51	NM_00726	Homo sapi	STXBP3,P	1.34043	5.89E-07	1.34043	2889.037	3888.855
A_23_P33	NM_02054	Homo sapi	ADCY2,HE	1.34029	4.64E-08	1.34029	617.525	827.5364
A_23_P11	NM_01506	Homo sapi	JMJD2C,G	1.3399	0.00035	1.3399	1097.172	1471.905
A_24_P67	NM_17856	Homo sapi	NM_17856	1.33953	0.00017	1.33953	2511.677	3379.185
A_32_P82	BC034583	Hypothetic	LOC40065	1.33944	0.00019	1.33944	309.943	415.2416
A_24_P21	3396	Unknown	A_24_P21	1.33932	2.95E-08	1.33932	49236.14	65746.03
A_32_P11	723	Unknown	ENST0000	1.33914	2.25E-11	1.33914	48324.6	64680.4
A_23_P10	NM_00688	Homo sapi	ZFP36L2,E	1.33893	9.79E-10	1.33893	4736.597	6336.262
A_23_P20	NM_01662	Homo sapi	WAC,Wwp	1.3388	3.45E-10	1.3388	10732.25	14351.66
A_23_P65	NM_00418	Homo sapi	WARS,IFI5	1.33875	4.63E-12	1.33875	5951.148	7966.96
A_23_P12	NM_01652	Homo sapi	UBAP1,UA	1.33873	1.01E-06	1.33873	2719.916	3633.481
A_23_P20	NM_03286	Homo sapi	TNS4,CTE	1.33852	0.00022	1.33852	2319.056	3115.201
A_23_P84	NM_01664	HDCMA18	HDCMA18	1.33847	0.00025	1.33847	2953.354	3995.389
A_23_P14	NM_03221	Homo sapi	ANKRD17,	1.3384	4.38E-08	1.3384	4634.65	6216.777
A_23_P11	NM_01826	Homo sapi	SEC3L1,SI	1.33837	0.00057	1.33837	839.5432	1128.589
A_23_P15	NM_00379	Homo sapi	CTNNAL1,	1.33812	9.69E-07	1.33812	14336.23	19130.3
A_24_P17	NM_01799	hypothetic	FLJ10081,I	1.33769	2.60E-12	1.33769	14550.78	19461.51
A_23_P82	NM_02086	Homo sapi	ZBTB2,ZNI	1.33769	6.05E-07	1.33769	689.2516	921.1932
A_23_P21	NM_02406	Homo sapi	ZNF655,VI	1.33747	0.00001	1.33747	1512.409	2025.158
A_23_P63	NM_14466	Homo sapi	SAMD8,FL	1.33742	0.00009	1.33742	539.3074	721.4584
A_23_P43	NM_17370	Homo sapi	NM_17370	1.33693	2.28E-07	1.33693	7632.672	10203.88
A_23_P18	NM_02213	Homo sapi	MCCC2,M	1.33651	1.75E-07	1.33651	976.3372	1305.959
A_23_P76	AF173389	Early endo	EEA1	1.33634	0.00094	1.33634	2072.941	2796.447
A_32_P14	AL049969	PDZ and L	PDLIM5	1.33631	0.00004	1.33631	2306.213	3105.478
A_24_P18	NM_03246	aspartate b	ASPH,BAF	1.33571	3.35E-09	1.33571	10187.94	13580.98
A_24_P37	NM_00445	fatty-acid-C	ACSL3,AC	1.3353	0.00061	1.3353	3283.711	4424.054
A_23_P34	NM_00145	Homo sapi	FOXO3A,A	1.33509	6.83E-09	1.33509	833.5491	1113.135
A_23_P45	BC017665	Chromosom	CXorf39	1.33503	7.64E-06	1.33503	6075.251	8159.884
A_23_P11	NM_00629	Homo sapi	VRK2	1.33441	1.40E-08	1.33441	4392.081	5867.397
A_23_P75	NM_01473	Homo sapi	ZFYVE16,I	1.33429	8.66E-06	1.33429	1356.946	1805.32
A_23_P20	NM_17792	Homo sapi	H2AFJ,MG	1.33428	7.18E-12	1.33428	20462.76	27309.88
A_23_P15	NM_00422	Homo sapi	STK17B,DI	1.33426	0.00006	1.33426	1804.314	2406.907
A_23_P73	AK124904	FYVE, Rho	FGD6	1.33408	1.54E-06	1.33408	1519.487	2028.797
A_23_P38	NM_01621	Homo sapi	POLK,DINI	1.33356	3.98E-10	1.33356	969.0479	1292.208
A_23_P35	NM_00539	Homo sapi	PPP1R3C,	1.33339	2.29E-07	1.33339	3239.851	4316.448
A_23_P31	NM_18165	Homo sapi	NCOA3,AC	1.33333	0.00011	1.33333	424.1435	565.5776
A_23_P99	NM_19821	Homo sapi	ING1,p33,r	1.33314	6.42E-07	1.33314	1237.732	1654.069
A_32_P15	NM_01527	Homo sapi	KIAA1033	1.33286	6.11E-08	1.33286	4460.627	5953.976
A_24_P64	AK054814	Meis1, myc	MEIS4	1.33284	0.00034	1.33284	1846.411	2484.746
A_24_P37	5573	Unknown	ENST0000	1.33262	0.00002	1.33262	5929.046	7875.254
A_23_P36	NM_03351	Homo sapi	LIMS3	1.33247	0.00003	1.33247	907.1675	1204.999
A_23_P40	NM_14496	RNA (guan	RG9MTD3	1.33243	0.00034	1.33243	363.2094	484.1368
A_32_P58	029	Unknown	A_32_P58	1.3323	0.00001	1.3323	492.3641	655.7681

A_23_P12	NM_01506	Homo sapi	PPRC1,PR	1.3319	7.12E-07	1.3319	5040.396	6733.931
A_23_P69	NM_03271	hypothetic	MGC1132	1.33158	0.00004	1.33158	7832.459	10526.68
A_23_P13	AK026695	Paxillin	PXN	1.33131	1.14E-10	1.33131	122769	163294.3
A_23_P45	NM_01208	Homo sapi	GLUD2,GL	1.33123	0.00028	1.33123	1176.018	1571.468
A_24_P152793		Unknown	ENST0000	1.33105	7.20E-09	1.33105	23434.68	31229.59
A_23_P13	NM_01394	chloride int	CLIC4,H1,I	1.3309	8.13E-06	1.3309	1886.391	2528.144
A_24_P577289		Unknown	A_24_P57	1.3308	0.00002	1.3308	354.1431	471.2794
A_23_P65	NM_00344	Homo sapi	ZBTB17,M	1.33073	2.95E-06	1.33073	900.0258	1196.193
A_32_P58	NM_02471	hypothetic	C15orf29,F	1.33056	3.04E-06	1.33056	4983.12	6619.885
A_23_P54	NM_01326	transcriptio	HCNGP,H	1.32909	6.89E-08	1.32909	712.7861	947.698
A_23_P41	NM_01225	zinc finger	ZNF212,M	1.32888	3.23E-06	1.32888	938.366	1245.58
A_24_P10	NM_02087	Homo sapi	SH3MD2,F	1.32886	0.00004	1.32886	1713.218	2285.124
A_24_P95	NM_00602	Tax1 (hum:	TAX1BP1,	1.32877	3.36E-06	1.32877	23255.14	31028.63
A_32_P13	AL137304	Intersex-lik	IXL	1.32822	4.05E-08	1.32822	964.9424	1281.673
A_24_P89	NM_19795	Homo sapi	C9orf90,KI	1.32788	0.00077	1.32788	1885.474	2539.301
A_24_P41309		Unknown	ENST0000	1.32767	1.28E-11	1.32767	2351.262	3121.587
A_23_P21	NM_01608	palladin	KIAA0992,	1.32702	0.00023	1.32702	6472.759	8655.582
A_32_P14	NM_00698	nuclear por	NPPIP	1.32677	6.87E-10	1.32677	33295.55	44219.07
A_24_P33	NM_18153	Homo sapi	BTN2A2,B	1.32663	5.59E-06	1.32663	597.5101	792.5389
A_23_P33	NM_15247	Homo sapi	ZNF578,FL	1.32652	4.41E-06	1.32652	538.4827	714.3882
A_24_P89	NM_00344	zinc finger	TRIM26,AF	1.32645	4.77E-11	1.32645	4202.642	5575.151
A_23_P21	NM_00659	Homo sapi	POLQ,POL	1.32609	4.65E-09	1.32609	1357.985	1800.23
A_32_P13	NM_19893	Homo sapi	SS18L1,CF	1.32581	1.17E-06	1.32581	887.8827	1177.677
A_23_P30	NM_00123	Homo sapi	CCNH,CA	1.3247	6.37E-09	1.3247	5546.615	7359.646
A_24_P22	NM_01781	hypothetic	RAB20,FL	1.32361	0.00046	1.32361	524.4957	693.3095
A_23_P30	NM_14713	Homo sapi	NFX1,NFX	1.32315	2.89E-10	1.32315	3760.671	4975.375
A_23_P50	NM_14519	Homo sapi	LIPT1,MGC	1.32276	0.0001	1.32276	449.7584	595.0693
A_32_P123589		Unknown	A_32_P12	1.32213	6.34E-06	1.32213	900.5848	1190.068
A_23_P201521		Unknown	A_23_P20	1.32197	5.93E-06	1.32197	131989.3	173555.6
A_24_P18	NM_01538	Nipped-B h	NIPBL	1.32196	0.0003	1.32196	1031.138	1372.542
A_32_P40	BC047332	Zinc finger	ZEC	1.32187	7.95E-07	1.32187	978.4361	1294.433
A_32_P22989		Unknown	A_32_P22	1.32184	0.00033	1.32184	131832.5	173002.6
A_24_P502660		Unknown	A_24_P50	1.32136	0.00004	1.32136	420.9603	556.2221
A_23_P86	NM_00567	Homo sapi	ZNF239,M	1.32118	4.92E-08	1.32118	1630.861	2156.828
A_32_P10	NM_00377	UDP-Gal:b	B4GALT4,I	1.32094	1.68E-06	1.32094	871.6884	1153.551
A_23_P38	AK055693	Zinc finger	FLJ35036	1.32073	3.71E-11	1.32073	4820.948	6368.903
A_23_P40	NM_17588	hypothetic	FLJ36031	1.32071	4.75E-08	1.32071	1471.221	1945.192
A_23_P25	NM_00550	Homo sapi	DMXL1	1.32054	0.00088	1.32054	547.7094	724.9379
A_24_P13	AK025245	KIT ligand	KITLG	1.32044	3.08E-06	1.32044	2379.11	3150.725
A_24_P479065		Unknown	A_24_P47	1.32042	4.49E-08	1.32042	1478.666	1952.579
A_32_P87	AB046850	Homo sapi	KIAA1630	1.3202	0.00023	1.3202	1038.776	1370.54
A_24_P33	AB107355	FLJ16231	FLJ16231	1.31992	4.59E-07	1.31992	883.3005	1166.353
A_23_P66	NM_01565	Homo sapi	NAT9,EBS	1.31989	5.33E-06	1.31989	7678.469	10095.51
A_32_P171921		AB017004	THC18654	1.31981	0.00013	1.31981	378.8735	500.0009
A_23_P14	AK021784	Acidic (leuc	ANP32A	1.31978	1.57E-07	1.31978	49091.8	64951.2
A_23_P14	NM_00456	Homo sapi	PCK2,PEP	1.31956	8.78E-10	1.31956	15347.19	20231.25
A_23_P15	NM_00439	Homo sapi	DDX5,p68,	1.31937	3.10E-09	1.31937	40068.05	52947.65
A_24_P20	NM_02472	MICAL-like	MICAL-L2	1.31919	0.00004	1.31919	1433.594	1891.858
A_23_P15	NM_01488	Homo sapi	FAM3C,GS	1.31901	1.85E-09	1.31901	4556.443	6004.914
A_23_P20	NM_01830	hypothetic	FLJ11029	1.31888	6.88E-06	1.31888	27855.29	36559.71
A_23_P38	NM_01463	KIAA0336	GCC2,GCC	1.31885	1.52E-06	1.31885	509.6915	672.2296
A_24_P666795		Unknown	A_24_P66	1.31826	0.00009	1.31826	1864.417	2445.141

A_23_P92	NM_00434	Homo sapi	CASP3,CP	1.3181	3.36E-09	1.3181	6773.043	8936.277
A_23_P36	NM_20736	Homo sapi	NM_20736	1.31802	0.00079	1.31802	780.8832	1029.632
A_23_P50320		Unknown	A_23_P50	1.31765	1.27E-11	1.31765	62339.71	82140.31
A_23_P50	NM_17586	Homo sapi	CBFA2T2,I	1.31753	0.00093	1.31753	339.3862	447.3696
A_23_P56	NM_03275	hypothetic	MGC15631	1.31674	1.48E-10	1.31674	90937.38	119670.6
A_23_P32	NM_01904	hypothetic	FLJ20054,I	1.31669	0.00035	1.31669	924.0523	1222.521
A_23_P43	NM_02478	Homo sapi	ZBTB3,FL	1.31636	0.00007	1.31636	1094.582	1439.22
A_23_P46	NM_02119	Homo sapi	PTBP2,PTI	1.31607	8.20E-07	1.31607	1441.219	1901.152
A_23_P14	NM_00174	Homo sapi	CANX,CN	1.31601	0.00003	1.31601	20761.48	27485.78
A_24_P41	NM_01814	Sec61 alph	SEC61A2,I	1.31601	5.06E-09	1.31601	1636.093	2151.323
A_24_P31	NM_00574	protein dist	PDIA6,P5,I	1.31592	4.19E-06	1.31592	8448.797	11158.37
A_23_P59	NM_00038	cyclin-depe	CDKN1A,F	1.31583	2.14E-08	1.31583	7263.936	9566.186
A_32_P68	NM_17592	hypothetic	LOC28563	1.3157	0.00006	1.3157	4131.838	5423.858
A_24_P23	NM_00380	tumor necr	TNFSF14,I	1.31551	2.06E-06	1.31551	494.2853	650.1943
A_32_P215900		Unknown	A_32_P21	1.31544	0.00035	1.31544	536.5212	704.5668
A_24_P32	NM_00036	tuberous sc	TSC1,LAM	1.31532	0.00017	1.31532	859.4998	1137.189
A_24_P24	AK025794	PI-3-kinase	SMG1	1.31527	0.00014	1.31527	1559.45	2062.952
A_24_P724886		Unknown	A_24_P72	1.31483	0.00037	1.31483	346.3288	455.3191
A_24_P47	BX118521	BX118521	BX118521	1.31478	3.61E-10	1.31478	9975.357	13117.3
A_23_P53	NM_17850	Homo sapi	DTX3,RNF	1.31439	0.00038	1.31439	4932.823	6523.188
A_24_P11	NM_00698	nuclear pot	NPIP	1.31423	2.46E-09	1.31423	24741.66	32554.96
A_32_P37	NM_14531	Homo sapi	LACE1	1.31396	0.00041	1.31396	518.855	682.5882
A_23_P55	NM_00379	Homo sapi	RNMT,ME	1.31394	0.00014	1.31394	1254.396	1653.802
A_23_P27	NM_01833	Homo sapi	ZNF444,E2	1.31388	3.64E-06	1.31388	6288.17	8309.276
A_24_P12	NM_03341	hypothetic	MGC13138	1.31382	0.00016	1.31382	1050.055	1382.809
A_23_P99	NM_00345	zinc finger	ZNF198,FI	1.31319	2.33E-08	1.31319	1435.945	1886.763
A_23_P14	NM_03145	Homo sapi	C10orf45,F	1.31317	7.09E-06	1.31317	5412.416	7133.452
A_24_P63	NM_01477	KIAA0141	KIAA0141	1.31288	6.04E-06	1.31288	2552.864	3362.008
A_23_P21	NM_01404	DKFZP564	CHMP2B,C	1.31269	4.77E-06	1.31269	2880.109	3803.89
A_23_P13	NM_00574	YY1 assoc	YAF2	1.31257	0.00014	1.31257	1329.374	1754.375
A_32_P15	NM_00362	protein phc	PPM1D,WI	1.31229	0.00016	1.31229	2074.615	2723.719
A_23_P84	NM_02407	hypothetic	SECISBP2	1.31207	0.00007	1.31207	624.3018	820.4475
A_24_P366465		Unknown	ENST0000	1.31163	5.15E-09	1.31163	3474.691	4560.43
A_23_P35	NM_17851	Homo sapi	ANKK1,PK	1.31146	0.00001	1.31146	160781.1	209878.1
A_24_P23	NM_20747	FLJ42953	FLJ42953	1.31146	1.76E-06	1.31146	678.8185	889.7672
A_23_P13	NM_00638	HIV-1 Tat i	HTATIP,TII	1.31121	1.58E-09	1.31121	1052.168	1379.555
A_24_P76	AK097724	Ariadne ho	ARIH2	1.31111	0.00003	1.31111	381.9552	500.769
A_24_P58	NM_00381	tumor necr	TNFSF9,4-	1.31104	6.56E-06	1.31104	1970.285	2596.018
A_32_P51	BC015790	Hypothetic	LOC40057	1.3109	0.00002	1.3109	2486.096	3237.974
A_23_P34	NM_01492	KIAA0971	KIAA0971	1.31084	7.93E-08	1.31084	4178.72	5492.099
A_23_P16	NM_01651	Pipelicolic a	PIPOX	1.3108	2.29E-08	1.3108	4218.306	5525.769
A_32_P20	AK096653	Hypothetic	LOC91664	1.31045	4.38E-08	1.31045	2938.943	3848.898
A_23_P20	NM_02245	Homo sapi	NOC3L,AD	1.31022	4.05E-06	1.31022	4399.616	5805.903
A_32_P22	NM_03202	CDA11 pro	TATDN1,C	1.31002	1.81E-10	1.31002	4834.794	6331.496
A_24_P12	NM_00534	Heat shock	HSPA1A	1.30943	4.42E-09	1.30943	8931.879	11687.45
A_23_P13	NM_00501	Homo sapi	NUCB2,NE	1.3091	0.00007	1.3091	3404.814	4493.199
A_24_P58	NM_03062	Homo sapi	CPEB4,KI	1.30909	0.00014	1.30909	658.0826	859.736
A_23_P11	NM_01206	activating t	ATF5,ATF	1.30903	1.11E-06	1.30903	710.7648	929.9735
A_23_P21	NM_01487	KIAA0354	ZBTB5	1.30903	0.0002	1.30903	2796.294	3645.821
A_32_P11	NM_15228	Homo sapi	ZNF561,M	1.30886	0.00013	1.30886	924.4623	1207.78
A_32_P55987		Unknown	THC19826	1.30815	0.00018	1.30815	352.8459	461.611
A_23_P15	NM_00343	Homo sapi	ZNF133,ZI	1.30781	3.96E-07	1.30781	2175.333	2849.51

A_24_P289043	Unknown	ENST0000	1.30773	0.00092	1.30773	1914.705	2525.82	
A_23_P79	BC004943	LAG1 long	LASS1	1.30756	1.17E-08	1.30756	374255.9	488680.3
A_24_P17	AB002321	KIAA0323	KIAA0323	1.30741	0.00005	1.30741	697.4764	912.2087
A_23_P13	NM_01601	Homo sapi	PHF20L1,C	1.30741	0.00001	1.30741	1926.17	2517.765
A_23_P13	NM_02182	Homo sapi	C6orf75,MI	1.30637	5.10E-06	1.30637	491.561	642.1266
A_23_P51	NM_01239	Homo sapi	PFDN2,PF	1.30631	4.89E-08	1.30631	40614.39	52924.57
A_32_P53	NM_00580	putative tra	EIF1,A121	1.30625	2.30E-08	1.30625	256945.7	335997.9
A_23_P13	NM_01609	Homo sapi	RBM7,FLJ	1.30622	0.00016	1.30622	1097.315	1438.738
A_24_P10	NM_03212	implantatio	DKFZp564	1.3059	0.00001	1.3059	3684.394	4799.97
A_23_P16	NM_00419	Homo sapi	GBF1,KIAA	1.30569	3.08E-07	1.30569	4775.581	6220.451
A_32_P166693	Unknown	THC19104		1.30563	8.21E-06	1.30563	5364.491	7002.964
A_24_P18	NM_01834	hypothetic	FLJ11171	1.30507	0.00001	1.30507	1126.787	1471.628
A_23_P75	NM_02489	Homo sapi	PDZK7	1.30505	0.00003	1.30505	121065.5	157136.3
A_23_P14	NM_02411	hypothetic	MGC4504	1.30504	9.47E-07	1.30504	13210.38	17287.3
A_24_P84	M27126	Major histo	HLA-DRB1	1.30409	0.00003	1.30409	502.7695	655.8606
A_24_P49860	Unknown	A_24_P49		1.30382	8.21E-07	1.30382	717.0684	935.2377
A_23_P10	NM_13911	Homo sapi	YY1AP1,Y	1.30378	3.96E-08	1.30378	4369.401	5707.521
A_32_P12	NM_03098	small GTP-	RAB1B	1.3037	3.01E-09	1.3037	10578.47	13793.46
A_24_P68	NM_00534	Heat shock	HSPA1A	1.30352	0.00047	1.30352	4334.627	5666.624
A_23_P43	NM_17383	hypothetic	YIPF6,FinC	1.30343	0.00034	1.30343	9035.727	11879.9
A_23_P39	NM_01490	Homo sapi	GLS,GLS1	1.30325	0.00021	1.30325	3358.136	4399.524
A_24_P41530	Unknown	A_24_P41		1.30279	6.86E-06	1.30279	3049.456	3961.218
A_23_P32	NM_03320	Homo sapi	C16orf45,F	1.30271	0.00002	1.30271	1531.77	2002.542
A_24_P25	NM_02093	Homo sapi	ZNF317,KI	1.3023	0.00009	1.3023	1277.441	1670.813
A_23_P12	NM_01243	Homo sapi	SETDB1,E	1.30186	0.00005	1.30186	549.986	716.0113
A_23_P13	NM_01568	Homo sapi	ARRDC2,F	1.3016	0.00002	1.3016	1016.2	1317.854
A_23_P16	NM_00187	Homo sapi	CPM	1.30145	0.00025	1.30145	529.5037	688.4868
A_23_P13	NM_02196	Homo sapi	ZNF148,BE	1.30078	0.00096	1.30078	3479.94	4571.263
A_32_P15	AK127081	Ubiquitin s	USP54	1.30011	0.00005	1.30011	1162.868	1518.354
A_24_P31	AK026491	Homo sapi	RPL12	1.30003	1.11E-07	1.30003	40198.94	52128.06
A_23_P34	NM_13849	Homo sapi	ZNF655,VI	1.30002	0.0001	1.30002	1787.272	2326.459
A_23_P60753	TRUA_HUI	THC20026		1.29961	2.32E-10	1.29961	244341.6	317615.2
A_23_P11	NM_00499	Homo sapi	MECP2,RT	1.29918	2.96E-06	1.29918	5101.608	6627.076
A_23_P13	NM_02500	Thrombos	TSRC1	1.29896	9.18E-08	1.29896	3735.843	4850.058
A_23_P14	NM_01895	Homo sapi	HLA-F,HLA	1.29886	3.93E-06	1.29886	4877.125	6328.487
A_24_P32	BC006474	U2-associa	SR140	1.29821	0.00004	1.29821	6391.592	8332.881
A_23_P86	NM_00452	Homo sapi	KIF5B,KN5	1.29732	0.00073	1.29732	622.6592	809.7568
A_23_P59	AB018259	Dedicator c	DOCK4	1.29723	0.00002	1.29723	824.8256	1069.386
A_23_P34	NM_00311	Homo sapi	SP3,SPR-2	1.297	8.53E-08	1.297	4226.754	5473.929
A_23_P31	NM_17827	Homo sapi	PILRA,FD	1.29699	0.00017	1.29699	596.7608	773.6652
A_23_P46	NM_14527	kinesin ligh	KLC3,KLC	1.29657	0.00003	1.29657	257023.1	331332.9
A_24_P493100	Unknown	A_24_P49		1.29638	0.0002	1.29638	990.9985	1283.265
A_23_P12	NM_02101	Homo sapi	KCNJ12,IR	1.29614	0.00098	1.29614	895.1544	1151.202
A_24_P247576	Unknown	ENST0000		1.29603	8.31E-09	1.29603	4402.003	5701.043
A_23_P32	BC007811	Similar to g	FLJ22795	1.29588	6.23E-07	1.29588	674.6248	874.3666
A_24_P221475	Unknown	A_24_P22		1.29586	6.49E-06	1.29586	2914.562	3786.377
A_23_P76	NM_01344	Homo sapi	BAZ1A,AC	1.29538	1.39E-06	1.29538	2462.129	3205.871
A_24_P14	NM_00472	DEAD (As	DDX21	1.29535	0.00009	1.29535	15271.76	19925.34
A_23_P21	NM_01826	Homo sapi	RSAFD1,F	1.29527	6.25E-06	1.29527	2497.011	3243.133
A_23_P40	NM_00676	Homo sapi	MYST3,MC	1.29515	0.00077	1.29515	1625.167	2124.058
A_23_P78	NM_01770	hypothetic	FBXL12,Ft	1.2951	0.00037	1.2951	541.0021	700.4343
A_23_P11	NM_02197	Homo sapi	MAP2K1IP	1.29481	3.48E-08	1.29481	2155.736	2794.565

A_23_P41	NM_15246 Homo sapi	EVER2, EV	1.29468	0.00009	1.29468	737.9551	953.2941
A_23_P12	NM_02065 Homo sapi	PELI1, pelli	1.29465	0.00005	1.29465	1254.518	1613.933
A_23_P48	NM_00117 Homo sapi	ARHGAP5	1.2944	1.49E-07	1.2944	703.3422	910.4166
A_32_P34	AK056119 CDNA FLJ	AK056119	1.29423	0.00097	1.29423	299.4373	387.5505
A_23_P90	NM_15252 hypothetic	FLJ40432	1.29387	1.12E-06	1.29387	1611.273	2087.51
A_24_P40	NM_01465 KIAA0446	KIAA0446,	1.29341	3.69E-06	1.29341	3288.218	4263.478
A_32_P64	AK026852 Similar to h	ZNF694, ZS	1.29326	9.33E-07	1.29326	973.7066	1259.913
A_23_P94	AY260738 Zinc finger	ZKSCAN1	1.29323	9.21E-08	1.29323	99884.2	129326.1
A_23_P38	NM_20733 zinc finger	ZFPL, DKF	1.29302	5.51E-06	1.29302	567.6802	734.0738
A_24_P40	AK057725 Matrilin 1,	MATN1	1.29295	0.00067	1.29295	894.0439	1157.916
A_23_P71	BC034271 Fanconi an	FANCC	1.29276	1.75E-08	1.29276	14421.81	18654
A_24_P27	NM_19904 Homo sapi	NSUN4, MC	1.29246	0.00004	1.29246	1004.347	1296.956
A_24_P41	AK093442 Zinc finger	ZNF596	1.29173	5.75E-06	1.29173	7919.088	10293.35
A_23_P16	NM_14472 hypothetic	FLJ31951, I	1.29168	0.00012	1.29168	3654.661	4752.119
A_23_P21	AB014535 Homo sapi	KIAA0635	1.29131	2.08E-09	1.29131	3718.092	4799.471
A_24_P36	NM_01617 hypothetic	LOC51149	1.29094	5.38E-07	1.29094	3581.956	4616.793
A_32_P20884	ALU7_HUM	THC19762	1.29079	0.00088	1.29079	594.4699	767.8296
A_23_P32	NM_00343 Homo sapi	ZNF134, p	1.29062	0.00031	1.29062	1367.771	1755.121
A_24_P21	NM_02475 modulator	(FLJ13213, I	1.29044	8.89E-10	1.29044	3270.191	4219.909
A_23_P20	NM_00325 Tight juncti	TJP1	1.29042	3.40E-06	1.29042	1495.921	1932.999
A_23_P76	NM_00142 Homo sapi	EMP1, TMF	1.29037	0.00001	1.29037	2948.259	3804.05
A_23_P97	NM_02215 Rag C prot	RRAGC, G	1.29021	0.00002	1.29021	3829.646	4925.572
A_23_P12	NM_02482 Homo sapi	C9orf82, FL	1.29016	2.13E-06	1.29016	1013.436	1308.593
A_23_P17	NM_00218 Homo sapi	IL1RAP, IL1	1.28979	6.16E-07	1.28979	653.2115	842.5768
A_24_P13	NM_15228 Homo sapi	ZFP276, ZN	1.28975	8.30E-06	1.28975	1671.135	2161.579
A_32_P11	AK091308 NMDA rec	NARG1	1.28945	2.98E-06	1.28945	660.1144	851.3978
A_23_P45	NM_00580 putative tra	EIF1, A121	1.28943	2.80E-07	1.28943	311765.4	402869.4
A_24_P572974	Unknown	A_24_P57	1.28934	0.00004	1.28934	530.3701	683.5947
A_23_P12	M21539 Small proli	SPRR2C	1.28902	0.00021	1.28902	4372.502	5570.128
A_23_P35	U68019 SMAD, mo	SMAD3	1.28817	0.00009	1.28817	1660.483	2146.856
A_24_P21	NM_03281 hypothetic	TBRG1, TB	1.28764	0.00017	1.28764	920.5256	1189.426
A_23_P11	NM_19431 FRBZ1 pro	FRBZ1	1.2875	0.00004	1.2875	546.9985	704.3596
A_24_P26	AK055669 Transportir	TNPO1	1.28728	0.00002	1.28728	5107.287	6605.785
A_24_P13	NM_00333 ubiquitin-c	UBE2D1, S	1.28662	0.0001	1.28662	5250.372	6797.6
A_23_P41	NM_01436 Homo sapi	GNL3, NS, E	1.28659	1.83E-07	1.28659	21759.74	27915.7
A_24_P33	NM_03212 implantatio	DKFZp564	1.28614	4.54E-07	1.28614	12706.53	16373.68
A_23_P39	NM_00235 Homo sapi	MAFG, MG	1.28611	5.68E-10	1.28611	19146.42	24624.77
A_32_P17	AB002293 Zinc finger	ZNF609	1.28578	0.00068	1.28578	9223.809	11948.53
A_24_P22	NM_00407 cysteine ar	CSRP1, CF	1.28542	3.76E-07	1.28542	822.3028	1057.192
A_24_P332471	Unknown	A_24_P33	1.28529	2.76E-06	1.28529	37847.43	48842.98
A_23_P89	AL080108 KIAA0753	KIAA0753	1.28464	0.00002	1.28464	1870.074	2408.373
A_23_P12	NM_00626 Homo sapi	DPF2, REC	1.28453	4.71E-08	1.28453	4405.774	5652.042
A_23_P20	AL353936 Pleckstrin	PSD3	1.28423	0.00004	1.28423	652.4528	837.57
A_23_P57	NM_00613 Homo sapi	TMEM50B	1.28417	0.00031	1.28417	1284.198	1652.521
A_23_P14	AK026323 Alpha-kina	ALPK1	1.2841	2.67E-09	1.2841	160543.4	206243
A_23_P39	BC037316 Zinc finger	ZNF630	1.28403	0.00062	1.28403	4315.569	5495.141
A_24_P29	NM_15320 Heat shock	HSPA8	1.28354	0.00001	1.28354	9848.959	12635.58
A_23_P93	AK098134 Similar to	AK098134	1.28331	9.38E-06	1.28331	5069.791	6522.459
A_24_P25	NM_00189 casein kina	CSNK1A1,	1.28301	0.0004	1.28301	4017.897	5128.415
A_23_P15	AK021957 Collagen, t	COL27A1	1.28254	0.0002	1.28254	2416.706	3086.241
A_32_P16	NM_03225 hypothetic	ANKRD20,	1.28241	0.00062	1.28241	374.9847	480.9252
A_32_P22	NM_18178 Homo sapi	SPRED2, S	1.28224	0.00067	1.28224	519.0997	666.2019

A_24_P913005	AF124394	THC18472	1.28217	0.0001	1.28217	1109.068	1426.985	
A_23_P15	(AK027372	Hypothetic	FLJ14466	1.28196	2.72E-07	1.28196	133737	171156.8
A_23_P20	NM_00609	Homo sapi	NDRG1,G	1.28158	0.00001	1.28158	5445.292	6964.266
A_23_P21	NM_01774	Homo sapi	ANKHD1,M	1.28145	0.00005	1.28145	1615.054	2077.297
A_23_P16	NM_01835	hypothetic	FLJ11184	1.28138	0.0001	1.28138	3843.594	4936.703
A_23_P15	NM_02303	ankyrin rep	ANKRA2,A	1.28111	4.88E-07	1.28111	2263.122	2899.137
A_32_P146450	Unknown	A_32_P14		1.2804	4.19E-07	1.2804	7738.759	9924.258
A_23_P15	NM_00284	Homo sapi	PTPRM,RF	1.28038	5.61E-09	1.28038	1635.75	2094.444
A_23_P54	NM_00594	Homo sapi	MT1A,MT1	1.28025	0.00003	1.28025	72072.63	92574.56
A_24_P37	NM_01894	cytochrom	CYCS,CYC	1.28019	0.00003	1.28019	31074.22	39666.04
A_23_P15	(AF338191	Homo sapi	FKSG62	1.28007	0.00006	1.28007	6744.516	8674.066
A_23_P38	NM_15258	hypothetic	DKFZp762	1.27975	0.00038	1.27975	1075.471	1376.076
A_24_P41	NM_01343	potential tu	LRP12,ST	1.2797	0.00013	1.2797	1582.952	2040.54
A_23_P15	NM_01607	Bcl-2 inhibi	BIT1,CGI-1	1.27947	1.87E-09	1.27947	16700.99	21370.87
A_23_P17	(NM_00694	stress 70 p	STCH	1.27932	0.00047	1.27932	444.0095	568.0571
A_23_P23	(NM_01605	CGI-115 pr	CGI-115	1.2793	4.62E-07	1.2793	8098.307	10381.56
A_23_P90	(NM_14896	Homo sapi	OTOS,OTC	1.27917	0.00045	1.27917	2765.538	3562.701
A_23_P63	(NM_00720	Homo sapi	DDX20,DP	1.27911	0.00021	1.27911	3212.884	4099.209
A_23_P32	(NM_15324	Homo sapi	NPHP3,NF	1.27888	0.00001	1.27888	1085.954	1387.173
A_23_P77	(AK024389	C-Maf-indu	CMIP	1.2787	3.82E-08	1.2787	3045.203	3896.969
A_23_P31	(NM_00409	Homo sapi	EMX1	1.27859	3.84E-08	1.27859	13943.77	17824.07
A_32_P81	(NM_01436	putative nu	GNL3,NS,E	1.27844	0.00006	1.27844	30490.11	38832.16
A_23_P13	(NM_02456	hypothetic	FLJ21616,I	1.27767	1.99E-06	1.27767	3107.567	3973.62
A_23_P21	(NM_00342	Homo sapi	ZNF35,HF	1.27724	0.00003	1.27724	487.6641	622.8716
A_23_P21	(NM_01494	Homo sapi	UBOX5,Ull	1.27676	1.98E-07	1.27676	263758.9	337339.9
A_23_P11	(NM_00484	Homo sapi	ATG5,ASP	1.2766	6.13E-06	1.2766	3189.14	4081.498
A_23_P26	(NM_02475	modulator	(FLJ13213,I	1.27641	9.43E-06	1.27641	2959.434	3785.959
A_23_P11	(NM_01637	hydroxyste	HSD17B7,I	1.27604	0.001	1.27604	825.4517	1056.623
A_23_P21	(NM_00207	guanine nu	GNAZ	1.27571	1.51E-08	1.27571	6593.481	8416.037
A_24_P49	(AB037853	KIAA1432	KIAA1432	1.27569	0.0003	1.27569	940.5531	1198.852
A_23_P98042	Unknown	A_23_P98		1.27566	0.00007	1.27566	15369.44	19690.91
A_24_P12	(NM_00166	ADP-ribosy	ARF4L,AR	1.2755	7.47E-06	1.2755	698.1168	890.469
A_24_P36	(NM_01794	hypothetic	NOL8,Nop	1.27542	6.26E-06	1.27542	9350.615	11959.61
A_24_P10	(NM_01591	Thioredoxir	TXNDC11	1.27539	0.00012	1.27539	4265.741	5451.861
A_32_P28	(NM_17216	Homo sapi	NASP,FLB	1.27531	9.02E-06	1.27531	6673.458	8483.616
A_23_P33	(XM_37095	similar to n	LOC38823	1.27529	2.50E-07	1.27529	15108.16	19292.29
A_23_P16	(AF151059	Formin bin	FNBP3	1.27527	0.00058	1.27527	10485.03	13456.59
A_23_P99	(NM_00531	Homo sapi	PDIA3,P58	1.27508	8.31E-06	1.27508	13748.27	17593.27
A_32_P125558	Unknown	A_32_P12		1.27498	0.00051	1.27498	3282.285	4206.799
A_24_P487766	Unknown	A_24_P48		1.27486	1.05E-06	1.27486	1876.847	2395.592
A_24_P33	(D31888	REST core	RCOR1	1.27485	2.80E-06	1.27485	2678.839	3426.815
A_23_P80	(NM_00680	Homo sapi	BTG3,ANA	1.27483	0.00003	1.27483	6742.735	8645.48
A_32_P214969	Unknown	A_32_P21		1.27471	0.00007	1.27471	786.8256	1002.606
A_24_P16	(NM_00016	GTP cyclo	GCH1,GCH	1.27465	1.95E-07	1.27465	921.9121	1175.06
A_23_P21	(NM_02297	Homo sapi	ACSL4,AC	1.27407	0.00035	1.27407	1546.177	1972.979
A_24_P37	(NM_00097	ribosomal	rRPL12	1.27345	1.67E-06	1.27345	72983.67	92655.45
A_23_P15	(NM_15335	hypothetic	MGC33214	1.27322	0.00018	1.27322	3741.071	4768.196
A_23_P10	(AF077204	GTP bindin	GTPBP1	1.27319	0.00025	1.27319	1596.063	2036.281
A_23_P15	(NM_02214	leucine zip	FKSG14,P	1.27301	0.00078	1.27301	2873.13	3691.327
A_24_P30	(NM_00344	zinc finger	ZNF155,pf	1.27263	0.00002	1.27263	830.6883	1056.057
A_23_P11	(NM_14521	Homo sapi	TRIM11,BI	1.27262	0.00016	1.27262	1704.85	2174.028
A_23_P13	(NM_00626	Homo sapi	PROP1	1.27249	0.00094	1.27249	21403.7	26956.19

A_24_P281853	Unknown	A_24_P281853	1.27205	0.00005	1.27205	3390.195	4332.317
A_23_P201	NM_00204	Homo sapiens GABPB2, E	1.272	7.92E-09	1.272	4323.982	5498.97
A_23_P201	NM_19879	Homo sapiens MAP4K5, K	1.27179	0.0005	1.27179	1029.157	1315.374
A_23_P201	NM_00076	Homo sapiens CYP2B6, C	1.27175	0.00026	1.27175	7187.839	9173.77
A_32_P671	BC045163	Homo sapiens BC045163	1.27161	0.00023	1.27161	1174.658	1493.04
A_24_P371	XM_37271	Homo sapiens XM_37271	1.27154	0.00004	1.27154	41766.95	53463.37
A_24_P851	AI791206	MRNA (fetus: APG12L, AI)	1.2708	0.00047	1.2708	384.1207	488.11
A_24_P101	NM_00531	glucose receptor PDIA3, P58	1.27058	9.68E-08	1.27058	15794.55	20084.84
A_23_P141	NM_02198	Homo sapiens OPTN, NRF	1.27019	4.83E-08	1.27019	2003.106	2545.011
A_24_P341	XM_37181	Homo sapiens XM_37181	1.26992	3.44E-08	1.26992	129910.2	164826
A_24_P471	AK093640	CDNA FLJ: AK093640	1.26987	0.00048	1.26987	3668.459	4697.468
A_23_P391	NM_02500	Thrombospondin TSRSC1	1.26978	2.89E-06	1.26978	862.4924	1095.554
A_23_P121	NM_00167	activating transcription factor ATF4, CRE	1.26963	8.17E-07	1.26963	17026.98	21643.86
A_23_P111	AF249273	BCL2-associated protein BCLAF1	1.26941	0.00086	1.26941	5038.934	6442.979
A_23_P141	NM_01479	Homo sapiens ZFHX1B, S	1.26937	0.0001	1.26937	910.0101	1152.486
A_23_P811	NM_08060	Homo sapiens TJAP1, PIL	1.26872	0.00003	1.26872	1355.774	1719.99
A_23_P201	NM_01221	Homo sapiens MGEA5, MI	1.26861	4.75E-06	1.26861	10695.42	13588.19
A_23_P971	NM_01222	Homo sapiens NT5C2, GM	1.26809	0.00015	1.26809	982.3973	1249.166
A_24_P651	AK125790	Hypothetical LOC40135	1.26793	0.0001	1.26793	6810.076	8577.2
A_32_P611	AF290475	Round specter RSBN1L	1.2678	2.77E-07	1.2678	879.4421	1114.935
A_24_P765053	ALU5_HUMAN	THC18911	1.26763	0.00052	1.26763	406.2131	514.9318
A_24_P261	NM_00586	Bet1 (S. cerevisiae) BET1, hbet	1.26688	0.00035	1.26688	1525.573	1941.617
A_23_P711	NM_01794	Homo sapiens NOL8, Nop	1.26668	1.64E-08	1.26668	10628.7	13458.19
A_32_P151	BC033829	A kinase (F) AKAP12	1.26658	1.03E-06	1.26658	2599.842	3289.615
A_24_P110541	Unknown	A_24_P110541	1.26644	1.41E-06	1.26644	50061.02	63581.93
A_32_P711	AL049987	FLJ40092 FLJ40092	1.26635	0.00003	1.26635	1546.502	1960.397
A_23_P161	NM_00313	signal receptor SRP54	1.26633	0.00065	1.26633	2267.857	2890.318
A_23_P421	NM_02020	Homo sapiens C9orf102, S	1.2663	1.43E-06	1.2663	1446.14	1830.549
A_24_P381	NM_00171	carcinoembryonic antigen CEACAM1	1.26499	0.00008	1.26499	1527.748	1922.522
A_23_P501	NM_00715	Homo sapiens ZXDA	1.26494	0.0009	1.26494	470.2337	595.0486
A_23_P336854	DJ999L4.1	RPL12L3	1.26466	1.87E-06	1.26466	125946.1	159780.9
A_23_P136013	Unknown	A_23_P136013	1.26452	0.00079	1.26452	5461.702	6959.69
A_24_P261	NM_15375	Homo sapiens C21orf88	1.26443	2.24E-07	1.26443	5817.241	7355.478
A_32_P171	D86974	KIAA0220- KIAA0220	1.26419	2.20E-07	1.26419	40389.42	51035.8
A_23_P511	NM_00724	Homo sapiens DUSP12, Y	1.26414	0.00031	1.26414	9265.242	11782.29
A_24_P341	NM_01499	Homo sapiens DAAM1, KI	1.26412	0.0007	1.26412	942.8012	1189.227
A_23_P131	NM_00636	nuclear ribonucleoprotein RNXF1, TAP	1.26404	0.00039	1.26404	1342.606	1697.148
A_23_P141	NM_01793	Homo sapiens PHIP, ndr	1.26373	8.99E-09	1.26373	5289.284	6684.321
A_24_P261	NM_13325	Homo sapiens LRPPRC, L	1.26368	0.00003	1.26368	8752.191	11104.96
A_24_P602348	ALU6_HUMAN	THC19585	1.26366	0.00011	1.26366	10611.14	13343.32
A_24_P401	XM_37231	Homo sapiens XM_37231	1.26359	0.00002	1.26359	47997.49	60350.25
A_24_P461	AK022083	Hypothetical FLJ12750	1.26347	0.00014	1.26347	460.256	581.4894
A_23_P501	NM_17237	Homo sapiens IL4I1, FIG1	1.26346	0.00008	1.26346	745.1948	941.7758
A_23_P991	NM_02256	Homo sapiens MESDC1, M	1.26341	0.00001	1.26341	2124.64	2690.094
A_32_P451	AL049265	Interleukin IL6ST	1.26249	1.35E-08	1.26249	3971.108	5013.483
A_23_P271	NM_19924	multidrug resistance MGC13170	1.2621	0.00002	1.2621	10241.84	12908.4
A_23_P481	NM_00643	Homo sapiens NPC2, HE1	1.2617	9.83E-06	1.2617	19731.48	24938.55
A_32_P222060	Unknown	THC18266	1.26164	0.00068	1.26164	961.848	1215.121
A_32_P76853	AF263742	THC20011	1.26137	0.00061	1.26137	966.4484	1221.198
A_24_P731	BI041703	Cisplatin resistance CRR9	1.26089	0.00029	1.26089	16737.27	21135.53
A_23_P211	NM_01988	Homo sapiens CYP26B1, I	1.26077	0.0005	1.26077	4768.853	6035.374
A_23_P111	NM_01226	DnaJ (Hsp) DNAJB5	1.26076	0.00091	1.26076	816.5429	1025.93

A_23_P10	NM_17592	Homo sapi	ILDR1,MGI	1.26045	4.16E-06	1.26045	160651.6	201961.8
A_24_P35	XM_37331	Homo sapi	XM_37331	1.25999	0.00048	1.25999	13334.46	16907.61
A_23_P79	NM_05284	Homo sapi	EMILIN3,E	1.25965	0.00025	1.25965	6416.209	8045.17
A_23_P30	NM_01447	Chromosor	C10orf28	1.25928	0.00004	1.25928	540.9055	681.0898
A_23_P25	NM_01339	Homo sapi	USP25,US	1.25925	0.00017	1.25925	2152.159	2721.608
A_23_P47	NM_01533	Homo sapi	THRAP2,M	1.25899	1.17E-06	1.25899	1621.064	2042.422
A_23_P20	NM_01480	KIAA0435	PCNXL2,F	1.25898	0.00055	1.25898	2081.738	2625.187
A_32_P11	NM_01569	Homo sapi	PDZK6,PD	1.25786	0.00042	1.25786	864.1372	1082.56
A_23_P14	NM_00359	Homo sapi	TPST1	1.25764	1.98E-07	1.25764	1661.112	2089.634
A_24_P17	NM_00525	GA-binding	GABPB2,E	1.25696	0.00011	1.25696	588.9392	740.0704
A_32_P80	532	Unknown	A_32_P80	1.25694	0.00062	1.25694	754.2685	947.885
A_23_P10	BX647159	TSPY-like	TSPYL3	1.25688	0.00045	1.25688	10879.63	13650.65
A_24_P57	NM_01849	COBW-like	CBWD1	1.25645	0.0003	1.25645	1096.434	1382.954
A_23_P21	NM_01826	Homo sapi	ASXL2,AS	1.2564	0.00027	1.2564	3727.923	4703.478
A_23_P48	NM_00474	Homo sapi	CCPG1,CF	1.25616	0.00032	1.25616	4663.408	5887.068
A_23_P21	NM_03256	Homo sapi	KIAA2010,	1.25509	2.77E-08	1.25509	6693.362	8400.766
A_23_P25	NM_00084	Homo sapi	GSTA3,GT	1.25484	1.02E-06	1.25484	1742.589	2187.923
A_23_P21	AK096769	Lanosterol	LSS	1.25468	0.00032	1.25468	818.4016	1024.609
A_32_P13	AF086472	Homo sapi	AF086472	1.2545	0.00029	1.2545	486.1108	609.7803
A_24_P24	XM_29357	similar to	HLOC34474	1.25442	0.00014	1.25442	1741.745	2191.163
A_24_P26	NM_02065	sentrin/SUI	SENTP7,KI	1.25415	0.00003	1.25415	1703.376	2134.646
A_23_P16	NM_01630	Homo sapi	DNAJB11,I	1.25415	3.44E-06	1.25415	10962.18	13718.41
A_23_P25	NM_02510	hypothetic	CCDC2,CM	1.25396	7.35E-06	1.25396	1148.824	1442.237
A_32_P23	NR_00143	Homo sapi	HLA-H	1.2539	0.00083	1.2539	3795.838	4750.211
A_23_P32	NM_14458	Zinc finger,	ZFYVE27	1.25385	0.00003	1.25385	6441.473	8101.479
A_23_P12	NM_00127	Homo sapi	CHKA,CH	1.25356	0.00001	1.25356	3653.137	4581.288
A_23_P15	NM_02458	Homo sapi	EFTUD1,F	1.25351	2.89E-07	1.25351	2880.013	3611.021
A_24_P26	NM_00456	6-phosphol	PFKFB3,PI	1.25344	0.00018	1.25344	2339.161	2926.006
A_32_P95	NM_17213	Homo sapi	WFDC10B	1.25317	6.69E-07	1.25317	21775.81	27286.64
A_23_P87	NM_02072	Homo sapi	FAM62B,C	1.25299	0.00058	1.25299	4557.401	5720.251
A_23_P50	NM_00178	cell divisior	CDC2L1,p	1.25296	0.0008	1.25296	2987.046	3737.592
A_23_P20	1998	AF320996	THC19433	1.25274	0.00004	1.25274	7686.512	9666.883
A_23_P15	NM_00052	Homo sapi	HEXB,ENC	1.2527	7.96E-07	1.2527	7798.663	9758.925
A_23_P68	NM_02074	KIAA1271	KIAA1271	1.25193	0.00092	1.25193	4037.164	5073.138
A_23_P21	11603	Unknown	A_23_P21	1.25189	2.88E-08	1.25189	13146.02	16457.51
A_24_P9	14208	Unknown	A_24_P9	1.2518	0.00033	1.2518	7250.987	9027.988
A_24_P18	NM_02452	hypothetic	FLJ20986	1.25169	5.70E-07	1.25169	6642.616	8326.394
A_23_P97	NM_02215	Homo sapi	RRAGC,G	1.25169	0.00059	1.25169	2165.566	2724.705
A_23_P11	NM_00726	suppressor	HSGT1	1.25136	1.99E-07	1.25136	5822.681	7289.323
A_24_P17	0309	Unknown	A_24_P17	1.25064	0.00006	1.25064	3420.922	4292.249
A_23_P87	NM_00653	Homo sapi	YEATS4,G	1.25045	0.00004	1.25045	4014.515	5031.884
A_32_P22	AB014574	KIAA0674	KIAA0674	1.24981	0.00001	1.24981	1540.314	1928.704
A_24_P33	NM_01909	nudix (nucl	NUDT4,DIF	1.24922	0.00012	1.24922	5534.222	6936.308
A_23_P11	NM_00092	Homo sapi	PDE4C,DF	1.24911	0.00028	1.24911	44673.99	55535
A_23_P42	NM_18178	Homo sapi	HKR1	1.2491	0.00071	1.2491	410.887	513.2265
A_23_P12	NM_00271	Homo sapi	PPP2R2A,	1.24905	5.59E-06	1.24905	2815.608	3519.348
A_32_P21	AL049250	Homo sapi	DKFZp547	1.24881	0.0002	1.24881	1332.803	1669.724
A_23_P88	NM_00683	Homo sapi	PLEKHC1,	1.2486	1.33E-06	1.2486	1560.809	1947.662
A_23_P11	NM_01773	hypothetic	THUMPD1	1.24834	0.00005	1.24834	2953.33	3700.995
A_32_P15	NM_14571	Homo sapi	ATXN2L,A	1.2479	0.00047	1.2479	18346.01	22831.63
A_23_P42	NM_00374	Homo sapi	SOCS1,JA	1.24777	0.00002	1.24777	4034.191	5010.635
A_23_P11	NM_01804	hypothetic	BSDC1,FL	1.24765	0.00017	1.24765	4259.99	5320.666

A_23_P75:	NM_02122	Homo sapi	ARHGAP2	1.24741	0.00016	1.24741	1365.386	1703.085
A_24_P49687		Unknown	ENST0000	1.24707	6.95E-07	1.24707	131061.3	163214.3
A_23_P15:	NM_02213	golgi phosph	GOLPH3,C	1.24706	0.00074	1.24706	2773.029	3475.716
A_24_P25:	NM_00376	phosphopri	PEA15,PEI	1.2466	0.0003	1.2466	6552.839	8195.818
A_23_P16:	NM_01571	Misshapen	MINK1	1.24643	1.77E-06	1.24643	6730.908	8388.564
A_23_P11:	NM_00239	Homo sapi	MDK,MK,N	1.24643	0.00028	1.24643	12372.28	15475.74
A_32_P14:	AB029033	IQ motif an	IQSEC3	1.24631	0.00019	1.24631	7036.368	8723.545
A_23_P11:	NM_17492	hypothetic	LOC20189	1.24621	0.00006	1.24621	2681.597	3343.651
A_23_P15:	NM_01869	elaC (E.col	ELAC1,D2:	1.24619	0.00031	1.24619	901.797	1125.762
A_32_P53713		CLN3_HU	THC19788	1.246	0.00014	1.246	3598.694	4472.897
A_32_P73:	NM_17073	Homo sapi	BDNF,MG	1.24553	0.0006	1.24553	363.8953	453.2383
A_32_P122494		Unknown	A_32_P12:	1.24553	0.00006	1.24553	661.9334	824.4517
A_23_P20:	NM_00321	Homo sapi	TEGT,BI-1	1.2454	0.00002	1.2454	21430.38	26759.46
A_24_P10:	AK026491	Homo sapi	RPL12	1.24528	0.00024	1.24528	59029.62	73033.92
A_32_P34:	AB051482	Formin hor	FHOD3	1.24521	2.32E-06	1.24521	1436.177	1789.344
A_23_P30:	NM_00551	Major histo	HLA-E	1.24512	0.00014	1.24512	2046.825	2541.272
A_23_P11:	NM_01774	Homo sapi	TEX10,FLJ	1.24485	0.00021	1.24485	6029.214	7547.745
A_23_P72583		Unknown	A_23_P72:	1.24483	0.00003	1.24483	39124.73	48622.04
A_23_P66:	NM_00125	Homo sapi	CDC27,AP	1.24483	0.00022	1.24483	2656.942	3326.299
A_24_P135830		Unknown	A_24_P13:	1.24442	0.00014	1.24442	833.3579	1038.362
A_23_P10:	AK027211	Dynein, ax	DNAH3	1.24408	5.38E-06	1.24408	27050.41	33714.59
A_23_P11:	NM_00426	cofactor re	CRSP6,ME	1.24393	0.00075	1.24393	1207.672	1506.479
A_24_P92:	AK026155	Chromosom	C2orf17	1.24373	0.00004	1.24373	3215.311	3976.257
A_32_P11:	AK022044	Ribosomal	RPL10A	1.24341	5.53E-06	1.24341	1419.688	1764.014
A_32_P85:	AK056739	CDNA FLJ:	AK056739	1.24293	0.00046	1.24293	1998.974	2499.13
A_24_P323715		Unknown	ENST0000	1.24283	9.12E-07	1.24283	38799.01	48178.59
A_24_P35:	AK025305	CDNA: FL:	AK025305	1.24276	0.00062	1.24276	796.3401	991.9773
A_23_P37:	NM_14529	similar to	RLOC19733	1.24261	0.00071	1.24261	504.6578	627.2639
A_24_P71:	NM_20733	hypothetic	LOC15356	1.24245	0.00007	1.24245	1457.412	1815.464
A_32_P27209		Unknown	THC19258	1.2424	0.00097	1.2424	598.5309	742.7811
A_23_P40:	NM_02019	Homo sapi	C7orf36,GI	1.24215	0.0005	1.24215	3105.567	3871.754
A_32_P99:	NM_00227	karyopherin	TNPO1,MII	1.24208	2.72E-06	1.24208	17315.17	21533.91
A_23_P40:	NM_17856	Homo sapi	NM_17856	1.24193	0.00009	1.24193	655.0886	813.53
A_24_P40:	NM_01742	cat eye syr	CECR1,IDI	1.24137	0.00024	1.24137	10654.57	13207.67
A_24_P31:	NM_01475	mastermin	MAML1,Me	1.24134	0.00011	1.24134	6994.022	8675.75
A_23_P25:	NM_00708	Homo sapi	WDHD1,AI	1.24083	0.00033	1.24083	2451.988	3060.283
A_32_P19:	AB033066	KIAA1240	KIAA1240	1.24081	0.00004	1.24081	2528.864	3133.432
A_32_P61:	BC036622	CDNA clon	BC036622	1.24049	0.0002	1.24049	960.8993	1194.168
A_23_P16:	NM_01807	Homo sapi	RBM28,FL	1.2402	0.00033	1.2402	4905.522	6106.601
A_24_P358215		Unknown	A_24_P35:	1.24004	0.0008	1.24004	1553.107	1932.732
A_23_P11:	NM_00336	Homo sapi	USP1,UBP	1.23938	1.83E-07	1.23938	15177.21	18807.4
A_24_P15:	NM_00060	serine (or	SERPINE1	1.23924	8.77E-06	1.23924	1816.904	2245.561
A_24_P30:	NR_00003	Homo sapi	RAB9P1	1.23912	3.44E-06	1.23912	1058.807	1311.738
A_23_P33:	NM_17382	hypothetic	MGC3951E	1.23896	0.00083	1.23896	758.1141	940.9594
A_23_P10:	NM_03144	Homo sapi	C18orf21,X	1.23878	0.00086	1.23878	2876.821	3577.465
A_23_P13:	NM_05295	Homo sapi	DNTTIP1,T	1.23832	8.58E-07	1.23832	5311.099	6574.354
A_24_P50:	NM_01748	telomeric r	TERF1,TR	1.23782	1.47E-06	1.23782	3854.235	4774.417
A_24_P192338		AF167441	THC18072	1.23777	5.56E-06	1.23777	922.9686	1142.524
A_23_P21:	NM_03137	heterogene	HNRPDL,F	1.23776	0.00004	1.23776	10125.92	12499.95
A_23_P13:	NM_19858	Homo sapi	SLC27A1,F	1.23768	5.97E-07	1.23768	37973.34	46979.43
A_23_P65:	NM_02248	BTB/POZ	ENC2,FLJ:	1.23671	0.00016	1.23671	3984.189	4920.907
A_32_P219554		Unknown	A_32_P21:	1.23623	0.00002	1.23623	878.0188	1085.234

A_23_P15(XM_37243 Olfactory r	ENST0000	1.23584	0.00071	1.23584	3414.094	4198.227
A_23_P50(NM_13863 Homo sapi	BCL2L12,1	1.23579	0.00004	1.23579	10315.62	12776.78
A_23_P78(NM_00705 Homo sapi	SFRS16,C	1.23559	0.00029	1.23559	5725.962	7077.937
A_32_P11(BX537851 Hypothetic	RP11-19J3	1.23553	2.23E-06	1.23553	2165.838	2674.274
A_24_P50157	Unknown A_24_P50	1.23536	0.00073	1.23536	95707.45	117457.5
A_23_P11(NM_08086 Homo sapi	ASB16,FL	1.23512	0.00003	1.23512	10056.2	12441.81
A_23_P11(AF116688 Homo sapi	PRO2133	1.23479	0.00013	1.23479	6051.973	7479.311
A_23_P30(AK023110 HSPC182	HSPC182	1.23473	6.12E-06	1.23473	2794.222	3454.902
A_23_P16(NM_15228 Homo sapi	ZFP276,Z	1.23408	0.00012	1.23408	1133.31	1397.384
A_23_P12(AK090619 KIAA0241	KIAA0241	1.23376	0.00023	1.23376	758.1617	935.5721
A_23_P40(NM_15337 Homo sapi	LN2,PDZ	1.23368	0.00058	1.23368	1041.246	1289.737
A_23_P17(NM_00628 Homo sapi	TFPI,EPI,L	1.23351	0.00003	1.23351	13029.74	16120.41
A_23_P14(NM_00318 Homo sapi	ADAM17,T	1.23323	0.00002	1.23323	2815.129	3469.56
A_23_P49(NM_00450 homeo box	HOXB7,HC	1.23306	0.00012	1.23306	3110.868	3822.835
A_32_P92(AK092715 Homo sapi	AK092715	1.233	0.00001	1.233	4186.706	5150.838
A_23_P32(NM_02023 x 009 prote	DTWD1,MI	1.23275	7.42E-07	1.23275	2734.588	3370.305
A_23_P32(NM_00243 Homo sapi	MNAT1,M	1.23262	0.00055	1.23262	4298.577	5320.722
A_23_P15(NM_03341 Homo sapi	ZNF622,ZF	1.23257	0.00015	1.23257	4092.402	5060.44
A_23_P48(NM_01540 Homo sapi	DKFZP586	1.23229	0.0004	1.23229	1273.031	1571.073
A_32_P98(NM_02248 hypothetic	FLJ22405,I	1.23211	2.72E-07	1.23211	5138.757	6331.545
A_32_P82(BC029651 CTD-bindir	KIAA1542	1.23204	0.00005	1.23204	21780.51	26897.98
A_24_P10(NM_01836 LIN-7 prote	LIN7C,VEL	1.23201	0.00024	1.23201	1654.518	2043.018
A_23_P78(NM_00557 lectin, man	LMAN1,MF	1.23128	0.00003	1.23128	3742.839	4613.722
A_23_P39(NM_01211 CCR4-like	CCRN4L,N	1.23125	0.00019	1.23125	15037.32	18558.18
A_23_P94(NM_17503 Homo sapi	ST6GALN/	1.23117	2.22E-06	1.23117	2784.102	3430.812
A_23_P13(NM_02277 Homo sapi	TBC1D15,I	1.23108	0.00098	1.23108	3130.292	3863.605
A_23_P37(NM_00656 cell growth	CGRRF1,C	1.23005	0.001	1.23005	1534.278	1897.754
A_24_P10(XM_37757 similar to	6 LOC40194	1.22995	3.95E-07	1.22995	17750.4	21828.86
A_24_P23(NM_00614 dynein, cy	DNCLI2,LI	1.22942	0.00002	1.22942	956.0804	1175.129
A_23_P31(NM_00608 tubulin, bet	TUBB2	1.22915	2.57E-06	1.22915	15913.21	19540.94
A_24_P14(NM_01632 Homo sapi	NUP98,AD	1.22844	0.0005	1.22844	705.0817	865.5949
A_23_P96(NM_00614 Homo sapi	DNCLI2,LI	1.228	0.00003	1.228	8225.055	10121.77
A_23_P10(NM_14525 LOC12449	LOC12449	1.22784	0.00001	1.22784	2801.675	3434.886
A_23_P95(NM_00211 Major histo	HLA-C	1.22771	0.00004	1.22771	14305.08	17567.38
A_24_P11(NM_14529 Homo sapi	ZNF626	1.22749	0.00006	1.22749	1124.359	1380.37
A_24_P25(NM_19855 Homo sapi	NM_19855	1.2267	0.0002	1.2267	698.3608	857.0636
A_23_P35(AK057602 Ribosomal	RPL12	1.22657	0.00018	1.22657	229408.3	280163.1
A_24_P66(NM_00660 stromal ant	STAG2,SA	1.22642	0.00008	1.22642	3828.553	4708.049
A_24_P88(AK026213 Homo sapi	WDR42A,F	1.22626	0.00002	1.22626	1225.133	1503.13
A_24_P74(AI140519 LIM domai	LMO2	1.22572	0.00072	1.22572	1670.746	2053.312
A_23_P34(NM_01504 Homo sapi	KIAA0261,	1.22506	0.00036	1.22506	1941.498	2385.603
A_24_P28(BC008094 Retinal G	p RGR	1.22505	0.00002	1.22505	1354.147	1657.605
A_24_P19(NM_18256 hypothetic	MGC29814	1.22448	0.00031	1.22448	5679.899	6970.346
A_23_P69(NM_00024 Homo sapi	MLH1,FCC	1.2241	0.0003	1.2241	16569.81	20237.61
A_24_P56(AK023803 ADP-ribosy	ARF1	1.22396	0.00058	1.22396	3384.857	4127.514
A_23_P27(NM_00157 Homo sapi	IRF3	1.22383	3.39E-06	1.22383	1126.772	1378.978
A_23_P77(NM_01246 TERF1 (TF	TINF2	1.22375	9.73E-06	1.22375	6813.138	8333.987
A_24_P585004	Unknown A_24_P58	1.2237	5.32E-06	1.2237	2498.057	3059.19
A_24_P92(XM_29283 Homo sapi	ZNF66,FL	1.22361	0.00089	1.22361	1516.837	1848.126
A_23_P41(NM_01517 Homo sapi	BAT2D1,X	1.22359	0.00015	1.22359	5817.808	7102.077
A_23_P26(NM_01442 copine VII	CPNE7,MC	1.2235	8.71E-06	1.2235	4175.589	5106.964
A_23_P25(NR_00154 Homo sapi	TTY14	1.22309	1.45E-06	1.22309	3217.911	3936.026

A_24_P75: NM_03354 myosin reg MRLC2,ML	1.22298	0.00008	1.22298	27939.2	34084.33
A_23_P37: AF147723 Homo sapi LSR68	1.22283	0.0002	1.22283	1078.14	1319.394
A_23_P25: NM_00510 Homo sapi ENDOGL1	1.2228	0.00006	1.2228	754.0508	922.1739
A_23_P43: NM_05819 Homo sapi CDKN2A,A	1.2225	0.00017	1.2225	6990.925	8572.706
A_23_P30: AF211119 Cyclin-dep CDKN2A	1.22209	0.00055	1.22209	2207.465	2707.389
A_23_P31: NM_00629 ubiquinol-c UQCRB,QI	1.22209	0.00014	1.22209	8635.535	10592.62
A_24_P38: NM_03287 hypothetic TNFRSF19	1.22202	0.00018	1.22202	14654	17892.44
A_24_P25: XM_37331 similar to ri LOC39238	1.22175	0.00008	1.22175	10590.02	12985.81
A_23_P87: BC025767 Rhophilin, LRHPN1	1.22171	0.00005	1.22171	8262.138	10067.99
A_24_P51: NM_02011 hypothetic LARS,LEU	1.22157	0.00062	1.22157	8840.313	10792.04
A_23_P14: NM_03189 Homo sapi FTHL17	1.22138	0.00091	1.22138	97111.98	119050.5
A_23_P13: NM_00531 granzyme IGZMM,ME	1.22119	0.00012	1.22119	5522.726	6716.112
A_23_P21: NM_15285 Homo sapi WTAP,MG	1.22009	0.00038	1.22009	2282.282	2795.265
A_24_P34: NM_00287 RAB intera RABIF,MS	1.2198	0.00062	1.2198	3380.15	4135.155
A_23_P39: NM_00627 Homo sapi SFRS7,9G	1.21979	6.01E-06	1.21979	36965.29	45126.61
A_23_P22: NM_15257 hypothetic FLJ38564,I	1.21961	0.00039	1.21961	870.7285	1062.698
A_23_P79: NM_15363 Homo sapi PANK2,HS	1.2196	0.00006	1.2196	2962.165	3618.365
A_32_P20: AK128772 PI-3-kinase KIAA0220	1.2196	0.00021	1.2196	20339.19	24847.74
A_23_P36: NM_01480 KIAA0528 KIAA0528	1.2191	0.00065	1.2191	2248.896	2752.443
A_23_P69: NM_01856 hypothetic C4orf16,GI	1.21899	0.00004	1.21899	1305.778	1591.169
A_23_P21: AB018307 SPTF-assc STAF65(ga	1.21883	0.00005	1.21883	6614.727	8078.334
A_32_P18523 Unknown A_32_P18:	1.21878	0.00013	1.21878	17627.66	21455.77
A_23_P57: NM_01780 Homo sapi CKLFSF6,I	1.21857	0.00041	1.21857	11796.43	14424.26
A_23_P21: NM_15287 Homo sapi DGKD,dgk	1.21847	0.00017	1.21847	5689.604	6918.24
A_24_P69: XM_37170 hypothetic LOC38922	1.21787	0.00053	1.21787	1055.661	1283.697
A_23_P44: NM_03305 Homo sapi KRTAP4-1	1.21764	0.00095	1.21764	5669.46	6862.353
A_24_P102043 Unknown ENST0000	1.21708	0.00017	1.21708	8859.743	10831.62
A_23_P25: NM_02478 Homo sapi ZDHHC11,	1.21667	0.00002	1.21667	2567.775	3121.077
A_23_P21: NM_00097 Homo sapi RPL12	1.21661	0.00006	1.21661	349194.6	423934.2
A_23_P20: NM_08042 Homo sapi PTPN2,PT	1.21622	6.44E-06	1.21622	3047.761	3707.716
A_24_P29: NM_17380 Homo sapi BLOC1S2,	1.21551	0.00022	1.21551	10860.6	13241.88
A_23_P15314 PPRB_HUI THC19795	1.21518	0.00004	1.21518	1886.4	2289.205
A_23_P69: NM_00115 Homo sapi ANXA5,PP	1.21501	9.34E-06	1.21501	23366.05	28416.86
A_23_P12: NM_01790 PAK1 inter: PAK1IP1	1.21498	0.00033	1.21498	3794.088	4636.158
A_23_P34: NM_17216 Homo sapi NASP,FLB	1.2145	0.00001	1.2145	5699.686	6916.51
A_23_P31: NM_00391 G10 proteir G10,EDG2	1.21444	0.00001	1.21444	5277.439	6403.116
A_23_P25: NM_00289 Homo sapi RBBP8,RIP	1.2141	0.00016	1.2141	4765.625	5802.359
A_32_P10: AX721252 Sequence :AX721252	1.21373	0.00005	1.21373	51533.07	62406.93
A_24_P17: NM_00484 APG5 (aut: ATG5,ASP	1.21333	0.0005	1.21333	627.1144	760.7257
A_23_P21: NM_01829 Homo sapi NGLY1,PN	1.21306	0.00022	1.21306	12430.97	15118
A_23_P41: AK027663 Stanniocalc STC2	1.2128	0.00005	1.2128	1052.543	1276.073
A_23_P61: NM_00445 Homo sapi ETFDH,MA	1.21273	7.88E-06	1.21273	4616.62	5595.206
A_24_P32: NM_00552 heterogene HNRPH1,H	1.2127	0.00011	1.2127	20717.13	25160.82
A_23_P13: NM_01396 Neuregulin NRG1	1.21251	0.00028	1.21251	985.8541	1196.531
A_23_P74: NM_18263 hypothetic FLJ39963	1.21247	0.00002	1.21247	56674.84	68764.97
A_23_P38: NM_15270 Homo sapi MGC26597	1.21243	0.00017	1.21243	1927.066	2336.771
A_23_P35: NM_00076 Homo sapi CYP3A7,C	1.21174	0.00018	1.21174	1033.253	1253.376
A_24_P21: NM_03194 oculospani TSPAN10,I	1.21168	3.13E-06	1.21168	11035.25	13370.22
A_24_P94: NM_19808 hypothetic LOC28400	1.21139	0.0006	1.21139	3556.158	4284.909
A_24_P16: AK096509 ADP-ribosy ARFIP1	1.21134	0.00017	1.21134	4225.157	5120.556
A_23_P16: NM_00567 Homo sapi TAF1C,SL	1.21123	0.00003	1.21123	3681.063	4462.598
A_32_P21: NM_14462 Homo sapi ZBTB8,BO	1.21113	0.00059	1.21113	1023.652	1238.075

A_23_P21	NM_03066	Homo sapi	SERPINB1	1.21067	0.00002	1.21067	15231.98	18431.9
A_32_P20	NM_05293	similar to h	PCMTD1,F	1.21061	0.0007	1.21061	580.4616	702.7333
A_23_P82	NM_00101	Homo sapi	RPS12	1.21045	0.00009	1.21045	270219.7	327961.7
A_23_P16	AK026132	PHD finger	PHF17	1.21038	0.00003	1.21038	4072.865	4925.765
A_23_P13	NM_15276	Homo sapi	C1orf62,M	1.21008	0.00005	1.21008	82195.55	99342.38
A_32_P92	NM_00678	SMA3	SMA3,b55	1.20996	0.00007	1.20996	1966.915	2380.393
A_24_P936282	Unknown	A_24_P93		1.20984	0.00014	1.20984	3307.227	3999.533
A_23_P58	NM_02289	Homo sapi	RANBP17,	1.20979	0.00018	1.20979	651.2404	787.9013
A_24_P82	AK024177	Hypothetic	AK024177	1.20912	0.00028	1.20912	753.6445	910.9575
A_24_P93	BC005231	Serine/thre	STK4	1.2091	0.00057	1.2091	1574.469	1903.331
A_23_P5512	Unknown	A_23_P55		1.20866	0.00098	1.20866	532.8326	643.9624
A_24_P76	AY358240	IQ motif an	IQSEC1	1.20842	0.00008	1.20842	10530.38	12695.32
A_24_P11	NM_00734	zinc finger	ZNF236,Z	1.20823	0.00002	1.20823	1219.931	1474.025
A_24_P925334	Unknown	A_24_P92		1.20806	7.38E-06	1.20806	3595.01	4345.037
A_23_P21	NM_00334	Homo sapi	UBE2H,UE	1.20784	0.00001	1.20784	3516.066	4247.416
A_23_P14	D79984	Suppressor	SUPT6H	1.20674	0.00035	1.20674	4143.161	5003.734
A_24_P36	NM_18188	Homo sapi	UBE2D3,U	1.20603	0.00001	1.20603	29664.28	35753.67
A_23_P24	NM_00347	Homo sapi	CSRP3,CL	1.20593	0.00021	1.20593	892.3276	1076.596
A_23_P16	NM_19845	Homo sapi	ZNF600,KF	1.20562	0.00006	1.20562	2315.384	2795.894
A_32_P22	NM_17589	hypothetic	FLJ37266	1.20423	0.00064	1.20423	4272.352	5136.917
A_23_P39	AK021860	Homo sapi	FBXO17,FI	1.20406	0.00036	1.20406	10563.49	12782.22
A_23_P36	AK024423	FLJ00012	FLJ00012	1.204	0.0001	1.204	859.4909	1034.967
A_24_P39	AW302767	xr55h07.x1	AW302767	1.20391	0.00019	1.20391	72045.35	86479.42
A_23_P96	NM_00253	Homo sapi	NVL	1.2038	0.00058	1.2038	9669.787	11665.45
A_32_P94	BC042589	Homo sapi	BC042589	1.20347	0.00012	1.20347	3893.75	4698.506
A_23_P11	NM_00272	Homo sapi	PRB4,Po	1.20311	0.00022	1.20311	5200.997	6278.455
A_23_P213468	Unknown	A_23_P21		1.20305	0.00007	1.20305	4816.312	5802.194
A_23_P34	NM_03191	Homo sapi	SETDB2,C	1.20304	0.00013	1.20304	2153.459	2589.336
A_24_P17	NM_02019	GK001 pro	GK001	1.20247	0.00017	1.20247	6688.69	8028.409
A_32_P11	NM_00203	Ferritin, he	FTH1	1.2024	0.00004	1.2024	351399.8	422096.7
A_23_P23	NM_02481	Homo sapi	C1orf82,FL	1.20235	0.0002	1.20235	4201.488	5033.43
A_32_P20	AK025669	Hypothetic	KIAA1991	1.202	0.00069	1.202	885.4993	1065.417
A_23_P13	NM_01807	hypothetic	FLJ10374	1.20187	0.00006	1.20187	1202.693	1445.146
A_32_P31	NM_01243	splicing fac	SF3B1,PR	1.20164	0.00006	1.20164	12055.12	14507.89
A_23_P25	NM_00295	Homo sapi	RSN,CLIP,	1.20152	0.00006	1.20152	6598.479	7920.634
A_23_P89	NM_02215	Homo sapi	FN3K	1.20151	0.00014	1.20151	1626.565	1957.975
A_23_P25	NM_00109	ATP-bindin	ABCF1,AB	1.20124	0.00045	1.20124	6773.071	8116.197
A_24_P912818	AJ251787	THC18588		1.20084	0.00023	1.20084	1014.439	1217.527
A_23_P12	NM_00417	Homo sapi	SREBF1,S	1.20013	0.00037	1.20013	51954.91	62414.17
A_23_P59	NM_00667	Homo sapi	TPBG,5T4,	1.20007	0.00002	1.20007	2866.804	3439.268
A_23_P97	NM_01459	Estrogen r	ERBP	1.19999	0.00009	1.19999	6561.331	7899.633
A_23_P33	AX721087	Sequence	AX721087	1.19933	0.00011	1.19933	3769.997	4516.111
A_24_P20	NM_02407	hypothetic	TMEM38A,	1.19908	0.00017	1.19908	8988.941	10786.2
A_24_P290314	Unknown	ENST0000		1.1988	0.00005	1.1988	38845.47	46516.85
A_32_P47	BX393727	Transcribe	BX393727	1.19864	0.00082	1.19864	725.4096	869.204
A_23_P25	NM_00361	Homo sapi	OFD1,71-7	1.19833	0.00022	1.19833	865.0039	1036.84
A_23_P14	NM_01642	Homo sapi	ABI3,NESt	1.19826	0.00047	1.19826	3702.709	4426.479
A_24_P152753	Unknown	A_24_P15		1.19805	0.00002	1.19805	67496.32	80927.25
A_23_P33	AK097598	Olfactory r	OR4N4	1.19801	0.00074	1.19801	783.479	939.2437
A_23_P208579	Unknown	A_23_P20		1.19677	0.00079	1.19677	695.6289	832.6361
A_24_P28	NM_08091	Homo sapi	DGUOK,dC	1.19646	0.00002	1.19646	6395.949	7649.852
A_23_P36	NM_00563	Homo sapi	SYT1,P65,	1.19591	0.00008	1.19591	975.1401	1166.117

A_23_P83:	NM_14505	Homo sapi	RNF183,M	1.19581	0.00004	1.19581	3395.003	4058.425
A_23_P370707		Unknown	A_23_P370	1.19512	0.00031	1.19512	3377.982	4040.832
A_32_P31355		Unknown	A_32_P31:	1.19494	0.00024	1.19494	1455.279	1740.74
A_24_P606663		Unknown	A_24_P60:	1.19446	0.00002	1.19446	71518.37	85398.5
A_23_P47220		Unknown	A_23_P47:	1.19443	0.0001	1.19443	23020.15	27460.63
A_23_P24:	NM_03212	hypothetic	DKFZP566	1.19364	0.00089	1.19364	6076.917	7240.511
A_23_P64184		Unknown	A_23_P64:	1.19337	0.00088	1.19337	19532.84	23359.49
A_23_P40:	NM_00025	Homo sapi	MUT,MCM	1.19335	0.00098	1.19335	1648.535	1969.3
A_23_P12:	NM_02015	Homo sapi	SARA1,SA	1.19304	0.00016	1.19304	23743.78	28367.26
A_24_P92771		Unknown	A_24_P92:	1.19284	0.0002	1.19284	250422.8	298155.4
A_23_P33:	NM_01414	CD274 ant	PDCD1LG	1.19192	0.00031	1.19192	4178.147	4989.021
A_23_P48:	NM_08074	Homo sapi	RNF36,Trif	1.19127	0.00022	1.19127	9055.041	10777.45
A_23_P20:	NM_02400	Homo sapi	GJB3,CX3	1.19121	0.00006	1.19121	3710.331	4417.617
A_32_P82:	NM_00203	Ferritin, he	FTH1	1.19084	0.0006	1.19084	275701.8	329096.4
A_23_P21:	NM_00200	Homo sapi	FGF2,BFG	1.18933	0.00005	1.18933	2395.497	2848.07
A_23_P70:	NM_01462	Trace amin	GPR58	1.18867	0.00007	1.18867	2930.292	3479.754
A_24_P306527		Unknown	A_24_P30:	1.18862	0.00005	1.18862	65027.52	77344.73
A_23_P20:	NM_02462	Homo sapi	ZC3HAV1,	1.18854	0.00045	1.18854	2435.878	2891.242
A_32_P79152		Unknown	THC19758	1.18814	0.00028	1.18814	30257.97	35940.11
A_32_P13:	NM_01654	Ras-associ	RBJ,RabJ	1.18805	0.00006	1.18805	4217.597	5006.711
A_23_P85:	NM_00064	interleukin	IL13RA2,IL	1.18784	0.00006	1.18784	47707.44	56630.27
A_23_P47:	NM_00428	Homo sapi	BAG3,BIS,	1.1865	0.0009	1.1865	1798.748	2135.99
A_32_P39:	U79275	Hypothetic	HSU79275	1.18645	0.00009	1.18645	6022.542	7156.11
A_23_P35:	NM_00428	Homo sapi	BAG2,BAG	1.18611	0.00094	1.18611	10282.24	12236.27
A_23_P59:	NM_01621	Homo sapi	HECA,HDC	1.18609	0.00038	1.18609	1944.551	2304.844
A_32_P32:	BC045677	Hypothetic	FLJ25477	1.18582	0.00049	1.18582	5408.351	6440.217
A_23_P15:	NM_00314	Homo sapi	SSB,LARP	1.18543	0.00026	1.18543	47353.25	56248.95
A_24_P10:	NM_03278	Homo sapi	RSPO3,PV	1.18424	0.0002	1.18424	1849.168	2189.442
A_23_P25:	NM_05306	Homo sapi	UBQLN1,D	1.18403	0.00036	1.18403	5961.551	7069.567
A_23_P29:	NM_03328	Homo sapi	MRPS36,D	1.18359	0.00006	1.18359	17799.16	21056.83
A_23_P76:	NM_00596	Homo sapi	NAB2,MAC	1.18199	0.00008	1.18199	13238.02	15644.63
A_23_P12:	NM_14841	Homo sapi	ATXN2L,A:	1.18198	0.00034	1.18198	8486.458	10041.68
A_23_P31:	NM_15334	Ataxin 7-lik	ATXN7L2	1.18184	0.0004	1.18184	2518.056	2973.158
A_24_P11:	NM_01513	KIAA0246	STAB1,FEI	1.18184	0.00065	1.18184	2645.479	3119.314
A_23_P16:	NM_00037	Homo sapi	VDR,NR1:	1.1812	0.00028	1.1812	3762.121	4448.106
A_24_P41:	NM_00736	non-POU-c	NONO,P54	1.17969	0.00026	1.17969	6534.626	7715.51
A_23_P10:	NM_02192	Homo sapi	SCT	1.17948	0.00046	1.17948	979.651	1155.385
A_24_P14:	NM_02210	G-protein s	GPSM3	1.17932	0.0002	1.17932	19696.61	23219.82
A_23_P10:	NM_01432	Homo sapi	ORC6L,OF	1.17868	0.00045	1.17868	4359.237	5132.782
A_23_P15:	AL834362	GTPase ac	GARNL1	1.17865	0.00056	1.17865	1524.553	1798.89
A_23_P14:	NM_03229	Homo sapi	ANKRD32,	1.17804	0.0001	1.17804	3430.284	4041.001
A_23_P15:	NM_01402	LR8 proteir	LR8,FLJ26	1.17708	0.00008	1.17708	2493.008	2934.507
A_23_P30:	NM_19850	HGS_RE4(UNQ1912,I	1.17707	0.00018	1.17707	2465.363	2903.183
A_23_P70:	NM_01763	Homo sapi	FAM46A,X	1.17701	0.00012	1.17701	14671.88	17276.89
A_23_P25:	NM_01832	Homo sapi	BXDC2,BR	1.17679	0.00006	1.17679	14943.34	17585.85
A_24_P93:	M64259	Major histo	HLA-B	1.17634	0.00087	1.17634	1789.623	2107.47
A_23_P14:	NM_00193	membrane	MPP3,DLC	1.17633	0.00044	1.17633	9644.222	11318.67
A_23_P13:	NM_00067	Homo sapi	ADRA1D,C	1.17633	0.00044	1.17633	11209.35	13158.14
A_24_P25:	AK092418	Fc fragmer	FCGR1A	1.17591	0.00081	1.17591	2725.43	3201.292
A_23_P31:	NM_17371	Homo sapi	NM_17371	1.1756	0.00007	1.1756	9251.738	10875.25
A_23_P93:	NM_00543	Homo sapi	ARHGEF5	1.17556	0.00026	1.17556	1998.738	2350.781
A_32_P15:	AB037847	Chromosor	C16orf34	1.17519	0.00099	1.17519	2883.9	3394.087

A_23_P34	NM_00100 Homo sapi	RPL39	1.1749	0.00035	1.1749	131666.7	154956.9
A_24_P17	NM_00667 MHC class HCP5,P5-1		1.17462	0.00022	1.17462	5666.73	6646.897
A_24_P22	AK057676 RNA bindir	RBM18	1.17346	0.00018	1.17346	1557.498	1827.714
A_32_P12	AW938928 PM4-DT00	AW938928	1.17245	0.00057	1.17245	1092.321	1280.653
A_32_P20	NM_13346 Homo sapi	CITED4	1.17202	0.00015	1.17202	8042.854	9426.036
A_24_P11	XM_20818 similar to l	LOC28423	1.17128	0.00063	1.17128	100548.2	117561.4
A_23_P86	BC048281 Zinc finger,	ZRANB1	1.17107	0.00022	1.17107	4743.642	5554.133
A_23_P129258	Unknown A_23_P12		1.17103	0.00061	1.17103	139613.8	163695.2
A_23_P36	NM_13326 WIRE prot	WIRE,WIC	1.17097	0.00063	1.17097	3008.38	3524.697
A_23_P25	NM_00628 Homo sapi	SSR4,TRA	1.17019	0.00049	1.17019	22355.39	26117.64
A_24_P91	L79986 Homo sapi	nuclear aut	1.16988	0.00017	1.16988	15825.17	18513.07
A_32_P16	AL833530 Nuclear fac	NFE2L1	1.16974	0.00036	1.16974	9318.053	10908.8
A_24_P10	BC022822 Immunoglo	IGLC2	1.16962	0.00019	1.16962	26960.39	31532.93
A_24_P17	NM_01799 hypothetic	FLJ10081,I	1.16938	0.00049	1.16938	1333.324	1559.614
A_23_P34	NM_15275 hypothetic	FLJ23834	1.1692	0.00015	1.1692	3815.358	4460.347
A_32_P71437	ALU5_HUM	THC19106	1.16875	0.00011	1.16875	17239.94	20149.63
A_24_P20	NM_00590 MAD (motr	SMAD2,JV	1.16846	0.00052	1.16846	1761.556	2059.312
A_24_P30	NM_01326 transcriptio	HCNGP,H	1.16837	0.00053	1.16837	1253.542	1464.862
A_24_P56	BC035751 Homo sapi	TAF4A5	1.16837	0.00093	1.16837	21618.73	25264.99
A_23_P36	NM_01525 Homo sapi	UBR2,C6o	1.1679	0.00039	1.1679	2643.88	3089.355
A_24_P25	NM_00608 tubulin, bet	TUBB3,MC	1.16784	0.0003	1.16784	22995.7	26870.91
A_32_P127492	Unknown A_32_P12		1.16759	0.00024	1.16759	1809.344	2112.438
A_23_P25	NM_00699 Homo sapi	SCGN,SEC	1.16712	0.00016	1.16712	4040.962	4716.668
A_23_P55	NM_15235 Homo sapi	ZNF440,FL	1.16659	0.00073	1.16659	2301.037	2685.649
A_24_P362100	SKI_HUM	THC18331	1.16483	0.00019	1.16483	2463.303	2869.337
A_24_P38	NM_02039 eukaryotic	EIF5A2,EIF	1.16475	0.00054	1.16475	1307.338	1522.707
A_32_P9707	Unknown A_32_P97		1.16468	0.00016	1.16468	14419.38	16794.02
A_23_P11	NM_01541 Homo sapi	LETMD1,H	1.16462	0.00051	1.16462	4389.43	5110.823
A_32_P172198	Unknown A_32_P17		1.16401	0.0007	1.16401	27536.37	32091.8
A_23_P37	NM_02466 hypothetic	FLJ11506,I	1.16394	0.00039	1.16394	13137.09	15290.84
A_23_P16	NM_03193 Homo sapi	TBC1D10A	1.16332	0.00059	1.16332	3152.176	3670.343
A_23_P38	NM_00655 Homo sapi	C18orf43,F	1.16316	0.00063	1.16316	1591.759	1852.029
A_32_P125820	Unknown THC	18366	1.16281	0.00061	1.16281	1533.495	1783.248
A_23_P14	NM_00674 Homo sapi	RBM3,RNF	1.16184	0.00086	1.16184	19921.68	23120.88
A_32_P14	BX092209 BX092209	BX092209	1.16071	0.00092	1.16071	5000.543	5799.284
A_23_P58	NM_00482 Homo sapi	HAND1,Hx	1.16058	0.00079	1.16058	5028.965	5825.74
A_32_P11	AW901958 Protein kin	PRKCB1	1.15939	0.001	1.15939	1178.986	1366.72
A_24_P72	AK123446 CDNA FLJ	AK123446	1.15866	0.00091	1.15866	6223.57	7205.825
A_24_P91	AK022038 Similar to b	na,ARVP6	1.15859	0.00085	1.15859	14752.77	17074.58
A_23_P17	NM_00123 Homo sapi	CASQ1,CA	1.1584	0.00048	1.1584	7649.783	8856.745
A_24_P28	NM_17586 Homo sapi	ELYS	1.15826	0.00069	1.15826	2553.656	2958.033
A_24_P24	BC029274 CXYorf1 pr	CXYORF1	1.15801	0.00081	1.15801	8580.609	9930.959
A_23_P43	NM_03279 hypothetic	MFSD2,FP	1.15792	0.00096	1.15792	2672.863	3090.918
A_23_P34	NM_00218 Homo sapi	IL9R	1.15422	0.00038	1.15422	11865.55	13695.85
A_24_P50509	Unknown A_24_P50		1.15174	0.00061	1.15174	7913.179	9115.389
A_23_P15	NM_00015 Homo sapi	GAA,LYAC	1.15132	0.00061	1.15132	8245.911	9494.855
A_32_P187143	ALU1_HUM	THC18956	1.14935	0.001	1.14935	6193.877	7114.35
A_23_P42	NM_01421 homeo box	HOXD9,HC	1.14889	0.00068	1.14889	3545.056	4073.066
A_24_P32	BC010562 Homo sapi	MGC1541	1.14877	0.00096	1.14877	3904.793	4483.77
A_24_P358534	Unknown A_24_P35		1.14858	0.00073	1.14858	5472.298	6284.732
A_23_P13	NM_00646 Homo sapi	IVNS1ABP	1.14809	0.00064	1.14809	8244.716	9465.346
A_23_P39	AK022001 Latrophilin	LPHN1	1.14787	0.00079	1.14787	11826.34	13578.6

A_23_P26:	NM_01601	Homo sapi	NDUFAF1,	-1.14732	0.00098	0.8716	10326.88	9003.784
A_24_P27:	BC040043	Steroid rec	SRA1	-1.14807	0.00087	0.87103	2695.905	2348.084
A_24_P67408		Unknown	ENST0000	-1.14913	0.00077	0.87022	12418.33	10804.63
A_23_P21:	NM_00408	Homo sapi	TIMM8A,D	-1.14955	0.00065	0.86991	6471.866	5630.303
A_23_P80:	NM_15229	kleisin beta	hCAP-H2,3	-1.14958	0.00099	0.86988	4445.065	3865.969
A_23_P14:	NM_01434	Homo sapi	MTCH1,PS	-1.15021	0.00078	0.86941	22259.67	19346.82
A_23_P81:	NM_00186	Cytochrom	COX7A2	-1.15106	0.00091	0.86877	70561.9	61257.97
A_23_P20:	NM_00275	Homo sapi	MAP2K6,M	-1.15131	0.00071	0.86857	4509.938	3917.78
A_23_P21:	M74817	Tropomyos	TPM2	-1.15201	0.00069	0.86805	30787.71	26715.95
A_23_P30:	NM_00280	Homo sapi	PSMC2,S7	-1.15228	0.00054	0.86784	18511	16067.72
A_23_P15:	NM_02466	hypothetic	VPS37B,FI	-1.15295	0.00099	0.86734	1518.136	1316.727
A_23_P21:	NM_02212	Homo sapi	PERP,THV	-1.15313	0.00074	0.8672	8966.575	7779.428
A_24_P41:	NM_08066	similar to	RMGC19604	-1.15331	0.00068	0.86707	5561.586	4821.989
A_24_P15:	NM_00615	neural prec	NEDD8,Ne	-1.15432	0.00052	0.86631	13665.92	11834.86
A_23_P29:	NM_02068	Homo sapi	C3orf14,H1	-1.15446	0.00054	0.86621	7965.047	6897.053
A_23_P16:	XM_37482	Homo sapi	PDCD11	-1.15493	0.00037	0.86585	10128.98	8769.939
A_32_P13:	NM_20700	Homo sapi	FAM45A	-1.15504	0.00084	0.86577	2161.8	1872.147
A_23_P29:	NM_01217	Homo sapi	FBXO7,FB	-1.15549	0.00059	0.86543	6036.999	5225.54
A_32_P15:	NM_00098	CDNA seq	RPL17	-1.15564	0.00038	0.86532	72444.81	62699.15
A_23_P46:	NM_01834	Homo sapi	SLC29A3,E	-1.15695	0.00048	0.86434	2760.656	2386.178
A_23_P53:	NM_02089	Homo sapi	OSBPL5,O	-1.15696	0.00053	0.86433	4182.857	3614.98
A_24_P152845		Unknown	A_24_P15:	-1.15698	0.0004	0.86432	19269.59	16651.5
A_23_P63:	BC025793	Chromosor	C1orf31	-1.15711	0.00082	0.86422	59656.1	51522.96
A_24_P118341		Unknown	A_24_P11:	-1.15821	0.00068	0.8634	1560.76	1347.595
A_23_P61:	NM_01645	brain prote	C8orf30A,F	-1.15861	0.00076	0.86311	2049.899	1769.019
A_24_P41250		Unknown	A_24_P41:	-1.15869	0.00027	0.86304	168061.8	145055.8
A_23_P63:	NM_01544	DKFZP586	C1orf43,NI	-1.15871	0.00037	0.86303	20340.61	17551.02
A_24_P25:	NM_05286	hypothetic	C20orf72,F	-1.15912	0.00071	0.86272	1596.438	1377.317
A_23_P14:	NM_00186	cytochrom	COX4I1,CC	-1.15995	0.0008	0.86211	32176.04	27713.64
A_23_P74:	NM_01871	RCC1-like	RCC2,TD-(-1.15997	0.00039	0.86209	4061.747	3501.321
A_24_P370484		Unknown	A_24_P37:	-1.15997	0.00056	0.86209	1883.407	1623.516
A_23_P51:	NM_05294	Homo sapi	LRRC42,M	-1.1601	0.00097	0.86199	7328.24	6319.104
A_23_P14:	NM_00686	Homo sapi	RAB31,Ral	-1.16042	0.001	0.86175	2679	2307.178
A_23_P12:	NM_01604	Exosome c	EXOSC3	-1.16089	0.00065	0.86141	9356.082	8052.153
A_24_P19:	NM_00165	ADP-ribosy	ARF1	-1.16101	0.00048	0.86132	55006.73	47403.2
A_32_P16:	NM_15329	Homo sapi	FAM10A5	-1.16117	0.00083	0.8612	8566.85	7381.471
A_23_P54:	NM_02124	Homo sapi	SNX6,TFA	-1.16136	0.0008	0.86106	5981.224	5155.798
A_32_P49:	BC036909	Hypothetic	LOC28488	-1.16156	0.00062	0.86091	4583.277	3942.924
A_24_P21:	NM_00273	Homo sapi	PRKAR1B,	-1.16161	0.00084	0.86088	1477.281	1271.746
A_24_P109973		Unknown	A_24_P10:	-1.16236	0.00073	0.86032	130132.3	112113.9
A_23_P14:	BC030119	Chromosor	C14orf109	-1.16384	0.0003	0.85922	2280.17	1959.041
A_32_P99:	NM_14459	hypothetic	MGC29937	-1.16456	0.0002	0.85869	4392.595	3771.652
A_32_P21:	BC019241	Hypothetic	FLJ33297	-1.16476	0.00077	0.85854	1768.973	1517.816
A_24_P37:	NM_00455	nuclease s	YBX1,YB1	-1.16549	0.00018	0.85801	15177.15	13024.23
A_32_P67:	BX647764	Keratin ass	KRTAP4-7	-1.16633	0.00034	0.85739	8178.746	7016.421
A_32_P88:	NM_03226	hypothetic	AE2,DKFZ	-1.16652	0.00061	0.85725	1586.431	1359.797
A_32_P10:	NM_00098	Ribosomal	RPL24	-1.16804	0.00017	0.85614	119694.4	102513.7
A_23_P39:	NM_00023	Homo sapi	LIG1	-1.16805	0.00016	0.85613	4058.318	3474.783
A_23_P10:	NM_00141	Homo sapi	EIF2B1,EIF	-1.16851	0.00018	0.85579	11302.71	9670.723
A_23_P13:	NM_01773	GPI7 prote	GPI7,PRO	-1.16999	0.00023	0.85471	4037.448	3451.78
A_24_P16:	NM_03293	DC32	TMEM60,C	-1.17006	0.00012	0.85466	4195.8	3585.913
A_23_P25:	NM_00319	Homo sapi	TBCC,CFC	-1.17015	0.00039	0.85459	3615.96	3090.521

A_23_P11	NM_00514	DnaJ (Hsp	DNAJA3,TI	-1.17076	0.00019	0.85415	16475.8	14076.55
A_23_P74	D50923	KIAA0133	KIAA0133	-1.17082	0.00053	0.8541	2739.369	2340.627
A_23_P36	NM_17490	hypothetic	LOC14796	-1.17123	0.00047	0.85381	1823.581	1556.868
A_23_P16	NM_00453	Homo sapi	MYBPC2,M	-1.17124	0.00084	0.8538	2171.15	1853.686
A_32_P78	BC043390	Glucagon-I	GLP2R	-1.17153	0.00085	0.85358	8573.442	7325.044
A_24_P15	AK092522	Zinc finger	ZNF545	-1.17157	0.00094	0.85355	1100.136	939.0916
A_23_P78	NM_00143	Homo sapi	FBL,FIB,FL	-1.1716	0.00072	0.85354	83306.5	70983.98
A_23_P15	NM_03234	Homo sapi	C6orf125,M	-1.17233	0.00012	0.853	11400.46	9727.172
A_23_P52	NM_00271	Homo sapi	PPP2R1B,	-1.17287	0.00031	0.85261	2944.601	2511.313
A_23_P88	NM_00194	Homo sapi	DUT,dUTP	-1.17311	0.00051	0.85243	29752.28	25402.23
A_32_P37	AB051443	KIAA1656	KIAA1656	-1.17351	0.00066	0.85214	1363.892	1162.554
A_24_P27	NM_01551	DKFZP566	DKFZp566	-1.17357	0.00019	0.8521	1791.519	1526.521
A_23_P25	BC015230	Chromosom	C9orf115	-1.17357	0.00079	0.8521	1877.698	1598.878
A_32_P99	NM_00450	heat shock	HSF2,MGC	-1.17368	0.00058	0.85202	1592.932	1357.637
A_23_P38	AL833405	Transmem	TMEM16F	-1.17413	0.00025	0.85169	4355.307	3712.223
A_24_P33	595	AJ249551	THC18320	-1.17414	0.00026	0.85169	2379.938	2027.481
A_23_P13	NM_00290	reticulocal	RCN1,RCN	-1.17496	0.00072	0.85109	3886.646	3313.097
A_23_P15	NM_01535	Homo sapi	SCRIB,CR	-1.17503	0.00006	0.85104	26989.57	22969.59
A_23_P19	NM_03128	Homo sapi	SF3B5,SF	-1.17513	0.00093	0.85097	42719.02	36259
A_23_P98	NM_00639	Homo sapi	SSSCA1,p	-1.1752	0.00082	0.85092	6275.169	5335.815
A_24_P43	NM_01847	TPA regula	TPARL,TM	-1.17533	0.00015	0.85083	2942.169	2502.991
A_24_P12	NM_00248	NADH deh	NDUFA2,B	-1.17606	0.00033	0.8503	15438.43	13148.1
A_23_P25	NM_00460	Homo sapi	STX4A,p3	-1.17631	0.00023	0.85012	3396.287	2885.68
A_23_P31	NM_00327	transmemb	TSPAN4,N	-1.17633	0.00015	0.8501	7918.488	6729.545
A_24_P86	NM_01895	chromosom	C9orf9	-1.17654	0.00091	0.84995	2934.546	2492.332
A_23_P92	NM_00456	Homo sapi	PET112L,F	-1.17665	0.00027	0.84987	1191.995	1013.048
A_23_P12	NM_00281	Homo sapi	PSMD4,AF	-1.17684	0.00065	0.84973	14174.88	12036.31
A_23_P61	NM_13078	Homo sapi	RAB24	-1.17788	0.00028	0.84898	3610.916	3064.122
A_23_P14	NM_03148	Homo sapi	GRWD1,G	-1.17797	0.00024	0.84892	3236.33	2749.552
A_24_P32	NM_00185	solute carri	SLC31A1,(-1.17805	0.00083	0.84886	1173.78	995.8412
A_23_P10	NM_01649	hypothetic	LOC51255	-1.17829	0.00021	0.84869	17172.39	14581.08
A_23_P34	NM_03142	Homo sapi	MRPL9,L9	-1.17867	0.00005	0.84842	29235.89	24802.73
A_32_P16	AK021744	CDNA FLJ	AK021744	-1.17876	0.00011	0.84835	2269.846	1925.932
A_24_P29	NM_13080	hypothetic	LOC13361	-1.17887	0.00013	0.84827	1473.409	1249.86
A_23_P15	NM_01787	Homo sapi	ASB6,MGC	-1.17913	0.00034	0.84808	1869.885	1585.824
A_23_P20	NM_01795	Homo sapi	CDCA4,HE	-1.17915	0.00072	0.84807	1038.193	880.5883
A_23_P14	NM_00186	cytochrom	COX4I1,C	-1.17946	0.00045	0.84785	63448.04	53899.71
A_23_P11	NM_02023	Homo sapi	TNFSF5IP	-1.17947	0.00018	0.84784	4141.449	3510.514
A_23_P17	BC059410	Hypothetic	LOC28514	-1.17961	0.00039	0.84774	12154.61	10285.79
A_23_P27	NM_01554	DKFZP564	DKFZP564	-1.17994	0.00036	0.8475	1377.627	1167.858
A_23_P78	NM_03283	hypothetic	FLJ14775	-1.18002	0.00015	0.84744	5102.006	4326.66
A_24_P16	6789	ALU6_HUM	THC19528	-1.18012	0.00062	0.84737	1854.302	1570.552
A_23_P67	NM_00174	Homo sapi	CAPNS1,3	-1.18057	0.00097	0.84705	5792.069	4898.554
A_23_P71	NM_01752	Homo sapi	WBSCR22	-1.18087	0.00028	0.84683	7529.209	6374.661
A_23_P70	NM_00630	JTV1 gene	JTV1,P38,I	-1.18088	0.00018	0.84683	14523.93	12312.57
A_23_P31	NM_00591	malate deh	MDH2,MDI	-1.18102	0.0002	0.84673	19284.57	16311.27
A_23_P56	NM_00128	cytoskeletc	CKAP1,CC	-1.18148	0.00031	0.84639	7559.739	6405.683
A_23_P40	NM_13840	Homo sapi	C6orf51,bA	-1.18156	0.00007	0.84634	30916.71	26154.46
A_24_P34	NM_01500	SMART/HI	SPEN,MIN	-1.18248	0.00099	0.84568	1199.958	1014.601
A_23_P16	NM_01636	LPAP for Iy	ACP6,LPA	-1.18256	0.00095	0.84562	834.7864	705.9418
A_23_P41	NM_00489	Homo sapi	SEC22L1,E	-1.18267	0.00049	0.84554	3520.567	2974.052
A_24_P15	NM_01209	peroxiredo	PRDX5,PL	-1.18282	0.00046	0.84544	16077.57	13619.97

A_23_P21	NM_00474	cell cycle p TBRG4,CF	-1.18305	0.00021	0.84527	5806.168	4900.846
A_32_P202708	Unknown	A_32_P20	-1.18337	0.00033	0.84505	1210.134	1022.733
A_23_P36	NM_00354	Homo sapi HYAL3,LU	-1.18344	0.00015	0.845	1781.919	1505.495
A_23_P11	NM_01773	hypoxia-inc PH-4	-1.18347	0.00007	0.84497	14288.73	12072.58
A_24_P84	BC032295	Ribosomal RPL22	-1.18352	0.00008	0.84493	2880.713	2433.558
A_23_P45	NM_01216	Homo sapi FBXO2,FB	-1.18376	0.00072	0.84477	1780.315	1505.313
A_23_P13	NM_00418	Homo sapi UCHL1,PA	-1.18376	0.00012	0.84477	237551.9	200860.2
A_23_P50	NM_02157	breakpoint BCR,ALL,C	-1.18376	0.00019	0.84477	1376.427	1162.86
A_32_P166152	Unknown	A_32_P16	-1.18413	0.00094	0.8445	973.3724	822.082
A_23_P15	NM_03290	hypothetic MGC1428	-1.18445	0.00031	0.84427	42325.08	35678.43
A_23_P83	NM_00270	Homo sapi PPP1CB,P	-1.18447	0.00039	0.84426	2153.85	1817.101
A_23_P19	NM_00222	Homo sapi ITPR3,IP3	-1.18484	0.00019	0.84399	1584.85	1337.71
A_23_P13	NM_01401	Homo sapi ASCC3L1,I	-1.18498	0.00004	0.84389	14019.15	11833.18
A_23_P12	NM_00013	fumarylce FAH	-1.18509	0.00009	0.84382	5227.599	4412.695
A_23_P72	NM_00676	Homo sapi YWHAE,M	-1.18561	0.00074	0.84344	9679.113	8187.829
A_24_P37	NM_00215	Homo sapi HSPA4,RY	-1.18567	0.00027	0.8434	9507.084	8033.697
A_23_P35	NM_00313	Homo sapi SRI,SCN	-1.18608	0.0001	0.84311	33108.69	27889.7
A_23_P97	NM_02021	Homo sapi RNPEP,Dk	-1.18614	0.00006	0.84307	2144.451	1808.177
A_23_P21	BM805607	Caspase re CARD9	-1.18645	0.00033	0.84285	3796.857	3205.918
A_24_P75	AF274938	PNAS-13 PNAS-13	-1.1865	0.00099	0.84281	1759.844	1487.262
A_23_P44	NM_01438	linker for at LAT,LAT1,	-1.18679	0.00096	0.84261	847.4983	714.1739
A_24_P39	NM_02023	x 003 prote TNFSF5IP	-1.1869	0.00024	0.84253	2950.154	2486.567
A_23_P40	NM_13877	Homo sapi C9orf112,F	-1.18701	0.00009	0.84245	5160.242	4346.431
A_32_P73507	Unknown	THC18821	-1.18702	0.00073	0.84245	2599.73	2192.325
A_24_P12	NM_13880	hypothetic ZFAND2B	-1.18703	0.00011	0.84244	1432.776	1207.118
A_23_P16	NM_00035	Homo sapi ABCA4,FF	-1.18725	0.00003	0.84228	10831.75	9121.17
A_32_P70	NM_01233	transmemb TSPAN15,I	-1.18728	0.0007	0.84226	1251.607	1054.211
A_32_P222335	Unknown	A_32_P22	-1.18743	0.00008	0.84216	8420.281	7089.174
A_23_P11	AK025895	Hypothetic FLJ32731	-1.18755	0.00012	0.84207	2348.441	1976.557
A_23_P32	AB037814	KIAA1393 KIAA1393	-1.1878	0.00036	0.84189	57708.82	48672.7
A_24_P10	NM_14586	Homo sapi ANXA11,A	-1.18787	0.00066	0.84184	1366.479	1150.928
A_24_P32	NM_17851	hypothetic FLJ39421	-1.18798	0.00009	0.84177	2531.852	2131.071
A_23_P44	NM_00425	Homo sapi COX5A,VA	-1.18803	0.00003	0.84173	26957.82	22681.62
A_23_P16	NM_00368	KH-type sp KHSRP	-1.18809	0.00017	0.84169	27221.55	22901.35
A_32_P15	NM_00551	high-mobili HMG2,HI	-1.18827	0.00066	0.84156	78004.03	65785.01
A_23_P37	AK000815	Homo sapi C9orf63,FL	-1.18843	0.00062	0.84145	1160.954	977.6033
A_23_P12	NM_00564	Homo sapi TAF6,TAF	-1.18854	0.00014	0.84137	16062.67	13507.96
A_23_P68	NM_01338	ubiquinol-c UCRC,HSF	-1.1888	0.00004	0.84118	40265.8	33858.48
A_24_P13	NM_00429	ring finger j RNF14,AR	-1.18895	0.00007	0.84108	1710.418	1438.701
A_23_P81	NM_03057	hypothetic NDFIP1,N	-1.18951	0.00024	0.84068	5880.578	4933.632
A_32_P10100	Unknown	A_32_P10	-1.18961	0.00033	0.84061	66752.96	55966.32
A_24_P18	NM_01401	mitogen-ac MAPBPIP,I	-1.18964	0.00059	0.84059	11046.4	9256.894
A_23_P10	NM_13828	Homo sapi NUP35,MF	-1.18993	0.00098	0.84038	5933.77	4993.563
A_23_P91	NM_03277	Homo sapi MINA,MDI	-1.18995	0.00024	0.84037	1952.406	1641.745
A_24_P25	NM_01585	unknown p LOC51035	-1.19008	0.00024	0.84028	5460.542	4581.733
A_23_P13	NM_01791	Homo sapi C19orf24,F	-1.1901	0.00088	0.84027	5113.913	4303.657
A_24_P17	BC013295	Homo sapi HK2	-1.19011	0.00055	0.84026	947.2731	796.1603
A_23_P392897	Unknown	A_23_P39	-1.19015	0.00029	0.84023	3523.035	2960.156
A_23_P14	NM_00375	eukaryotic EIF3S5,eIF	-1.19029	0.00004	0.84013	4503.994	3782.881
A_23_P10	NM_02066	Homo sapi DECR2,PC	-1.19055	0.00052	0.83995	6274.057	5277.718
A_23_P55	AF217993	Hypothetic MGC1552	-1.1906	0.00004	0.83991	7331.702	6160.502
A_24_P358474	Unknown	A_24_P35	-1.19065	0.00023	0.83988	2637.262	2214.544

A_24_P12	NM_01418 HSPC160	ORMDL2, M	-1.19074	0.00011	0.83982	1110.537	932.6392	
A_24_P724106	Unknown	A_24_P72	-1.19084	0.0007	0.83974	31197.18	26136.67	
A_24_P707543	Unknown	A_24_P70	-1.19085	0.00015	0.83973	101004.2	84665.81	
A_24_P54	AK026687	Signal seq	SSR1	-1.1909	0.00036	0.8397	1320.267	1108.842
A_23_P25	NM_01838	hypothetic	FLJ11305, I	-1.1911	0.00003	0.83956	4235.522	3556.101
A_24_P16	NM_19933	similar to	RLOC37439	-1.19136	0.00016	0.83938	1387.958	1164.922
A_23_P20	NM_02429	leukocyte r	LENG4, BB	-1.19172	0.00009	0.83912	12190.63	10224.02
A_23_P39	AK124717	Purkinje ce	PCP2	-1.19199	0.00006	0.83893	14265.26	11976.98
A_23_P20	NM_00556	Homo sapi	STMN1, La	-1.19272	0.00098	0.83842	24939.82	20962.96
A_23_P34	AK090472	Kelch-like	KLHL21	-1.19284	0.00003	0.83833	6186.182	5187.165
A_23_P46	NM_00472	BUB3 (bud	BUB3, BUB	-1.19303	0.00004	0.8382	7701.639	6452.39
A_23_P10	NM_00608	Homo sapi	BPNT1, PIF	-1.19313	0.0003	0.83813	3817.971	3200.243
A_24_P57	NM_00068	S-adenosy	AHCY, SAH	-1.19321	0.00032	0.83807	1137.81	953.7639
A_24_P42	NM_00280	proteasom	PSMD2, S2	-1.19392	0.00005	0.83758	4098.144	3430.61
A_24_P38	NM_01603	CGI-92 pro	COQ4, CGI	-1.19394	0.0002	0.83756	1331.336	1114.751
A_24_P238896	Unknown	A_24_P23		-1.194	0.00003	0.83752	3323.289	2782.488
A_24_P152792	Unknown	A_24_P15		-1.19408	0.00035	0.83746	5208.745	4369.631
A_23_P20	NM_00263	Homo sapi	SERPINB5	-1.19415	0.00044	0.83742	875.147	732.7839
A_24_P234871	Unknown	A_24_P23		-1.19431	0.00002	0.8373	2281.032	1909.884
A_23_P42	NM_17360	Homo sapi	C15orf21, L	-1.19435	0.00049	0.83728	149807.8	125691.9
A_23_P34	NM_00311	Homo sapi	SP100, FLJ	-1.19472	0.00024	0.83701	108167	90751.63
A_23_P50	NM_02198	ubiquitin-cc	UBE2V1, C	-1.19476	0.00041	0.83699	6288.339	5270.533
A_23_P11	NM_02526	hypothetic	MGC2744	-1.19484	0.00005	0.83693	2787.207	2331.754
A_23_P36	NM_14579	Homo sapi	MGST1, MC	-1.19486	0.00033	0.83692	53624.7	44739.41
A_32_P17	AK057740	Laminin, al	LAMA5	-1.19487	0.00006	0.83691	1795.919	1502.928
A_23_P14	NM_02257	Homo sapi	VPS16, hVf	-1.19506	0.00016	0.83678	1823.883	1526.594
A_23_P35	NM_01902	Homo sapi	TMCO1, PC	-1.19506	0.00019	0.83678	17106.94	14285.64
A_32_P29	NM_00551	high-mobili	HMGN2, HI	-1.19515	0.00005	0.83671	52379.41	43793.33
A_32_P98	NM_00425	cytochrom	COX5A, VA	-1.19518	0.00001	0.83669	41497.37	34714.11
A_23_P93	NM_00591	Homo sapi	MCM7, MC	-1.19528	0.00006	0.83663	61267.41	51206.59
A_24_P30	NM_02068	SCY1-like	SCYL1	-1.19538	0.00079	0.83655	11301.61	9435.445
A_23_P72	NM_00020	Homo sapi	IF, FI, factor	-1.1954	0.00029	0.83654	2077.437	1734.828
A_23_P10	NM_01822	Homo sapi	MOBK1B, M	-1.19543	0.00015	0.83652	1069.947	895.1279
A_23_P50	NM_03215	Williams B	WBSCR20	-1.19545	0.00004	0.83651	27574.76	23052.11
A_23_P14	NM_00544	Homo sapi	PAPSS1, S	-1.19561	0.00053	0.83639	2372.044	1987.34
A_24_P14	NM_00538	protein-L-is	PCMT1	-1.19593	0.00071	0.83617	4900.488	4108.439
A_24_P16	NM_01437	replication	REPIN1, AF	-1.19641	0.00002	0.83584	3470.396	2900.926
A_23_P22	NM_01906	Homo sapi	GNL3L, FL	-1.19649	0.00016	0.83578	2021.174	1688.06
A_24_P18	NM_01824	hypothetic	TMEM30A	-1.19692	0.00025	0.83548	1042.491	870.8465
A_24_P39	NM_01323	like mouse	ATXN10, E	-1.19692	0.00035	0.83548	3760.849	3147.421
A_24_P229833	Unknown	A_24_P22		-1.19711	0.00001	0.83534	323082.6	269836.2
A_23_P99	NM_00398	Homo sapi	SLC7A7, LF	-1.19726	0.00002	0.83524	2850.812	2380.656
A_24_P19	NM_00267	promyeloc	PML, MYL, I	-1.19739	0.00009	0.83515	2913.77	2433.979
A_32_P11	AB029025	KIAA1102	KIAA1102	-1.19747	0.00004	0.83509	10142.6	8475.902
A_23_P70	NM_00720	Homo sapi	MRPL3, MF	-1.1976	0.00008	0.835	5867.918	4903.286
A_23_P21	NM_01602	Ubc6p hor	UBE2J1, UI	-1.19764	0.00014	0.83497	1001.199	836.0194
A_32_P91	BE779749	CDNA clon	BE779749	-1.19782	0.00011	0.83485	1193.748	996.7386
A_24_P35	AK057071	Homo sapi	AK057071	-1.19842	0.00006	0.83443	34178.77	28562.48
A_23_P20	D21262	Nucleolar	NOLC1	-1.1986	0.0001	0.83431	6116.085	5109.066
A_32_P69	NM_19427	protein con	MOPT, BLC	-1.19864	0.0006	0.83428	4874.621	4080.853
A_24_P161074	Unknown	A_24_P16		-1.19872	0.00002	0.83422	6716.486	5606.095
A_23_P60	AF152351	Homo sapi	TIMM13a	-1.1992	0.00001	0.83389	9486.654	7913.09

A_32_P47616	Unknown	A_32_P47616	-1.19927	0.0002	0.83384	1664.771	1388.361
A_23_P12	NM_00567	RNA bindir RBM10,MC	-1.19943	0.00001	0.83373	11247.13	9379.158
A_32_P19460	AP003240	THC19590	-1.19943	0.00009	0.83373	2566.357	2139.668
A_24_P80	AK024496	Kelch dom: JPH3	-1.19944	0.00041	0.83372	1119.089	933.2578
A_23_P20	NM_00229	Homo sapi LAMC1,LA	-1.19999	8.74E-06	0.83334	5177.166	4314.085
A_24_P32	BC067760	Mitogen-ac LOC40783	-1.20011	0.00083	0.83326	1498.539	1250.533
A_24_P20	NM_00645	multifunctic PAICS,AIR	-1.20061	0.00088	0.83291	10125.52	8455.686
A_23_P35	NM_14586	Homo sapi ANXA11,A	-1.20121	0.00015	0.83249	11097.5	9220.85
A_23_P21	NM_00488	ATP synth: ATP5J2,A1	-1.20143	0.0005	0.83234	28156.21	23508.68
A_23_P31	NM_17812	Homo sapi CXorf40A,I	-1.20152	0.00005	0.83228	6888.32	5736.266
A_32_P17	AK091332	CDNA FLJ: AK091332	-1.20165	0.00021	0.83219	3982.217	3319.253
A_23_P16	NM_02405	Homo sapi C17orf39,1	-1.20179	0.00014	0.83209	1219.002	1014.026
A_23_P13	NM_03232	K+ channe BTBD10,G	-1.20218	0.00052	0.83182	1518.35	1263.43
A_23_P41	NM_14460	Homo sapi NOXO1,P4	-1.20238	0.00041	0.83168	1361.869	1133.951
A_23_P15	NM_01810	Homo sapi ZDHHC4,Z	-1.20241	0.00002	0.83167	13146.14	10936.82
A_32_P82251	Unknown	A_32_P82251	-1.20248	8.92E-06	0.83162	2322.567	1931.496
A_24_P10	NM_00644	Homo sapi DRAP1,NC	-1.2026	0.00025	0.83153	2536.919	2112.173
A_23_P30	NM_00139	Homo sapi ECM2,MG	-1.20287	0.00026	0.83135	877.1785	729.3451
A_24_P11	AK022629	Homo sapi FLJ12567	-1.20294	0.0002	0.8313	1373.71	1142.195
A_23_P79	NM_00018	Homo sapi HADHB,H/	-1.20295	0.00095	0.83129	15503.57	12932.88
A_24_P63	NM_01783	nucleolar p NOLA2,NH	-1.20308	0.00022	0.8312	16187.07	13420.69
A_24_P280873	Unknown	ENST0000	-1.20335	0.00001	0.83102	2945.736	2447.966
A_23_P25	NM_01494	KIAA0843 ABLIM3,HI	-1.20345	0.00003	0.83094	1477.912	1227.933
A_24_P20	NM_00723	dynactin 3 DCTN3,DC	-1.20346	7.76E-06	0.83094	5529.427	4593.575
A_23_P20	NM_00240	Homo sapi MAP3K3,M	-1.20348	0.00012	0.83093	1048.147	871.0436
A_32_P41	NM_00551	high-mobili HMGN2,HI	-1.20359	5.09E-06	0.83085	164045.8	136307.8
A_32_P86009	Unknown	A_32_P86009	-1.20376	0.00075	0.83073	2548.436	2118.803
A_24_P27	NM_01902	Homo sapi TMCO1,PC	-1.20421	0.00022	0.83042	12307.54	10209.73
A_23_P13	AL832747	Guanine nt GNAQ	-1.20429	0.00021	0.83037	3954.652	3290.105
A_23_P11	NM_00634	Homo sapi ZNHIT1,CC	-1.20448	0.00074	0.83023	9117.958	7594.074
A_24_P93	NM_02475	hypothetic: AGMAT,FL	-1.2045	0.00003	0.83022	2696.287	2238.95
A_23_P10	NM_00458	Homo sapi RAB5C,RA	-1.20463	0.00025	0.83013	5311.399	4405.143
A_23_P46	NM_01607	Homo sapi BOLA1,CC	-1.20474	0.00016	0.83005	7274.688	6039.247
A_23_P32	NM_14516	Homo sapi SFT2D1,pF	-1.20477	0.00008	0.83003	13388.32	11118.62
A_23_P25	NM_00637	Homo sapi STK25,SO	-1.20478	0.00077	0.83003	3434.39	2858.598
A_23_P86171	FXD2_HU	THC18578	-1.20481	0.00001	0.83	2358.837	1957.617
A_24_P14	NM_00370	RuvB (E cc RUVBL1,R	-1.20495	0.0009	0.82991	1002.334	831.3518
A_24_P930707	Unknown	A_24_P930707	-1.20512	0.00089	0.82979	808.2143	670.8765
A_24_P94	NM_00632	ariadne (D: ARIH2,ARI	-1.2052	0.00009	0.82974	11298.59	9388.023
A_23_P65	NM_03285	Homo sapi ZFYVE19,I	-1.20546	0.00037	0.82956	3779.516	3133.596
A_24_P50639	Unknown	A_24_P50639	-1.20573	0.00014	0.82937	10059.64	8320.123
A_24_P27	AB032986	Homo sapi KIAA1160	-1.20577	0.00016	0.82935	1564.503	1296.969
A_23_P12	NM_00084	Homo sapi GSTM3,G	-1.20585	6.01E-06	0.82929	72601.35	60197.55
A_23_P18	NM_00699	polymerase POLS,POL	-1.20585	0.00014	0.82929	2349.591	1950.781
A_32_P26443	AF010144	THC19676	-1.20598	9.29E-06	0.8292	2021.726	1676.378
A_23_P13	NM_14456	Homo sapi RPIA,RPI	-1.20614	0.00003	0.82909	8234.115	6829.555
A_32_P19	BC035417	Homo sapi BC035417	-1.20616	0.0001	0.82907	1557.824	1291.197
A_23_P17	NM_00138	Homo sapi DPAGT1,C	-1.20656	0.00003	0.8288	1817.704	1506.82
A_23_P25	NM_01324	Homo sapi NXT1,P15,	-1.20658	8.03E-06	0.82879	15109.64	12526.54
A_23_P15	NM_00629	X-ray repai XRCC1,RC	-1.20684	0.00044	0.82861	4432.399	3680.179
A_24_P17	NM_00532	hydroxyacy HAGH,GLC	-1.20684	0.00003	0.82861	3393.298	2809.419
A_23_P38	NM_00642	Homo sapi RABAC1,P	-1.20692	0.00025	0.82855	10788.02	8952.97

A_23_P94	NM_01231	Homo sapi	LETM1	-1.20714	0.00004	0.8284	3439.155	2847.076
A_32_P60	NM_00300	succinate c	SDHD,PGL	-1.20744	0.00002	0.8282	3868.545	3206.817
A_24_P350136		Unknown	A_24_P350136	-1.2076	0.00042	0.82809	30048.04	24822.71
A_23_P91	NM_00195	Homo sapi	ECGF1,TP	-1.20774	0.00012	0.82799	3303.062	2737.593
A_24_P17	NM_00166	ras homolo	RHOA,ARF	-1.2078	0.00004	0.82795	3674.844	3045.161
A_32_P50005		C40201 art	THC18368	-1.20793	0.00085	0.82786	788.4055	652.5898
A_24_P18	NM_17812	hypothetic	LOC16242	-1.20797	0.00052	0.82784	980.2956	811.4502
A_24_P91	NM_03230	hypothetic	SPATA11,1	-1.20805	0.00001	0.82778	2144.06	1775.053
A_23_P16	NM_01536	S-phase 2	DERP6,HS	-1.20815	4.05E-06	0.82771	4721.926	3908.222
A_32_P12	AK001369	Chromosom	C6orf70	-1.20819	0.00007	0.82769	1653.068	1367.656
A_24_P23	BC009463	CDNA FLJ	BC009463	-1.2082	0.00016	0.82768	3130.372	2596.122
A_24_P83	NM_00327	transmemb	TSPAN6,T	-1.20852	0.00033	0.82745	2413.759	2002.167
A_23_P31	NM_02080	Homo sapi	WDR56,KI	-1.20884	0.00015	0.82724	936.6403	774.9332
A_23_P16	NM_02113	TNF recept	TRAF2,TR	-1.20887	0.00002	0.82722	1499.916	1240.915
A_23_P15	NM_01403	Homo sapi	BZW2,MS1	-1.20895	0.00031	0.82716	32651.96	26930.84
A_23_P31	AB095939	Chromosom	C14orf78	-1.20928	0.00015	0.82694	1272.674	1052.09
A_32_P34201		Unknown	A_32_P34201	-1.2096	3.08E-06	0.82672	52319.14	43249.55
A_24_P12	NM_17688	TR4 orpha	TRA16	-1.21023	4.98E-06	0.82629	2901.546	2397.767
A_24_P22	NM_01909	UDP glyco	UGT1A3,U	-1.21038	0.00008	0.82619	1405.06	1161.292
A_24_P75	XM_16725	similar to tr	LOC22187	-1.21065	0.00011	0.826	1919.104	1586.638
A_24_P178663		Unknown	A_24_P178663	-1.21069	0.00007	0.82597	25503.57	21089.73
A_24_P23	AB007963	KIAA0494	KIAA0494	-1.21071	0.00002	0.82596	1967.45	1624.802
A_23_P16	BC044842	Tetratricop	TTC19	-1.21091	0.00011	0.82582	1587.714	1312.404
A_23_P20	NM_00271	Homo sapi	PPP1CC,P	-1.21092	2.39E-06	0.82582	25743.47	21260.91
A_23_P32	NM_03190	SVH protei	SVH,MGC	-1.21122	0.00024	0.82561	6224.053	5150.03
A_23_P13	NM_00097	Homo sapi	RPL7A,TR	-1.21134	0.00002	0.82554	449363.3	370404.2
A_23_P90	NM_01857	amyotrophi	ALS2CR2,1	-1.21166	0.00006	0.82531	2320.802	1916.412
A_23_P5171		Unknown	A_23_P5171	-1.21188	0.0001	0.82516	2669.524	2203.24
A_23_P25	NM_00501	phosphate	PCYT1A,C	-1.21196	0.00004	0.82511	1218.055	1005.027
A_32_P19	NM_00182	CDC28 prc	CKS1B,CK	-1.2123	0.00009	0.82488	84807.18	70102.19
A_32_P2354		ALU8_HUM	THC19686	-1.21245	0.0004	0.82477	742.046	611.9635
A_23_P86	NM_00403	annexin A7	ANXA7,SN	-1.21275	6.63E-06	0.82457	7257.375	5981.371
A_23_P20	NM_00686	Homo sapi	RAB35,RA	-1.21278	6.09E-06	0.82455	5852.36	4823.934
A_23_P10	NM_01645	hypothetic	LOC51234	-1.21282	0.00011	0.82453	15014.44	12409.16
A_23_P59	NM_00309	Homo sapi	SNRPC,FL	-1.21282	1.94E-06	0.82452	29337.54	24188.34
A_23_P11	NM_17853	Homo sapi	LOC33879	-1.21303	0.00004	0.82438	4348.59	3589.89
A_23_P52	NM_01544	deleted in	RP11-529I	-1.21326	0.00015	0.82422	3094.273	2547.393
A_32_P19	AA167144	zp06e02.r1	AA167144	-1.21334	0.00054	0.82417	1346.224	1110.324
A_24_P19	NM_00682	tyrosine 3-	YWHAQ,1C	-1.21348	0.00008	0.82408	62050.92	50995.38
A_24_P575336		Unknown	A_24_P575336	-1.21365	0.00018	0.82396	27673.65	22862.25
A_23_P37	NM_03223	hypothetic	FLJ22875	-1.21384	0.00008	0.82383	42095.02	34739.43
A_23_P31	NM_01244	six transm	STEAP1,S	-1.21389	0.00098	0.82379	3312.064	2737.625
A_23_P20	NM_01233	Homo sapi	MYCBP,AM	-1.21389	0.00064	0.8238	5623.816	4646.878
A_24_P24	BC006177	Metastasis	MTA1	-1.21393	0.0002	0.82377	1036.147	854.0801
A_24_P418386		Unknown	ENST0000	-1.21394	0.00015	0.82376	1221.856	1006.847
A_23_P11	NM_02402	Homo sapi	MRP63,bM	-1.21431	2.80E-06	0.82352	14348.94	11813.55
A_23_P32	NM_17561	Homo sapi	NDUFA11,	-1.21459	0.00028	0.82333	15188.42	12553.31
A_23_P83	NM_00156	Homo sapi	ING2,ING1	-1.21472	0.00059	0.82324	1988.067	1639.971
A_23_P71	NM_01824	hypothetic	CCDC25,F	-1.21474	0.00028	0.82322	703.6284	579.2231
A_23_P20	NM_00002	presenilin 1	PSEN1,AD	-1.21488	0.00005	0.82313	1434.79	1181.187
A_23_P33	NM_00286	Homo sapi	PYGB,MGC	-1.21509	2.10E-06	0.82298	2983.369	2455.265
A_23_P36	NM_00165	Homo sapi	ARF3	-1.21539	0.00084	0.82278	750.5266	617.2995

A_24_P33	NM_18250	hypothetic	LOC14450	-1.21562	0.00003	0.82262	1415.234	1164.592
A_32_P94	BC059396	Hypothetic	MGC71735	-1.21562	0.00051	0.82263	1183.275	975.2603
A_32_P17	XM_37988	similar to H	LOC40256	-1.21565	0.00002	0.8226	78635.41	64582.34
A_23_P14	NM_01663	Homo sapi	SPG21,MA	-1.21593	5.91E-06	0.82242	15238.96	12534.32
A_24_P71	NM_17371	Homo sapi	NM_17371	-1.21608	0.00001	0.82232	7509.791	6178.313
A_23_P63	NM_00614	stratifin	SFN	-1.2161	0.00029	0.8223	1164.298	958.1334
A_32_P15	NM_01781	hypothetic	CHCHD3,F	-1.21612	4.67E-06	0.82228	3902.068	3209.567
A_23_P16	NM_02440	Homo sapi	NDUFS7,P	-1.21613	1.46E-06	0.82228	7708.552	6338.882
A_24_P74	NM_01537	chromosom	PGEA1,CE	-1.21623	0.00032	0.82222	661.1069	543.5917
A_24_P65	AK098360	Homo sapi	PKM2	-1.21635	0.00049	0.82213	1713.377	1408.509
A_24_P50489	Unknown	A_24_P50		-1.21643	0.00069	0.82207	26288.31	21709.34
A_23_P40	NM_15239	hypothetic	FLJ30834	-1.21646	0.00001	0.82206	2737.934	2252.397
A_23_P87	NM_00517	Homo sapi	ATP5G2	-1.21653	0.00001	0.82201	75605.48	62082.89
A_32_P75	BC034247	F-box prote	FBXO10	-1.21658	0.00005	0.82198	35773.01	29461.18
A_23_P43	D83781	Nucleopor	NUP160	-1.21675	0.00002	0.82186	3505.797	2881.255
A_23_P32	NM_00153	Homo sapi	HSBP1,NP	-1.21699	3.61E-06	0.8217	39387.55	32351.13
A_23_P56	BC015995	General tra	GTF3C3	-1.2171	0.00006	0.82163	2884.445	2369.573
A_23_P32	NM_00327	Homo sapi	TMPO,TP,I	-1.21735	0.00043	0.82146	2331.735	1921.8
A_23_P41	AK024480	Hypothetic	LOC12691	-1.21747	2.67E-06	0.82137	1993.183	1637.154
A_24_P73	NM_00168	ATPase, C	ATP2A2,DI	-1.21768	0.00039	0.82123	2553.417	2105.238
A_32_P36	BX436400	BX436400	BX436400	-1.21773	0.00003	0.8212	2327.791	1908.76
A_24_P93	NM_01782	hypothetic	COMMD4,I	-1.21783	0.00001	0.82113	3641.98	2988.796
A_23_P10	NM_01812	hypothetic	FLJ10534,I	-1.21786	0.00011	0.82111	3235.261	2652.584
A_23_P13	NM_00286	Homo sapi	ALDH18A1	-1.21787	0.00026	0.82111	6631.594	5443.089
A_23_P16	NM_00556	Lysosomal	LAMP1	-1.21794	0.0004	0.82106	40792.49	33278.14
A_23_P14	NM_01863	Homo sapi	WSB2,SBA	-1.218	0.00002	0.82101	1656.522	1359.985
A_23_P45	NM_00182	Homo sapi	CKS1B,CK	-1.21807	0.00002	0.82097	97239.55	79694.3
A_24_P61	AL833755	Chromosom	C6orf89	-1.21808	0.00002	0.82096	4537.875	3719.683
A_24_P41	NM_00094	protein phc	PPP3CA,C	-1.21817	0.00002	0.8209	3555.345	2917.922
A_23_P13	NM_01475	KIAA0097	CKAP5,TO	-1.21831	0.00008	0.82081	2790.406	2293.342
A_23_P20	NM_00645	Homo sapi	SPFH1,KE	-1.2184	0.00079	0.82075	8080.361	6664.051
A_23_P16	NM_00639	Homo sapi	RNASEH2	-1.21854	1.98E-06	0.82066	16258.16	13343.07
A_23_P41	NM_03235	Homo sapi	MON1A,SA	-1.2187	9.08E-06	0.82055	3241.817	2660.167
A_32_P157671	Unknown	A_32_P15		-1.21874	0.00017	0.82052	11270.33	9248.061
A_24_P229807	Unknown	A_24_P22		-1.21876	4.47E-06	0.8205	22509.61	18455.17
A_23_P15	NM_05716	Homo sapi	KLHDC3,P	-1.21901	0.00066	0.82034	747.4137	613.1616
A_23_P32	AL110257	Hypothetic	MGC26690	-1.21917	0.0004	0.82023	1278.184	1049.532
A_23_P88	NM_01475	Homo sapi	DLG7,DLG	-1.21927	0.00017	0.82016	5892.402	4815.367
A_32_P12	BE018697	Similar to F	BE018697	-1.2194	0.00025	0.82008	1802.568	1479.749
A_24_P32	NM_03121	Chromosom	C19orf27	-1.2198	5.54E-06	0.8198	3883.548	3183.331
A_23_P77	NM_01834	hypothetic	FLJ11151,I	-1.21993	0.00074	0.81972	807.6688	662.5212
A_23_P31	NM_00030	Homo sapi	PON2	-1.22014	0.00009	0.81958	2161.196	1774.207
A_24_P75	XM_37330	Homo sapi	XM_37330	-1.2202	0.00019	0.81954	3623.585	2963.276
A_24_P80	NM_01784	hypothetic	FLJ20507,I	-1.22034	5.03E-06	0.81944	1977.242	1620.276
A_23_P11	NM_00012	Homo sapi	ERCC5,XF	-1.22034	0.00022	0.81944	2793.154	2288.29
A_32_P14	NM_18247	Pyruvate ki	PKM2	-1.22049	0.00013	0.81935	6857.551	5600.929
A_24_P406591	AP003453	THC19028		-1.22058	0.00032	0.81928	749.2963	613.883
A_23_P12	NM_02272	Homo sapi	FANCF,FA	-1.22063	0.00027	0.81925	1263.832	1033.67
A_32_P13	BC009510	Chromosom	C9orf123	-1.22074	5.92E-06	0.81917	13252.2	10845.13
A_24_P204095	Unknown	A_24_P20		-1.22079	0.00007	0.81914	90780.54	74538.53
A_32_P107833	AB072039	THC18587		-1.22085	0.00008	0.8191	2231.255	1828.469
A_23_P14	NM_00169	ATPase, H	ATP6V1E1	-1.22088	0.00022	0.81908	1172.04	959.0731

A_23_P64	AB029016 Trinucleotic TNRC6B	-1.22095	0.00021	0.81904	1043.172	854.8524
A_23_P42	NM_00094 Homo sapi PPIC,CYP	-1.22101	0.0002	0.819	6153.591	5033.389
A_24_P17	BC044624 Pre-B-cell IPBX1	-1.22105	0.00002	0.81897	16991.59	13892
A_24_P94	NM_01812 hypothetical FLJ10534,I	-1.22107	0.00002	0.81895	4950.303	4058.035
A_23_P20	NM_00721 Homo sapi GALNT6,G	-1.22134	0.00054	0.81877	638.0023	522.418
A_23_P14	NM_00227 Homo sapi KRT8,K8,K	-1.22135	0.00009	0.81877	89937.86	73720.28
A_23_P94	NM_02521 G kinase a GKAP1	-1.22144	0.00013	0.81871	1349.691	1103.743
A_32_P19	NM_03115 heterogene HNRPA1,H	-1.22156	0.00005	0.81863	139045	113563.3
A_24_P34	NM_00660 unactive pr PTGES3,P	-1.22162	0.00028	0.81859	10292.93	8422.268
A_23_P21	NM_00539 Homo sapi PODXL,PC	-1.22171	0.00004	0.81853	8616.193	7039.242
A_23_P20	NM_01440 Homo sapi TRAPPC3,	-1.22176	0.00094	0.81849	2950.826	2412.174
A_24_P93	111 Unknown A_24_P93	-1.22178	0.00036	0.81848	19734.16	16124.38
A_23_P21	NM_19882 Ras-relate RAC1	-1.22182	0.00017	0.81845	15722.58	12912.54
A_23_P13	NM_00126 cyclin-depe CDK8,K35	-1.222	0.0004	0.81833	998.2407	816.4956
A_23_P15	NM_00469 Homo sapi SLC16A5,I	-1.22204	0.00004	0.8183	15810.8	12957.2
A_23_P37	AK027341 KIAA1856 KIAA1856	-1.22225	0.00013	0.81816	4612.098	3776.246
A_24_P47	AL117621 ADP-ribosy ARF6	-1.22232	0.0008	0.81812	1966.396	1612.435
A_32_P12	NM_15337 hypothetical LYSD2,M	-1.22237	0.00056	0.81808	5270.548	4323.989
A_24_P13	NM_00078 cytochrom CYP51A1,I	-1.22237	0.00026	0.81808	6334.388	5189.064
A_23_P89	NM_00646 Homo sapi SPAG5,MA	-1.22254	0.00042	0.81797	9997.639	8215.637
A_23_P39	NM_00666 Homo sapi RUVBL2,R	-1.22276	9.04E-06	0.81782	3828.479	3131.804
A_24_P47	8940 Unknown A_24_P47	-1.22284	0.00001	0.81777	128153.6	104671.2
A_23_P77	NM_00485 Homo sapi PIGB,MGC	-1.22284	0.00047	0.81777	1441.154	1179.317
A_23_P58	NM_13442 Homo sapi HPCAL1,B	-1.22284	9.00E-06	0.81777	2472.098	2020.849
A_23_P30	NM_00415 Homo sapi PPP2CB,P	-1.22302	4.69E-06	0.81765	3419.845	2796.292
A_23_P39	Z24727 Tropomyos TPM1	-1.22306	0.00005	0.81762	4731.721	3877.76
A_23_P33	NM_00282 Homo sapi PTPN1,PT	-1.22312	0.00014	0.81758	1588.625	1300.659
A_23_P34	NM_00469 Homo sapi FAM50A,9I	-1.2232	0.00002	0.81753	5231.043	4273.419
A_24_P14	NM_17795 Homo sapi DNCL2A,B	-1.22337	9.33E-07	0.81742	4299.389	3514.655
A_23_P25	NM_01897 Homo sapi DPM3,MG	-1.22367	0.00027	0.81722	6433.009	5266.009
A_23_P42	NM_00645 Mad4 homi MXD4,MAI	-1.22383	0.00002	0.81711	1352.835	1105.197
A_23_P25	NM_02246 endoplasm SIL1,BAP	-1.22396	0.0005	0.81702	6967.796	5676.08
A_23_P31	NM_01336 Homo sapi ANAPC4,A	-1.22399	0.00048	0.817	3668.904	3008.096
A_32_P104	263 AJ318213 THC19216	-1.22405	0.00012	0.81696	3771.222	3077.64
A_32_P15	NM_00682 tyrosine 3-YWHAQ,1C	-1.2241	0.00003	0.81693	56606.84	46168
A_24_P186	354 Unknown ENST0000	-1.22419	0.00074	0.81686	11729.58	9602.527
A_23_P34	NM_01554 Homo sapi RPAP1,MC	-1.2242	0.00008	0.81686	6400.725	5218.627
A_24_P39	NM_00660 putative ho PHTF1,PH	-1.22448	0.0001	0.81667	972.8378	794.8106
A_23_P35	NM_01816 NAD synth NADSYN1,	-1.2246	0.00069	0.81659	1152.151	939.9231
A_24_P26	NM_00482 potassium KCNK6,TC	-1.22475	3.50E-06	0.81649	2312.075	1888.089
A_23_P38	NM_00096 Homo sapi RARG,RAF	-1.22491	0.00059	0.81638	833.6984	680.0437
A_24_P69	AK027214 CCR4-NOI CNOT6L	-1.22506	0.00091	0.81629	1709.442	1396.943
A_24_P40	S80864 cytochrom cytochrom	-1.22515	0.00035	0.81622	1812.452	1482.293
A_23_P11	NM_03078 cisplatin re CRR9,DKF	-1.2255	0.00002	0.81599	12721.07	10395.38
A_24_P15	XM_37792 similar to CLOC40224	-1.22554	0.00017	0.81597	3103.502	2536.697
A_32_P11	AL137938 PX domain P XK	-1.2256	0.00033	0.81593	4936.346	4043.124
A_23_P15	NM_03237 hypothetical MGC4251	-1.22567	5.62E-06	0.81588	1754.87	1431.408
A_24_P91	NM_00084 glutathione GSTM3,G	-1.22568	0.00011	0.81587	25916.29	21078.77
A_23_P20	NM_00179 Homo sapi CDC42,G2	-1.22568	0.0006	0.81588	13283.5	10884.51
A_23_P12	NM_00455 Homo sapi NDUFS6	-1.22599	9.63E-06	0.81566	46087.4	37544.32
A_24_P16	AK025511 ATPase, C ATP11A	-1.22601	0.0002	0.81565	1159.793	946.1481
A_23_P65	NM_00638 Homo sapi CIB2,KIP2	-1.2261	0.00018	0.81559	3151.733	2570.373

A_24_P315654	Unknown	A_24_P315654	-1.22625	0.00002	0.81549	128162	104249.3
A_24_P662972	Unknown	THC18135	-1.22627	0.00039	0.81548	750.4422	612.1939
A_24_P911	NM_00415	Ornithine d OAZ1	-1.22639	0.00004	0.8154	421698.4	344437.1
A_23_P49	NM_00518	Calmodulin CALM3	-1.22652	5.30E-06	0.81532	3527.557	2874.319
A_23_P11	NM_01810	Homo sapi PAPD1,mtl	-1.22661	1.67E-06	0.81526	1986.035	1619.233
A_32_P14	BC028735	Hypothetic LOC40143	-1.22661	5.98E-06	0.81525	1376.3	1122.033
A_24_P17	NM_03092	DNA replic CDT1,DUF	-1.22662	9.04E-06	0.81525	2905.685	2367.313
A_24_P23	NM_00392	RAB7, mer RAB7L1,R	-1.22669	0.00003	0.8152	1066.477	869.5527
A_23_P27	NM_01554	DKFZP564 DKFZP564	-1.22677	0.00017	0.81515	2396.854	1956.027
A_23_P37	BC001774	Platelet-act PAFAH1B2	-1.22678	1.54E-06	0.81514	4736.021	3859.27
A_23_P59	NM_00564	Homo sapi TAF11,TAF	-1.22688	0.00018	0.81507	2557.107	2079.766
A_23_P16	NM_02274	Homo sapi HIPK2,PRC	-1.2273	1.21E-06	0.8148	3966.803	3231.381
A_23_P37	NM_20329	Homo sapi CHCHD1,C	-1.22743	0.00002	0.81471	26631.38	21655.38
A_23_P44	NM_00131	Homo sapi CRIP1,CRI	-1.22757	0.00024	0.81461	9086.157	7431.926
A_23_P11	NM_20330	Homo sapi C9orf132,E	-1.22768	0.00034	0.81454	10062.4	8224.311
A_23_P88	AY359110	Mesoderm MESDC2	-1.22772	2.91E-06	0.81452	5650.399	4601.528
A_24_P85	NM_01223	sirtuin (sile SIRT3,SIR	-1.22777	1.59E-06	0.81448	2534.022	2063.647
A_24_P28	NM_01869	HSVI bindi ARL6IP4,S	-1.2278	0.00024	0.81447	58642.08	47632.16
A_23_P49	NM_01650	Homo sapi MRPL27,L	-1.22783	2.82E-06	0.81444	11239.83	9144.548
A_23_P21	NM_01224	sirtuin (sile SIRT5,SIR	-1.22787	0.00001	0.81442	2563.686	2086.251
A_23_P14	NM_00169	Homo sapi ATP5O,AT	-1.228	0.00002	0.81433	58515.3	47741.29
A_23_P53	NM_01605	CGI-116 pr CGI-116	-1.22803	2.81E-06	0.81431	3429.892	2794.469
A_24_P45	AY007126	Hypothetic LOC33922	-1.22805	9.79E-06	0.8143	7817.064	6358.88
A_24_P29	NM_00334	ubiquitin-cc UBE2G1,U	-1.22814	0.00018	0.81424	1859.521	1514.3
A_24_P418498	Unknown	ENST0000	-1.22815	0.00036	0.81424	594.2901	483.886
A_24_P21	NM_01743	prostate tu PTOV1,MC	-1.22818	0.00025	0.81421	2585.591	2108.147
A_23_P74	NM_17845	Hypothetic MGC54289	-1.22824	7.74E-06	0.81417	1874.35	1526.456
A_23_P56	BC022086	Troponin T TNNT1	-1.22831	0.00001	0.81413	1033.101	841.0542
A_23_P25	NM_17336	LOC31767 LOC31767	-1.22832	0.00007	0.81412	1303.834	1060.875
A_24_P32	NM_00079	dihydrofola DHFR	-1.22861	0.00024	0.81393	2744.353	2240.719
A_23_P13	NM_01640	hypothetic HSPC138,I	-1.22884	4.02E-06	0.81378	10678.85	8683.049
A_24_P52	NM_13078	Homo sapi RAB24	-1.22916	7.66E-06	0.81356	2324.141	1890.048
A_32_P138178	Unknown	A_32_P138178	-1.22922	0.00027	0.81353	694.2012	564.6905
A_23_P21	NM_00396	Homo sapi UBE1C,UE	-1.22931	0.00014	0.81346	2099.258	1707.177
A_23_P31	NM_19888	Homo sapi NUP43,p42	-1.22934	0.00015	0.81344	4177.609	3385.758
A_23_P16	NM_00657	Homo sapi RRAGA,R/	-1.22957	0.00028	0.81329	9187.47	7447.21
A_32_P22	NM_00589	membrane M11S1,GP	-1.22958	0.00027	0.81328	689.0571	560.4493
A_24_P32	BC014228	Transcripti TFCP2	-1.2296	0.00021	0.81327	2288.876	1863.491
A_23_P43	NM_01493	Homo sapi PLEKHA6,	-1.22992	0.00006	0.81306	1928.829	1569.646
A_23_P30	NM_01462	Homo sapi MEA1,HYS	-1.23007	0.00007	0.81296	8055.876	6534.052
A_24_P11	NM_17074	Homo sapi ALDH5A1,;	-1.2301	0.00075	0.81294	1178.31	959.0571
A_23_P13	NM_02407	Homo sapi TSEN34,LI	-1.23011	5.25E-06	0.81294	11139.52	9063.109
A_24_P84	XM_29392	similar to p LOC34564	-1.23045	0.00019	0.81271	8370.27	6824.52
A_23_P43	NM_14501	Homo sapi MRGPRF,I	-1.23048	0.00011	0.81269	1137.32	925.4572
A_24_P59	NM_02198	ubiquitin-cc UBE2V1,C	-1.23049	0.00094	0.81269	2470.197	2013.425
A_23_P21	NM_00654	S-phase re SPHAR,MC	-1.23067	0.00074	0.81257	1724.083	1404.636
A_24_P75708	Unknown	ENST0000	-1.23074	0.0004	0.81252	1253.874	1016.582
A_32_P115451	ALU7_HUM	THC19333	-1.23079	0.00017	0.81249	873.8729	709.6491
A_23_P27	NM_03268	Homo sapi ZNF607,FL	-1.23083	1.73E-06	0.81246	5385.337	4378.217
A_24_P34	NM_02484	duodenal c CYBRD1,C	-1.23087	0.0004	0.81243	669.1509	543.553
A_23_P49	NM_19934	hypothetic LOC37492	-1.23089	0.00018	0.81242	3261.844	2654.964
A_24_P25	NM_13878	HEPIS LOC11971	-1.23103	0.00013	0.81233	1169.323	951.1373

A_32_P28	NM_00675	transaldola TALDO1,T	-1.23112	0.0003	0.81227	54798.14	44698.24
A_23_P12	NM_00297	Homo sapi SCP2,NLT	-1.23113	0.00064	0.81226	4681.919	3818.836
A_23_P61	NM_00460	syntaxin 4/ STX4A,p3	-1.23113	0.00066	0.81226	518.1838	420.8945
A_23_P64	NM_00641	Homo sapi HTATIP2,C	-1.2312	0.00002	0.81222	19808.89	16059.26
A_23_P12	NM_01769	Homo sapi LRRC49,FI	-1.23125	6.30E-06	0.81218	1801.485	1462.725
A_32_P18	NM_00675	transaldola TALDO1,T	-1.23141	0.00009	0.81207	96395.25	78449.55
A_23_P210285	Unknown	A_23_P210	-1.23144	0.00002	0.81206	240315.3	194364.7
A_23_P57	NM_00544	Homo sapi CHAF1B,C	-1.23163	6.54E-08	0.81193	4171.918	3394.255
A_23_P94	NM_03292	hypothetic MGC14141	-1.23166	0.00065	0.81191	10395.82	8406.936
A_23_P98	NM_02121	transposon LOC58486	-1.2317	0.00021	0.81189	5603.821	4534.964
A_32_P175098	Unknown	ENST0000	-1.23178	0.00003	0.81184	33756.48	27468.14
A_32_P21	NM_00079	dihydrofola DHFR	-1.23178	0.00003	0.81183	6315.058	5115.969
A_24_P39	NM_02064	chromoson TMEM9B,C	-1.23209	9.01E-06	0.81163	1301.967	1056.633
A_23_P37	NM_15321	Homo sapi ZNF524,M	-1.23251	6.52E-06	0.81135	1670.293	1354.937
A_24_P38	NM_00601	CD164 ant CD164,MC	-1.23276	0.00004	0.81119	1184.769	961.6067
A_23_P13	NM_01773	GPI7 prote GPI7,PRO	-1.23283	1.41E-06	0.81114	1526.521	1238.248
A_23_P20	NM_00640	Acidic (leuc ANP32B	-1.23286	0.0001	0.81112	11266.86	9165.902
A_24_P212565	Unknown	A_24_P212	-1.23305	0.00015	0.811	1495.586	1214.072
A_24_P13	NM_01812	elaC (E. co ELAC2,EL	-1.23315	1.24E-06	0.81093	2831.153	2295.677
A_23_P25	NM_00090	natriuretic (NPR3,NPF	-1.23319	0.00004	0.8109	951.3124	771.356
A_23_P90	AK127147	MBC3205 UNQ501	-1.23321	4.70E-07	0.81089	16989.06	13777.6
A_23_P48	NM_01247	Homo sapi UCK2,UK,I	-1.23326	6.04E-06	0.81086	26453.87	21427.9
A_24_P7470	Unknown	A_24_P747	-1.23327	0.00002	0.81086	1376.776	1116.201
A_23_P38	NM_01472	Homo sapi STARD8,K	-1.23331	0.00004	0.81082	954.2961	773.583
A_23_P32	NM_00208	Homo sapi GOT2,FLJ	-1.23346	0.00016	0.81073	4019.61	3265.358
A_24_P18	NM_01840	Homo sapi LPTM4B,	-1.23371	0.00009	0.81056	9882.092	7995.947
A_23_P11	NM_00666	Homo sapi PGRMC1,M	-1.23375	6.50E-06	0.81054	10340.35	8368.301
A_23_P15	NM_00279	proteasom PSMB1,HC	-1.23379	0.00013	0.81051	30656.82	24951.48
A_23_P85	NM_02490	hypothetic DENND2D	-1.23386	0.00018	0.81046	778.4773	631.0958
A_23_P38	NM_02483	Homo sapi ZNF403,DI	-1.2339	9.94E-06	0.81044	3099.531	2510.787
A_23_P50	NM_00168	Homo sapi ATP5D	-1.2339	0.00013	0.81044	5761.138	4653.962
A_24_P76	AK056691	Germ cell GSG2	-1.23398	4.56E-06	0.81039	4091.273	3316.227
A_32_P36	NM_00117	ADP-ribosy ARL1,ARF	-1.23398	0.00056	0.81039	3850.403	3132.586
A_23_P81	NM_01328	Homo sapi MAT2B,TC	-1.23434	5.32E-07	0.81015	3958.353	3207.235
A_23_P12	NM_00433	Homo sapi BUB1,BUB	-1.23461	0.00011	0.80997	2806.2	2279.995
A_23_P74	NM_00660	nuclear dis NUDC,HNI	-1.23485	0.00044	0.80981	8394.35	6766.968
A_24_P93	AK058065	G protein-c GPR133	-1.23513	0.00004	0.80963	929.0082	752.3401
A_32_P19608	AF247786	THC18468	-1.23513	0.00011	0.80963	986.4462	798.4719
A_24_P10	BC013255	Hypothetic MGC1048	-1.23519	0.00002	0.80959	1848.714	1496.995
A_23_P86	NM_00375	Homo sapi EIF3S10,E	-1.23538	0.00063	0.80946	59925.2	48681.66
A_32_P53	BG327427	Full-length BG327427	-1.23555	5.92E-07	0.80936	2196.708	1778.012
A_23_P70	U83115	Absent in n AIM1	-1.23569	7.59E-07	0.80927	2605.049	2107.721
A_24_P17	NM_01458	solute carri SLC2A8,G	-1.23573	1.93E-06	0.80924	1420.891	1149.754
A_23_P14	NM_03095	Homo sapi RNF170,DI	-1.23577	0.00069	0.80921	1201.317	973.0391
A_32_P19	NM_01599	Kruppel-lik KLF13,BTE	-1.23586	0.00004	0.80915	6052.711	4898.069
A_23_P46	NM_01225	Homo sapi SLC27A5,/	-1.236	0.00094	0.80906	5994.844	4826.423
A_23_P20	NM_01617	Calcium bir Cab45	-1.23605	0.00022	0.80903	11881.23	9662.438
A_32_P23	NM_03340	Homo sapi DOCK7,KI	-1.2362	0.00091	0.80893	2034.698	1645.739
A_23_P47	NM_00601	Homo sapi CLPP	-1.2363	0.00068	0.80886	18200.16	14645.71
A_24_P17	NM_00435	cell divisior CDC34,UB	-1.2363	0.0002	0.80887	1782.467	1446.31
A_24_P80	NM_01436	protein tyrc PTPN18,BI	-1.23634	0.00004	0.80884	1568.382	1270.209
A_23_P53	NM_03327	Ku70-bindi KUB3	-1.23653	1.19E-06	0.80872	2733.599	2211.355

A_23_P12	NM_02120	E-1 enzym	MASA,MS	-1.23659	0.00013	0.80867	4100.88	3325.261
A_23_P20	NM_00293	Homo sapi	RNH1,RAI	-1.23659	0.00036	0.80867	51727.36	41922.11
A_23_P26	NM_02067	HSCARG	HSCARG	-1.23661	5.93E-07	0.80866	5426.857	4387.334
A_23_P129405	Unknown	A_23_P12		-1.23678	1.46E-07	0.80855	12160.08	9832.082
A_23_P49	NM_01438	Proline-, gl	PELP1	-1.23691	0.00002	0.80847	5248.895	4239.031
A_24_P14	NM_01220	glyoxylate	GRHPR,GI	-1.23697	6.35E-06	0.80842	3974.499	3215.589
A_24_P170203	Unknown	A_24_P17		-1.23709	0.0004	0.80835	2507.28	2031.653
A_23_P14	NM_01474	Family with	LOC34818	-1.23719	0.00003	0.80828	2225.202	1795.985
A_23_P11	NM_00489	Homo sapi	MINPP1,M	-1.23742	0.00001	0.80813	1723.599	1391.878
A_23_P80	NM_00640	Homo sapi	PROCR,C	-1.2375	0.00084	0.80808	10221.93	8215.186
A_32_P99	AK022020	MRNA; cD	AK022020	-1.23753	0.00025	0.80806	1666.384	1345.337
A_32_P35	NM_00464	poly(A)-bin	PABPN1,C	-1.23778	4.89E-07	0.8079	9398.972	7590.805
A_24_P13	NM_00409	eukaryotic	EIF2S1,EIF	-1.23782	3.54E-06	0.80787	7861.317	6357.184
A_23_P99	NM_01799	Homo sapi	RNF31,ZIE	-1.23795	0.00052	0.80779	1574.092	1271.901
A_24_P24	NM_00693	SMT3 (sup	SUMO3,SM	-1.23803	9.15E-06	0.80773	1706.928	1378.192
A_23_P42	NM_17355	Homo sapi	TYSND1,M	-1.23804	0.00084	0.80773	965.381	780.0378
A_23_P50	NM_13362	Homo sapi	RAD51L3,	-1.23804	0.00004	0.80773	747.1567	603.5112
A_23_P41	NM_00585	Homo sapi	PURA,PUF	-1.23808	0.00003	0.8077	5049.768	4077.79
A_32_P86	BC032913	Hypothetic	LOC38902	-1.23815	0.00061	0.80766	1687.447	1364.326
A_23_P10	NM_00455	Homo sapi	NDUFS5	-1.23817	9.83E-06	0.80764	113544.8	91875.47
A_23_P21	NM_00145	Homo sapi	FLNB,AOI,	-1.23818	0.00013	0.80764	3863.87	3124.207
A_23_P31	NM_20749	Homo sapi	FLJ44896	-1.23836	0.00037	0.80752	1999.417	1616.834
A_23_P11	NM_00454	NADH deh	NDUFB10,	-1.23839	0.00034	0.8075	50853.74	40826.55
A_24_P213256	Unknown	A_24_P21		-1.2384	0.00094	0.80749	1056.73	854.1596
A_23_P13	NM_01223	Homo sapi	SIRT3,SIR	-1.23853	0.00002	0.80741	828.7648	669.1943
A_23_P20	NM_00296	Homo sapi	SAFB,HAP	-1.23862	0.00013	0.80735	3210.842	2585.676
A_23_P14	NM_03282	Homo sapi	ATOH8,HA	-1.23965	0.00007	0.80668	827.812	667.795
A_24_P54	AK096279	Homo sapi	AK096279	-1.23967	0.00014	0.80667	986.027	796.303
A_32_P22	NM_00691	Homo sapi	RPE,RPE2	-1.23967	0.00002	0.80666	928.0569	748.7321
A_23_P169766	AF208291	THC18182		-1.23977	3.20E-06	0.8066	4416.389	3566.958
A_23_P13	NM_00682	Homo sapi	YWHAQ,1	-1.23988	1.34E-06	0.80653	31898.52	25700.66
A_23_P78	NM_07847	Homo sapi	MYO18A,M	-1.23992	0.00008	0.80651	1319.293	1063.315
A_23_P37	BC006267	Asparagin	ASRGL1	-1.24003	0.00004	0.80643	2298.27	1851.212
A_24_P37	NM_00716	zinc finger	ZMYM6,M	-1.24017	1.84E-06	0.80634	1926.101	1553.019
A_23_P38	NM_00589	Homo sapi	ACAT2	-1.24019	0.00021	0.80633	2155.921	1734.111
A_23_P36	NM_00681	coated ves	TMED2,P2	-1.24023	0.00006	0.8063	7552.717	6083.158
A_23_P31	NM_00314	single-strar	SSBP1,SS	-1.24025	0.00013	0.80629	29507.5	23664.5
A_23_P13	NM_01811	Homo sapi	DALRD3,F	-1.24027	0.00002	0.80628	1836.964	1481.778
A_24_P921264	ALU7_HUM	THC19452		-1.24041	4.31E-06	0.80619	1001.411	807.3231
A_23_P68	NM_00385	Homo sapi	DPM1,MPI	-1.24044	0.00007	0.80616	20788.34	16818.12
A_23_P14	NM_00319	transcriptio	TCEA2,TFI	-1.24051	1.57E-06	0.80612	1656.403	1335.639
A_23_P13	NM_03317	Homo sapi	BAT4,G5,C	-1.24055	1.29E-06	0.80609	5621.196	4533.57
A_24_P278375	Unknown	A_24_P27		-1.24058	0.00017	0.80608	2157.933	1741.007
A_23_P10	AB020637	KIAA0830	KIAA0830	-1.24079	0.00096	0.80594	1859.191	1492.207
A_24_P73	NM_18250	hypothetic	MTERFD2,	-1.24087	0.0002	0.80589	593.7794	478.5106
A_23_P42	NM_01405	Homo sapi	TMEM14A,	-1.24097	0.00002	0.80582	8457.273	6820.991
A_23_P28	NM_00456	Homo sapi	PIGH,GPI-	-1.24097	6.15E-06	0.80582	1302.955	1050.358
A_23_P12	NM_14589	Homo sapi	PFDN5,MM	-1.24104	0.00005	0.80577	13399.38	10747.63
A_24_P38	NM_00210	H2A histon	H2AFX,H2	-1.24107	0.00006	0.80576	17151.64	13836.4
A_23_P91	NM_14471	Homo sapi	FNDC6,MC	-1.24135	1.59E-07	0.80558	6187.11	4985.011
A_24_P22	NM_14461	hypothetic	MGC3212	-1.24137	0.00004	0.80556	828.5987	667.5582
A_24_P76	AB039791	Actin-relate	ARP3BET/	-1.24146	0.00052	0.8055	498.1334	401.2443

A_23_P22	NM_00207	Homo sapi	GNB1	-1.24147	8.73E-06	0.8055	3312.653	2673.039
A_23_P25	NM_01791	hypothetic	FLJ20647	-1.24149	0.00004	0.80548	2251.45	1812.337
A_23_P89	NM_01521	Homo sapi	EXOC7,EX	-1.24155	3.26E-07	0.80544	4592.016	3697.597
A_23_P21	NM_02451	Homo sapi	FYCO1,RL	-1.24157	0.00004	0.80543	887.5957	714.8052
A_24_P17	NM_01897	dolichyl-ph	DPM3,MG	-1.24176	1.29E-06	0.80531	6048.517	4866.095
A_24_P41	BC014384	CDNA FLJ	BC014384	-1.24196	0.00083	0.80518	4284.642	3473.181
A_24_P627678	Unknown	A_24_P62		-1.24201	0.00001	0.80515	9340.41	7531.696
A_23_P78	NM_01404	HSPC023	HSPC023	-1.24207	0.00093	0.80511	3616.789	2919.994
A_24_P22	NM_01232	microtubul	MAPRE1,E	-1.24235	0.00009	0.80493	6244.32	5040.316
A_23_P10	NM_01479	Homo sapi	LCMT2,MC	-1.24238	1.52E-06	0.80491	3005.19	2421.262
A_24_P27	AK126559	Ash1 (abse	ASH1L	-1.24242	0.00043	0.80488	1947.102	1568.935
A_24_P29	NM_02394	hypothetic	MGC3234	-1.2425	0.00035	0.80483	594.0749	478.126
A_24_P18	NM_00347	protein tyrc	PTP4A2,HI	-1.24253	2.14E-07	0.80481	2659.922	2140.587
A_23_P16	NM_01243	Homo sapi	SNAPAP,S	-1.24255	4.88E-06	0.8048	5947.9	4793.175
A_32_P19	NM_00067	alcohol def	ADH5,FDH	-1.2426	2.16E-07	0.80476	2921.057	2350.912
A_23_P71	NM_00652	Homo sapi	SLBP,HBP	-1.24265	0.00062	0.80473	11038.43	8907.61
A_23_P53	NM_00234	lymphotoxi	LTBR,CD1	-1.24269	0.00028	0.8047	1115.196	898.0707
A_24_P46	NM_01786	Homo sapi	PIGX,FLJ2	-1.24294	1.24E-06	0.80454	3452.029	2778.408
A_24_P68	BC044933	Hypothetic	LOC14690	-1.24305	1.25E-06	0.80447	1745.796	1404.905
A_24_P28	NM_00417	solute carri	SLC1A3,E	-1.24351	0.00011	0.80417	1099.45	884.5642
A_23_P29	NM_00730	Synuclein,	SNCA	-1.2436	3.54E-06	0.80412	5326.414	4285.723
A_32_P14	AY026351	Similar to	AFKSG73	-1.24361	0.00046	0.80411	593.4341	477.236
A_23_P32	NM_00013	Homo sapi	FANCC,FA	-1.2439	8.76E-06	0.80392	1737.619	1396.192
A_24_P77	AK093903	HBV preS1	PS1TP4	-1.24421	0.00019	0.80373	1003.754	807.0133
A_23_P40	NM_01239	Homo sapi	PFAS,PUR	-1.24431	0.00006	0.80366	1044.88	839.6838
A_24_P608790	UCRH_HU	THC18041		-1.24437	0.00058	0.80362	64837.39	51759.41
A_23_P30	NM_03341	IMP4, U3 s	IMP4	-1.2445	0.00084	0.80354	4588.694	3671.34
A_23_P58091	BC011341	THC19337		-1.24455	0.00002	0.8035	4429.134	3559.96
A_24_P418536	Unknown	A_24_P41		-1.24461	0.00009	0.80346	44164.39	35681.63
A_23_P35	NM_01829	Homo sapi	CWF19L1,	-1.24461	0.00002	0.80346	2153.274	1732.45
A_23_P13	NM_00256	prostatic bi	PBP,HCNF	-1.24474	9.67E-08	0.80338	74248.7	59640.5
A_24_P22	NM_00505	RAD23 (S.	RAD23A,H	-1.24475	0.00003	0.80338	1742.262	1400.652
A_24_P87	BC061590	GTF2I repe	GTF2IRD2	-1.24506	0.00036	0.80318	1880.787	1513.9
A_32_P43	NM_00574	accessory	BCAP31,C	-1.24533	0.00026	0.803	47367.61	37769.21
A_24_P25020	Unknown	ENST0000		-1.24545	0.00002	0.80293	2433.26	1954.988
A_24_P19	NM_01614	PTD008 pr	PTD008	-1.24545	2.90E-07	0.80292	2702.855	2170.79
A_24_P674924	ALU8_HU	THC19984		-1.24549	6.13E-06	0.8029	2189.996	1757.326
A_23_P15	NM_00567	reproductic	UBXD6,RE	-1.24558	4.63E-07	0.80284	6072.118	4876.727
A_23_P39	NM_01344	Homo sapi	DSCR1L2,	-1.2456	0.00022	0.80282	624.485	501.2964
A_23_P21	NM_00671	Homo sapi	FASTK,FA	-1.24565	0.0002	0.80279	6101.451	4894.887
A_24_P16230	Unknown	ENST0000		-1.24571	0.00002	0.80276	20786.95	16655.21
A_32_P14	BQ447618	Polymeras	PTRF	-1.24585	0.00016	0.80266	4130.45	3326.248
A_23_P38	NM_15230	Homo sapi	C20orf158,	-1.2459	3.51E-06	0.80263	3848.247	3087.321
A_24_P255914	Unknown	A_24_P25		-1.24621	0.00003	0.80243	1784.253	1432.993
A_23_P50	NM_00332	Homo sapi	TUB	-1.24629	0.00007	0.80238	1486.499	1193.074
A_23_P15	NM_00392	Homo sapi	MBD2,DM	-1.24641	0.00035	0.80231	5751.333	4629.763
A_23_P69	NM_00160	acetyl-Coe	ACAA1,AC	-1.24641	3.99E-06	0.8023	4531.074	3628.598
A_23_P13	NM_15241	hypothetic	TMEM68,F	-1.24649	0.00009	0.80226	1500.351	1201.927
A_32_P12388	Unknown	A_32_P12		-1.24674	0.00005	0.80209	840.1189	674.0434
A_32_P16	NM_18262	Homo sapi	FAM33A,F	-1.24677	0.00023	0.80207	2812.135	2259.698
A_24_P98	NM_01332	sorting nex	SNX10,MC	-1.24679	4.60E-06	0.80206	1429.461	1146.672
A_23_P29	NM_00305	Homo sapi	SLC19A1,C	-1.24696	0.00007	0.80195	1615.917	1297.283

A_23_P46	AK126751	NHL repea	NHLRC2	-1.24717	0.00022	0.80181	896.3236	719.3543
A_23_P35	NM_01618	60S acidic	C1orf33,dJ	-1.2472	8.54E-07	0.8018	4807.001	3849.791
A_24_P16	NM_00727	lung cance	TUSC2,PA	-1.24738	0.00002	0.80168	4151.028	3325.31
A_24_P68	2601	Unknown	A_24_P68:	-1.2476	0.00002	0.80154	2188.4	1755.81
A_23_P13	NM_03290	phosphatid	PIG-Y	-1.24774	0.00001	0.80145	11584.84	9305.172
A_23_P31	AF356877	Acetyl-Coe	ACAT2	-1.24774	0.00003	0.80145	8278.782	6617.192
A_32_P17	BC014113	Hypothetic	AFAP	-1.24778	5.75E-06	0.80142	1470.343	1177.549
A_23_P40	NM_00409	E2F transc	E2F2,E2F-	-1.24806	3.61E-06	0.80125	4176.877	3348.386
A_24_P18	AK002097	Hypothetic	FLJ11235	-1.2481	0.00002	0.80122	2527.879	2023.604
A_23_P16	NM_17071	Homo sapi	DAZAP1,M	-1.24817	5.30E-07	0.80117	8482.635	6800.971
A_23_P14	NM_03242	Homo sapi	LZTS2,LAF	-1.24824	4.10E-07	0.80113	2091.888	1675.348
A_23_P14	NM_01809	Homo sapi	NLE1,Nle,F	-1.2483	8.89E-08	0.80109	8691.907	6964.407
A_23_P10	NM_01206	Homo sapi	AKR7A3,A	-1.24849	0.00004	0.80097	2667.229	2137.984
A_24_P35	NM_01516	Homo sapi	ARL6IP,All	-1.2486	4.51E-06	0.8009	33271.93	26696.6
A_32_P25	AK022962	Pre-B-cell	IPBX1	-1.24869	0.00022	0.80084	4964	3988.844
A_24_P47	1121	Unknown	A_24_P47:	-1.24884	0.00012	0.80074	994.3202	796.9824
A_23_P51	NM_00381	Homo sapi	PABPC4,A	-1.24887	0.00056	0.80073	516.7195	413.7374
A_23_P25	NM_01609	HSPC038	JZNF706,H	-1.24922	6.85E-08	0.8005	21709.56	17381.38
A_24_P79	NM_01430	SEC11-like	SEC11L1	-1.24928	0.00002	0.80046	45042.68	35953.84
A_23_P65	NM_00222	Homo sapi	ITPKA	-1.24949	0.00014	0.80033	2755.375	2205.072
A_23_P27	NM_00727	Homo sapi	GABARAP	-1.25003	0.00022	0.79998	4043.952	3221.801
A_23_P21	NM_01822	hypothetic	FLJ10803	-1.25017	0.00011	0.79989	4635.883	3719.551
A_24_P16	NM_01754	Homo sapi	EPDR1,UC	-1.25025	0.00007	0.79984	12835.3	10298.38
A_23_P11	NM_00635	Homo sapi	UBE2E3,U	-1.25032	6.79E-07	0.7998	8767.361	7015.72
A_23_P13	NM_03298	Homo sapi	TBL2,WBS	-1.25038	3.55E-07	0.79976	9779.414	7827.173
A_23_P88	NM_00003	aldolase A,	ALDOA,AL	-1.25048	0.00031	0.79969	68841	55278.52
A_24_P34	0866	Unknown	A_24_P34:	-1.25048	0.00058	0.7997	2166.705	1742.246
A_32_P22	BC067856	Succinate	SDHAL2	-1.25053	0.00057	0.79966	2580.606	2066.482
A_24_P26	NM_00316	sulfotransf	SULT1A3,S	-1.25062	2.69E-07	0.7996	10962.44	8772.529
A_23_P50	NM_01549	Homo sapi	ANKRD25,	-1.25068	1.80E-06	0.79957	10597.47	8476.871
A_23_P11	NM_00048	Homo sapi	APRT,AMF	-1.25081	0.00035	0.79948	42018.07	33416.43
A_24_P49	421	Unknown	ENST0000	-1.25082	0.00011	0.79948	13018.29	10457.59
A_23_P16	NM_00362	Homo sapi	PKP4,p007	-1.25094	0.00018	0.7994	814.6187	651.0949
A_23_P13	NM_02107	Homo sapi	NDUFV2	-1.25145	6.12E-07	0.79907	22386	17868.15
A_24_P16	NM_18173	Homo sapi	PRDX2,PR	-1.2515	0.00043	0.79904	808.6399	646.8066
A_23_P75	NM_00455	NADH deh	NDUFS3	-1.25162	0.00004	0.79897	13197.64	10512.37
A_23_P10	AK027315	Homo sapi	PPIL3	-1.25185	2.55E-06	0.79882	6785.14	5415.927
A_23_P24	NM_03225	hypothetic	FLJ37970,l	-1.25193	8.60E-06	0.79877	2360.916	1885.811
A_23_P14	NM_01599	Homo sapi	ATP6V1D,'	-1.25194	0.00035	0.79876	5529.238	4434.066
A_23_P88	NM_01221	Homo sapi	MLYCD,M	-1.25198	3.22E-06	0.79873	1268.459	1013.516
A_23_P10	NM_00393	Adaptor-rel	AP3D1	-1.25227	1.09E-07	0.79855	35004.28	27942.22
A_32_P14	243	AE016951	THC20045	-1.25231	2.75E-06	0.79852	1508.269	1203.925
A_23_P89	NM_00611	acetyl-Coe	ACAA2,DS	-1.25245	9.35E-06	0.79844	5277.376	4223.106
A_32_P18	9034	K1CR_HUI	THC19529	-1.25269	7.88E-08	0.79828	14510.93	11587.57
A_23_P94	NM_00331	Homo sapi	TSTA3,FX,	-1.25269	0.00015	0.79828	1401.586	1119.12
A_23_P24	NM_01767	Homo sapi	OTUB1,OT	-1.25275	0.00056	0.79824	1707.074	1355.65
A_23_P13	NM_02207	hypothetic	MANBAL	-1.25279	0.00074	0.79822	3278.323	2621.739
A_24_P94	NM_00630	SMC1 (str	SMC1L1,S	-1.25291	0.00055	0.79814	896.2628	716.0858
A_23_P40	NM_00340	Homo sapi	XPO1,CRM	-1.25295	3.83E-08	0.79812	12898.06	10295.45
A_24_P94	NM_01461	C9orf10 pr	C9orf10,DI	-1.25296	0.00008	0.79811	8561.055	6863.195
A_32_P11	NM_15226	hypothetic	FLJ38663	-1.25302	0.00001	0.79807	3228.874	2578.292
A_23_P40	NM_00415	Ornithine d	OAZ1	-1.25306	2.04E-06	0.79804	246633.8	197042.9

A_23_P14	NM_00580 Homo sapi	PRDX2,PR	-1.25306	1.11E-07	0.79805	9841.838	7858.084
A_24_P40	NM_18130 Homo sapi	MRPL52	-1.25332	0.00007	0.79788	9553.025	7648.106
A_23_P20	NM_01961 WDR45-lik	WDR45L	-1.25365	0.00002	0.79767	12620.75	10097.12
A_24_P14	NM_02278 likely ortho	ARV1	-1.25374	2.67E-06	0.79762	1581.218	1260.707
A_23_P42	NM_01518 KIAA0746	KIAA0746,	-1.25378	3.59E-07	0.79759	5915.452	4713.568
A_23_P80	NM_00318 Homo sapi	TAF4,TAF2	-1.25391	0.00002	0.7975	1218.813	972.5275
A_23_P20	AB007925 SLIT-ROB	SRGAP2	-1.25398	4.07E-08	0.79746	3993.391	3184.368
A_23_P42	NM_03092 Homo sapi	KAZALD1,	-1.25411	0.00015	0.79738	1769.975	1413.785
A_23_P18	NM_00720 mitochondr	MRPL3,MF	-1.25411	0.00062	0.79738	15525.78	12293.31
A_32_P17	BC059174 TNF recept	TRAF3IP1	-1.2543	0.00009	0.79726	1836.27	1464.3
A_32_P20	NM_00682 tyrosine 3-t	YWHAQ,10	-1.25459	1.01E-06	0.79707	57935.24	46230.32
A_24_P38	NM_03094 HESB like	HBLD2	-1.25469	0.00007	0.79701	616.012	490.9765
A_24_P23	BC000821 Hypothetic	LOC14960	-1.25472	2.76E-08	0.79699	6307.354	5026.9
A_23_P66	NM_01653 sirtuin (sile	SIRT7,SIR	-1.25478	0.00002	0.79695	1366.58	1090.084
A_23_P40	NM_02092 Homo sapi	BCOR,MA	-1.25488	7.89E-06	0.79689	903.3779	719.9193
A_23_P39	NM_20330 Homo sapi	RKHD1,TIF	-1.2549	0.00006	0.79687	1279.958	1020.279
A_23_P77	NM_00480 Homo sapi	WDR39,C1	-1.25495	4.43E-06	0.79684	23030	18388.83
A_24_P35	BC034573 Chromosom	C6orf12	-1.2551	0.00014	0.79675	837.5596	667.9095
A_23_P11	NM_00588 Membrane	MARCH-VI	-1.25514	0.00006	0.79673	3305.946	2633.464
A_23_P29	NM_01570 CGI-96 pro	CGI-96,BK	-1.25515	1.07E-06	0.79672	1366.485	1088.49
A_23_P21	NM_03362 Homo sapi	TREX1,DR	-1.25518	5.46E-06	0.7967	2168.804	1726.046
A_23_P15	NM_01847 Homo sapi	ACTR10,A	-1.25531	3.92E-07	0.79662	9620.623	7673.24
A_23_P45	NM_01785 hypothetic	C1orf109,F	-1.25554	1.23E-06	0.79647	2438.115	1942.654
A_23_P20	NM_00686 RAB35, me	RAB35,RA	-1.25556	6.44E-08	0.79645	9477.139	7546.492
A_32_P109777	PHB_HUM	THC19411	-1.25558	0.00015	0.79644	2170.606	1724.653
A_23_P50	NM_00585 Homo sapi	RAMP1	-1.25562	0.00005	0.79642	1798.137	1434.127
A_23_P83	NM_03255 hypothetic	HIATL1,FL	-1.2558	0.00006	0.79631	3388.669	2689.544
A_23_P13	NM_00464 Homo sapi	CDK2AP1,	-1.25582	6.13E-06	0.79629	19416.94	15429.6
A_24_P29	NM_00680 chromobox	CBX1,CBX	-1.2559	6.16E-07	0.79624	11167.45	8900.38
A_23_P10	NM_00458 RAB5C, m	RAB5C,RA	-1.25595	0.00027	0.79621	1161.236	924.7366
A_23_P30	NM_00733 Wolf-Hirsch	WHSC1,N	-1.25598	0.00003	0.79619	814.0186	648.1273
A_23_P20	NM_00550 Homo sapi	SCARB1,C	-1.25598	3.01E-07	0.79619	99682.99	79288.92
A_32_P17	NM_01857 amyotrophi	ALS2CR2,	-1.25628	0.00033	0.796	618.0281	492.0445
A_24_P41	NM_00564 transducin	TBL1X,EBI	-1.25631	7.43E-07	0.79598	2398.526	1908.284
A_23_P15	NM_01246 Homo sapi	C20orf14,T	-1.25637	0.00005	0.79594	7539.256	5981.343
A_23_P73	NM_00058 Homo sapi	SPP1,OPN	-1.25653	1.32E-06	0.79585	209297.6	165931.3
A_23_P16	NM_15235 Homo sapi	CPT1C,CA	-1.25656	2.25E-06	0.79583	1435.678	1141.734
A_23_P23	NM_00597 papillary re	PRCC,TPF	-1.25659	0.00082	0.79581	792.7795	631.8641
A_23_P37	NM_14460 hypothetic	HEXIM2,L	-1.2566	1.34E-06	0.7958	4100.879	3260.646
A_24_P20	NM_00420 vesicle traf	SEC22L3,S	-1.25662	0.00045	0.79578	2511.118	2007.628
A_24_P13	NM_02289 likely ortho	ACTR8,AR	-1.25692	0.00004	0.79559	723.9154	575.9859
A_24_P16	NM_00168 ATP synth	ATP5D	-1.25693	0.00009	0.79559	7221.936	5714.663
A_24_P25	NM_01841 hypothetic	DKFZp762	-1.25727	3.42E-06	0.79537	1849.38	1470.434
A_23_P124823	SPC4_HUI	THC19785	-1.25731	0.00028	0.79535	10543.94	8432.955
A_23_P20	NM_01418 HSPC163	CNIH4,HSI	-1.25736	0.00003	0.79531	36601.76	29005.98
A_24_P25	NM_02410 hypothetic	MGC2747	-1.2574	3.90E-08	0.79529	2939.542	2337.896
A_23_P32	NM_02040 CGI-203 pr	C6orf149,C	-1.25747	1.03E-06	0.79524	3954.217	3146.928
A_24_P26	NM_00282 prothymosi	PTMA,TM	-1.25752	1.55E-07	0.79522	268512.6	213764.8
A_23_P20	NM_02513 melanoma/	ARMC9,AF	-1.25771	0.00001	0.7951	1497.773	1192.929
A_23_P11	NM_00611 HS1 bindin	HAX1,HS1	-1.25782	0.00014	0.79503	18693.03	14786.69
A_32_P14	AI204246 Transmem	TMED4	-1.25785	0.00049	0.79501	2326.55	1845.66
A_23_P75	NM_01539 Homo sapi	BRMS1,Dk	-1.25786	5.11E-07	0.795	2459.976	1954.188

A_23_P81	NM_00651	Homo sapi	TCTEL1,C	-1.25789	6.08E-06	0.79498	33602.87	26649.21
A_23_P30	NM_00128	chloride int	CLIC1,G6,I	-1.25795	0.00006	0.79494	20069.95	15898.98
A_24_P25	X64990	Olfactory r	OR5L2	-1.25797	0.00098	0.79493	3515.988	2802.735
A_24_P97	NM_01662	hypothetic	MRPL35	-1.25802	1.91E-07	0.7949	7654.012	6079.814
A_32_P11	NM_03247	Homo sapi	MRPL41,B	-1.25811	0.00018	0.79485	37121.83	29398.36
A_23_P88	NM_00323	Homo sapi	TGFB3,TG	-1.25819	9.31E-06	0.79479	754.4924	599.6871
A_32_P20	AK090739	CDNA FLJ	AK090739	-1.25821	0.00016	0.79478	6101.293	4857.579
A_24_P71	NM_01614	dynein ligh	DNCLI1	-1.25821	0.00066	0.79478	1552.795	1227.103
A_24_P41	NM_08066	similar to	RMGC1960	-1.25827	0.00001	0.79474	10019.6	7938.251
A_24_P118472		Unknown	A_24_P11	-1.25842	0.00056	0.79465	6938.539	5545.104
A_23_P11	NM_00056	glutathione	GSTM1,ML	-1.25887	1.18E-06	0.79437	5416.201	4306.352
A_24_P15	BC036012	Tripartite r	TRIM37	-1.2591	0.00048	0.79422	779.7376	619.4628
A_23_P33	NM_00466	Homo sapi	HERC2,jdf	-1.25921	8.56E-06	0.79415	1642.698	1306.053
A_23_P40	NM_00123	carbonyl re	CBR3,hCB	-1.25923	3.89E-06	0.79413	1179.97	936.8054
A_23_P69	NM_02297	small EDR	SERF1B,F	-1.25942	0.00006	0.79402	14767.04	11776.19
A_23_P12	NM_01537	Homo sapi	BRI3,I3	-1.25943	6.80E-06	0.79401	47122.56	37311.41
A_23_P41	AL122036	Three prim	TREX2	-1.25944	0.00012	0.794	2920.677	2307.924
A_23_P14	NM_01502	KIAA0251	KIAA0251,	-1.25945	0.00004	0.794	3706.673	2951.426
A_23_P75	NM_03279	Homo sapi	RPUSD4,F	-1.25994	0.00002	0.79369	3520.37	2800.414
A_32_P16	AK075484	Homo sapi	AK075484	-1.26001	9.74E-08	0.79365	3974.758	3153.006
A_23_P20	AK027500	Leucine ric	LRRN6A	-1.26022	3.38E-06	0.79352	4675.142	3714.014
A_23_P14	NM_14458	Homo sapi	C14orf149,	-1.26026	0.00076	0.79349	2588.301	2059.853
A_23_P20	BC032831	G protein-c	GPR135	-1.26046	0.00018	0.79336	1467.159	1164.759
A_23_P16	NM_00275	Homo sapi	MAPK9,JN	-1.26047	0.00003	0.79335	858.1698	680.6199
A_23_P32	NM_01229	Homo sapi	ESPL1,ES	-1.26048	3.24E-06	0.79335	5169.078	4107.12
A_23_P21	NM_01592	Homo sapi	C6orf82,Cl	-1.26053	7.60E-08	0.79331	22034.2	17474.16
A_23_P38	NM_00411	Homo sapi	FDXR,AD	-1.26061	0.00035	0.79327	10190.42	8076.681
A_24_P94	NM_01494	spastic par	SPAST,FS	-1.26061	0.00091	0.79327	1276.429	1014.684
A_23_P58	NM_02115	Homo sapi	RAP1GDS	-1.26061	0.0004	0.79326	1096.618	868.7727
A_32_P12	AK002014	Chromosor	C6orf70	-1.26061	2.37E-06	0.79327	1716.335	1363.025
A_24_P91	NM_00099	ribosomal	RPL28	-1.26065	1.61E-06	0.79324	1198.159	950.6555
A_24_P84970		Unknown	A_24_P84	-1.26079	2.91E-06	0.79315	14532.53	11546.78
A_24_P94	AK091153	AHNAK nu	MGC5395	-1.26105	0.00005	0.79299	6859.948	5453.548
A_23_P37	NM_01324	FH1/FH2 d	FHOD1,FH	-1.26109	0.00036	0.79296	4723.81	3733.25
A_23_P10	NM_15286	Homo sapi	ARPC2,AR	-1.2611	5.27E-07	0.79296	33951.61	26881.88
A_23_P20	NM_00473	Homo sapi	MTA2,PID,	-1.26115	3.00E-06	0.79293	1917.919	1521.233
A_23_P11	NM_01778	Homo sapi	SEMA4C,S	-1.26116	8.57E-06	0.79292	3073.793	2437.503
A_23_P211899		Unknown	THC19455	-1.26127	2.17E-06	0.79285	2270.635	1799.418
A_23_P426279		AB046417	THC18444	-1.26134	6.28E-06	0.79281	2310.956	1827.728
A_24_P90	NM_00546	heterogene	HNRPDL,I	-1.26141	0.00039	0.79276	4409.474	3500.419
A_23_P44	NM_01407	Homo sapi	MRPL13,L	-1.26161	2.55E-08	0.79264	15038.23	11924.85
A_23_P80	NM_00336	Homo sapi	USP4,UNF	-1.26181	0.00009	0.79251	632.9733	501.6593
A_23_P77	NM_15233	Homo sapi	SLC25A29	-1.26187	0.00019	0.79248	621.0015	492.0988
A_23_P61	NM_00578	Tyrosine ki	TNK2	-1.2619	0.00012	0.79246	2213.211	1759.935
A_23_P12	NM_00464	Homo sapi	PABPN1,C	-1.2622	0.00001	0.79227	14095.04	11197.84
A_24_P10	NM_00600	ubiquinol-c	UQCRH	-1.26233	0.00009	0.79218	97407.11	76731.17
A_23_P36	NM_17883	similar to	CLOC20116	-1.26243	0.00067	0.79213	805.943	638.6276
A_24_P74	NM_05818	Homo sapi	C21orf51	-1.26243	6.24E-07	0.79212	2680.295	2124.585
A_24_P18	NM_01988	glycogen s	GSK3A,DK	-1.2625	5.21E-07	0.79208	8888.654	7032.549
A_23_P25	NM_03279	hypothetic	FLJ14466	-1.26294	0.00006	0.7918	666.726	527.9578
A_23_P70	NM_00563	Homo sapi	SMO,Gx,S	-1.26298	0.00094	0.79178	2448.7	1943.719
A_23_P11	NM_03250	Homo sapi	RBM13	-1.26301	7.21E-06	0.79176	3803.491	3013.69

A_24_P15	XM_37299 similar to c	LOC39154	-1.26303	0.00066	0.79175	1081.25	857.9496
A_24_P18	NM_00119 BH3 intera	BID,MGC1	-1.26305	1.54E-06	0.79174	11780.64	9311.848
A_24_P12	XM_37292 similar to C	LOC39137	-1.26309	0.0002	0.79171	1458.951	1155.033
A_24_P358406	Unknown	A_24_P35	-1.26321	0.00062	0.79164	18051.39	14199.04
A_23_P10	NM_00191 Homo sapi	CYB5,MCE	-1.26363	4.89E-08	0.79137	18437.48	14581.7
A_24_P68	BC041926 Homo sapi	BC041926	-1.26366	0.00003	0.79135	660.526	522.6904
A_23_P66	NM_14468 hypothetic	MGC23280	-1.26376	0.00025	0.79129	1713.255	1356.182
A_23_P34	NM_00548 Homo sapi	PIGK,GPI8	-1.26387	0.00018	0.79122	2436.186	1927.975
A_24_P34	NM_00192 dCMP dea	DCTD,MG	-1.26402	2.52E-08	0.79113	2451.598	1939.437
A_24_P25	NM_14504 alpha tubul	MGC16703	-1.26425	0.00008	0.79098	3630.122	2871.094
A_23_P44	NM_00039 Homo sapi	ABCC2,DJ	-1.26434	3.13E-08	0.79093	8933.227	7066.56
A_23_P63	NM_00037 uroporphyr	UROD,PC	-1.26435	1.62E-07	0.79092	9739.632	7712.25
A_23_P84	NM_03351 Homo sapi	SLC38A5,S	-1.26436	0.00003	0.79091	3008.473	2379.127
A_23_P21	NM_01790 hypothetic	FLJ20625	-1.26437	3.41E-07	0.79091	4388.696	3468.707
A_24_P38	NM_00112 adaptor-rel	AP1G1,AD	-1.26441	2.04E-07	0.79088	4872.085	3853.499
A_23_P11	NM_00640 Homo sapi	HBXIP,XIP	-1.26445	1.06E-06	0.79086	26104.2	20689.63
A_23_P53	NM_00379 Homo sapi	EED,HEEC	-1.26459	2.07E-07	0.79077	7411.349	5867.709
A_23_P91	NM_00288 Homo sapi	RANBP1,M	-1.26469	0.00073	0.79071	69209.95	54252.03
A_23_P31	NM_01988 Homo sapi	CHST7,C6	-1.26486	0.00039	0.7906	1169.417	925.0855
A_23_P38	NM_01430 SEC11-like	SEC11L1	-1.2652	4.25E-07	0.79039	56778.2	44794.31
A_23_P35	NM_00408 Homo sapi	TIMM8A,D	-1.26526	0.00004	0.79035	2200.757	1744.003
A_24_P17	NM_01504 KIAA0676	KIAA0676	-1.26543	0.00015	0.79024	5345.879	4236.352
A_23_P12	NM_00542 tumor prote	TP53BP2,S	-1.26548	0.00033	0.79022	1535.277	1216.868
A_23_P74	AB020714 KIAA0907	KIAA0907	-1.26549	0.00009	0.79021	2292.947	1818.187
A_23_P23	NM_01428 Homo sapi	ITGB3BP,N	-1.26554	1.27E-06	0.79018	2798.966	2222.529
A_23_P12	NM_00483 Homo sapi	GSTO1,P2	-1.26558	0.00008	0.79015	28831.88	22660.75
A_24_P87	AK093346 CDNA FLJ	AK093346	-1.2656	8.64E-06	0.79014	2895.014	2292.263
A_23_P36	NM_13841 hypothetic	MGC15416	-1.26585	4.73E-08	0.78999	34437.66	27215.84
A_23_P71	NM_00071 Homo sapi	BLVRA,BL	-1.26586	0.00007	0.78998	24995.6	19697.96
A_32_P203300	IF4E_HUM	THC18793	-1.26593	0.00035	0.78993	988.1337	779.9327
A_23_P14	NM_03232 hypothetic	MGC13102	-1.26601	0.00002	0.78989	1316.354	1039.844
A_23_P64	NM_01245 translocase	TIMM10,TI	-1.26619	0.00019	0.78977	16300.77	12964.58
A_24_P118512	Unknown	A_24_P11	-1.2662	0.00002	0.78977	950.2813	750.1879
A_23_P13	NM_00417 Homo sapi	SNRPD3,S	-1.26635	0.00003	0.78967	38357.16	30176.17
A_24_P32	NM_01549 DKFZP434	ANKRD25,	-1.2664	0.00015	0.78964	937.1141	740.4636
A_24_P36	NM_01787 hypothetic	HRASLS2,	-1.26642	0.00024	0.78963	1220.083	964.7119
A_23_P76	NM_00632 Homo sapi	TUBGCP3,	-1.26658	0.00046	0.78953	507.3796	400.5804
A_24_P315594	Unknown	A_24_P31	-1.26663	0.00021	0.78949	979.7675	773.468
A_23_P25	NM_00429 Homo sapi	RNF14,AR	-1.26687	0.0001	0.78935	2591.153	2041.856
A_23_P45	NM_00682 hypothetic	IFI44L,GS	-1.26705	0.00033	0.78923	2680.037	2114.735
A_23_P34	NM_01430 Homo sapi	NIFUN,ISC	-1.26716	1.37E-08	0.78917	17300.34	13657.74
A_32_P19	BC006406 Chromosor	CXorf44	-1.26726	3.64E-06	0.7891	4661	3683.743
A_23_P12	NM_01328 Homo sapi	RBM15B,H	-1.26729	9.13E-09	0.78908	4241.728	3347.019
A_23_P10	NM_00213 Homo sapi	HMOX2,HC	-1.26732	1.36E-07	0.78907	6300.978	4972.966
A_24_P18	NM_01663 SRp25 nuc	ARL6IP4,S	-1.26734	0.00037	0.78905	9086.171	7133.473
A_24_P878419	Unknown	A_24_P87	-1.26743	0.00096	0.789	8490.621	6724.512
A_23_P60	NM_00731 V-abl Abela	ABL1	-1.2675	0.00075	0.78896	2004.846	1582.599
A_24_P13	NM_00386 carboxyles	CES2,iCE,	-1.26759	8.85E-08	0.7889	10880.05	8577.952
A_23_P13	NM_01508 Homo sapi	MRPS27,M	-1.26771	0.00011	0.78883	5422.117	4294.125
A_32_P46	NM_08062 Homo sapi	C20orf135,	-1.26772	0.00095	0.78882	1948.939	1542.25
A_23_P98	NM_00490 Homo sapi	PRDX6,PR	-1.26773	3.77E-06	0.78881	125767.7	99493.14
A_23_P13	NM_00177 Homo sapi	CD58,LFA	-1.26815	1.78E-06	0.78855	2838.983	2236.259

A_24_P383660	Unknown	A_24_P383660	-1.26819	3.42E-06	0.78853	19330.89	15219.51
A_23_P12	NM_00182	Homo sapiens CLCN3, CL	-1.26835	6.00E-06	0.78843	1417.114	1118.108
A_23_P19	NM_01432	Homo sapiens HEBP2, PP	-1.2685	0.00009	0.78833	10506.39	8302.234
A_24_P42	NM_00022	keratin 18 KRT18, K18	-1.26856	0.00017	0.7883	38822.18	30454.85
A_32_P20	NM_00182	CDC28 protein kinase CKS1B, CK	-1.26861	0.00015	0.78826	12635.98	10012.69
A_24_P230057	Unknown	ENST00000230057	-1.26901	1.77E-06	0.78802	26925.32	21177.09
A_24_P777185	Unknown	A_24_P777185	-1.26931	1.01E-06	0.78783	3803.787	2998.449
A_23_P10	NM_00693	Homo sapiens SNRPD1, S	-1.26932	0.00065	0.78782	66736.25	52165.55
A_23_P40	NM_00134	Homo sapiens DGKG, DA	-1.26954	7.56E-07	0.78769	1882.943	1482.459
A_24_P35	NM_00728	GABA(A) receptor subunit GABARAP	-1.26966	0.0003	0.78762	3487.71	2768.254
A_24_P39	NM_00208	growth factor receptor GRB2, AS	-1.26982	0.00004	0.78751	914.9697	719.8854
A_23_P20	NM_00280	proteasome activator PSMD1, S1	-1.26993	3.63E-07	0.78744	23240.16	18274.27
A_23_P42	NM_15282	Homo sapiens SNX3, SDP	-1.26995	0.0001	0.78743	17830.67	13967.5
A_23_P33	BC024231	Hypothetical protein MGC27345	-1.27019	0.00003	0.78728	1355.144	1067.599
A_23_P44	NM_00201	Homo sapiens FLII, FLI, FL	-1.27025	9.44E-09	0.78724	5122.184	4031.645
A_24_P10	BC031469	MRNA similar to LOC55420	-1.27029	0.00006	0.78722	2345.753	1852.039
A_24_P401601	Unknown	ENST000002401601	-1.27031	1.53E-06	0.78721	10916.08	8581.328
A_24_P418687	Unknown	A_24_P418687	-1.27038	0.00028	0.78716	20743.27	16265.3
A_23_P32	NM_17843	Homo sapiens LCE3B, LEI	-1.27054	0.00017	0.78707	1082.792	852.9136
A_23_P97	NM_02512	hypothetical protein C10orf57, F	-1.27057	0.00003	0.78705	1246.099	979.5779
A_23_P26	AK026008	WD-repeat domain HAN11	-1.2707	1.59E-07	0.78697	3591.241	2826.676
A_23_P90	NM_01619	Homo sapiens LSM7, YNL	-1.271	3.51E-09	0.78678	26739.94	21038.96
A_23_P16	NM_00293	ring finger protein RNF4, SNU	-1.27102	5.73E-07	0.78677	4956.085	3903.368
A_24_P74	M21191	Aldolase A ALDOAP2	-1.27105	0.00004	0.78675	13758.19	10796.44
A_24_P51	NM_00715	Upstream of C10orf57 UNR	-1.2712	1.15E-07	0.78666	6848.866	5395.419
A_23_P58	NM_02409	hypothetical protein MGC5297,	-1.27125	1.65E-06	0.78663	2991.309	2349.004
A_24_P30	NM_00110	A disintegrin-like and repeats ADAM8	-1.27145	0.00001	0.7865	2692.407	2118.04
A_24_P98	NM_01847	uncharacterized protein FAM45B, H	-1.27157	0.00005	0.78643	579.1798	455.4865
A_24_P14	AK125007	WIPI49-like protein WIPI-2	-1.27166	0.0003	0.78637	520.3386	409.1566
A_24_P213144	Unknown	A_24_P213144	-1.27202	0.0003	0.78615	5001.209	3945.983
A_23_P89	NM_00614	LIM and domain protein LASP1, ML	-1.27204	2.17E-08	0.78614	8179.126	6428.658
A_24_P161733	Unknown	A_24_P161733	-1.27208	2.68E-06	0.78611	34884.36	27366.04
A_23_P36	NM_17506	Homo sapiens DDX51, MC	-1.27212	0.00002	0.78609	2525.806	1988.343
A_32_P13	NM_00651	ret finger protein RFP, RNF7	-1.27217	0.00005	0.78606	5530.234	4359.806
A_23_P13	NM_03265	Homo sapiens DHX37, DD	-1.27222	0.00007	0.78603	3531.026	2773.092
A_24_P18	NM_00262	prefoldin 5 PFDN5, MM	-1.27227	1.18E-06	0.786	38533.3	30201.48
A_23_P31	NM_18324	hypothetical protein LOC28625	-1.27235	6.63E-06	0.78594	4430.044	3472.715
A_23_P67	NM_13880	Homo sapiens GALM, BLC	-1.27277	8.77E-06	0.78569	906.2842	711.8683
A_32_P14	NM_00693	small nuclear protein SNRPD1, S	-1.27305	0.00003	0.78552	20333.46	15886.74
A_23_P13	NM_00136	Homo sapiens DKC1, DKC	-1.27308	5.29E-07	0.78549	19081.21	15002.13
A_23_P50	NM_00424	Homo sapiens TRIP10, CII	-1.27308	0.00007	0.7855	853.061	669.7733
A_24_P230466	Unknown	A_24_P230466	-1.27321	0.00003	0.78542	35232.59	27585.91
A_23_P83	NM_02403	hypothetical protein MGC5242,	-1.27323	4.31E-06	0.7854	1411.873	1109.334
A_32_P92	AK023269	Homo sapiens LYCAT, AL	-1.27323	0.00002	0.78541	5905.182	4655.797
A_23_P10	NM_01554	Homo sapiens LRIG1, LIG	-1.27325	0.00004	0.78539	2477.816	1948.513
A_23_P12	NM_00123	Homo sapiens CAV2, CAV	-1.27354	0.00025	0.78521	2161.24	1701.579
A_24_P24	NM_00538	protein-L-isomerase PCMT1	-1.27392	3.80E-06	0.78498	5936.292	4644.785
A_23_P38	NM_14499	Homo sapiens LRRC45, M	-1.27406	0.00075	0.78489	1995.853	1569.977
A_23_P15	NM_00412	G protein-coupled receptor GPS1	-1.27409	4.81E-08	0.78487	3690.055	2894.714
A_23_P27	NM_00293	ribonucleoprotein RNASEH1,	-1.27438	0.00036	0.78469	4641.893	3669.009
A_23_P21	NM_01593	CGI-09 protein CGI-09, MC	-1.27441	0.00005	0.78468	3737.114	2942.323
A_24_P60	BC048794	Hypothetical protein LOC33904	-1.27483	0.00014	0.78442	1155.069	907.1686

A_24_P83	NM_00566	Wolf-Hirsch WHSC2,NI	-1.275	2.85E-06	0.78432	1003.704	786.984
A_32_P11	NM_02040	CGI-203 pr C6orf149,C	-1.27511	3.64E-07	0.78425	3839.682	3008.381
A_23_P50	NM_03301	FAST kina: FASTK,FA	-1.27512	1.00E-06	0.78424	19406.56	15211.71
A_24_P28	AL833099	CGI-14 pro CGI-14	-1.27525	0.00001	0.78416	2464.213	1942.68
A_23_P28	NM_01608	apoptosis r C2orf28,p1	-1.27531	2.06E-06	0.78412	8252.054	6487.617
A_23_P26	NM_02484	Homo sapi PCNT1,PC	-1.27565	6.30E-07	0.78392	6347.954	4985.266
A_24_P20	NM_02099	B-cell CLL/ BCL7A,BC	-1.27568	6.21E-07	0.7839	1223.283	959.0156
A_24_P410246	Unknown	A_24_P410	-1.27588	0.00023	0.78377	6461.784	5053.165
A_23_P11	NM_00640	Homo sapi PRDX4,AC	-1.27591	0.00094	0.78375	66532.55	51710.02
A_23_P15	NM_00465	Homo sapi PNPLA4,G	-1.27594	0.00009	0.78373	7430	5847.084
A_23_P33	NM_05816	Homo sapi TRIM6,RNI	-1.27605	0.00015	0.78367	931.7526	730.8325
A_24_P67748	Unknown	A_24_P677	-1.27613	2.93E-07	0.78362	2247.633	1761.33
A_23_P65	NM_01837	Homo sapi SYNJ2BP,,	-1.27617	9.41E-09	0.78359	1713.175	1342.431
A_24_P83	AK026647	CDNA: FL: AK026647	-1.27627	8.72E-07	0.78354	4704.89	3687.92
A_23_P15	BC021714	PTPRF int: PPFIBP2	-1.27635	1.37E-08	0.78349	1923.561	1507.17
A_23_P30	NM_14502	Homo sapi C6orf136,Λ	-1.27658	0.00002	0.78334	1585.477	1244.156
A_24_P19	NM_02046	KIAA1387 SMEK2,KI/	-1.27679	4.54E-06	0.78321	1551.928	1217.425
A_23_P50	NM_15273	Homo sapi HOXA9,HC	-1.27697	9.25E-08	0.7831	1572.162	1231.373
A_24_P195164	Unknown	ENST0000	-1.27719	0.00008	0.78297	35926.13	27990.61
A_24_P93	AK095167	CDNA FLJ: AK095167	-1.27747	0.0002	0.7828	940.2734	736.4806
A_24_P92	AF233453	Protein kin: PRKCBP1	-1.2775	0.00002	0.78278	2654.603	2080.768
A_24_P33	BC007880	KIAA1970 KIAA1970	-1.2775	0.00004	0.78278	6792.802	5306.338
A_24_P307126	Unknown	A_24_P307	-1.27763	1.71E-06	0.7827	1554.318	1217.227
A_23_P20	NM_00398	Homo sapi PRC1,MGC	-1.27773	6.61E-07	0.78264	12503.1	9797.646
A_24_P35	AK129956	Nuclear fac NFIC	-1.27774	0.00002	0.78263	6360.189	4997.374
A_24_P10	NM_19885	Homo sapi TRIM50C	-1.27778	0.00003	0.78261	1433.001	1117.656
A_23_P18	NM_01590	Homo sapi LAP3,LAP,	-1.2779	0.00071	0.78253	5124.136	4032.89
A_24_P58	NM_03270	tubulin alpt TUBA6,MC	-1.27793	0.00062	0.78252	32922.02	25661.18
A_23_P10	NM_02037	Homo sapi AVEN,PDC	-1.27795	0.00009	0.7825	7178.103	5637.76
A_23_P15	NM_08088	Homo sapi DBN1,D0S	-1.27809	1.99E-06	0.78242	2579.497	2017.087
A_24_P25	NM_01835	hypothetic: ABCF3,FL	-1.2781	8.88E-06	0.78241	1699.813	1332.245
A_23_P25	NM_03146	Homo sapi CD99L2,M	-1.27831	2.17E-07	0.78228	21156.38	16575.46
A_23_P33	NM_00238	Homo sapi MAZ,PUR1	-1.27833	8.62E-07	0.78227	977.4992	764.5201
A_32_P54	NM_00600	ubiquinol-c UQCRH	-1.27834	2.17E-06	0.78226	127429.4	99326.44
A_23_P23	NM_00681	Homo sapi MLLT11,AI	-1.27837	4.74E-06	0.78225	25654.41	20139.75
A_32_P10	AK056836	Hypothetic: LOC15016	-1.27844	0.00061	0.78221	571.171	446.8554
A_24_P35	AF161353	Chromosor C9orf126	-1.27845	2.11E-07	0.7822	1622.508	1269.18
A_32_P180265	Unknown	THC18073	-1.27854	2.29E-07	0.78214	1424.68	1114.034
A_23_P35	NM_03081	hypothetic: MGC11335	-1.27859	4.43E-06	0.78211	2362.198	1846.41
A_23_P21	NM_02471	Chromosor C9orf86	-1.2787	7.91E-07	0.78204	14315.51	11206.78
A_32_P35	AF343078	Homo sapi TOB3	-1.27886	0.00011	0.78195	1729.563	1355.397
A_23_P16	NM_00414	Homo sapi NINJ1,NIN	-1.27899	0.00015	0.78187	7752.335	6030.262
A_24_P30	NM_00483	glutathione GSTO1,P2	-1.27903	0.00032	0.78184	20746.46	16102.04
A_32_P15	NM_00300	succinate c SDHD,PGL	-1.27908	0.00015	0.78181	5648.483	4442.465
A_32_P50	NM_00632	RAN, mem RAN,TC4,/	-1.27934	7.42E-07	0.78165	341420.9	267476.1
A_23_P13	NM_01649	Homo sapi MRPL37,M	-1.27946	9.34E-09	0.78158	9260.01	7241.008
A_24_P409115	ITBA2 prot: ENST0000		-1.27956	2.89E-06	0.78152	4620.474	3606.854
A_24_P91	AV760375	Megakaryo MKL1	-1.27958	0.00001	0.78151	7218.354	5658.354
A_23_P80	NM_00522	Homo sapi E2F1,RBP:	-1.27959	0.0003	0.7815	2671.19	2083.297
A_23_P51	NM_00452	Homo sapi MGST3,G5	-1.27967	0.00002	0.78145	54799.57	42580.91
A_24_P76	XM_29266	similar to NLOC34265	-1.27973	0.00074	0.78142	11429.48	9026.748
A_23_P13	NM_01605	Homo sapi PPIL1,CYF	-1.27977	0.0005	0.78139	6478.05	5088.02

A_23_P10	NM_00606	Homo sapi	COX4NB,N	-1.27989	0.00006	0.78132	15572.51	12220.7
A_24_P36	NM_05284	hypothetic	MMAB,ATF	-1.28011	8.58E-07	0.78118	6146.016	4809.424
A_23_P13	NM_01657	PTX1 prote	PTX1	-1.28018	0.00005	0.78114	2952.845	2325.334
A_23_P13	NM_00655	SNARE pr	YKT6	-1.2804	0.00001	0.78101	900.8272	703.6993
A_24_P55	AK094583	Hypothetic	LOC33974	-1.28041	0.00075	0.781	761.7712	594.8893
A_23_P14	NM_00455	Homo sapi	NDUFS2	-1.28047	2.39E-06	0.78096	3171.436	2481.693
A_23_P15	NM_01232	Homo sapi	LSM4,YER	-1.28089	0.00001	0.78071	16757.57	13123.15
A_23_P79	NM_00124	Homo sapi	ENTPD6,C	-1.28091	0.00005	0.78069	3961.238	3107.123
A_23_P20	NM_00325	Homo sapi	THRA,AR7	-1.28097	0.00008	0.78066	3181.137	2479.776
A_23_P33	NM_02429	Homo sapi	DERL1,DE	-1.28115	0.00083	0.78055	2046.672	1603.977
A_23_P10	NM_00504	PWP2 (per	PWP2H,PV	-1.28138	5.94E-07	0.78041	1546.807	1206.124
A_32_P18	NM_01450	ubiquitin c	UBE2S,E2	-1.28142	9.23E-07	0.78039	18129.7	14152.21
A_23_P21	NM_01428	homolog of	EXOSC2,p	-1.28149	1.04E-07	0.78034	1392.462	1086.858
A_24_P375360	Unknown	A_24_P37		-1.28162	3.45E-07	0.78026	15207.45	11889.07
A_23_P20	NM_00235	mannose-6	M6PR,SMF	-1.2819	1.85E-07	0.78009	1516.87	1183.493
A_23_P10	NM_00279	Homo sapi	PSMB3,HC	-1.28201	0.00001	0.78003	13152.86	10212.86
A_23_P15	NM_00381	a disintegri	ADAM15,M	-1.28207	0.00083	0.77999	1307.323	1022.525
A_23_P16	NM_01605	Homo sapi	MRPL48,C	-1.28213	1.14E-08	0.77995	5840.106	4552.779
A_23_P25	NM_02215	Homo sapi	DUS1L,DU	-1.28213	5.83E-09	0.77995	4195.298	3271.946
A_23_P20	NM_01404	DKFZP564	LRP10,MS	-1.28219	0.00014	0.77991	4994.936	3919.628
A_23_P10	NM_00253	ornithine d	OAZ2	-1.2822	4.04E-07	0.77991	3201.381	2494.169
A_24_P39	D87447	KIAA0258	KIAA0258	-1.28242	3.35E-08	0.77977	1333.102	1039.464
A_24_P21	NM_02291	hypothetic	NOL6,NRA	-1.28266	0.00003	0.77963	688.1828	536.6419
A_23_P26	NM_01788	Homo sapi	HCFC1R1,	-1.28276	1.20E-06	0.77957	5982.062	4651.932
A_23_P12	NM_00445	Homo sapi	ERH,DROI	-1.28288	0.00061	0.7795	31226.54	24147.79
A_24_P30	AK054966	Ankylosis,	ANKH	-1.28292	0.00018	0.77947	500.6047	390.1863
A_24_P144084	Unknown	A_24_P14		-1.28323	5.87E-08	0.77928	6343.916	4949.414
A_24_P936252	Unknown	THC18329		-1.28345	8.13E-09	0.77915	4480.273	3492.868
A_23_P69	NM_17804	Homo sapi	ACTL6A,A	-1.28346	6.06E-07	0.77914	12018.87	9378.479
A_23_P13	NM_03228	Leucine zip	LDOC1L	-1.2836	0.00016	0.77906	3580.637	2804.573
A_32_P112546	Unknown	A_32_P11		-1.28364	0.00014	0.77903	50003.61	38756.16
A_32_P27706	L10908	Gc THC19450		-1.28369	1.79E-08	0.77901	1971.276	1535.946
A_23_P78	NM_00446	phenylalan	FARSLA,F	-1.28386	3.04E-07	0.7789	1793.186	1395.537
A_24_P73	NM_00209	G1 to S ph	GSPT1,GS	-1.28419	4.16E-07	0.7787	2175.939	1694.595
A_23_P14	NM_00259	programme	PDCD2,RF	-1.28423	6.16E-07	0.77868	2828.507	2206.315
A_23_P12	NM_01418	Homo sapi	MRPL22,R	-1.28431	0.00014	0.77863	37908.71	29652.24
A_24_P21	NM_03231	hypothetic	WBSCR18	-1.28448	0.001	0.77853	1124.856	877.7966
A_23_P10	NM_03143	hypothetic	THAP2,DK	-1.28468	0.00088	0.77841	623.1968	485.2213
A_23_P30	NM_13913	Homo sapi	NUP98,AD	-1.28476	3.54E-06	0.77836	2317.636	1807.548
A_32_P16	AK023018	Homo sapi	AK023018	-1.28481	0.00028	0.77832	2171.822	1692.08
A_23_P11	AK023379	Splicing fac	SRP46	-1.28481	4.44E-08	0.77832	1801.417	1402.274
A_32_P69	AK023933	BTB (POZ)	HSPC063	-1.28494	0.00092	0.77825	1321.908	1031.463
A_24_P48	BC045718	LOC44046	SH3GLP3,	-1.285	0.00012	0.77821	3788.397	2962.125
A_23_P11	NM_00138	Homo sapi	DPH1,DPI	-1.28505	5.78E-08	0.77818	13537.55	10518.7
A_23_P36	NM_00662	Homo sapi	AHCYL1,D	-1.2856	0.00001	0.77785	21586.5	16731.8
A_23_P87	NM_00310	Homo sapi	SORL1,LR	-1.28562	1.38E-07	0.77784	999.2098	777.251
A_23_P50	NM_01761	Homo sapi	DONSON,I	-1.28564	0.00017	0.77782	5806.815	4542.484
A_24_P65	XM_21136	hypothetic	LOC28418	-1.2857	0.00022	0.77778	36468.2	28104.41
A_23_P89	NM_00444	Homo sapi	ERBB2,NE	-1.28601	9.73E-06	0.7776	1617.775	1259.406
A_23_P57393	Unknown	THC18373		-1.28608	0.00067	0.77756	2100.562	1646.814
A_23_P11	NM_00258	Homo sapi	PARN,DAN	-1.28612	0.00056	0.77753	1066.602	831.1918
A_32_P18	AK095472	Hypothetic	DKFZp762	-1.28622	0.00006	0.77747	576.6297	448.3251

A_23_P51	NM_02410	Homo sapi	WDR18, M	-1.28622	0.00007	0.77747	8253.215	6436.391
A_24_P69	NM_00208	glutamic-o	GOT2, FLJ	-1.28626	0.00031	0.77745	1235.964	961.8092
A_23_P11	NM_00652	Homo sapi	WNT6	-1.2869	0.00035	0.77706	1697.042	1320.109
A_24_P401124		Unknown	A_24_P40	-1.28698	0.00016	0.77701	6909.465	5352.117
A_24_P20	NM_00138	diphtheria to	DPH1, DPH	-1.28701	0.00031	0.777	5594.182	4365.619
A_23_P14	NM_00662	Homo sapi	ZNF271, H	-1.28708	1.47E-07	0.77695	1427.018	1108.837
A_24_P19	NM_01595	eNOS inter	NOSIP, CG	-1.28714	4.90E-06	0.77692	999.4581	776.1506
A_32_P30	NM_00660	inactive pr	PTGES3, P	-1.28721	4.38E-07	0.77687	138155	107068.4
A_23_P21	NM_03238	Homo sapi	GFM2, EFC	-1.28723	0.00061	0.77686	8874.387	6931.473
A_24_P92	BC041879	Homo sapi	LOC39070	-1.28728	2.00E-08	0.77683	4368.626	3392.023
A_23_P12	NM_00530	Homo sapi	GRK5, GPF	-1.28742	8.15E-07	0.77675	2523.391	1959.746
A_24_P23	BC009369	Hypothetic	LOC33922	-1.28747	2.84E-08	0.77671	5990.956	4658.059
A_23_P12	NM_14477	Homo sapi	APOA1BP,	-1.28751	0.00043	0.77669	17288.29	13349.48
A_24_P306704		Unknown	A_24_P30	-1.28768	0.00055	0.77659	36559.95	28188.16
A_24_P21	NM_00437	COX11 (ye	COX11, CC	-1.28792	8.56E-07	0.77645	3854.594	2999.198
A_24_P26	XM_06142	similar to	SLOC11935	-1.28803	0.00021	0.77638	44008.01	33936.14
A_23_P28	NM_00324	Homo sapi	THOP1, MF	-1.28807	2.09E-06	0.77635	26284.04	20454.88
A_32_P16	XM_37067	similar to	h LOC38783	-1.28828	4.29E-08	0.77623	5399.231	4188.023
A_23_P25	NM_00270	protein phc	PPP1CB, P	-1.2883	0.00006	0.77622	990.8624	769.4628
A_24_P358131		Unknown	ENST0000	-1.28834	0.00002	0.7762	13787.74	10662.95
A_23_P25	NM_00671	activator of	ASK, DBF4	-1.28839	0.00003	0.77617	1892.994	1481.391
A_23_P91	NM_08092	Homo sapi	LOC91219	-1.2885	2.27E-07	0.77609	983.9351	763.6469
A_23_P11	NM_01630	protein x 0	(ATPBD1C,	-1.28856	0.00099	0.77606	5359.222	4188.842
A_24_P206848		ALU1_HUM	THC18433	-1.28857	1.25E-09	0.77606	2614.381	2028.886
A_24_P17	NM_01568	B9 protein	EPPB9, B9	-1.28865	0.00002	0.77601	1541.323	1194.844
A_23_P12	NM_00314	Homo sapi	SSR2, TLA	-1.28895	0.00037	0.77582	65728.3	50534.66
A_24_P79	XM_37307	similar to	h LOC39176	-1.28905	0.00075	0.77577	139580.8	107126.2
A_23_P11	NM_01608	Homo sapi	STYXL1, D	-1.28908	0.0004	0.77574	1360.842	1057.719
A_24_P87	NM_01240	protein kin	PRKCBP1,	-1.28909	1.72E-06	0.77574	1243.37	964.09
A_23_P25	NM_01793	hypothetic	FLJ20718	-1.28925	9.05E-07	0.77565	10662.98	8281.39
A_23_P54	NM_00458	RAB27A, n	RAB27A, G	-1.28939	5.06E-06	0.77556	2533.272	1960.065
A_23_P16	NM_00421	fibroblast g	FIBP, FGFI	-1.28943	0.00007	0.77554	1231.035	953.433
A_23_P25	NM_01540	Homo sapi	HDAC7A, F	-1.28945	2.59E-09	0.77553	2116.406	1641.398
A_24_P45	NM_02183	hypothetic	FLJ21839	-1.28955	4.22E-07	0.77546	1038.424	805.4456
A_23_P15	NM_00039	ATP-bindin	ABCC2, DJ	-1.28966	0.00004	0.7754	9055.494	6990.518
A_24_P10	AF217993	Hypothetic	MGC1552	-1.28975	0.00002	0.77534	1297.998	1007.334
A_32_P10	NM_01900	Homo sapi	ARMCX6, F	-1.28977	6.59E-06	0.77533	9747.37	7588.242
A_23_P20	NM_00015	glucosidas	GBA, GCB,	-1.28977	1.34E-06	0.77533	1767.871	1369.571
A_23_P40	BC021714	PTPRF int	PPFIBP2	-1.28978	0.00004	0.77532	1137.094	883.8558
A_23_P37	NM_13879	Homo sapi	OACT2, FL	-1.28982	0.00014	0.7753	948.7835	736.5659
A_23_P79	NM_02431	hypothetic	PLEKHF1,	-1.29016	7.14E-08	0.7751	3828.586	2970.718
A_23_P26	NM_00637	membrane	VAT1, VAT	-1.29025	4.50E-06	0.77504	16798.45	12979.44
A_23_P15	NM_01814	hypothetic	FAM82C, p	-1.29026	2.88E-06	0.77504	6863.314	5339.52
A_23_P40	NM_17858	Homo sapi	HM13, H13	-1.29035	1.36E-07	0.77498	1958.737	1519.154
A_24_P56	NM_01657	GMPR2 fo	GMPR2, M	-1.2904	0.00017	0.77496	12794.22	9975.092
A_24_P186746		Unknown	A_24_P18	-1.29057	8.00E-06	0.77485	34916.8	26951.96
A_23_P16	NM_01406	Homo sapi	DBNL, SH3	-1.29067	0.00011	0.77479	734.5496	569.54
A_32_P34	AK125119	Chromosor	C6orf68	-1.29089	0.00003	0.77466	2116.207	1640.152
A_23_P92	NM_02192	Homo sapi	FGFRL1, F	-1.2909	0.00001	0.77465	617.5158	478.3734
A_24_P894763		Unknown	A_24_P89	-1.29096	8.67E-08	0.77462	4711.521	3654.96
A_24_P88	NM_01650	putative	S1PS1D, pNO	-1.29097	0.00082	0.77461	3376.791	2632.14
A_23_P14	NM_00132	Homo sapi	CSTF2, Cst	-1.29102	8.22E-07	0.77458	5394.811	4189.997

A_23_P43	NM_00282	prothymosi	PTMA, TMS	-1.29105	0.00064	0.77456	52249.04	40707.86
A_24_P23	NM_00279	proteasom	PSMB1, HC	-1.29109	1.27E-06	0.77454	74131.42	57229.34
A_23_P11	NM_01602	Homo sapi	DHRS7, ret	-1.29129	2.52E-06	0.77442	3165.068	2443.946
A_24_P20	BC029662	Homo sapi	C20orf142,	-1.29133	0.00002	0.77439	904.3215	700.8695
A_23_P43	BC039468	Nucleopor	NUP50	-1.29137	6.29E-09	0.77437	3721.071	2882.313
A_24_P93	BC001027	Transmem	TMEM4	-1.29142	0.00047	0.77434	5365.459	4175.273
A_24_P35	NM_00578	proteasom	PSME3, Ki	-1.29147	8.73E-06	0.77431	2081.156	1613.229
A_24_P94	AB033058	Discs, larg	DLG3	-1.29164	0.00012	0.77421	1266.614	982.549
A_24_P12	NM_00270	phosphorib	PPAT, GPA	-1.29187	1.18E-06	0.77407	3510.438	2714.756
A_24_P67	NM_02483	C3HC4-typ	ZNF403, DI	-1.29223	2.80E-06	0.77385	3807.978	2937.635
A_23_P20	NM_02518	lin-10 prote	LIN10, lin-1	-1.29223	4.01E-07	0.77386	1764.975	1364.999
A_23_P97	NM_01797	Homo sapi	MRPL20, L	-1.29226	1.93E-08	0.77384	28715.87	22257.23
A_24_P33	NM_01900	hypothetic	FLJ20232, I	-1.29229	0.00002	0.77382	3253.519	2511.665
A_23_P69	NM_00506	Homo sapi	SLAH2, hSi	-1.29236	0.00002	0.77378	757.7466	586.4066
A_23_P73	NM_00601	DNA segm	DXS9879E	-1.29257	3.47E-08	0.77365	31356.79	24239.44
A_32_P20	BG178211	Mesenchyr	BG178211	-1.2926	0.00002	0.77364	3198.648	2479.661
A_23_P16	NM_00628	Homo sapi	TESK1	-1.29277	0.00058	0.77353	2826.007	2198.54
A_23_P10	NM_01430	RNA bindir	RBM9, RTA	-1.29282	0.00044	0.7735	588.4047	455.1906
A_23_P11	NM_01596	cell death-r	NDUFA13,	-1.29295	2.93E-10	0.77343	13823.9	10692.25
A_24_P73	AK093246	Ribosomal	RPL13	-1.29302	5.85E-09	0.77338	1493.522	1155.111
A_23_P65	NM_01630	60S riboso	C15orf15, L	-1.29304	1.07E-06	0.77337	13833.42	10724.08
A_24_P264004	Unknown	A_24_P26		-1.29317	5.85E-07	0.77329	2033.515	1571.73
A_23_P42	NM_03201	Homo sapi	MRPS24, H	-1.29317	0.0003	0.77329	69445.59	53242.54
A_32_P59	NM_00300	succinate c	SDHD, PGL	-1.29322	0.0001	0.77326	3071.735	2386.328
A_32_P18	NM_02283	hypothetic	C1orf80, FL	-1.29331	0.0006	0.77321	679.7023	525.6088
A_23_P11	NM_00701	Homo sapi	DDX52, RC	-1.29334	7.70E-07	0.77319	1990.013	1541.689
A_23_P34	BC036837	BA299N6.3	LOC19843	-1.29352	0.00037	0.77308	530.1266	409.8348
A_32_P53	AF060511	Homo sapi	My016 pro	-1.29356	3.38E-07	0.77306	57714.31	44490.84
A_23_P15	NM_00251	Homo sapi	NME3, DR-	-1.29363	5.37E-09	0.77302	20798.31	16085.42
A_23_P15	NM_01210	Homo sapi	ARL2BP, B	-1.29379	0.00016	0.77292	10964.17	8499.916
A_23_P36	NM_01327	Homo sapi	ANKRD11,	-1.2939	1.74E-06	0.77286	7725.342	5982.258
A_23_P24	NM_15226	Homo sapi	SLC39A13	-1.29407	0.00002	0.77276	3706.867	2870.194
A_24_P37	NM_00449	hepatoma-	HDGF, HM	-1.29411	0.00088	0.77273	21150.89	16479.29
A_23_P16	NM_17060	Homo sapi	MLX, MAD	-1.2943	3.66E-08	0.77262	1900.832	1467.464
A_23_P14	NM_00485	Homo sapi	STX8, CAR	-1.29431	0.00004	0.77261	2746.126	2126.783
A_23_P31	AJ277796	Solute carr	SLC16A1	-1.29455	1.53E-06	0.77247	793.8899	613.3183
A_32_P61	NM_00600	ubiquinol-c	UQCRFS1	-1.29474	4.74E-08	0.77235	24654.99	19079.79
A_23_P42	NM_14463	Homo sapi	CHCHD4, F	-1.29501	0.00005	0.77219	7000.365	5429.177
A_23_P13	NM_01814	Homo sapi	RNMTL1, H	-1.29507	0.00013	0.77216	4601.819	3574.258
A_24_P20	NM_00673	homeo box	HOXA2, HC	-1.29531	5.50E-08	0.77202	1520.113	1173.603
A_23_P41	NM_01534	Homo sapi	BRRN1, CA	-1.29563	0.00007	0.77182	2726.698	2094.073
A_24_P264293	Unknown	A_24_P26		-1.29565	7.49E-09	0.77182	12103.49	9335.303
A_23_P46	NM_15237	hypothetic	C1orf131, C	-1.29571	0.00026	0.77178	5131.805	3989.17
A_23_P33	NM_02112	Homo sapi	POLR2L, R	-1.296	0.00043	0.7716	108509.5	82852.95
A_23_P15	NM_01774	Homo sapi	BCOR, MA	-1.29605	6.41E-06	0.77157	1667.683	1286.296
A_23_P95	NM_00435	Homo sapi	CD151, GP	-1.29608	0.00026	0.77156	2287.566	1757.656
A_23_P29	NM_00331	Homo sapi	TST, RDS, I	-1.29616	8.70E-07	0.77151	22465.17	17310.88
A_23_P30	NM_18183	Homo sapi	UBE2D2, U	-1.29627	6.50E-08	0.77145	4894.14	3781.099
A_24_P127098	Unknown	A_24_P12		-1.29668	0.00019	0.7712	756.2617	582.7078
A_24_P17	NM_02519	Homo sapi	GRPEL1, H	-1.29669	0.00006	0.77119	1975.519	1530.661
A_23_P16	NM_00534	Homo sapi	HSPCA, HS	-1.29703	0.00027	0.77099	105279.8	80502.16
A_23_P26	NM_02510	hypothetic	FLJ13909	-1.2972	7.97E-07	0.77089	5411.582	4173.228

A_32_P168727	Unknown	A_32_P168727	-1.29737	0.00013	0.77079	2662.127	2072.025
A_24_P247464	Unknown	A_24_P247464	-1.29771	1.41E-06	0.77059	1999.67	1540.679
A_23_P15	NM_19427 HESB like	HBLD1	-1.2978	1.49E-08	0.77053	7055.929	5442.452
A_24_P354488	ASAL_HUM	THC19567	-1.2978	0.00001	0.77054	1357.081	1045.966
A_23_P15	NM_01805 homolog of	ARHGAP1	-1.29786	0.00009	0.7705	2954.518	2287.248
A_32_P177937	ALU7_HUM	THC19694	-1.29791	0.00031	0.77047	523.9334	403.6841
A_23_P12	NM_01900 Spindlin	far SPIN2	-1.29795	0.00007	0.77045	2273.215	1747.363
A_23_P33	NM_01592 Androgen-i	APRIN	-1.29797	0.00003	0.77043	853.4316	657.8389
A_23_P10	NM_01640 Ufm1-conj	UFC1,HSP	-1.29805	2.97E-08	0.77039	6084.412	4696.224
A_23_P72	AK095357 ELMO	dorr ELMOD2	-1.29813	3.61E-06	0.77034	2210.245	1704.953
A_32_P19	NM_17588 Homo sapi	PRPS1L1,I	-1.29828	7.10E-09	0.77025	1501.301	1156.519
A_24_P22	NM_02274 Homo sapi	ATPAF1,A	-1.29845	6.11E-08	0.77015	1750.673	1349.146
A_23_P92	NM_01240 pp32 relate	ANP32C,P	-1.29863	0.00099	0.77004	766.3395	591.7076
A_24_P471242	Unknown	A_24_P471242	-1.29874	0.00001	0.76997	14155.32	10849.68
A_23_P26	NM_02237 G protein b	GBL,MGC	-1.29879	0.00029	0.76995	735.3809	565.3494
A_23_P42	AB028986 Ubiquitin	s1 USP22	-1.29895	1.57E-06	0.76985	2738.775	2111.853
A_24_P34	XM_37771 Homo sapi	XM_37771	-1.29898	0.00053	0.76984	2332.062	1780.431
A_23_P29	NM_00233 low density	LRPAP1,R	-1.29912	0.00034	0.76975	964.1787	742.9248
A_23_P14	AL161962 Zinc finger,	ZDHHC9	-1.29918	1.11E-08	0.76972	3551.012	2734.1
A_24_P25	NM_02115 hypothetic	TXNDC13,	-1.29936	2.59E-06	0.76961	2404.136	1852.359
A_24_P38	NM_01631 polymerase	POLR3K,C	-1.29939	7.56E-07	0.76959	15302.34	11740.19
A_23_P11	NM_01598 Homo sapi	HEBP1,HB	-1.29967	0.0002	0.76942	13776.46	10676.28
A_32_P166356	Unknown	THC19174	-1.29994	2.46E-07	0.76927	1132.706	871.7227
A_23_P16	NM_03129 hypothetic	CDCA3,GF	-1.30014	1.13E-07	0.76915	2119.569	1631.547
A_32_P20	AL569368 Similar to	t1 FKBP1A	-1.30024	0.00018	0.76909	2898.783	2223.724
A_23_P25	NM_00262 Homo sapi	PFN2,PFL	-1.30032	0.00018	0.76904	3039.178	2350.749
A_24_P39	NM_00437 COX11 (ye	COX11,CC	-1.30059	1.42E-06	0.76888	2227.261	1712.163
A_23_P40	NM_02393 hypothetic	C20orf116,	-1.3006	7.00E-10	0.76887	4879.01	3752.273
A_23_P30	NM_00104 Homo sapi	SRD5A1	-1.30064	7.96E-07	0.76885	881.9437	678.1872
A_23_P42	NM_00717 Homo sapi	CIT,CRIK,S	-1.30066	1.28E-07	0.76884	3764.041	2889.034
A_23_P12	NM_01866 Homo sapi	PXMP2,PM	-1.30069	1.69E-07	0.76882	6495.917	4998.729
A_24_P74	NM_17754 Homo sapi	SNRPD2,S	-1.3007	0.00006	0.76882	51002.26	38941.45
A_24_P13	NM_01793 hypothetic	FLJ20718	-1.30077	0.00042	0.76878	808.9797	623.0544
A_32_P46	NM_17349 Homo sapi	LSM11,FL	-1.3009	0.0002	0.7687	842.1786	647.4885
A_23_P92	NM_02209 hypothetic	CIDEC,Fsp	-1.30108	3.10E-06	0.76859	1194.825	919.7983
A_32_P15	NM_00022 keratin 18	KRT18,K18	-1.30108	0.00004	0.76859	209541.2	160131.4
A_24_P12	NM_00474 discs, large	DLG5,PDL	-1.30117	0.00002	0.76854	3934.43	3037.827
A_23_P40	NM_14566 Dbf4-relate	DRF1,ASK	-1.30158	0.00013	0.76829	4920.44	3814.392
A_32_P16	NM_01539 Homo sapi	ANAPC13,	-1.3016	0.00005	0.76828	3384.898	2606.879
A_23_P37	AK097804 Hypothetic	LOC28480	-1.30167	6.99E-07	0.76824	1167.578	896.4037
A_23_P67	NM_00225 Homo sapi	KCNN4,IK	-1.30172	0.00001	0.76822	2807.311	2158.264
A_23_P40	NM_01476 Homo sapi	DAZAP2,M	-1.3019	4.40E-07	0.76811	23779.16	18210.7
A_23_P14	NM_00337 Homo sapi	VDAC1,PC	-1.30196	0.00064	0.76807	58191.35	44165.2
A_32_P15	AK056140 CDNA	FLJ:AK056140	-1.30198	0.00076	0.76806	679.9822	522.3179
A_24_P83	NM_19884 Homo sapi	C9orf10OS	-1.30224	6.92E-10	0.76791	12108.21	9293.863
A_23_P41	NM_02210 CDW92 an	CDW92	-1.30233	3.10E-07	0.76785	865.4894	664.6287
A_24_P83	NM_00309 small nucle	SNRPC,FL	-1.30248	4.88E-07	0.76777	1013.486	777.7186
A_23_P80	NM_00368 DNA segm	D21S2056I	-1.30269	1.06E-10	0.76765	7045.332	5408.506
A_24_P82	NM_01868 zinc finger	ZNF313,RI	-1.30272	1.54E-09	0.76762	2521.945	1935.517
A_24_P365015	Unknown	THC18585	-1.3028	8.38E-06	0.76758	2203.514	1690.017
A_23_P43	NM_18169 Homo sapi	C10orf9,CF	-1.30291	0.00009	0.76751	890.4317	684.5789
A_24_P161827	Unknown	A_24_P161827	-1.30292	0.00017	0.76751	8093.869	6177.428

A_24_P58:	BC039246 Homo sapi	BC039246	-1.30308	0.00083	0.76741	4160.5	3209.23
A_24_P247303	Unknown A_24_P24		-1.30364	8.61E-06	0.76708	13459.66	10284.38
A_23_P24:	NM_01425 Homo sapi	TMEM5,HF	-1.30382	1.38E-06	0.76698	13875.03	10688.41
A_24_P23:	AB018310 Death-indu	KIAA0767	-1.30402	0.00004	0.76686	1071.941	822.5034
A_23_P16:	NM_00191 Homo sapi	DCI	-1.30419	0.00023	0.76676	37764.61	28719.44
A_23_P14:	NM_00349 Homo sapi	ARD1A,TE	-1.30424	0.00009	0.76673	4312.47	3328.739
A_24_P29:	NM_01542 aminoadip	AASDHPP	-1.30426	4.08E-06	0.76672	8707.648	6702.898
A_23_P82:	NM_00310 SRY (sex c	SOX4,EVI1	-1.30431	0.00002	0.76669	624.6627	478.9088
A_24_P82:	NM_00619 proliferati	PA2G4,EB	-1.30432	0.00024	0.76668	9586.507	7372.978
A_23_P50:	NM_00340 Homo sapi	YWHAB,H	-1.30443	0.00004	0.76662	73508.44	55937.16
A_23_P14:	NM_00063 Homo sapi	GSR,MGC	-1.30451	0.00006	0.76657	5518.077	4207.296
A_24_P115621	ALU8_HUM	THC19007	-1.30467	7.08E-07	0.76648	3420.77	2628.615
A_24_P32:	NM_01620 hematopoi	HSPC176,I	-1.3049	9.21E-07	0.76634	3078.128	2354.289
A_23_P64:	NM_03201 Homo sapi	RNF26,MC	-1.30533	0.00004	0.76609	4936.758	3794.932
A_23_P20:	NM_02513 melanoma/	ARMC9,AF	-1.30539	0.0006	0.76606	374.0417	286.5096
A_32_P11:	NM_01441 Siah-intera	CACYBP,S	-1.30547	2.53E-07	0.76601	29115.81	22236.52
A_23_P60:	NM_02100 Homo sapi	PTTG3	-1.30582	0.00058	0.7658	2185.24	1668.675
A_32_P84:	BC041995 Homo sapi	BC041995	-1.30599	4.54E-07	0.7657	957.3494	732.7677
A_23_P16:	NM_17686 Homo sapi	PSME3,Ki,	-1.30603	8.13E-06	0.76568	2419.676	1850.933
A_24_P17:	NM_03194 c-Myc targ	CDCA7,JP	-1.30607	4.63E-08	0.76566	986.4855	755.3681
A_23_P13:	NM_00272 Homo sapi	PREP,PE,I	-1.30608	3.77E-10	0.76565	3928.017	3007.546
A_24_P21:	NM_01817 hypothetic	FLJ10661,I	-1.3061	1.61E-10	0.76564	2725.527	2086.802
A_32_P10:	BC052334 Similar to	BC052334	-1.30632	2.95E-08	0.76551	1098.592	841.0096
A_32_P11:	BF686720 Transcribe	BF686720	-1.30653	0.00079	0.76539	513.0294	392.6484
A_24_P39:	NM_00138 diptheria to	DPH2,DPH	-1.30655	1.87E-09	0.76537	2217.424	1696.823
A_23_P42:	NM_13871 Homo sapi	PPT2,G14,	-1.30658	0.00012	0.76536	757.9528	580.3834
A_23_P64:	NM_00481 Homo sapi	DDX23,prp	-1.3072	2.12E-09	0.76499	9613.556	7345.708
A_23_P122650	Unknown A_23_P12:		-1.30721	2.89E-06	0.76499	202520.4	154236
A_23_P70:	NM_01234 Homo sapi	NNT	-1.30756	0.00005	0.76478	945.9929	724.8232
A_23_P15:	NM_00136 Homo sapi	DHODH,DI	-1.30758	3.20E-06	0.76477	1937.302	1484.303
A_23_P25:	NM_08065 similar to	RMGC13017	-1.30776	1.92E-06	0.76467	3961.483	3018.98
A_24_P82:	BC009950 LOC44066	BC009950	-1.30792	2.63E-09	0.76458	1965.863	1502.297
A_23_P20:	NM_00271 protein phc	PPP1CC,P	-1.30798	7.68E-06	0.76454	7011.367	5330.579
A_23_P15:	NM_03211 Katanin p6	KATNAL1	-1.30831	0.00037	0.76434	632.8963	483.8756
A_23_P27:	D50928 Scaffold at	SAFB2	-1.30865	0.00002	0.76414	2953.41	2255.044
A_23_P71:	NM_00623 polymerase	POLD2	-1.30878	1.11E-08	0.76407	1812.611	1385.253
A_23_P29:	NM_00501 Homo sapi	OXA1L	-1.309	1.23E-07	0.76394	4436.468	3384.238
A_24_P272713	Unknown A_24_P27:		-1.30929	0.00061	0.76377	4213.167	3243.235
A_23_P31:	NM_00328 Homo sapi	TPD52L1,I	-1.30933	6.06E-06	0.76375	15327.49	11767.65
A_24_P94:	NM_00622 phosphatid	PITPNA,PI	-1.30937	2.04E-10	0.76373	7864.333	6004.087
A_23_P11:	NM_03263 Homo sapi	GATA2,NF	-1.30953	0.00031	0.76363	1935.987	1480.588
A_24_P14:	NM_01558 DKFZP586	POLDIP2,F	-1.30971	0.00006	0.76353	10394.06	7976.973
A_23_P16:	NM_01567 sentrin/SUI	SEN3,SS	-1.30994	5.55E-07	0.76339	1578.252	1203.85
A_23_P20:	NM_00430 Rho GDP c	ARHGDI A,	-1.31002	1.09E-10	0.76335	3387.288	2585.416
A_24_P22:	XM_37635 Homo sapi	ENST0000	-1.31013	0.0002	0.76328	5837.22	4473.861
A_23_P15:	NM_00600 Homo sapi	UQCRFS1	-1.31024	7.63E-08	0.76322	31779.16	24193.05
A_24_P213336	Unknown A_24_P21:		-1.31025	1.51E-09	0.76321	7972.551	6080.609
A_23_P13:	NM_00473 eukaryotic	ETF1,ERF,	-1.31032	9.86E-06	0.76317	4732.295	3614.627
A_23_P21:	NM_03281 Homo sapi	C9orf100,F	-1.31033	3.95E-08	0.76316	798.3301	609.2561
A_32_P18:	BC007606 Baculoviral	BIRC5	-1.31036	7.69E-09	0.76315	8493.879	6482.376
A_23_P25:	AB037813 Storkhead	DKFZp762	-1.31046	4.03E-06	0.76309	671.246	512.2619
A_24_P854758	Unknown THC19657		-1.31051	0.00054	0.76306	5381.612	4114.101

A_32_P65395	AB019038	THC18668	-1.31054	2.62E-09	0.76305	2658.297	2028.5	
A_24_P123720	Unknown	A_24_P12:	-1.31055	0.00074	0.76304	530.6516	404.7961	
A_24_P25:	NM_03223	Ubiquitin s	USP48	-1.31065	4.43E-06	0.76298	2708.026	2067.06
A_24_P19:	NM_20711	Homo sapi	GTF2H5,TI	-1.31075	7.75E-08	0.76292	5099.218	3899.391
A_23_P21:	NM_00551	Homo sapi	GTF2E1,FI	-1.31076	0.00001	0.76291	3615.672	2742.67
A_24_P27:	NM_00637	serine/thre	STK25,SO	-1.31076	0.00049	0.76292	1497.369	1140.063
A_23_P74:	NM_01441	Homo sapi	CACYBP,S	-1.31094	4.41E-07	0.76281	55546.23	42184.51
A_24_P36:	NM_00430	Rho GDP c	ARHGDI A,	-1.31104	0.00013	0.76275	519.0839	395.9401
A_23_P63:	NM_03233	Homo sapi	C10orf58,N	-1.31109	8.04E-11	0.76273	8320.611	6345.842
A_23_P64:	NM_20127	Homo sapi	MTMR2,C	-1.31115	8.87E-08	0.76269	2532.575	1935.506
A_23_P40:	NM_02469	Homo sapi	SLC25A22	-1.31117	0.00003	0.76268	7861.479	6010.789
A_23_P75:	NM_17593	Homo sapi	PSMD13,p	-1.31118	0.00034	0.76267	2317.955	1767.662
A_24_P90:	BC064610	Selenopho	SEPHS1	-1.31125	5.33E-06	0.76263	1403.806	1072.993
A_24_P34:	NM_15252	Homo sapi	SLC16A14	-1.3113	0.00042	0.7626	1839.413	1411.044
A_24_P384200	Unknown	A_24_P38:		-1.31131	7.65E-10	0.7626	7407.4	5653.141
A_24_P384379	Unknown	ENST0000		-1.31134	0.00041	0.76258	3624.011	2778.12
A_24_P16:	NM_01829	hypothetic	CWF19L1,	-1.31139	2.36E-09	0.76255	1728.781	1318.688
A_23_P13:	NM_01466	Homo sapi	LRRC14,K	-1.31141	0.00005	0.76254	1061.61	812.8682
A_23_P78:	NM_00413	Homo sapi	KRTHA3A,	-1.31147	0.00011	0.7625	548.9131	418.5687
A_24_P187174	Unknown	A_24_P18:		-1.31165	5.08E-06	0.7624	1787.418	1363.885
A_24_P230486	Unknown	A_24_P23:		-1.31167	0.00058	0.76238	5915.076	4478.631
A_23_P21:	NM_01221	Homo sapi	ILF3,DRBF	-1.31174	0.00027	0.76234	7201.354	5499.787
A_23_P66347	Unknown	A_23_P66:		-1.31188	7.11E-06	0.76227	1174.796	895.7806
A_23_P21:	NM_00220	integrin, al	ITGAE,CD	-1.312	5.66E-06	0.7622	8051.408	6159.563
A_24_P41:	NR_00003	Homo sapi	ESTRRA	-1.31205	6.75E-10	0.76217	5096.755	3886.68
A_23_P47:	NM_01784	Homo sapi	MRPL16,L	-1.31213	6.75E-07	0.76212	18781.02	14273.7
A_24_P37939	Y08307	rib	THC18779	-1.31215	8.73E-06	0.76211	924.9813	704.6428
A_23_P32:	NM_05281	caspase re	CARD14,B	-1.31235	0.00064	0.76199	407.6905	310.5724
A_24_P68:	AK123675	Lix1 homol	MGC46719	-1.31249	7.78E-06	0.76191	1265.142	964.6834
A_23_P10:	AB014592	KIAA0692	KIAA0692	-1.31256	5.96E-06	0.76187	1129.367	861.5037
A_23_P73:	NM_01636	clone HQ0	LOC51204	-1.31282	5.30E-06	0.76172	1719.584	1314.043
A_23_P25:	NM_01404	Homo sapi	UNC50,UR	-1.31294	4.01E-06	0.76165	8519.167	6507.773
A_24_P62:	NM_00044	presenilin 2	PSEN2,AD	-1.31304	0.00003	0.76159	1767.948	1351.229
A_23_P52:	NM_00347	Homo sapi	RASSF7,H	-1.31311	0.00002	0.76155	2690.302	2044.604
A_23_P21:	NM_00002	Homo sapi	ADA	-1.31312	2.86E-07	0.76155	10959.38	8332.784
A_23_P62:	NM_00031	Homo sapi	PPT1,PPT	-1.31312	9.64E-09	0.76155	14182.89	10819.54
A_32_P103131	ALU1_HUM	THC20038		-1.3135	3.21E-07	0.76133	5372.369	4098.168
A_23_P51:	NM_00274	Homo sapi	PRKCZ,PK	-1.31367	3.68E-07	0.76123	2791.146	2119.281
A_32_P103464	Unknown	A_32_P10:		-1.31368	0.00002	0.76122	3533.466	2701.686
A_23_P50:	NM_03121	Chromosor	C19orf27	-1.31399	5.07E-09	0.76104	4818.467	3667.19
A_23_P12:	NM_02404	Homo sapi	CUEDC2,N	-1.31418	7.67E-10	0.76093	2896.453	2205.252
A_23_P34:	NM_01444	gamma tub	76P,GCP4	-1.31468	0.00052	0.76064	1093.894	828.7047
A_24_P87:	NM_01797	mitochondr	MRPL20,L	-1.31475	7.82E-07	0.7606	27929.3	21168.66
A_23_P24:	NM_17381	hypothetic	TTC9C,MC	-1.31477	0.00023	0.76059	2001.84	1518.77
A_23_P38:	NM_00644	Homo sapi	PRPF8,PR	-1.31479	0.00011	0.76058	6373.908	4836.1
A_24_P40:	AB030654	Adaptor-rel	AP4S1	-1.31489	0.00058	0.76052	383.6254	291.7469
A_23_P63:	NM_00517	ATP synth	ATP5C1,A	-1.31491	1.25E-07	0.76051	5177.413	3945.985
A_24_P57:	NM_01503	Homo sapi	ZHX3,TIX1	-1.31492	0.00092	0.7605	909.0602	688.9648
A_23_P11:	NM_00643	ADP-ribosy	PARP4,PH	-1.31508	0.00005	0.76041	3722.079	2843.456
A_24_P17:	NM_02429	hypothetic	DERL1,DE	-1.31516	9.10E-07	0.76037	3335.398	2541.916
A_24_P37:	NM_00549	SUMO-1 at	UBA2,SAE	-1.31526	1.58E-06	0.76031	4927.354	3753.848
A_23_P11:	NM_00280	Homo sapi	PSMB9,LV	-1.31573	5.70E-11	0.76004	5974.294	4541.588

A_24_P490704	Unknown	A_24_P490704	-1.31585	0.00097	0.75997	1794.688	1366.258
A_32_P14	NM_20348	hypothetical MGC70870	-1.31612	1.34E-07	0.75981	1438.107	1092.397
A_23_P12	NM_00562	S100 calcit S100A11, N	-1.31636	0.00003	0.75967	9771.361	7472.718
A_23_P37	NM_02032	ATP-binding ABCD4, P7	-1.31641	0.00045	0.75964	2786.253	2128.172
A_24_P28	NM_00314	signal seq SSR2, TLA	-1.31651	4.54E-06	0.75958	9779.422	7411.952
A_23_P11	NM_01782	Homo sapi COMMD4, l	-1.31667	9.03E-07	0.75949	1765.054	1336.13
A_23_P41	NM_00480	Homo sapi WDR39, Cl	-1.31669	0.00013	0.75948	4123.96	3123.373
A_24_P31	NM_14530	similar to RLOC20245	-1.31687	3.14E-07	0.75938	5629.013	4289.338
A_23_P29	NM_01404	Signal pep SPCS1	-1.31711	4.45E-06	0.75924	19445.3	14854.28
A_24_P409661	Unknown	A_24_P409661	-1.31713	1.65E-08	0.75922	10519.81	7997.595
A_32_P13	NM_17307	Homo sapi SLITRK4, C	-1.31724	8.38E-07	0.75916	804.0403	610.4902
A_32_P12	BC039468	Nucleoporin NUP50	-1.31739	0.00033	0.75908	2887.983	2197.031
A_23_P15	NM_01834	Homo sapi RSAD1, FL	-1.31747	5.62E-06	0.75903	2696.48	2053.116
A_23_P27	NM_02023	Homo sapi PPAN, SSF	-1.31752	0.00049	0.759	1872.853	1414.417
A_23_P20	NM_03277	Homo sapi ZNF503, M1	-1.31766	4.70E-06	0.75892	1666.159	1265.046
A_23_P10	AK026518	Myosin he MYOHD1	-1.31778	1.55E-10	0.75885	4051.234	3075.203
A_23_P41	NM_03066	Homo sapi RAI1, SMS,	-1.31831	4.30E-08	0.75855	3594.383	2730.805
A_23_P15	NM_01883	Homo sapi SULF2, HS	-1.31836	1.13E-08	0.75852	2186.261	1657.271
A_23_P35	NM_00461	Homo sapi TK2	-1.31857	6.50E-06	0.7584	679.1392	515.0753
A_23_P94	NM_01502	Homo sapi POP1, KIAA	-1.31857	0.00002	0.7584	1602.595	1215.619
A_23_P32593	Unknown	A_23_P32593	-1.3186	0.00012	0.75838	4175.46	3193.684
A_23_P30	NM_17812	Homo sapi CXorf40A, l	-1.31862	4.08E-07	0.75837	4196.708	3191.352
A_23_P54	NM_00731	kinesin-like KIF22, KID,	-1.31875	7.34E-09	0.7583	3637.73	2759.531
A_24_P41	NM_00293	ribonucleas RNH1, RAI,	-1.31892	3.77E-09	0.7582	5820.156	4415.668
A_24_P13	NM_15232	Homo sapi PPIL5, LRR	-1.31892	5.73E-09	0.75819	3440.999	2605.447
A_24_P19	NM_07847	Homo sapi TM2D3, BL	-1.31897	0.00002	0.75817	1110.966	842.8301
A_23_P15	NM_03275	hypothetical MGC15429	-1.31907	6.12E-08	0.75811	1277.719	968.0712
A_23_P11	NM_00461	Homo sapi TSPAN7, A	-1.31909	0.00009	0.7581	876.8018	664.3423
A_23_P11	NM_02023	Homo sapi INCENP, FI	-1.31914	1.35E-09	0.75807	1199.612	909.4148
A_23_P68	NM_00470	U3 snoRNA RNU3IP2, U	-1.31931	3.02E-10	0.75797	4190.502	3178.173
A_23_P64	NM_02474	Homo sapi ALG9, DIB1	-1.31931	2.58E-10	0.75797	1660.462	1258.634
A_32_P102252	Unknown	A_32_P102252	-1.3194	3.54E-08	0.75792	6879.91	5205.337
A_23_P37	NM_00728	Homo sapi OIP5, 5730	-1.31944	2.45E-10	0.7579	11712.71	8870.416
A_23_P13	NM_01847	TPA regula TPARL, TM	-1.31967	5.35E-07	0.75777	6083.501	4620.011
A_24_P64	NM_01540	histone de HDAC7A, H	-1.31979	0.00003	0.7577	5733.612	4377.558
A_24_P17	NM_19428	DnaJ homc DNAJA5	-1.31992	1.62E-09	0.75762	1223.96	927.3029
A_23_P38	NM_17492	hypothetical MGC31967	-1.31994	0.0001	0.75761	1064.693	804.4397
A_23_P25	NM_00209	Homo sapi GSK3B	-1.31996	3.43E-07	0.7576	1182.343	897.0474
A_32_P59811	U51990	hF THC18043	-1.32009	0.00053	0.75752	1236.761	932.6165
A_23_P54	NM_15342	Homo sapi TRADD, Hs	-1.32022	2.05E-06	0.75745	9107.133	6874.613
A_23_P40	NM_17221	Homo sapi CAMKK2, C	-1.32024	0.00021	0.75744	530.0302	401.4613
A_23_P25	NM_00675	Homo sapi UGP2, UDF	-1.32028	2.75E-06	0.75742	9061.075	6893.932
A_23_P14	NM_00370	Homo sapi SLC25A12	-1.32033	0.00012	0.75738	3214.526	2450.375
A_23_P25	NM_02473	calponin lik CLMN, caln	-1.32039	4.89E-06	0.75735	955.9523	724.5051
A_32_P17635	JC7219	nu THC19529	-1.32049	8.39E-10	0.7573	1543.45	1169.043
A_24_P281605	Unknown	A_24_P281605	-1.3205	1.03E-06	0.75729	4030.09	3042.184
A_23_P19	NM_02508	hypothetical ASRGL1, A	-1.32073	0.00066	0.75716	928.9502	703.4236
A_23_P10	NM_15228	hypothetical MGC13024	-1.32083	0.00004	0.7571	618.595	468.3945
A_24_P38	NM_00583	Rab9 effec RABEPK, p	-1.32093	3.08E-09	0.75704	6659.283	5047.566
A_23_P12	NM_02459	steroid 5 al SRD5A2L	-1.32108	7.34E-10	0.75696	5053.757	3827.37
A_23_P41	BC010736	Translocas TIMM50	-1.32141	2.34E-11	0.75677	10463.58	7917.611
A_24_P686014	Unknown	A_24_P686014	-1.32142	4.26E-09	0.75676	6425.711	4865.429

A_23_P56:	NM_00623	Homo sapi	POLR2I,RF	-1.32144	3.74E-07	0.75675	15186.55	11532.68
A_24_P14:	NM_00467	mitogen-ac	MAP3K6,A	-1.32149	0.00082	0.75672	2954.031	2234.483
A_32_P78:	NM_00457	phosphose	PSPH,PSF	-1.3215	2.29E-07	0.75672	2199.911	1667.085
A_24_P900721	Unknown	A_24_P90		-1.32153	0.00001	0.7567	3733.959	2836.534
A_24_P12:	NM_00374	dynein, cyt	DNCL1,LC	-1.32158	0.00001	0.75667	68654.26	51591.58
A_24_P255473	Unknown	A_24_P25:		-1.32215	7.94E-12	0.75634	12152.58	9191.715
A_23_P30:	NM_15242	hypothetic	FLJ39827,	-1.32237	2.06E-07	0.75622	881.419	666.4407
A_23_P50:	NM_00416	Homo sapi	SHMT1,SH	-1.32241	1.22E-10	0.75619	3094.861	2340.938
A_23_P33:	AK023854	KIAA0657	KIAA0657	-1.32254	0.00006	0.75612	457.6013	346.0008
A_23_P34:	NM_14461	hypothetic	MGC3212:	-1.32261	0.00001	0.75608	2117.068	1606.282
A_23_P13:	NM_00583	Homo sapi	POP7,RPF	-1.32272	0.00071	0.75602	54248.35	40658.95
A_23_P92:	NM_00235	Homo sapi	MAD2L1,M	-1.32282	1.03E-10	0.75596	5708.439	4313.219
A_32_P13:	NM_01774	hypothetic	ZCCHC2,F	-1.32298	0.00005	0.75587	2105.653	1580.046
A_23_P38:	BC009872	Branched c	BCKDK	-1.32308	1.51E-08	0.75581	4272.59	3228.86
A_24_P28:	NM_02431	MGC4170	GNPTAB,	-1.32309	2.11E-10	0.75581	1763.801	1333.264
A_23_P16:	NM_00372	Homo sapi	RFXANK,E	-1.32321	6.76E-06	0.75574	5788.442	4360.412
A_23_P23:	NM_19904	Homo sapi	NUDT4,DII	-1.32329	1.55E-10	0.75569	3401.083	2569.378
A_24_P36:	NM_00388	developme	DDEF2,PA	-1.32339	0.00003	0.75563	633.8799	478.8671
A_24_P25:	NM_01847	uncharacte	ACTR10,A	-1.32345	1.92E-10	0.7556	6246.908	4723.234
A_23_P13:	NM_00467	Homo sapi	PIAS2,miz,	-1.32346	8.76E-07	0.75559	963.8108	728.311
A_23_P94:	NM_01417	Homo sapi	MRPL15,R	-1.32356	8.64E-12	0.75554	13236.79	10001.71
A_24_P22746	Unknown	THC18471		-1.32363	2.13E-09	0.7555	1233.502	932.0288
A_32_P11181	P2G4_HU	THC18164		-1.32383	4.28E-08	0.75538	20130.71	15168.47
A_23_P10:	NM_00340	Homo sapi	YWHAH,Y	-1.32387	0.00003	0.75536	2419.02	1831.637
A_23_P91:	NM_00577	Homo sapi	RCL1,RNA	-1.32401	0.00088	0.75528	5133.259	3859.495
A_23_P86:	NM_01758	NICE-5 prc	UBE2Q1,U	-1.32409	0.00006	0.75524	708.228	534.3801
A_23_P14:	NM_00379	membrane	MBTPS1,S	-1.32414	6.39E-07	0.75521	4284.325	3243.887
A_24_P341546	Unknown	A_24_P34:		-1.32416	5.60E-06	0.75519	3269.815	2461.867
A_23_P43:	NM_00144	Homo sapi	FABP6,ILB	-1.32422	0.00001	0.75516	1274.655	961.5487
A_23_P11:	NM_00619	proliferatio	PA2G4,EB	-1.32435	4.97E-11	0.75509	29216.09	22046.08
A_23_P12:	NM_02275	chromoson	C9orf12,IN	-1.32436	0.0003	0.75508	2730.074	2072.903
A_23_P21:	NM_01632	RAB14, m	RAB14,FB	-1.32446	1.01E-10	0.75503	2215.114	1672.102
A_24_P647163	Unknown	THC19914		-1.32449	0.00038	0.75501	2579.688	1973.402
A_23_P46:	NM_01333	Homo sapi	NME7	-1.32449	6.73E-10	0.75501	1978.98	1494.327
A_23_P40:	NM_03238	Homo sapi	HPS3,SUT	-1.32456	5.86E-13	0.75497	933.1146	703.8073
A_24_P34:	NM_00241	antigen ide	MKI67,KIA	-1.32456	0.00023	0.75497	441.178	333.1406
A_24_P17:	NM_02308	hypothetic	C8orf33,FL	-1.32464	1.01E-09	0.75492	7630.335	5754.664
A_24_P67681	Unknown	ENST0000		-1.32467	0.00009	0.75491	10698.73	8146.799
A_23_P38:	NM_15322	hypothetic	FLJ33318	-1.3249	6.09E-08	0.75477	1174.459	887.2065
A_24_P221724	Unknown	ENST0000		-1.32502	0.00003	0.75471	9256.969	7014.445
A_24_P201323	AF176008	THC18154		-1.32503	0.00002	0.7547	1312.404	989.8228
A_24_P32:	NM_00737	bromodom	BRD3,ORF	-1.3254	2.24E-10	0.75449	1418.246	1070.082
A_23_P27:	NM_03234	hypothetic	MGC1309:	-1.32548	0.00004	0.75444	5033.388	3807.495
A_23_P18:	NM_00660	Homo sapi	PTTG2	-1.32561	0.00019	0.75437	13904.51	10545.82
A_23_P43:	NM_17752	Homo sapi	SULT1A1,I	-1.32573	2.25E-07	0.7543	3011.787	2275.969
A_23_P10:	NM_00141	Homo sapi	EIF4G2,p9	-1.32576	1.17E-07	0.75428	28489.54	21541.85
A_23_P21:	NM_00737	Homo sapi	BRD3,ORF	-1.3258	0.00015	0.75426	2595.71	1944.579
A_24_P153003	Unknown	A_24_P15:		-1.3258	6.61E-07	0.75426	6403.729	4837.478
A_32_P16:	NM_13845	hypothetic	LOC11614	-1.32604	0.00001	0.75413	2193.322	1663.447
A_23_P20:	NM_01611	Homo sapi	TRPV2,VR	-1.32612	1.63E-11	0.75408	5549.871	4185.696
A_23_P58:	NM_01910	DNA methy	DMAP1,DM	-1.32618	1.24E-09	0.75405	2684.571	2025.781
A_24_P33:	NM_00676	tyrosine 3-	YWHAE,M	-1.32644	0.00048	0.7539	1145.004	865.1757

A_23_P26	NM_01542 DKFZP564C16orf51,C	-1.32647	0.00006	0.75388	2271.892	1720.942
A_23_P13	NM_00421 Homo sapi FIBP,FGFI	-1.32651	1.88E-06	0.75386	4398.476	3325.787
A_23_P50	NM_02196 Homo sapi ABR,MDB	-1.32655	5.77E-07	0.75383	5961.2	4484.676
A_24_P56	NM_07942 Homo sapi MYL6,ESM	-1.32659	5.96E-08	0.75381	49622.61	37497.95
A_24_P30	NM_01473 KIAA0152 KIAA0152	-1.3267	2.71E-06	0.75375	994.6509	749.6895
A_23_P40	NM_15251 hypothetical FLJ30990	-1.32692	0.00022	0.75363	603.8976	455.3138
A_23_P22	NM_00426 Homo sapi SURB7,SR	-1.32711	0.00013	0.75352	2129.121	1618.922
A_24_P16	NM_01894 tubulin, alp TUBA8,TU	-1.32713	0.00018	0.7535	4293.36	3238.302
A_24_P57	NM_20348 Homo sapi DLL3,SCD	-1.32716	1.37E-08	0.75349	941.2282	709.1378
A_23_P36	NM_01560 Homo sapi C20orf22,A	-1.32729	4.88E-07	0.75342	1494.181	1127.32
A_23_P87	NM_00198 Homo sapi ESD	-1.32731	0.00018	0.7534	5926.133	4505.808
A_24_P28	NM_03291 hypothetical FAM86B1,I	-1.32739	1.06E-10	0.75336	5425.593	4085.055
A_23_P94	NM_01217 Homo sapi FBXO25,FI	-1.32752	3.39E-06	0.75328	1140.837	860.8524
A_24_P52	NM_03219 AYP1 prote AYP1,FLJ2	-1.32759	1.30E-10	0.75324	6687.821	5041.171
A_32_P12	NM_00310 sorbitol de SORD,SOI	-1.32768	4.61E-09	0.75319	6537.209	4932.111
A_23_P10	NM_00015 Homo sapi GAMT,PIG	-1.32776	6.53E-10	0.75315	2560.808	1928.24
A_23_P25	NM_00182 Homo sapi CKB,B-CK	-1.32785	6.98E-08	0.7531	1415.515	1065.267
A_23_P10	NM_00627 splicing fac SFRS6,B5	-1.32793	1.66E-06	0.75305	1613.765	1213.802
A_23_P50	NM_00014 Homo sapi FGFR3,AC	-1.32806	2.41E-08	0.75298	756.9457	569.9635
A_24_P40	NM_15248 Homo sapi SAMD11,M	-1.32806	0.00008	0.75298	3925.203	2974.768
A_24_P64	AK025909 CDNA: FL AK025909	-1.32871	1.61E-06	0.75261	655.1176	493.0868
A_23_P14	NM_19934 hypothetical SFT2D2,UI	-1.3288	8.89E-08	0.75256	1141.058	859.1631
A_24_P28	9818 Unknown A_24_P28	-1.32882	1.41E-06	0.75255	1124.406	847.6522
A_32_P18	BC067852 Similar to h LOC38896	-1.32889	3.39E-09	0.75251	3659.313	2749.644
A_23_P90	NM_02291 Homo sapi MRPL44,F	-1.32895	0.00002	0.75248	3964.653	3003.048
A_23_P16	NM_01475 Homo sapi PTDSS1,P	-1.32901	5.90E-07	0.75244	30600.63	23011.34
A_24_P18	6986 Unknown ENST0000	-1.32902	1.56E-09	0.75243	24940.34	18787.95
A_24_P33	2532 Unknown ENST0000	-1.32909	1.81E-06	0.75239	32398.03	24282.37
A_24_P75	5169 Unknown A_24_P75	-1.32919	0.00073	0.75234	9471.23	7227.434
A_23_P12	NM_00146 thyroid aut XRCC6,ML	-1.32919	1.31E-08	0.75234	121324.6	91076.92
A_32_P48	NM_03250 Homo sapi FAM11A,C	-1.32926	6.11E-10	0.7523	2973.973	2237.09
A_32_P77	783 Unknown A_32_P77	-1.32933	0.00003	0.75226	546.1409	410.8243
A_24_P65	1129 Unknown A_24_P65	-1.32936	3.90E-08	0.75224	4671.572	3510.484
A_32_P37	AK098081 Exportin, t XPOT	-1.32937	0.00008	0.75224	9764.266	7403.144
A_23_P11	NM_18280 Arsenate r ARS2	-1.32941	1.57E-07	0.75221	17070.74	12865.44
A_23_P78	NM_03129 hypothetical MGC2963	-1.32954	0.00008	0.75214	17144.64	12786.27
A_24_P25	NM_00624 protein phc PPP2R5A	-1.32961	7.10E-07	0.7521	1299.844	976.4396
A_23_P10	NM_02501 Homo sapi TUBA4,FL	-1.32968	2.89E-09	0.75206	24632.24	18494.75
A_32_P21	3210 Unknown ENST0000	-1.32978	9.42E-06	0.752	1095.979	825.3956
A_24_P20	NM_00044 regulatory I RFX5	-1.33008	1.46E-07	0.75184	2036.786	1532.582
A_23_P59	NM_01779 Homo sapi YTHDF1,C	-1.33039	4.34E-08	0.75166	13818.45	10411.3
A_23_P10	NM_00471 Homo sapi CTD1,FC	-1.33047	0.00008	0.75161	1038.765	779.7406
A_24_P92	BC017281 Homo sapi LOC15022	-1.3307	2.33E-07	0.75149	9382.354	7038.009
A_24_P23	NM_03091 hypothetical SNX27,MY	-1.3307	4.68E-11	0.75148	1738.47	1306.385
A_24_P34	9869 Unknown A_24_P34	-1.33092	3.54E-06	0.75136	25426.97	19084.71
A_32_P15	NM_02406 hypothetical PDCL3,VI	-1.33095	2.57E-07	0.75134	4915.061	3696.354
A_32_P15	5026 Unknown THC19899	-1.3313	0.00005	0.75115	1922.078	1443.682
A_23_P21	BC014315 Small nucle SNAPC5	-1.33136	4.78E-11	0.75111	5294.677	3978.717
A_23_P17	NM_02019 Homo sapi SMYD2,HS	-1.33149	7.79E-08	0.75104	7779.208	5856.231
A_23_P43	NM_17885 organic sol OSTbeta,M	-1.3316	3.07E-08	0.75098	792.329	595.0253
A_24_P23	NM_00307 SWI/SNF r SMARCD1	-1.3316	6.94E-08	0.75098	5085.32	3811.373
A_23_P73	NM_17678 Homo sapi MTMR1	-1.3317	1.44E-06	0.75092	1597.673	1203.014

A_24_P24982	Unknown	A_24_P24982	-1.33171	3.70E-08	0.75092	16643.55	12528.05
A_23_P95	NM_18157 Homo sapi	RFC5,RFC	-1.33175	3.58E-12	0.75089	9491.137	7127.613
A_23_P12	NM_00409 Homo sapi	DUSP3,VH	-1.33179	3.00E-09	0.75087	2384.854	1789.731
A_23_P21	NM_01955 Homo sapi	HOXD8,HC	-1.33186	1.11E-06	0.75083	620.3876	465.7883
A_24_P91	BC058041 Ribosomal	RPL23A	-1.33189	7.48E-06	0.75081	5724.359	4328.456
A_24_P32	NM_01561 WIPI49-like	WIPI-2	-1.33194	0.00004	0.75079	15111.5	11394.48
A_23_P12	NM_00702 Homo sapi	CYB561D2	-1.3321	2.94E-07	0.75069	3643.334	2726.09
A_23_P41	NM_15265 Homo sapi	ZNF553,FL	-1.33234	1.14E-06	0.75056	1461.765	1099.711
A_23_P15	NM_01865 bladder car	SLC35E3,E	-1.33242	4.12E-08	0.75052	950.5604	713.4425
A_23_P13	NM_01778 hypothetical	FLJ20366,I	-1.33244	9.45E-11	0.7505	2206.22	1655.487
A_23_P29	NM_14564 Homo sapi	APOL3,CC	-1.33261	4.80E-08	0.75041	991.1323	743.9702
A_23_P41	NM_02082 phosphatid	PREX1,KI	-1.33261	9.13E-07	0.75041	1487.347	1117.307
A_23_P20	NM_03066 mitogen-ac	MAP2K2,M	-1.33271	8.91E-08	0.75035	4055.526	3043.159
A_23_P25	NM_00457 Homo sapi	PSPH,PSF	-1.33278	1.03E-07	0.75031	3923.766	2940.514
A_32_P82515	Unknown	A_32_P82515	-1.33291	0.00002	0.75024	524.78	393.7123
A_23_P15	NM_00640 Homo sapi	TM9SF1,M	-1.333	1.21E-08	0.75019	2547.829	1909.228
A_23_P38	NM_02203 Homo sapi	GPRC5C,F	-1.33302	0.00007	0.75018	4197.338	3142.005
A_32_P14	BF897263 IL2-MT018	PTMA,TM	-1.33306	2.15E-11	0.75016	2269.202	1702.343
A_23_P25	NM_03194 Homo sapi	CDCA7,JP	-1.33313	7.15E-10	0.75011	7180.047	5393.199
A_23_P40	NM_03238 Hermansky	HPS3,SUT	-1.33317	0.00004	0.75009	1959.113	1478.37
A_24_P28	BX537360 Cleavage ε	CPSF5	-1.33322	0.00001	0.75006	4119.982	3099.598
A_24_P780609	Unknown	A_24_P780609	-1.33344	6.28E-06	0.74994	8982.77	6723.441
A_23_P31	NM_03143 Homo sapi	C7orf21,SE	-1.33347	1.20E-08	0.74992	7344.78	5512.98
A_23_P53	NM_00148 Homo sapi	BLOC1S1,	-1.33374	1.37E-09	0.74977	6753.812	5070.312
A_24_P647965	Unknown	A_24_P647965	-1.33387	1.90E-06	0.7497	549.681	412.1004
A_24_P49	BE081109 Activity-dep	ADNP	-1.3339	1.72E-07	0.74968	2462.699	1846.715
A_24_P39	NM_08082 Homo sapi	C20orf108,	-1.33402	7.68E-08	0.74961	1591.279	1195.557
A_24_P17	BC011136 Transformi	TGFBRAP	-1.33423	6.01E-08	0.74949	944.239	707.4723
A_24_P144303	Unknown	A_24_P144303	-1.33428	7.86E-09	0.74947	1207.829	904.7855
A_23_P98	AB020674 Mlx interac	MONDOA	-1.33447	5.64E-06	0.74936	3411.302	2559.896
A_24_P56	NM_00131 cAMP resp	CREBL2,M	-1.3345	5.54E-07	0.74935	696.3916	521.8735
A_23_P50	NM_14586 Homo sapi	EDARADD	-1.33453	0.00011	0.74933	36372.55	27068.41
A_23_P18	NM_00654 homologou	TUSC4,NP	-1.33457	2.89E-08	0.7493	2518.513	1889.506
A_24_P72	AK022015 Sel-1 supp	SEL1L	-1.33475	9.69E-08	0.7492	822.3386	616.1166
A_24_P48	NM_00007 cystathionii	CBS,HIP4	-1.33485	0.00098	0.74915	845.8574	636.5602
A_24_P771278	Unknown	A_24_P771278	-1.33496	1.62E-10	0.74909	5845.479	4375.638
A_23_P14	BX648086 Coronin, ar	CORO2A	-1.33508	0.00021	0.74902	2704.138	2032.762
A_23_P11	NM_01800 Homo sapi	TMEM45A,	-1.33518	2.98E-10	0.74896	1336.938	1001.213
A_23_P33	NM_15241 hypothetical	C8orf38,M	-1.3353	0.00006	0.74889	985.3419	739.7018
A_23_P11	NM_02300 Homo sapi	JMJD4,FL	-1.33534	3.09E-08	0.74887	2295.678	1719.73
A_23_P84	NM_00094 Homo sapi	PPIC,CYP	-1.33539	5.47E-06	0.74884	7049.251	5296.193
A_23_P11	NM_01754 NF-kappaE	NKRF	-1.33555	0.00016	0.74875	3267.387	2468.446
A_23_P21	NM_01521 KIAA0433	KIAA0433	-1.33576	9.85E-07	0.74864	3007.6	2267.19
A_23_P91	NM_00375 Homo sapi	EIF3S7,MC	-1.33578	7.28E-07	0.74863	34608.05	25814.68
A_24_P195400	Unknown	A_24_P195400	-1.33583	4.19E-06	0.7486	646.3828	483.8091
A_24_P854913	AF096870 THC18692		-1.3361	3.27E-11	0.74845	2011.429	1505.42
A_23_P64	NM_00686 putative G	RABL4,RA	-1.33622	6.93E-07	0.74838	1372.819	1027.573
A_23_P43	NM_01520 Homo sapi	BOP1,KIA	-1.33644	9.34E-08	0.74826	7901.416	5898.487
A_23_P95	NM_00468 Metastasis	MTA1	-1.33655	4.07E-12	0.74819	18177.71	13602.69
A_24_P18	NM_01850 hypothetical	PRO1855	-1.33682	2.81E-06	0.74804	16527.84	12423.51
A_23_P15	AB033092 Metastasis	MTA3	-1.33684	2.32E-10	0.74803	1542.702	1154.19
A_24_P36	XM_37258 Homo sapi	ENST0000	-1.3369	2.15E-06	0.748	1475.702	1100.294

A_24_P24685	Unknown	ENST0000	-1.33697	4.00E-10	0.74796	1769.465	1324.114	
A_24_P187448	Unknown	A_24_P187	-1.33699	8.09E-09	0.74795	1107.137	827.9962	
A_24_P15	NM_01563	Homo sapi	FGFR1OP	-1.33711	0.00002	0.74788	898.666	672.7284
A_23_P20	NM_01818	Homo sapi	ATAD3A,F	-1.33715	5.58E-07	0.74786	6642.788	4954.785
A_24_P169574	Unknown	ENST0000		-1.33715	4.30E-07	0.74786	780.536	583.5908
A_23_P28	NM_03262	Homo sapi	GTPBP3,M	-1.33723	5.33E-06	0.74781	1256.62	937.6444
A_23_P18	NM_02249	Homo sapi	MRPS25,R	-1.33737	5.27E-07	0.74774	22597.44	16843.89
A_23_P15	NM_01226	Homo sapi	C22orf3,P1	-1.33746	0.00072	0.74769	2080.437	1546.969
A_32_P157481	ALU7_HUM	THC18916		-1.33795	0.00011	0.74741	826.431	618.7682
A_23_P19	NM_00685	Homo sapi	KDELR2,E	-1.33855	2.91E-11	0.74708	12659.11	9460.832
A_23_P16	NM_00484	Homo sapi	EIF4E2,4E	-1.33861	3.96E-11	0.74704	18259.69	13658.04
A_23_P91	NM_01338	prolactin re	PREB,SEC	-1.33871	2.12E-07	0.74699	4156.41	3101.481
A_24_P409420	Unknown	A_24_P409		-1.33881	0.00042	0.74693	6178.805	4589.98
A_23_P15	NM_00632	Homo sapi	PGRMC2,I	-1.3391	3.03E-11	0.74677	3462.167	2583.99
A_23_P24	NM_01417	thymocyte	THY28,MY	-1.33924	1.66E-11	0.74669	2060.619	1538.472
A_23_P34	NM_07942	Homo sapi	MYL6,ESM	-1.33934	1.90E-12	0.74664	88678.53	66223.94
A_24_P32	AB051452	Polymerase	POLR3H	-1.33941	7.09E-06	0.7466	5548.979	4139.396
A_23_P14	NM_13839	Homo sapi	C1orf37	-1.33945	8.21E-09	0.74657	6961.683	5185.112
A_32_P10	NM_01341	adenylate	AK3L1,AK	-1.33975	0.00049	0.74641	9080.737	6823.716
A_24_P247616	Unknown	A_24_P247		-1.33987	0.00025	0.74634	2092.04	1571.638
A_23_P14	NM_01813	Homo sapi	C14orf104,	-1.3399	4.51E-10	0.74632	3947.372	2948.695
A_23_P16	NM_02465	Homo sapi	IPO4,Imp4	-1.33996	3.43E-11	0.74629	4855.687	3623.961
A_24_P19	XM_29150	Homo sapi	XM_29150	-1.33996	4.02E-06	0.74629	2966.469	2214.17
A_24_P17	NM_02043	Homo sapi	PHTF2,MC	-1.34	3.92E-06	0.74627	840.6429	626.5177
A_23_P12	NM_02100	deformed	ε DEAF1,SP	-1.34006	1.22E-07	0.74624	3469.649	2596.952
A_23_P15	NM_01528	Homo sapi	PHF15,JA	-1.34009	2.36E-08	0.74622	771.7023	575.8705
A_23_P25	NM_01804	hypothetic	AGGF1,VC	-1.34019	1.17E-06	0.74616	1528.556	1143.062
A_23_P25	NM_01210	Homo sapi	AUP1	-1.34048	0.00042	0.746	5093.76	3778.188
A_23_P92	NM_02114	Homo sapi	UNG2,UDC	-1.34058	1.03E-08	0.74594	2802.328	2086.323
A_23_P57	NM_01463	Homo sapi	PPM1F,FE	-1.34058	0.00036	0.74595	379.8727	283.4255
A_24_P28	NM_00156	interleukin	IL13RA1,N	-1.34081	7.46E-07	0.74582	10949.48	8198.32
A_23_P38	NM_01812	elaC (E. co	ELAC2,EL	-1.34094	2.95E-12	0.74575	34727.57	25905.39
A_23_P56	NM_01557	Homo sapi	FAM61A,C	-1.34096	4.37E-12	0.74573	3539.939	2639.178
A_24_P79	NM_00066	Homo sapi	TGFB1	-1.34107	0.00007	0.74568	400.4323	298.5939
A_23_P61	NM_02412	hypothetic	MGC4825	-1.34109	8.95E-13	0.74566	8186.006	6104.162
A_32_P40	NM_00644	U5 snRNP	PRPF8,PR	-1.34137	4.51E-10	0.74551	9730.887	7263.266
A_32_P16	NM_13845	hypothetic	LOC11614	-1.34139	0.00027	0.74549	1020.055	764.4574
A_24_P17	NM_00611	DNA segm	TOMM40,T	-1.34155	3.00E-06	0.7454	1524.126	1136.88
A_24_P83	XM_08836	Homo sapi	SEC11L2,S	-1.34158	1.03E-10	0.74539	4384.149	3271.065
A_24_P10	NM_14572	Homo sapi	DMN,SYN,	-1.34165	8.56E-09	0.74535	3694.186	2757.177
A_23_P72	NM_00422	Homo sapi	CRSP2,CS	-1.34175	4.08E-06	0.7453	1136.855	848.1696
A_24_P30	NM_01239	phosphotid	PITPNB,VI	-1.34184	6.01E-10	0.74525	6423.85	4779.033
A_24_P38	NM_00500	non-histon	NHP2L1,N	-1.34191	0.00005	0.74521	4429.955	3292.186
A_23_P42	NM_01594	Homo sapi	C7orf20,C	-1.34192	0.00064	0.7452	1521.317	1131.918
A_23_P47	NM_01402	Homo sapi	WDR37,FL	-1.342	0.00055	0.74516	863.4443	643.6486
A_23_P14	BC045739	Chromosor	C15orf23	-1.34213	4.17E-06	0.74508	11828.33	8774.883
A_23_P13	NM_03237	Homo sapi	ELOF1,MC	-1.34229	0.00028	0.745	1303.153	967.2114
A_24_P40	AK093744	TIA1 cytotc	TIA1	-1.34257	0.00082	0.74484	1461.474	1088.405
A_23_P41	NM_00164	N-acylamin	APEH,AP	-1.34267	1.95E-08	0.74479	1100.652	819.6382
A_23_P86	NM_01233	c-myc bind	MYCBP,AM	-1.34276	0.00017	0.74474	1618.922	1209.047
A_23_P45	NM_00719	Homo sapi	PROSC	-1.343	4.53E-06	0.7446	1518.203	1127.537
A_23_P35	NM_02236	popeye prc	POPDC3,F	-1.34302	0.00004	0.74459	3821.683	2826.427

A_24_P418216	Unknown	A_24_P418216	-1.34314	1.64E-12	0.74452	3292.178	2451.23
A_24_P14(NM_03290	phosphatid	PIG-Y	-1.34322	8.39E-06	0.74448	3309.908	2467.726
A_23_P21(NM_00287	Homo sapi	RAC2,Gx,E	-1.34323	4.80E-06	0.74447	5062.102	3782.033
A_23_P34(NM_02275	Homo sapi	SFXN1,FL	-1.34349	9.54E-09	0.74433	3111.165	2319.593
A_24_P28(NM_00563	small optic	SOLH,CAF	-1.34349	2.81E-11	0.74433	4655.357	3466.33
A_23_P10(NM_01782	Dual specif	DUSP23	-1.34352	5.90E-10	0.74431	8543.125	6378.53
A_24_P56(NM_15246	Homo sapi	C17orf32,M	-1.3439	1.43E-07	0.74411	1141.347	848.126
A_32_P73(AA441859	zw62c10.r1	AA441859	-1.34421	8.26E-07	0.74393	614.7439	457.3467
A_23_P20(NM_00290	radixin	RDX	-1.34445	6.84E-06	0.7438	3252.979	2426.374
A_24_P33(BC039449	Hypothetic	LOC33928	-1.34449	0.00016	0.74377	1644.463	1239.625
A_24_P44(NM_00036	tropomyosi	TPM1,TMS	-1.34461	0.0003	0.74371	718.2523	534.5466
A_23_P69(NM_00454	Homo sapi	NDUFB4,B	-1.34486	3.56E-13	0.74357	48206.14	35844.53
A_23_P40(NM_17370	Homo sapi	NM_17370	-1.34492	1.86E-12	0.74354	33870.48	25193.91
A_23_P40(NM_00415	Ornithine d	OAZ1	-1.34494	3.69E-11	0.74353	124726.1	92673.93
A_24_P39(NM_00319	tubulin-spe	TBCC,CFC	-1.34504	0.00016	0.74347	815.0311	607.1342
A_24_P101503	Unknown	A_24_P101503	-1.34506	4.25E-10	0.74346	2644.185	1965.943
A_23_P16(NM_02409	hypothetic	MGC2574,	-1.34507	1.48E-12	0.74345	14003.29	10412.25
A_32_P11(NM_00307	SWI/SNF r	SMARCC1	-1.34509	9.57E-10	0.74344	1041.021	773.9457
A_23_P22(AK022463	Homo sapi	FLJ12649,I	-1.34509	0.00002	0.74344	2767.408	2085.744
A_23_P14(NM_01460	Homo sapi	CYFIP1,ST	-1.34513	0.00014	0.74342	2692.669	2011.555
A_23_P36(AB046765	KIAA1545	KIAA1545	-1.34534	2.89E-06	0.74331	2760.969	2066.567
A_24_P43(NM_01406	src homolo	DBNL,SH3	-1.34543	6.09E-08	0.74326	686.5928	510.3143
A_23_P20(NM_02483	Homo sapi	C10orf119,	-1.34557	1.43E-07	0.74318	3628.166	2703.419
A_23_P31(NM_02400	DEAD/H (A	DDX3X,DE	-1.34573	0.00055	0.74309	654.9053	486.5917
A_23_P33(NM_00727	Homo sapi	FKBP9,FKI	-1.34574	0.00024	0.74308	4137.469	3097.967
A_24_P40(NM_03092	integral me	ITM2C,E25	-1.34591	0.00058	0.74299	3376.25	2523.223
A_24_P14(NM_01752	leptin rece	LEPROT,L	-1.34598	0.00066	0.74295	1327.612	993.5696
A_24_P15(NM_01457	hypothetic	MAC30	-1.34606	0.00006	0.74291	2534.595	1888.812
A_23_P20(NM_03223	Ubiquitin s	USP48	-1.34606	2.66E-08	0.74291	3985.016	2963.319
A_23_P13(NM_00210	histidyl-tR	N HARS,HR5	-1.34608	3.96E-07	0.7429	5568.905	4152.083
A_32_P75(NM_13880	hypothetic	MGC16471	-1.34623	4.21E-08	0.74282	6304.526	4691.228
A_23_P37(NM_00129	Homo sapi	CNGB1,G/	-1.34632	0.00003	0.74276	2072.102	1543.297
A_23_P10(NM_00119	Homo sapi	BCAT2,BC	-1.34639	2.79E-10	0.74273	2425.689	1802.203
A_23_P70(AK058186	Hypothetic	LOC22171	-1.34641	0.00003	0.74272	872.4578	648.4365
A_24_P143543	Unknown	ENST0000	-1.34645	0.00035	0.74269	4168.103	3118.377
A_24_P12(NM_00700	cleavage a	CPSF6,CF	-1.34656	9.68E-11	0.74263	4148.824	3084.419
A_24_P269315	ILEU_HUM	THC19557	-1.3466	2.44E-06	0.74261	2801.958	2073.009
A_23_P20(NM_00589	Homo sapi	GOLGA3,M	-1.3466	1.42E-08	0.74261	9276.82	6876.608
A_24_P925664	ALU1_HUM	THC18071	-1.34666	0.00096	0.74258	605.0295	449.4517
A_23_P20(NM_18323	Homo sapi	GSTO2,bA	-1.34684	2.29E-09	0.74248	2465.14	1834.705
A_24_P16(NM_18146	Homo sapi	MRPL55,A	-1.3469	1.41E-07	0.74245	25529.44	18856.71
A_23_P11(NM_01484	KIAA0196	KIAA0196	-1.34697	2.61E-11	0.74241	5133.167	3811.248
A_23_P37(NM_20153	Homo sapi	NDRG2,Sy	-1.34702	9.82E-08	0.74238	1980.612	1467.321
A_24_P31(NM_01500	Homo sapi	EXOSC7,p	-1.34717	7.29E-08	0.7423	7408.094	5512.416
A_32_P231302	AF059247	THC19800	-1.34721	0.00005	0.74228	1869.303	1393.101
A_23_P16(NM_00149	Homo sapi	GDI2,RAB	-1.34721	0.00003	0.74227	8162.347	6002.475
A_23_P41(AY423045	Vitamin K	ε VKORC1L	-1.34724	1.06E-06	0.74226	3346.315	2491.152
A_23_P80(NM_02197	Homo sapi	POLR2F,R	-1.34746	1.79E-09	0.74214	25092.75	18585.91
A_32_P96(NM_02474	hypothetic	SHCBP1,F	-1.34747	7.12E-09	0.74213	2354.428	1750.672
A_23_P59(NM_03282	Homo sapi	SLC35B4,)	-1.34749	2.55E-11	0.74212	1461.729	1084.785
A_23_P21(NM_03349	hexokinase	HK1,HKI,H	-1.34757	4.48E-06	0.74208	3801.561	2803.506
A_24_P67(NM_00399	CDC-like k	CLK2,hCLI	-1.34768	0.00094	0.74202	450.4464	334.1564

A_24_P76319	Unknown	ENST0000	-1.34774	0.00008	0.74198	1473.021	1088.2	
A_23_P88	NM_00287	Homo sapi	RAD51,RE	-1.34774	3.98E-10	0.74198	4195.517	3116.358
A_23_P65	NM_00374	Homo sapi	DNCL1,LC	-1.34786	1.99E-11	0.74191	92678.65	68683.11
A_24_P332595	Unknown	A_24_P33		-1.34787	1.39E-10	0.74191	5277.315	3917.957
A_23_P45	NM_01323	Homo sapi	PPP2R3B,	-1.34802	7.11E-09	0.74183	1920.854	1425.955
A_23_P66	NM_02107	Homo sapi	GCN5L2,G	-1.34833	1.52E-07	0.74166	6504.929	4809.475
A_23_P25	NM_00028	Homo sapi	PDHA1,PC	-1.34843	0.00002	0.7416	31013.68	22804.34
A_24_P24	NM_13904	Homo sapi	SMARCA3	-1.34843	0.0008	0.7416	2340.218	1752.516
A_23_P31	AK021933	Homo sapi	FLJ11871	-1.34906	0.00051	0.74125	405.9004	300.8426
A_32_P18	NM_02393	hypothetic	FUNDC2,C	-1.34919	1.20E-09	0.74119	8299.096	6166.858
A_23_P11	NM_00368	Homo sapi	AKR7A2,A	-1.3492	0.00007	0.74118	6653.078	4954.771
A_23_P96	NM_00690	Homo sapi	PRKDC,H)	-1.3494	6.37E-11	0.74107	3056.812	2266.609
A_23_P12	NM_03256	Homo sapi	EBPL,EBR	-1.34945	5.55E-09	0.74104	36958.84	27464.32
A_23_P41	NM_17354	Homo sapi	FUT11,MG	-1.34955	0.00003	0.74099	963.6021	714.3814
A_24_P22	NM_00619	proliferatio	PA2G4,EB	-1.34955	1.32E-06	0.74099	15949.67	11751.04
A_23_P37	NM_17846	Homo sapi	HMG4L	-1.34956	0.00004	0.74098	9501.801	7073.623
A_23_P10	NM_02275	sarcoma ar	C1orf149,F	-1.34976	6.66E-07	0.74087	3454.846	2567.852
A_24_P25	NM_00123	caveolin 3	CAV3,VIP2	-1.34981	4.19E-06	0.74084	2383.935	1764.813
A_32_P12	BG616663	602614865	BG616663	-1.34986	0.00019	0.74082	893.2875	662.9117
A_32_P15	NM_02406	hypothetic	PDCL3,VI	-1.35001	0.00001	0.74073	564.9052	418.4069
A_24_P21	NM_01789	hypothetic	C20orf11,T	-1.35006	0.00005	0.74071	4895.359	3615.565
A_24_P13	NM_03281	hypothetic	FLJ14627	-1.35011	3.81E-06	0.74068	4183.847	3113.253
A_24_P37	AK026976	Hypothetic	FLJ23323	-1.35016	4.34E-06	0.74065	2458.272	1823.063
A_23_P63	NM_02280	Homo sapi	CTBP2	-1.3505	3.14E-08	0.74047	29382.62	21694.14
A_32_P106553	Unknown	A_32_P10		-1.35067	1.82E-06	0.74038	6303.278	4657.307
A_23_P14	NM_00553	Homo sapi	IDH3A	-1.35071	2.06E-06	0.74035	1861.024	1376.786
A_24_P75	NM_01431	single-strar	SMUG1	-1.3508	7.02E-10	0.7403	3288.484	2430.576
A_24_P89	NM_00210	histidyl-tRN	HARS,HR	-1.35093	1.35E-06	0.74023	3778.365	2809.666
A_24_P98	NM_00735	guanine nu	GNA12,RM	-1.35096	7.24E-13	0.74021	2657.701	1967.291
A_24_P49	BC013389	LOC44015	BC013389	-1.35109	0.00002	0.74014	1365.291	1012.733
A_23_P12	NM_17574	Homo sapi	RHOC,ARI	-1.35118	5.91E-12	0.74009	8858.444	6560.364
A_23_P33	NM_02409	XTP3-trans	XTP3TPA,	-1.35119	1.52E-12	0.74009	12865.73	9517.232
A_23_P38	NM_00297	Homo sapi	ATXN2,AT	-1.3512	4.42E-10	0.74008	7673.601	5684.035
A_23_P96	NM_02523	Homo sapi	C8orf20,PF	-1.35128	1.38E-06	0.74004	847.1134	626.5906
A_24_P20	NM_01648	hypothetic	C6orf203,F	-1.35132	5.88E-07	0.74001	623.932	461.7126
A_23_P14	NM_00280	proteasom	PSMB10,L	-1.35145	1.45E-09	0.73995	14744.39	10886.04
A_23_P13	NM_01554	Homo sapi	PTCD1,KI	-1.35149	3.24E-11	0.73993	8085.683	5977.512
A_24_P23	AB004064	Homo sapi	tomoreguli	-1.3515	3.07E-11	0.73992	44089.26	32585.86
A_24_P23	NM_00645	Mad4 hom	MXD4,MAI	-1.35151	0.00048	0.73991	502.1661	371.495
A_24_P84	NM_01441	Siah-intera	CACYBP,S	-1.35156	3.34E-10	0.73988	15358.12	11346.81
A_32_P197432	Unknown	A_32_P19		-1.35176	0.00051	0.73978	2057.642	1525.153
A_23_P12	NM_00146	thyroid aut	XRCC6,ML	-1.35179	2.30E-08	0.73976	5720.001	4243.078
A_32_P129968	Unknown	A_32_P12		-1.35189	0.0002	0.7397	560.9226	414.8115
A_23_P16	NM_14471	Homo sapi	KLHL23,M	-1.35202	0.00074	0.73964	416.156	307.616
A_24_P55	NM_14580	Homo sapi	MTPN,V-1,	-1.35205	0.00032	0.73962	5845.042	4365.063
A_23_P21	NM_03313	Homo sapi	CNP,CNP1	-1.35217	4.77E-08	0.73955	19189.95	14227.31
A_23_P11	NM_18164	Homo sapi	CKLF,C32,	-1.35228	1.57E-08	0.73949	8886.752	6578.664
A_23_P37	NM_00121	Homo sapi	C1QBP,p3	-1.3523	2.73E-06	0.73948	150684.1	110727.5
A_23_P71	NM_01802	hypothetic	C8orf32,FL	-1.35235	4.10E-08	0.73945	1435.424	1060.619
A_23_P32	NM_08072	Homo sapi	SRXN1,Np	-1.35239	5.74E-08	0.73943	32114.59	23662.21
A_23_P13	NM_14572	Homo sapi	MRPL24,M	-1.35247	2.55E-09	0.73939	28130.07	20748.56
A_24_P28	NM_00538	protein-L-is	PCMT1	-1.35257	0.00003	0.73933	4045.717	3013.637

A_24_P332326	Unknown	A_24_P33:	-1.35269	9.57E-09	0.73927	64627.13	47855.22	
A_23_P28:	NM_00376	Homo sapi	VAMP8,EC	-1.35287	2.09E-06	0.73917	26111.88	19194.9
A_23_P61674	AF212224	THC19644		-1.35298	0.00002	0.73911	468.7129	346.3992
A_23_P15:	NM_02433	Homo sapi	IRX3,IRX-1	-1.35306	5.16E-08	0.73906	1870.127	1385.846
A_23_P35:	NM_17587	hypothetic	FLJ35119	-1.35317	4.32E-08	0.73901	1173.871	868.6218
A_23_P12:	NM_03098	hypothetic	ISG20L2,F	-1.35332	3.70E-09	0.73892	2251.946	1660.085
A_32_P200773	AF193342	THC18379		-1.35346	0.00005	0.73885	1173.726	865.4928
A_23_P14:	NM_00379	membrane	MBTPS1,S	-1.35357	7.54E-12	0.73879	3802.803	2810.854
A_23_P35:	NM_01812	hypothetic	C10orf6	-1.35374	1.22E-06	0.73869	765.7709	565.884
A_24_P35:	AK054681	Chromosom	C22orf13	-1.35383	0.00022	0.73864	3452.633	2569.99
A_32_P91156	AF030694	THC18524		-1.35396	7.91E-06	0.73857	968.0094	715.521
A_23_P25:	NM_13841	Homo sapi	FAM54A,D	-1.3541	5.38E-08	0.7385	1676.79	1236.415
A_24_P239017	Unknown	A_24_P23:		-1.35411	0.00033	0.73849	4676.153	3437.723
A_32_P27:	NM_01563	Chromosom	C10orf61	-1.35424	6.77E-12	0.73842	5523.32	4080.088
A_23_P29:	NM_05296	Homo sapi	RPL39L,RI	-1.35426	2.77E-07	0.73841	4085.459	3026.276
A_23_P38:	NM_17361	hypothetic	FLJ90652,I	-1.3544	0.00012	0.73833	5079.178	3730.552
A_23_P87:	NM_02206	mitochondr	MRPL17,R	-1.35464	0.00004	0.7382	11776.58	8774.427
A_23_P27:	NM_03204	Elastin mic	EMILIN2	-1.35465	0.00031	0.7382	568.667	419.7661
A_24_P57:	NM_18130	Homo sapi	MRPL52	-1.35511	0.00043	0.73795	1394.604	1034.752
A_23_P65:	NM_01327	Homo sapi	RACGAP1	-1.35515	6.82E-08	0.73793	7887.179	5838.841
A_23_P48:	NM_13855	Homo sapi	KIF23,CHC	-1.3552	9.19E-11	0.7379	7774.649	5738.984
A_23_P35:	NM_00037	Homo sapi	UROS	-1.35526	0.00011	0.73786	2137.049	1578.05
A_23_P84:	NM_02208	Homo sapi	GALNT11,I	-1.35548	0.00007	0.73775	5780.167	4276.194
A_24_P92:	NM_00638	DEAD/H (ADDX17,P7:		-1.35557	8.35E-10	0.7377	2673.435	1975.475
A_23_P15:	NM_00087	Homo sapi	ICAM2,CD	-1.35587	4.74E-12	0.73754	4734.655	3490.096
A_23_P37:	AK055367	Chromosom	C14orf126	-1.3559	3.15E-11	0.73751	2025.582	1494.429
A_23_P27:	NM_15229	Homo sapi	ATP1A3,RI	-1.35591	0.00039	0.73751	1490.222	1086.957
A_24_P122403	Unknown	THC18315		-1.3562	1.35E-11	0.73735	1322.339	975.0054
A_23_P35:	NM_00138	dihydropyri	DPYSL2,D	-1.35625	0.00004	0.73733	8705.026	6471.393
A_23_P21:	NM_00633	Homo sapi	LYPLA1,LF	-1.35625	8.78E-06	0.73733	5808.766	4306.216
A_23_P11:	NM_15333	Homo sapi	TCEAL8,M	-1.35628	4.02E-12	0.73731	8445.817	6231.649
A_24_P31:	NM_02100	nuclear rec	NR2F2,AR	-1.35634	0.00041	0.73728	5884.741	4374.028
A_24_P10:	NM_00639	nucleolar p	NOL5A,NC	-1.35637	0.00008	0.73726	53835.44	39343.93
A_32_P45:	NM_00589	isocitrate d	IDH1,IDH,I	-1.35639	3.24E-11	0.73725	27129.23	19972
A_24_P470754	Unknown	A_24_P47:		-1.35642	0.00002	0.73723	918.974	678.2146
A_23_P93:	NM_01899	DKFZp434	DKFZP434	-1.35662	0.00001	0.73713	956.923	706.0847
A_24_P11:	NM_03144	Homo sapi	C19orf12,Λ	-1.35664	4.85E-12	0.73712	2868.161	2113.363
A_23_P36:	NM_15246	Homo sapi	EME1,MM:	-1.35668	0.00046	0.73709	1781.879	1302.892
A_23_P64:	NM_00572	Homo sapi	TSFM,EF-	-1.35685	1.30E-06	0.737	26008.68	19079.14
A_23_P13:	NM_02017	Nicalin hon	NCLN	-1.35691	6.32E-08	0.73697	819.8195	604.3489
A_24_P13:	NM_02072	Homo sapi	FAM62B,C	-1.35692	3.27E-07	0.73697	1899.507	1403.167
A_24_P35:	BC022779	V-akt murir	AKT2	-1.35696	2.91E-09	0.73694	15597.92	11489.8
A_24_P101493	Unknown	A_24_P10:		-1.35707	1.29E-09	0.73688	3175.09	2339.965
A_23_P10:	NM_00710	Homo sapi	TCEB2,SIII	-1.35714	0.00034	0.73685	2641.143	1963.459
A_23_P91:	NM_01694	chromosom	C21orf6,GI	-1.35714	1.14E-09	0.73684	1113.321	820.0406
A_24_P79:	NM_00606	aldo-keto r	AKR1A1,A	-1.35725	2.16E-11	0.73678	5732.8	4227.653
A_23_P73:	NM_01478	armadillo r	ARMCX2,A	-1.35748	1.04E-07	0.73666	685.4643	504.9307
A_24_P73920	IMB1_HUM	THC18332		-1.35751	0.00023	0.73664	1165.666	861.0186
A_24_P14:	NM_01486	KIAA0475	FAM20B	-1.35752	3.80E-10	0.73664	16027.08	11780.36
A_32_P89:	NM_00314	signal seq	SSR1,TRA	-1.35753	4.31E-11	0.73663	1207.455	889.4843
A_23_P10:	NM_00186	Homo sapi	COX6B1,C	-1.35767	4.39E-08	0.73656	32439.46	23757.8
A_23_P25:	NM_13841	Homo sapi	CKLFSF7,I	-1.35771	4.65E-07	0.73654	4308.576	3172.831

A_23_P15	NM_00337	Homo sapi	VEGFB,VR	-1.35772	0.00013	0.73653	6392.547	4703.974
A_24_P30	NM_01899	Homo sapi	RIN2,RAS	-1.35792	7.47E-12	0.73642	2137.767	1573.912
A_23_P34	NM_18187	Homo sapi	DVL1,DVL	-1.35792	0.00001	0.73642	937.6813	691.0963
A_23_P13	NM_15239	Homo sapi	PQLC3,C2	-1.35792	0.00003	0.73642	653.1079	480.7833
A_24_P247596	Unknown	A_24_P24		-1.35805	8.49E-06	0.73635	4007.593	2969.563
A_23_P67	NM_00403	Homo sapi	ADCY3,AC	-1.3581	8.94E-13	0.73632	8814.783	6491.967
A_24_P28	BC066984	Mitogen-ac	MAP2K1	-1.3582	0.00031	0.73627	570.837	420.1494
A_24_P14	NM_02107	N-myristoy	NMT1,NM1	-1.35821	0.00084	0.73626	8475.995	6298.243
A_23_P13	NM_01424	Homo sapi	CELSR1,M	-1.35825	0.00051	0.73624	531.3202	391.108
A_23_P25	NM_00279	Homo sapi	PSMA6,IO	-1.35844	1.31E-09	0.73614	53563.37	39315.54
A_32_P86	BC035036	Aminoacyl	ACY1L2	-1.35866	2.98E-08	0.73602	1979.424	1459.437
A_24_P16	NM_05401	Homo sapi	FKBP1A,FI	-1.3587	1.01E-08	0.736	1327.234	976.6273
A_23_P41	NM_00132	Homo sapi	CTBP1,MC	-1.35881	7.33E-09	0.73594	9658.221	7097.472
A_24_P27	NM_00570	RNA, U tra	RNUT1,KP	-1.35889	0.00014	0.73589	690.2731	507.2312
A_24_P17	NM_19431	hypothetic	LOC33976	-1.3589	0.00001	0.73589	2128.343	1576.532
A_23_P12	NM_01405	feline leuke	FLVCR	-1.35913	1.88E-10	0.73577	2597.985	1914.306
A_23_P69	NM_01836	hypothetic	CNO,FLJ1	-1.35918	0.00003	0.73574	1958.651	1441.736
A_24_P10	NM_02459	hypothetic	FLJ13154	-1.35921	1.43E-10	0.73572	16217.35	11947.29
A_23_P38	NM_01812	Homo sapi	ELAC2,EL	-1.35938	5.65E-10	0.73563	17091.55	12597.42
A_23_P21	NM_01424	Homo sapi	RBX1,ROC	-1.35943	1.17E-08	0.7356	9804.896	7251.156
A_23_P47	NM_00556	Homo sapi	LDHA,LDH	-1.35944	3.01E-08	0.7356	51027.55	37408.46
A_24_P92	AK058024	Zinc finger	ZNF161	-1.35945	4.51E-07	0.73559	2258.54	1666.083
A_24_P884915	Unknown	A_24_P88		-1.35952	4.19E-13	0.73555	24378.29	17937.99
A_24_P11	BC014048	Kelch repe	KBTBD4	-1.35955	1.18E-07	0.73554	653.324	480.5602
A_23_P34	NM_00278	Homo sapi	PSMA2,ML	-1.35975	6.59E-11	0.73543	23971.68	17666.36
A_23_P82	NM_05818	Homo sapi	C21orf51	-1.35993	8.97E-06	0.73533	1939.787	1434.666
A_23_P82	NM_00041	Homo sapi	IFNGR1,CI	-1.35994	6.25E-13	0.73533	5831.315	4285.962
A_32_P19	NM_00711	transcriptio	TFDP1,DP	-1.36	1.11E-06	0.73529	1013.782	746.0639
A_24_P34	NM_01659	DHHC1 prc	ZDHHC3,C	-1.3601	3.85E-06	0.73524	1259.652	926.75
A_24_P87	BC039021	CDNA clon	BC039021	-1.36035	4.09E-06	0.73511	2185.614	1618.304
A_24_P79	AK056910	Hypothetic	FLJ11000	-1.36038	4.30E-10	0.73509	2211.659	1627.068
A_24_P58	XM_37805	similar to h	LOC40236	-1.36038	1.35E-06	0.73509	8274.311	6064.231
A_32_P34	NM_00141	eukaryotic	EIF4B,EIF	-1.36043	9.51E-10	0.73506	4604.567	3386.785
A_23_P32	NM_01221	interleukin	ILF3,DRBF	-1.36058	0.00005	0.73498	1214.532	893.5043
A_23_P10	NM_01785	hypothetic	FLJ20512	-1.36064	0.00002	0.73495	3162.272	2313.942
A_23_P15	NM_00600	Homo sapi	TUBA1,FL	-1.36082	0.00054	0.73485	25756.28	18674.83
A_23_P21	NM_01230	Homo sapi	AP2A2,HIF	-1.36099	0.00007	0.73476	2528.886	1867.87
A_24_P53	NM_13932	Homo sapi	ATR1N,MG	-1.36107	0.00011	0.73472	798.4711	587.4593
A_24_P26	NM_00011	dystonia 1,	TOR1A,DC	-1.36107	0.00005	0.73472	1875.62	1383.578
A_32_P20	NM_19827	Homo sapi	TMEM17,F	-1.36114	1.55E-06	0.73468	913.0269	669.9863
A_32_P20	AB032986	Homo sapi	KIAA1160	-1.36119	3.87E-11	0.73465	1248.866	917.3902
A_32_P16	BC013438	Hypothetic	LOC37529	-1.36125	8.53E-07	0.73462	2687.955	1969.348
A_23_P21	NM_00634	Homo sapi	TACC3,ER	-1.36125	1.13E-06	0.73462	1567.858	1149.113
A_23_P13	NM_02515	hypothetic	WDR71,PA	-1.36134	0.00018	0.73457	952.0931	700.1624
A_23_P13	NM_00503	Homo sapi	POLRMT,A	-1.36143	1.61E-09	0.73452	5033.333	3699.014
A_24_P42	NM_00685	KDEL (Lys	KDEL2,E	-1.36165	0.00007	0.7344	6567.436	4795.538
A_24_P91	AB062488	OK/SW-cl.	OK/SW-cl.	-1.36189	0.00019	0.73427	578.6574	424.8536
A_32_P23	NM_00556	lactate deh	LDHA,LDH	-1.36189	0.00012	0.73427	91525.34	66449.39
A_32_P145159	Unknown	THC18683		-1.36191	0.00017	0.73426	749.8003	551.2411
A_23_P21	NM_00411	Homo sapi	FGF12,FH	-1.36192	0.00026	0.73426	2816.192	2081.498
A_23_P14	NM_13880	similar to r	LOC13414	-1.36226	0.0001	0.73407	4712.08	3478.576
A_23_P10	NM_14506	pellino 3 al	PELI3,MG	-1.3624	1.77E-06	0.734	969.5905	712.2897

A_23_P21	NM_01842 Homo sapi	EPB41L4B	-1.36248	7.87E-06	0.73395	2562.062	1879.922
A_23_P11	NM_00395 Homo sapi	MPZL1,PZ	-1.36263	1.30E-07	0.73388	4240.621	3102.188
A_24_P26	NM_00660 unactive pr	PTGES3,P	-1.36272	0.0006	0.73383	4640.367	3400.061
A_32_P50	NM_00080 FK506-bin	FKBP1A,FI	-1.36338	0.00044	0.73347	2267.291	1678.915
A_23_P18	NM_00067 Homo sapi	ADH5,FDH	-1.36348	0.0001	0.73342	881.7296	648.3676
A_23_P12	NM_03228 hypothetic	TRG20,L6	-1.36362	6.99E-08	0.73334	16603.41	12129.27
A_23_P16	NM_13934 Homo sapi	BIN1,AMPI	-1.36366	2.78E-07	0.73332	1434.648	1055.042
A_23_P20	NM_01695 Homo sapi	CKLF,C32,	-1.36376	3.79E-13	0.73327	15798.95	11578.77
A_24_P39	NM_01784 tRNA seler	SECP43,P	-1.36376	0.00006	0.73327	653.669	479.7084
A_32_P23795	Unknown A_32_P23		-1.36409	6.60E-07	0.73309	5555.919	4072.889
A_23_P89	NM_01466 Homo sapi	NUP93,KI	-1.36423	2.20E-08	0.73301	3342.358	2453.061
A_23_P35	NM_00394 ribosomal	rRPS6KA4,	-1.36463	1.37E-07	0.7328	1614.56	1181.732
A_24_P12	NM_00676 tyrosine 3-	YWHAE,M	-1.36463	0.00005	0.7328	1752.326	1292.802
A_23_P39	NM_14531 Homo sapi	RASGEF1,	-1.36467	8.95E-08	0.73278	1430.738	1048.358
A_24_P40	NM_00451 integrin-lin	ILK,P59,DI	-1.36474	1.57E-06	0.73274	1454.681	1065.246
A_23_P15	NM_01431 integral inn	LEMD3,MA	-1.3648	0.00002	0.73271	878.4417	643.7231
A_24_P127425	Unknown ENST0000		-1.36486	3.37E-12	0.73267	2511.949	1839.886
A_24_P27	NM_00381 Homo sapi	CDS2	-1.36495	4.60E-06	0.73263	2024.355	1483.888
A_24_P25	NM_01653 apoptosis-r	FLJ39616	-1.36534	0.00008	0.73242	1846.905	1364.902
A_23_P12	NM_00290 Homo sapi	RDBP,RD,	-1.36573	2.99E-12	0.73221	12623.03	9236.154
A_24_P28	NM_03235 hypothetic	MGC1074	-1.36584	0.00002	0.73215	5782.201	4260.328
A_23_P83	NM_01416 Homo sapi	MRPL18,H	-1.36589	4.26E-07	0.73212	25693.97	18702.39
A_23_P20	NM_00526 Growth fac	GFER	-1.36602	1.53E-12	0.73205	2979.424	2181.997
A_23_P50	NM_01749 WD repeat	WDR1,AIP	-1.36605	2.17E-07	0.73204	4822.718	3545.736
A_23_P13	NM_00295 ribophorin	IRPN1,RBF	-1.36617	0.00014	0.73197	7185.499	5295.745
A_23_P59	NM_18172 hypothetic	FLJ12760	-1.36624	6.55E-12	0.73194	1621.356	1186.467
A_24_P93	NM_01434 mitochondr	MTCH2,HS	-1.36654	1.47E-07	0.73178	1469.573	1076.447
A_23_P28	NM_00376 syntaxin 1C	STX10,SYI	-1.36654	0.00049	0.73177	984.5071	718.3261
A_23_P25	NM_15274 Homo sapi	C7orf27,M	-1.36672	1.23E-06	0.73168	3501.953	2563.451
A_23_P16	NM_01785 Homo sapi	PARP16,C	-1.36676	0.00009	0.73166	828.3015	607.5263
A_23_P72	NM_00560 Homo sapi	PTK2,FAK	-1.36699	0.00005	0.73153	1253.503	917.2109
A_23_P53	NM_00488 Homo sapi	TP53I3,PIC	-1.36753	2.89E-08	0.73125	3044.78	2230.138
A_23_P20	AK027166 Lung canc	HLC-8	-1.36755	4.48E-11	0.73123	1778.638	1299.957
A_24_P22	NM_00627 splicing fac	SFRS7,9G	-1.36756	4.37E-08	0.73123	992.5532	725.2896
A_32_P45	BX648283 KIAA1171	KIAA1171	-1.36785	0.00011	0.73107	900.4096	660.1659
A_23_P31	NM_17850 NS5ATP13	NS5ATP13	-1.36788	1.04E-07	0.73106	637.3055	465.9197
A_23_P87	NM_00417 Homo sapi	TARBP2	-1.36791	1.73E-12	0.73104	1883.577	1377.162
A_24_P37	NM_00655 SNARE pr	YKT6	-1.36796	0.00004	0.73102	2792.657	2042.032
A_23_P67	NM_00274 Homo sapi	PKN1,DBK	-1.36802	0.00088	0.73098	519.7386	379.7362
A_23_P88	NM_01780 hypothetic	DUS2L,DU	-1.36803	0.00003	0.73098	580.9574	424.6458
A_24_P58759	Unknown A_24_P58		-1.36818	1.39E-12	0.7309	26375.12	19259.98
A_32_P204258	Unknown THC19199		-1.36825	1.75E-07	0.73086	1804.038	1321.561
A_23_P38	NM_14566 Homo sapi	APOL4,AP	-1.36832	0.00052	0.73082	461.5985	337.4688
A_23_P43	NM_00030 Homo sapi	PMM2,CD	-1.36846	1.18E-10	0.73075	2894.319	2112.149
A_32_P43	NM_00279 Homo sapi	PSMA6,IO	-1.36855	2.96E-06	0.7307	90501.05	65563.63
A_23_P15	AL832489 KIAA1970	KIAA1970	-1.36861	0.00016	0.73067	1438.806	1057.741
A_32_P14	XM_37220 similar to	RLOC38984	-1.36871	0.00099	0.73061	2378.955	1741.451
A_23_P16	NM_15342 Homo sapi	PITX2,RS,	-1.36898	2.56E-09	0.73047	1173.917	857.3015
A_23_P35	NM_03235 hypothetic	MGC14151	-1.36906	9.29E-08	0.73043	73322.59	53276.7
A_23_P21	NM_00047 adenylate	AK1	-1.36923	4.58E-14	0.73034	4566.78	3335.541
A_24_P99	NM_00227 RAN bindir	RANBP5	-1.3693	3.54E-08	0.7303	2669.972	1957.628
A_32_P84	NM_15237 hypothetic	C1orf93,M	-1.3695	0.00006	0.7302	699.0598	510.1368

A_23_P65	NM_03321	hypothetic	MGC10992	-1.36958	4.68E-11	0.73015	1084.004	791.4318
A_24_P10	XM_11662	Homo sapi	ENST0000	-1.36991	4.00E-08	0.72998	4684.979	3429.073
A_24_P12660		Unknown	THC18357	-1.36997	0.00046	0.72994	384.7533	280.6852
A_24_P161809		Keratin, ty	KRT18	-1.36997	4.89E-09	0.72994	4767.7	3486.083
A_23_P60	NM_00566	Homo sapi	VDAC3,HC	-1.37018	7.68E-10	0.72983	12999.64	9507.963
A_24_P118719		Unknown	A_24_P118	-1.37018	3.72E-07	0.72983	512.6705	374.1575
A_24_P84	NM_00290	RD RNA-bi	RDBP, RD,	-1.37072	4.36E-10	0.72954	29085.2	21262.39
A_23_P10	NM_00032	Homo sapi	ROM1,ROI	-1.37095	2.15E-12	0.72942	1433.01	1045.292
A_23_P43	NM_01786	hypothetic	C8orf52,FL	-1.37106	1.59E-07	0.72936	688.7555	502.3452
A_24_P75	AL049461	MRNA; cDI	ENST0000	-1.37108	7.63E-07	0.72935	1200.855	877.4495
A_23_P48	NM_01398	Homo sapi	DIO2,SeY	-1.37116	1.73E-10	0.72908	6256.448	4557.5
A_24_P27	NM_00599	tubulin-spe	TBCD,KIA	-1.37206	1.12E-09	0.72883	13224.8	9625.096
A_24_P195974		Unknown	A_24_P195	-1.37206	0.00054	0.72883	3909.78	2814.392
A_23_P10	AK026575	Heat shock	HSBP1	-1.37239	3.59E-09	0.72866	2140.277	1555.563
A_24_P43	NM_01799	hypothetic	C9orf40,FL	-1.37268	3.83E-08	0.7285	2160.075	1577.324
A_32_P1360		Unknown	A_32_P1360	-1.37277	0.00087	0.72845	1499.59	1091.198
A_23_P62	NM_02210	Homo sapi	MRPS14,M	-1.37287	4.62E-11	0.7284	13795.64	10069.6
A_24_P37	NM_00193	Homo sapi	DLAT,DLT	-1.37296	0.00001	0.72835	2076.568	1519.758
A_32_P60	NM_03232	Inhibitor of	ING5	-1.37322	1.11E-09	0.72822	2254.923	1641.768
A_23_P63	NM_03211	Homo sapi	MRPL43,bl	-1.3735	1.30E-12	0.72807	19092.07	13915.02
A_23_P56	NM_00131	casein kina	CSNK1G2,	-1.37354	1.66E-07	0.72805	887.0762	646.0302
A_32_P18	NM_01327	Rac GTPa	RACGAP1	-1.37355	5.20E-09	0.72804	9467.538	6912.964
A_24_P50	AK124185	Hypothetic	LOC33928	-1.37364	0.00034	0.72799	317.7738	231.3388
A_23_P13	NM_00264	Homo sapi	PIGF,MGC	-1.37374	5.61E-09	0.72794	3736.756	2724.929
A_23_P14	NM_00637	Homo sapi	VTI1B,VTI	-1.374	5.34E-07	0.7278	16593.54	12024.55
A_23_P14	NM_01750	Homo sapi	SURF2,SU	-1.37403	8.69E-15	0.72779	10613.18	7724.034
A_23_P15	NM_01840	hypothetic	HSA27219	-1.37418	4.47E-14	0.72771	33064.24	24054.78
A_23_P16	NM_00735	low density	COG2,LDL	-1.37419	5.89E-08	0.7277	5771.881	4213.853
A_24_P60	AF098968	LOC44033	AF098968	-1.37439	0.00086	0.7276	699.6918	508.5609
A_23_P25	NM_01207	Homo sapi	CCT5,CCT	-1.37443	7.33E-06	0.72757	141819.4	102292.9
A_24_P93	AF305817	Syntaxin 1	STX16	-1.37445	7.24E-13	0.72757	7206.446	5239.033
A_23_P10	NM_02021	Homo sapi	DKFZp547	-1.37447	8.98E-08	0.72755	685.2996	498.546
A_24_P19	NM_01900	Spindlin fa	SPIN2	-1.37462	8.99E-07	0.72747	1450.915	1051.932
A_23_P31	NM_02277	Homo sapi	RABL5,FL	-1.37467	4.73E-09	0.72745	7519.765	5485.115
A_23_P20	NM_01593	Homo sapi	KIAA0859,	-1.37483	9.98E-12	0.72736	10614.01	7729.482
A_24_P11	NM_01602	DORA rev	DREV1,DF	-1.37485	4.79E-08	0.72735	4188.517	3058.287
A_23_P93	NM_00670	Homo sapi	NUDT3,DI	-1.37488	2.02E-10	0.72734	1377.081	1002.129
A_23_P33	NM_17370	Homo sapi	NM_17370	-1.37491	0.00004	0.72732	25096.05	18297.91
A_24_P49	NM_18146	Homo sapi	MRPL55,A	-1.37491	4.99E-08	0.72732	10978.66	7944.785
A_24_P18	NM_03270	tubulin alp	TUBA6,MC	-1.37494	1.86E-07	0.7273	260961.8	190622.2
A_23_P12	NM_00029	Homo sapi	PGK1,PGK	-1.37495	1.47E-09	0.7273	126530.1	91823.45
A_23_P37	NM_01810	Homo sapi	CDCA8,BC	-1.37504	1.84E-11	0.72725	6854.894	4992.325
A_24_P489639		Unknown	A_24_P489	-1.37514	0.00002	0.7272	467.5875	340.0875
A_23_P74	NM_00319	Homo sapi	TCEB3,SII	-1.37524	1.09E-11	0.72715	2153.26	1566.833
A_23_P12	NM_15233	Homo sapi	TARSL2,FI	-1.37532	0.00011	0.7271	605.5865	440.1982
A_23_P67	NM_03249	Homo sapi	JAGN1,GL	-1.37546	2.48E-09	0.72703	14139.08	10249.5
A_24_P28	NM_03322	purine-rich	PURB,PUF	-1.37549	7.91E-08	0.72701	4879.419	3530.268
A_24_P29	NM_05401	Homo sapi	MGAT4B,C	-1.37553	0.00007	0.72699	2407.874	1754.135
A_23_P20	AB011134	Glycine-, g	KIAA0562	-1.37638	0.00039	0.72654	1100.031	799.7058
A_24_P306644		Unknown	A_24_P306	-1.37673	1.27E-07	0.72636	1708.792	1238.354
A_23_P35	NM_08060	Homo sapi	B3GALT6,I	-1.37674	1.09E-09	0.72635	3824.49	2779.002
A_23_P13	NM_00347	Homo sapi	STAM,STA	-1.3769	0.00002	0.72627	749.6293	544.9783

A_23_P40	AB033007	Endoplasm	KIAA1181	-1.37694	2.10E-12	0.72625	4383.699	3182.866
A_23_P12	NM_00340	Homo sapi	YWHAB,H	-1.37705	4.17E-11	0.72619	20180.78	14680.55
A_24_P35	AB011087	KIAA0515	KIAA0515	-1.37713	1.01E-08	0.72615	4287.603	3104.648
A_23_P10	NM_01884	Homo sapi	C20orf24,F	-1.37723	0.00007	0.72609	133283	95839.39
A_24_P56	NM_02110	mitochondr	MRPS12,R	-1.37726	4.52E-14	0.72608	9041.182	6566.412
A_23_P10	NM_00680	chromobox	CBX1,CBX	-1.37746	0.00005	0.72597	3815.869	2814.166
A_23_P20	NM_00317	Homo sapi	SURF1	-1.37747	6.57E-14	0.72597	3942.706	2861.518
A_24_P29	NM_01344	makorin, rir	MKRN1,R	-1.37749	5.52E-06	0.72596	706.8788	512.8484
A_24_P67	NM_00227	keratin 8	KRT8,K8,K	-1.3775	0.00002	0.72595	2668.546	1928.012
A_23_P21	NM_03143	Homo sapi	UCK1,URK	-1.3778	3.64E-09	0.7258	1234.825	896.0386
A_23_P39	NM_01597	mitochondr	MRPS7,M	-1.37796	5.85E-12	0.72571	4078.35	2956.15
A_23_P10	NM_02024	Homo sapi	CHPT1,CP	-1.37806	1.89E-10	0.72566	5225.733	3796.985
A_24_P38	NM_02067	AD023 pro	AD023,MG	-1.37815	0.00003	0.72561	412.2621	299.1818
A_23_P16	BC001097	Eukaryotic	EIF4B	-1.37828	7.72E-10	0.72554	32439.9	23586.09
A_24_P13	NM_01840	hypothetic	HSA27219	-1.37885	3.40E-12	0.72524	12031.12	8718.207
A_24_P93	6051	A40201 art	THC19143	-1.37893	2.02E-06	0.7252	565.5432	410.1323
A_23_P15	NM_00334	Homo sapi	UBE2I,P18	-1.37908	3.52E-09	0.72512	23038.37	16742.57
A_24_P28	1730	Unknown	A_24_P28	-1.37923	0.00006	0.72504	498.9249	361.7839
A_23_P37	BC028316	Tumor sup	TUSC1	-1.3796	3.86E-07	0.72485	1685.806	1221.953
A_23_P34	NM_02308	Homo sapi	CAPN10	-1.37962	0.00004	0.72484	1740.453	1264.084
A_24_P25	NM_020734	Homo sapi	ZDHHC24	-1.37998	4.71E-08	0.72465	2362.463	1709.071
A_23_P15	NM_02471	Homo sapi	ISOC2	-1.38048	6.42E-08	0.72438	4924.925	3559.583
A_23_P63	NM_03211	Homo sapi	MRPL43,bl	-1.38056	1.17E-11	0.72434	8721.779	6307.189
A_23_P34	NM_00012	Homo sapi	EPHX1,ME	-1.38087	7.58E-13	0.72418	2757.617	1998.005
A_24_P54	NM_00282	prothymosi	PTMA,TMS	-1.38089	2.60E-11	0.72417	181704.6	131351
A_24_P93	NM_00141	eukaryotic	EIF4B,EIF-	-1.38091	3.10E-10	0.72416	1162.794	842.1054
A_24_P17	AK091178	Hypothetic	LOC28540	-1.38104	0.00002	0.72409	2793.226	2013.523
A_23_P12	NM_03270	tubulin alpt	TUBA6,MC	-1.38151	0.00001	0.72385	136129.8	97390.43
A_23_P36	NM_02291	Homo sapi	VPS33A,FI	-1.38161	8.83E-11	0.72379	2072.067	1499.638
A_24_P20	4334	Unknown	A_24_P20	-1.38165	0.0001	0.72377	14560.96	10660.99
A_32_P16	NM_13845	hypothetic	LOC11614	-1.38176	1.95E-07	0.72372	1567.036	1136.748
A_24_P84	3309	Unknown	A_24_P84	-1.38186	1.01E-07	0.72366	1520.749	1102.476
A_32_P21	4005	Unknown	THC18565	-1.38208	3.85E-15	0.72355	10898.75	7885.468
A_24_P93	AK123264	CDNA FLJ	AK123264	-1.38218	0.00031	0.72349	768.5227	556.9122
A_24_P40	XM_29198	similar to h	LOC33875	-1.38253	4.27E-08	0.72331	15247.48	10986.11
A_23_P19	NM_02409	hypothetic	MGC2477	-1.38261	4.29E-10	0.72327	17000.32	12330.1
A_23_P25	NM_01604	Isochorism	CGI-111	-1.38293	8.75E-07	0.7231	6328.744	4555.376
A_23_P13	NM_01478	Homo sapi	ARHGAP1	-1.38294	0.00004	0.7231	1235.672	892.4054
A_23_P20	NM_00611	Homo sapi	VAV3	-1.38317	7.10E-07	0.72298	5069.796	3656.597
A_23_P29	NM_01649	mitochondr	MTP18,HS	-1.38333	9.25E-09	0.72289	3092.083	2227.886
A_23_P58	NM_00090	Homo sapi	NQO2,QR	-1.38334	6.52E-06	0.72289	11102.57	7953.357
A_24_P18	AB007935	Immunoglo	IGSF3	-1.38347	0.00005	0.72282	389.3535	281.4734
A_32_P17	BM701156	Transcribe	BM701156	-1.38355	3.87E-09	0.72278	857.5045	619.7605
A_23_P36	NM_15236	EID-2-like i	EID-3,FLJ	-1.38374	1.10E-08	0.72268	766.3694	553.7627
A_24_P14	NM_01792	hypothetic	PEX26,FL	-1.38388	2.01E-08	0.72261	5270.911	3818.379
A_23_P20	NM_01606	Homo sapi	TTC11,Fis	-1.38425	1.54E-06	0.72241	6076.137	4422.989
A_23_P20	NM_00572	actin relate	ARPC1B,A	-1.38443	0.00005	0.72232	9542.075	6830.564
A_23_P50	NM_01749	WD repeat	WDR1,AIP	-1.3846	6.98E-07	0.72223	3410.758	2472.98
A_24_P21	2715	Unknown	A_24_P21	-1.38462	6.25E-06	0.72222	730.787	527.8652
A_24_P26	NM_17845	Hypothetic	MGC5428	-1.3848	3.69E-06	0.72212	1483.435	1075.939
A_23_P13	NM_00404	Homo sapi	ATP6V0B,	-1.38487	1.43E-11	0.72209	12474.38	9021.242
A_23_P15	NM_00611	Homo sapi	TOMM40,T	-1.38523	2.16E-08	0.7219	4262.943	3085.12

A_23_P30	NM_03082	Homo sapi	PLA2G12A	-1.38525	7.40E-08	0.72189	2756.381	1997.887
A_24_P21	AF130080	Nuclear ub	NUCKS	-1.3854	6.28E-11	0.72181	996.5853	719.4327
A_23_P21	NM_00111	Homo sapi	ADD1,ADC	-1.38568	1.40E-14	0.72167	7772.901	5609.748
A_24_P19	NM_14460	hypothetic	FLJ31153	-1.38575	6.92E-06	0.72163	938.0109	678.6947
A_24_P94	NM_01824	hypothetic	TMEM34,F	-1.38581	3.14E-07	0.7216	1657.177	1198.453
A_23_P54	NM_01401	dexametha	DEXI,MYLI	-1.38605	1.74E-07	0.72148	7933.985	5735.094
A_24_P170103		Unknown	A_24_P17	-1.38618	1.32E-08	0.72141	116915.9	83912
A_23_P54	NM_03247	Homo sapi	MRPL38,R	-1.3862	0.00023	0.7214	16197.22	11538.6
A_24_P21	NM_00675	surfeit 6	SURF6,FL	-1.3868	7.86E-06	0.72108	714.9976	516.0429
A_23_P15	NM_02233	Homo sapi	ITGB1BP1	-1.38687	3.90E-12	0.72105	14626.31	10531.46
A_24_P13	NM_00430	Rho GDP c	ARHGDI A,	-1.387	6.40E-08	0.72098	639.1184	460.7776
A_24_P82	NM_01868	zinc finger	ZNF313,RI	-1.38701	1.14E-07	0.72098	8274.781	5987.841
A_23_P62	NM_00167	Homo sapi	ATP1B1,A	-1.38707	6.19E-11	0.72094	21256.3	15297.93
A_24_P94	NM_01485	Homo sapi	SLC35E2,I	-1.38709	9.29E-06	0.72094	2787.917	2006.484
A_32_P88	BC016882	Chromosor	C22orf9	-1.38715	4.90E-09	0.7209	9222.557	6642.768
A_23_P22	NM_00560	PTK2 prote	PTK2,FAK	-1.38716	8.70E-06	0.7209	896.7618	646.9915
A_24_P12	NM_02020	Nit protein	NIT2	-1.38718	1.26E-10	0.72089	14074.77	10166.65
A_23_P48	NM_03091	protein kin	CDADC1,A	-1.38729	0.00076	0.72083	366.2925	263.7261
A_23_P34	NM_00113	amino-terr	AES,GRG,	-1.38752	3.50E-09	0.72071	784.8007	565.6871
A_24_P288993		Unknown	ENST0000	-1.38759	4.28E-07	0.72067	7693.433	5576.34
A_23_P25	AL137321	RAB2, mer	RAB2	-1.38761	0.001	0.72066	656.8872	473.3725
A_24_P204257		Unknown	A_24_P20	-1.38767	0.00004	0.72063	3592.989	2610.094
A_24_P11	NM_02045	Homo sapi	C13orf1,Cl	-1.38785	2.44E-07	0.72054	1654.364	1195.352
A_23_P68	NM_00414	Homo sapi	DRG1,NEI	-1.38799	1.93E-08	0.72047	5409.845	3876.908
A_23_P35	NM_13834	hypothetic	KIAA2013,	-1.38844	1.69E-06	0.72023	5195.478	3734.257
A_23_P25	NM_02210	Homo sapi	C6orf79,FL	-1.38845	3.24E-06	0.72023	1992.868	1439.651
A_23_P20	NM_00029	Homo sapi	PHKB	-1.38849	6.88E-10	0.72021	7109.774	5138.415
A_23_P35	NM_01868	Homo sapi	ANLN,Scra	-1.3885	0.00007	0.7202	2911.104	2112.997
A_23_P47	NM_01818	Homo sapi	DHX32,DD	-1.38876	0.00037	0.72007	1263.452	908.54
A_24_P22	AF111845	3-phosphoi	PDPK1	-1.38888	1.30E-06	0.72001	1718.982	1241.094
A_23_P47	NM_00272	Homo sapi	HHEX,HEX	-1.38906	1.78E-10	0.71991	1183.706	852.3731
A_32_P13	NM_00566	Homo sapi	C5orf18,DF	-1.3891	1.32E-11	0.71989	10944.61	7894.363
A_23_P17	NM_02393	hypothetic	FUNDC2,C	-1.38928	2.77E-08	0.7198	1888.288	1355.805
A_24_P20	XM_21036	similar to	RLOC28428	-1.38929	4.76E-08	0.71979	1435.435	1034.776
A_32_P14	BC046178	Chromosor	C6orf173	-1.38956	9.73E-09	0.71965	8508.984	6149.211
A_24_P29	NM_00680	chromobox	CBX1,CBX	-1.38961	2.33E-08	0.71962	4173.943	3018.378
A_24_P46	NM_00487	etoposide-i	EI24,PIG8,	-1.38977	1.34E-07	0.71954	4248.784	3057.452
A_23_P13	NM_02112	Homo sapi	MPST,MS1	-1.38997	3.68E-06	0.71944	11066.13	7912.018
A_23_P97	NM_01222	Homo sapi	MSRB2,CE	-1.39112	1.71E-09	0.71885	7632.913	5503.572
A_32_P86	NM_00472	Ac-like tran	ZBED1,AL	-1.3912	0.00012	0.7188	3512.497	2523.592
A_23_P47	NM_02047	Homo sapi	YIF1A,54T	-1.39152	1.28E-10	0.71864	12326.85	8871.025
A_24_P19	NM_19948	Homo sapi	C20orf24,F	-1.3916	3.51E-15	0.7186	65461.27	47052.11
A_23_P38	NM_01563	Homo sapi	KIAA1279,	-1.39172	5.22E-10	0.71854	1795.258	1287.725
A_23_P30	NM_15234	hypothetic	FLJ30656	-1.39179	4.58E-09	0.7185	2038.54	1461.599
A_24_P81	BC044637	Hypothetic	LOC34892	-1.39179	4.03E-12	0.7185	5802.066	4174.571
A_23_P25	NM_01895	Homo sapi	HOXA10,P	-1.39203	0.00062	0.71837	297.946	214.017
A_24_P21	NM_03230	hypothetic	MGC10911	-1.39208	2.32E-11	0.71835	8124.653	5846.698
A_32_P13	NM_01741	ubiquitin sp	USP18,ISC	-1.39217	3.69E-15	0.71831	2756.266	1979.873
A_24_P713729		ALU8_HUM	THC19913	-1.3924	2.00E-06	0.71818	2571.063	1858.758
A_23_P26	AB037809	Zinc finger	ZNF319	-1.39262	0.00021	0.71807	846.6002	608.2903
A_23_P32	NM_00522	Homo sapi	EFNA4,EF	-1.39267	6.75E-06	0.71805	1036.701	745.8065
A_23_P65	NM_01641	Homo sapi	C14orf87,C	-1.39281	4.13E-09	0.71797	23943.32	17150.81

A_23_P59	NM_02117	U6 snRNA	LSM2,G7b	-1.39288	4.44E-12	0.71793	1393.813	1000.693
A_32_P54	BX537645	Hypothetic	LOC40139	-1.39289	7.53E-13	0.71793	10547.1	7567.462
A_24_P37	NM_02522	hypothetic	PRO2730,I	-1.39335	2.75E-06	0.7177	4200.131	3031.711
A_24_P41	NM_01840	Homo sapi	LAPTM4B,	-1.39344	5.10E-09	0.71765	2595.28	1860.152
A_32_P20	NM_00144	fatty acid b	FABP5,EF	-1.39347	1.99E-15	0.71763	11991.53	8607.587
A_23_P14	NM_00167	Homo sapi	ATP1B1,A	-1.39354	0.0002	0.7176	8087.817	5876.485
A_23_P14	NM_01437	Homo sapi	REPIN1,AF	-1.39365	6.41E-08	0.71754	15313.44	11012.26
A_23_P91	NM_00375	eukaryotic	EIF3S7,MC	-1.39402	6.39E-16	0.71735	70159.59	50326.43
A_23_P21	NM_01810	Homo sapi	EFHC1,EJ	-1.39405	0.00009	0.71733	679.1659	486.399
A_24_P934755		IFE3_HUM	THC20018	-1.39407	2.87E-12	0.71733	1363.453	977.7148
A_24_P358857		Unknown	A_24_P35	-1.39409	1.42E-07	0.71731	1681.11	1203.32
A_23_P14	NM_00189	Homo sapi	CSNK2A2,	-1.39439	9.32E-12	0.71716	1128.507	809.4611
A_24_P846755		Unknown	A_24_P84	-1.3944	0.00001	0.71715	665.0411	476.6223
A_32_P78285		Unknown	A_32_P78	-1.39471	0.00007	0.71699	7548.651	5447.423
A_23_P14	NM_01826	membrane	MTCBP-1,	-1.39473	9.02E-13	0.71699	4853.348	3477.459
A_23_P14	NM_00118	Homo sapi	BAK1,BAK	-1.39507	0.00003	0.71681	1744.986	1245.357
A_24_P247074		Unknown	A_24_P24	-1.39526	0.00022	0.71671	3720.549	2641.257
A_23_P48	NM_19916	Homo sapi	ADPRHL1,	-1.39529	1.18E-08	0.71669	2038.895	1461.429
A_32_P224449		Unknown	THC18364	-1.3953	7.85E-08	0.71669	660.8775	473.728
A_23_P50	NM_00039	Diaphorase	DIA1	-1.39531	1.08E-06	0.71669	53255.91	38395.41
A_23_P81	NM_01661	Homo sapi	PLAC8,C1	-1.39538	0.00066	0.71665	4137.549	3045.76
A_24_P11	NM_00200	farnesyl di	FDPS,FPS	-1.39542	1.65E-12	0.71663	33248.58	23852.63
A_23_P94	NM_00229	Homo sapi	LAMB1	-1.39545	2.61E-10	0.71661	12379.74	8895.854
A_24_P911648		Unknown	A_24_P91	-1.39557	1.17E-07	0.71656	3365.716	2421.824
A_23_P16	NM_00517	ATP synth	ATP5G1,A	-1.39582	4.56E-14	0.71642	26890.78	19247.93
A_24_P19	NM_03226	nerve injur	ZNRF1,NIF	-1.3959	0.00051	0.71638	440.6991	315.3276
A_24_P60	NM_00029	phosphogly	PGK1,PGK	-1.39598	0.00008	0.71634	15564.95	11039.2
A_24_P47	XM_20909	Homo sapi	LOC28426	-1.39598	1.15E-11	0.71634	1503.6	1077.637
A_23_P61	NM_01684	neuronal S	SHC3,NSF	-1.39633	2.15E-11	0.71616	2489.22	1780.836
A_24_P25	NM_20332	Homo sapi	SLC23A2,I	-1.39639	0.00001	0.71613	1054.654	755.5713
A_23_P16	AB002363	Homo sapi	KIAA0365	-1.39659	0.00013	0.71603	464.7773	332.5837
A_23_P25	NM_00216	Homo sapi	ID1,ID	-1.39674	4.34E-07	0.71595	77989.53	55503.61
A_32_P21	BU537444	Transcriber	BU537444	-1.3968	2.32E-07	0.71592	632.7943	453.0877
A_32_P48	BC004542	Plexin B2	PLXNB2	-1.39688	2.95E-09	0.71588	1714.147	1226.739
A_23_P10	NM_01742	Homo sapi	SPA17,SP	-1.39705	3.63E-07	0.71579	1644.461	1178.999
A_24_P15	NM_03262	hypothetic	SSBP4,MC	-1.39726	0.00097	0.71569	800.2402	572.249
A_23_P91	NM_00639	Homo sapi	NOL5A,NC	-1.39729	1.48E-12	0.71567	14455.08	10332.61
A_23_P25	NM_01762	hypothetic	FLJ20013	-1.39742	1.48E-08	0.71561	1767.841	1263.917
A_23_P15	NM_08064	Homo sapi	APEX1,AP	-1.39768	6.07E-09	0.71547	8208.123	5887.189
A_24_P42	NM_01233	acyl-Coenz	ACATE2,C	-1.39787	2.92E-12	0.71537	2959.017	2116.281
A_23_P42	NM_05304	Homo sapi	COMMD7,I	-1.39795	7.57E-07	0.71533	1550.311	1111.445
A_23_P13	NM_01767	Homo sapi	C20orf42,U	-1.39814	1.34E-10	0.71524	5019.495	3591.544
A_24_P22	NM_00377	nuclear loc	MRPL40,U	-1.39817	0.00013	0.71522	3980.711	2874.124
A_23_P28	AK022713	Homo sapi	C20orf27,F	-1.39837	5.23E-07	0.71512	807.958	577.5529
A_23_P53	NM_02064	Homo sapi	C11orf17,E	-1.39847	9.22E-09	0.71507	2992.302	2134.389
A_23_P24	NM_00547	potassium	KCNE3,HC	-1.39854	3.51E-07	0.71503	979.2521	701.7225
A_24_P74	NM_19432	hypothetic	MGC5201(C	-1.39858	7.68E-13	0.71501	9548.986	6817.488
A_23_P11	NM_03231	Homo sapi	NICN1,MG	-1.39882	4.69E-06	0.71489	1689.745	1212.546
A_23_P38	NM_02109	solute carri	SLC5A6,SI	-1.39911	0.00006	0.71474	920.7227	657.0997
A_23_P14	NM_00302	Homo sapi	SH3BGRL,	-1.39916	8.28E-13	0.71471	1814.105	1296.672
A_23_P98	NM_01245	Homo sapi	TIMM8B,D	-1.39925	1.28E-08	0.71467	31458.53	22359.18
A_24_P76	NM_00144	fatty acid b	FABP5,EF	-1.39936	7.51E-12	0.71461	21186.61	15176.02

A_23_P11	(NM_01534 Homo sapi	PPWD1,KI	-1.39939	8.58E-15	0.7146	3898.659	2784.73
A_23_P40	NM_02121 Homo sapi	PCTP,STA	-1.39942	1.58E-06	0.71458	3122.528	2245.207
A_23_P15	NM_00600 Homo sapi	TUBA1,FL	-1.39974	2.50E-10	0.71442	138850.3	98826.09
A_24_P213325	Unknown	A_24_P21	-1.39987	3.57E-07	0.71435	5513.147	3949.005
A_24_P48	NM_00300 secreted ar	SECTM1,K	-1.40012	0.00012	0.71423	1581.323	1132.724
A_23_P41	AF289571 Hypothetic	LOC28337	-1.40044	1.14E-11	0.71406	1514.371	1080.772
A_23_P66	NM_00261 3-phosphoi	PDPK1	-1.40069	1.19E-14	0.71393	1990.25	1420.769
A_23_P21	NM_00323 Homo sapi	TFRC,TFR	-1.40083	6.07E-12	0.71386	25903.45	18449.63
A_24_P144556	Unknown	A_24_P14	-1.40089	7.46E-13	0.71383	2179.688	1556.404
A_24_P410686	A42286 EF THC	18334	-1.40091	2.27E-06	0.71382	11521.44	8190.285
A_23_P38	NM_00604 Homo sapi	SNAPC5,S	-1.40112	4.49E-10	0.71371	2136.709	1526.453
A_23_P36	NM_00681 coated ves	TMED2,P2	-1.40116	1.24E-11	0.7137	3372.086	2402.91
A_23_P39	NM_05304 Homo sapi	HTRA3,Prs	-1.40123	9.07E-12	0.71366	25987.01	18511.76
A_24_P19	NM_01770 hypothetic	FLJ20186	-1.40127	2.73E-15	0.71364	3522.365	2514.257
A_23_P14	(NM_01214 solute carri	SLC25A10	-1.40129	3.23E-09	0.71363	1076.007	767.3753
A_24_P17	AL834528 Calmodulin	CAMSAP1	-1.40137	6.49E-06	0.71359	588.4841	419.8989
A_23_P11	NM_01601 Homo sapi	LCMT1,LC	-1.40154	4.81E-11	0.7135	4402.27	3147.114
A_23_P25	NM_01473 Homo sapi	KIAA0152	-1.40155	1.04E-11	0.71349	11592.77	8253.008
A_24_P27	NM_15282 Homo sapi	SNX3,SDP	-1.4016	1.32E-07	0.71347	6345.413	4539.713
A_24_P16	NM_00581 protein kin	PRKD3,EP	-1.40176	0.00031	0.71339	1289.498	925.4207
A_23_P25	NM_03275 Homo sapi	PHF5A,INI	-1.40207	0.00026	0.71323	1287.499	919.3664
A_23_P52	NM_00699 Homo sapi	NPM3	-1.40213	2.70E-07	0.7132	6652.814	4741.345
A_23_P78	NM_00238 Homo sapi	MCM3,HC	-1.40221	1.75E-10	0.71316	25157.04	17901.04
A_23_P14	(NM_01826 membrane	MTCBP-1,	-1.40235	3.79E-09	0.71309	3021.615	2159.038
A_24_P298744	Unknown	ENST0000	-1.40259	2.58E-07	0.71297	470.0876	335.1614
A_24_P238716	Unknown	A_24_P23	-1.40286	3.18E-12	0.71283	4272.476	3044.23
A_32_P49	AK023856 Hypothetic	LOC33980	-1.40298	5.49E-10	0.71277	1059.571	755.6083
A_32_P91	NM_00334 ubiquitin-cc	UBE2L3,E	-1.40311	4.41E-10	0.7127	4430.212	3166.143
A_23_P20	NM_18187 Homo sapi	DVL1,DVL	-1.40312	3.54E-15	0.7127	28541.42	20347.86
A_23_P76	NM_02083 KIAA1443	KIAA1443	-1.40322	1.03E-06	0.71265	653.2277	465.5312
A_23_P28	NM_17585 Homo sapi	DNMT3B,I	-1.40331	1.69E-10	0.7126	1172.847	835.7238
A_24_P97	(NM_03286 hypothetic	PRPF38A,I	-1.40354	0.00001	0.71248	2969.652	2116.972
A_23_P41	(NM_00207 Homo sapi	GNAZ	-1.40366	1.98E-08	0.71242	1245.335	887.1625
A_24_P12	NM_00707 adaptor-rel	AP4S1,AP	-1.40374	5.92E-06	0.71238	515.5793	367.2219
A_23_P15	NM_08064 Homo sapi	APEX1,AP	-1.40375	7.14E-12	0.71238	2050.558	1460.089
A_24_P32	(AK023593 Peroxisom	PEX3	-1.40392	1.67E-11	0.71229	2713.62	1935.552
A_23_P42	(BC031838 CGI-96 pro	CGI-96	-1.40434	8.40E-11	0.71208	1297.855	924.2419
A_23_P30	NM_15232 Homo sapi	ANKRD9,M	-1.4045	1.64E-06	0.712	2551.487	1811.365
A_23_P35	NM_00486 Homo sapi	LITAF,PIG	-1.40453	4.28E-08	0.71198	5652.49	4038.565
A_23_P12	(NM_01633 Selenoprot	SEPX1	-1.40513	3.17E-07	0.71168	7668.359	5430.67
A_32_P44	(NM_00556 lactate deh	LDHA,LDH	-1.4052	2.32E-07	0.71164	9976.117	7050.348
A_32_P13	(BC035377 Hypothetic	BC035377	-1.40543	0.00076	0.71153	564.4742	401.3248
A_23_P21	(D80008 DNA replic	KIAA0186	-1.40549	0.00003	0.71149	518.9253	369.2211
A_23_P11	(NM_01742 Homo sapi	COQ3,bA9	-1.40554	1.14E-11	0.71147	1778.511	1264.591
A_23_P90	(NM_01823 dudulin 2	STEAP3,2,	-1.40602	5.13E-09	0.71123	2042.495	1459.081
A_24_P170283	Unknown	ENST0000	-1.40613	1.16E-07	0.71117	1555.434	1105.758
A_23_P53	(NM_20161 IKK interac	IKIP,FLJ31	-1.40632	3.00E-08	0.71107	2204.679	1573.915
A_23_P53	(NM_03362 Homo sapi	FBXO21,FI	-1.40666	2.17E-14	0.7109	9115.384	6484.165
A_23_P11	(NM_03235 Homo sapi	MRPL45,M	-1.40675	2.32E-12	0.71086	19803.04	14045.05
A_24_P22	(NM_01232 U6 snRNA	LSM4,YER	-1.40688	8.12E-07	0.71079	3359.069	2396.236
A_23_P59	(NM_00144 fatty acid b	FABP5,EF	-1.40704	0.00003	0.71071	41980.52	29462.22
A_23_P54	(NM_03247 Homo sapi	MRPL38,R	-1.40705	0.00004	0.71071	1521.89	1079.5

A_23_P11	NM_03120	hypothetic	FAHD1, YIS	-1.40726	1.11E-13	0.7106	2194.473	1558.688
A_23_P42	NM_13857	Bcl-XL-bin	MGC5352,	-1.4073	2.72E-07	0.71058	537.2416	381.7606
A_23_P12	NM_00665	Homo sapi	TUBGCP2,	-1.40752	1.14E-09	0.71047	1339.204	951.7016
A_23_P11	NM_00579	Homo sapi	NUTF2,NT	-1.40759	6.11E-07	0.71043	5643.891	4016.249
A_23_P21	NM_03217	Src-associ	UTP15,FL	-1.4078	6.60E-08	0.71033	1584.481	1127.391
A_23_P14	NM_00015	Homo sapi	GALK1,GK	-1.4079	1.99E-11	0.71028	1678.673	1192.901
A_23_P81	NM_17850	Homo sapi	C6orf1,LB	-1.4081	0.00001	0.71018	6373.023	4515.119
A_23_P63	NM_01418	Homo sapi	SSU72,HS	-1.4081	5.51E-10	0.71018	11930.36	8448.636
A_23_P84	NM_01596	Homo sapi	THAP4,CG	-1.40813	1.87E-14	0.71016	6699.942	4755.558
A_32_P92	NM_03238	hypothetic	COG8,DOI	-1.40823	2.29E-16	0.71011	5807.218	4123.338
A_23_P15	NM_20701	Homo sapi	AP3M1,MC	-1.40828	8.22E-09	0.71009	1253.266	889.255
A_24_P497843		Unknown	THC18702	-1.40851	1.26E-12	0.70997	1497.417	1063.301
A_23_P15	NM_01594	Homo sapi	ATP6V1H,	-1.40857	2.29E-06	0.70994	3547.922	2532.882
A_24_P19	NM_01457	hypothetic	MAC30	-1.40857	2.95E-10	0.70994	3431.94	2431.432
A_23_P87	NM_01418	Homo sapi	ORMDL2,M	-1.40866	7.72E-15	0.70989	16642.03	11822.54
A_24_P12	NM_02077	Homo sapi	TBC1D14,I	-1.40871	0.0008	0.70987	874.8351	621.2166
A_24_P63	AK056973	CDNA FLJ	AK056973	-1.40872	0.00003	0.70987	340.3276	241.5931
A_23_P25	NM_01596	Homo sapi	MRPS17,R	-1.40876	3.63E-06	0.70984	5566.283	3968.298
A_23_P28	BC017237	Syntaxin 1	(STX10	-1.409	5.95E-10	0.70972	6628.348	4715.172
A_32_P4318		Unknown	A_32_P43	-1.40924	0.00033	0.7096	386.5319	274.3198
A_23_P105753		PWP1_HU	THC18059	-1.40951	1.81E-09	0.70946	2416.392	1719.844
A_32_P14	CA314936	Transcribe	CA314936	-1.40954	1.35E-08	0.70945	1424.632	1012.57
A_24_P41	NM_00532	L-3-hydrox	HADHSC,S	-1.40973	2.47E-11	0.70936	2560.919	1814.526
A_32_P14	NM_05402	Homo sapi	CNOT7,CA	-1.40979	6.81E-09	0.70932	1981.691	1402.642
A_23_P10	NM_02522	hypothetic	PRO2730,I	-1.40986	0.00007	0.70929	2406.818	1697.152
A_23_P97	NM_05283	secretory c	SCAMP3,C	-1.41001	0.00005	0.70921	3472.922	2452.37
A_32_P39	AL390079	Hypothetic	LOC58489	-1.4101	2.64E-10	0.70917	5695.455	4034.832
A_23_P33	AK055401	Dedicator c	DOCK7	-1.41031	0.00028	0.70907	528.2299	374.2669
A_23_P46	NM_00126	chromoson	RCC1,CHC	-1.41056	0.00011	0.70894	1624.203	1139.236
A_24_P11	NM_02193	likely ortho	RIC8A,RIC	-1.41058	1.71E-12	0.70893	1093.849	775.5573
A_23_P10	NM_02494	NEFA-inter	NIP30,CD	-1.41059	4.05E-09	0.70892	1399.298	993.6284
A_23_P16	NM_01595	Homo sapi	MRPL4,L4	-1.41085	8.39E-08	0.70879	2735.842	1933.24
A_24_P83	NM_17069	Homo sapi	dJ222E13.	-1.41105	2.00E-12	0.70869	1895.922	1342.699
A_23_P74	NM_03192	AAA-ATPa	ATAD3B,T	-1.41132	0.00014	0.70856	863.6067	611.1681
A_24_P786257		Unknown	A_24_P78	-1.4114	1.44E-06	0.70852	1160.472	819.9816
A_23_P21	NM_13083	Homo sapi	OPA1,NPC	-1.41175	0.00004	0.70834	2329.934	1645.348
A_32_P123629		ALU5_HU	THC20038	-1.41176	1.79E-15	0.70834	3806.679	2695.831
A_23_P42	NM_13846	hypothetic	LOC11623	-1.41193	8.39E-12	0.70825	8100.791	5738.085
A_32_P211752		Unknown	THC18206	-1.41199	3.30E-09	0.70822	1386.588	981.4623
A_23_P53	M10036	Human trio	TPI1	-1.41199	1.63E-07	0.70822	5769.368	4104.029
A_23_P10	NM_01900	hypothetic	FLJ20232,I	-1.41215	1.66E-09	0.70814	1465.64	1037.697
A_23_P41	NM_01403	DKFZP586	DKFZP586	-1.4124	1.30E-10	0.70802	4859.095	3453.336
A_23_P36	NM_01816	Homo sapi	C12orf11,F	-1.41244	6.54E-06	0.708	3169.114	2256.963
A_23_P36	NM_01222	myosin IB	MYO1B,my	-1.41254	1.55E-07	0.70795	1364.463	969.5992
A_24_P29	NM_02432	hypothetic	ZNF576,M	-1.41264	7.26E-08	0.70789	859.8456	609.1533
A_23_P10	BC008845	Homo sapi	MGC1438	-1.41268	1.43E-12	0.70788	6731.065	4767.296
A_24_P38	NM_00631	ring finger	PCGF3,RN	-1.41274	0.00002	0.70785	3023.787	2142.851
A_24_P56	AK024346	Peroxisom	PPARGC1	-1.41291	2.13E-10	0.70776	980.0465	693.2922
A_32_P89	AF110908	TNF recept	TRAF3	-1.41308	1.53E-10	0.70767	2472.343	1750.245
A_24_P187355		Unknown	A_24_P18	-1.41312	1.49E-12	0.70765	4155.93	2944.36
A_23_P30	NM_03056	Homo sapi	PRR7,MGC	-1.41313	1.92E-08	0.70765	4405.735	3108.31
A_23_P20	NM_01784	hypothetic	FLJ20489	-1.4133	5.05E-11	0.70757	9428.941	6668.738

A_24_P67	NM_00144 fatty acid b FABP5,EF	-1.41355	1.75E-07	0.70744	20499.68	14401.67
A_23_P16	NM_00358 Homo sapi DHX16,DB	-1.41359	4.71E-12	0.70742	1471.046	1040.357
A_32_P86	AB032986 Homo sapi KIAA1160	-1.41381	2.36E-06	0.70731	817.2956	578.7464
A_23_P41	NM_01804 Homo sapi AGGF1,VC	-1.41381	1.30E-10	0.70731	824.2687	583.0725
A_24_P12	NM_00712 valosin-cor VCP,p97,T	-1.41436	6.30E-06	0.70704	1498.7	1062.709
A_32_P31	NM_02520 FAD-synth FLAD1,FAI	-1.41474	1.08E-11	0.70684	1107.255	782.9132
A_23_P20	NM_03283 Homo sapi ALG10,KC	-1.41475	0.00029	0.70684	477.9298	337.3742
A_23_P56	BC051789 Hypothetic LOC12929	-1.4149	7.09E-07	0.70676	4826.404	3412.667
A_24_P84	NM_03256 empomamil t EBPL,EBR	-1.41496	2.89E-13	0.70673	6155.113	4353.301
A_24_P66908	Unknown A_24_P66	-1.41499	0.00003	0.70672	779.0968	550.4976
A_23_P17	(NM_17347 Homo sapi NTAN1,DK	-1.4151	3.77E-06	0.70666	896.3051	634.1965
A_23_P36	NM_17355 Sterile alph SAMD6	-1.41537	5.47E-06	0.70653	4053.353	2873.014
A_24_P70	BC004542 Plexin B2 PLXNB2	-1.41538	2.01E-08	0.70652	2105.51	1482.327
A_23_P50	NM_13364 Homo sapi SURF5,ME	-1.41548	9.43E-10	0.70647	9260.979	6563.199
A_24_P15973	Unknown A_24_P15	-1.41549	1.08E-09	0.70647	1355.528	956.6155
A_24_P83	BC041772 Hypothetic LOC12497	-1.41557	8.80E-14	0.70643	1983.743	1400.556
A_24_P246591	NADP+-sp ENST0000	-1.41572	4.29E-10	0.70635	7994.168	5634.32
A_24_P12	(NM_00280 proteasom PSMD1,S1	-1.41594	0.00001	0.70624	3235.459	2305.412
A_23_P15	(NM_00436 Homo sapi CDH7,CD	-1.41594	0.00002	0.70624	1793.92	1266.105
A_24_P48	(NM_19847 Homo sapi FLJ46111	-1.41626	4.46E-06	0.70608	956.0162	675.4618
A_24_P324506	Unknown A_24_P32	-1.41627	1.29E-09	0.70608	7872.365	5579.63
A_32_P36046	ALU8_HUM THC18442	-1.41644	0.00046	0.70599	572.8087	404.144
A_24_P263803	Unknown ENST0000	-1.41648	7.17E-10	0.70598	2477.962	1752.608
A_23_P11	(NM_17793 hypoxia-inc PH-4	-1.41674	2.70E-06	0.70584	1586.113	1114.857
A_32_P16	(NM_00337 voltage-dej VDAC1,PC	-1.4168	3.68E-15	0.70582	2788.811	1969.113
A_23_P16	(NM_00152 Homo sapi GTF3C2,K	-1.41682	2.26E-09	0.70581	5640.854	3978.43
A_32_P22	BC026043 Homo sapi BC026043	-1.41686	0.00042	0.70578	476.5943	336.6796
A_24_P41	(NM_15303 hypothetic FLJ32065,l	-1.41702	2.38E-09	0.7057	7850.576	5562.245
A_24_P272886	Unknown ENST0000	-1.41714	2.06E-10	0.70565	3715.628	2618.636
A_24_P59099	Unknown ENST0000	-1.41724	0.00006	0.7056	2170.087	1519.276
A_24_P49	(NM_18165 hypothetic LOC28401	-1.41731	0.00095	0.70556	656.3239	463.1761
A_23_P25	(NM_01486 chromoson CNAP1,KI/	-1.41744	2.97E-06	0.7055	3241.581	2274.666
A_24_P40	(NM_00437 COX11 (ye COX11,CC	-1.41797	6.70E-15	0.70523	2726.011	1921.849
A_23_P92	(NM_00676 interferon-r IFRD2,SM	-1.41807	1.98E-06	0.70518	11499.34	8085.206
A_32_P38	BC015133 CDNA clon BC015133	-1.41862	0.00015	0.70491	614.3984	432.8257
A_23_P10	(NM_02109 Homo sapi SLC5A6,SI	-1.41884	1.47E-09	0.7048	1326.806	935.4142
A_24_P32	(NM_00650 melanoma MCAM,CD	-1.41931	8.69E-08	0.70457	2188.858	1545.863
A_32_P52	(NM_01826 membrane MTCBP-1,;	-1.41938	3.36E-09	0.70453	3224.601	2275.193
A_23_P21	(AK025603 HSPC159 HSPC159	-1.41939	2.93E-09	0.70453	2641.785	1861.608
A_23_P17	(NM_01827 Homo sapi C20orf20,M	-1.41942	8.52E-09	0.70451	8103.82	5682.506
A_24_P12	AK021904 Hypothetic RPL7L1	-1.41978	3.32E-11	0.70433	1916.067	1347.221
A_23_P74	(NM_01871 RCC1-like RCC2,TD-(-1.41984	4.98E-06	0.70431	13520.02	9428.471
A_24_P12	(NM_02393 mitochondr MRPL34,L	-1.41985	2.71E-09	0.7043	18251.84	12899.92
A_23_P76	BC034962 Hypothetic 15E1.2	-1.42014	5.37E-09	0.70416	1293.753	909.6806
A_23_P21	(NM_01339 Homo sapi C6orf49,OI	-1.4202	3.16E-07	0.70413	19555.72	13868.76
A_23_P44	(NM_01623 anaphase- ANAPC7,A	-1.42026	0.00002	0.70409	1107.04	777.255
A_32_P72	(NM_01450 ubiquitin c UBE2S,E2	-1.42031	2.87E-09	0.70407	12698.09	8914.836
A_24_P126851	Unknown A_24_P12	-1.42062	4.62E-11	0.70392	1744.571	1229.099
A_23_P12	(NM_03270 tubulin alpt TUBA6,MC	-1.42064	1.66E-10	0.70391	26914.14	18887.18
A_23_P77	(NM_01867 mesoderm MESP1,IR	-1.42084	1.98E-09	0.70381	2937.385	2062.914
A_23_P17	(NM_00703 Tara-like p HRIHFB21	-1.42108	9.04E-12	0.70369	887.6835	624.6057
A_23_P43	(NM_01422 Homo sapi NDUFA8,P	-1.42111	7.29E-14	0.70368	55040.05	38669.16

A_24_P11	NM_03329	T-cell activ	MRFAP1,F	-1.42143	3.07E-14	0.70352	3105.246	2185.645
A_23_P56	NM_00168	ATP synth	ATP5G3,M	-1.42162	2.07E-10	0.70342	142839.2	100651.5
A_23_P11	NM_01221	Homo sapi	TRIM32,H1	-1.42164	4.86E-07	0.70342	648.1304	456.0148
A_23_P41	NM_01843	Homo sapi	RNF130,G	-1.42175	2.95E-06	0.70336	6002.556	4252.184
A_23_P50	NM_07847	Homo sapi	TM2D2,BL	-1.42212	2.40E-13	0.70318	4496.981	3163.017
A_23_P14	NM_00114	Homo sapi	AMFR,GP7	-1.42214	1.41E-10	0.70317	3240.273	2281.275
A_24_P7021	Unknown	A_24_P70		-1.42231	4.39E-13	0.70308	2105.374	1478.992
A_23_P16	NM_01630	Homo sapi	SS18L2,KI	-1.42225	2.29E-13	0.70299	12003.75	8450.822
A_24_P16	NM_00088	IMP (inosin	IMPDH2,IM	-1.42268	1.27E-07	0.7029	5629.078	3933.184
A_23_P43	NM_01642	Homo sapi	ZNF219,ZF	-1.42279	3.71E-08	0.70284	19800.8	13835.52
A_32_P39	BC039097	Homo sapi	BC039097	-1.42284	8.41E-11	0.70282	1013.533	712.5067
A_23_P48	NM_00028	Homo sapi	PCCA	-1.42291	1.05E-13	0.70279	1492.596	1048.711
A_23_P66	BC047010	Similar to	TLOC40050	-1.423	2.29E-06	0.70274	3339.528	2353.528
A_32_P22	BG283145	Hypothetic	LOC28602	-1.42327	0.00008	0.70261	1054.132	739.5183
A_24_P922120	ALU1_HU	THC18729		-1.42361	1.23E-06	0.70244	487.295	342.307
A_23_P21	NM_01333	protein trar	SEC61A1,1	-1.42409	0.00033	0.7022	2476.187	1715.481
A_24_P11	AK022628	Forkhead t	KIAA0415	-1.42413	4.69E-14	0.70218	2241.267	1574.293
A_23_P49	NM_05293	hypothetic	MGC20781	-1.42414	4.12E-06	0.70218	1071.959	752.9452
A_23_P16	NM_00736	Homo sapi	NID2	-1.42428	0.0002	0.70211	2125.019	1505.967
A_23_P25	NM_00192	Homo sapi	DDB1,XPE	-1.42434	0.00012	0.70208	7418.513	5169.711
A_23_P98	NM_00697	Homo sapi	ZNF32,KO	-1.42439	0.00007	0.70205	344.5828	241.8996
A_24_P25	NM_00683	ATP/GTP-t	HEAB,CLP	-1.42453	2.63E-07	0.70199	849.6666	596.5071
A_23_P10	NM_00260	Homo sapi	PDE6D,PD	-1.4246	0.00001	0.70195	1184.059	829.626
A_32_P12	AB011542	EGF-like-d	EGFL5	-1.4246	5.75E-08	0.70195	1252.696	879.3632
A_32_P169142	AF030694	THC18524		-1.42468	1.72E-13	0.70191	989.8185	694.7583
A_24_P39	NM_01544	DKFZP586	C1orf43,NI	-1.42476	1.16E-14	0.70187	4820.143	3378.574
A_23_P87	NM_03259	Homo sapi	FBXL10,C)	-1.42479	0.00002	0.70186	742.3269	520.9305
A_23_P99	NM_01591	homolog of	POP5,RPF	-1.42491	4.02E-10	0.7018	4314.611	3034.7
A_24_P18	NM_03121	Chromosor	C19orf27	-1.42499	3.30E-13	0.70176	5904.082	4147.84
A_23_P25	NM_00645	Homo sapi	MXD4,MAI	-1.42505	0.00009	0.70173	536.1228	376.1116
A_24_P29	NM_01461	expressed	E7EA,KIA/	-1.42516	0.00005	0.70168	2594.063	1831.936
A_23_P12	NM_00138	Homo sapi	DPH2,DPH	-1.4252	4.29E-14	0.70165	4089.997	2871.982
A_32_P14	AK001118	Ubiquitin p	UBR2	-1.42552	0.00048	0.7015	291.2433	204.2313
A_23_P21	NM_02276	Homo sapi	CERK,LK4	-1.4257	8.78E-06	0.70141	705.8637	495.0698
A_24_P14	NM_00477	P311 prote	C5orf13,P3	-1.42578	3.80E-09	0.70137	895.4677	627.6973
A_32_P194095	ALU5_HU	THC20037		-1.42592	0.00001	0.7013	2159.835	1512.302
A_23_P12	NM_00271	protein phc	PPP2CA,R	-1.42618	8.00E-14	0.70117	8668.441	6068.257
A_32_P50655	ALU5_HU	THC18078		-1.42619	5.46E-06	0.70117	1200.199	845.8035
A_24_P21	NM_00500	non-metas	NME4,NM2	-1.42634	1.28E-06	0.7011	3382.863	2357.887
A_24_P35	NM_00147	UDP-N-acc	GALGT,GA	-1.42651	2.36E-09	0.70101	855.5765	599.7453
A_23_P10	NM_01790	aurora-A ki	AURKAIP1	-1.42654	1.07E-14	0.701	18320.31	12860.75
A_23_P17	NM_02522	Homo sapi	ADPN,PNF	-1.42663	3.17E-09	0.70095	1121.056	786.8912
A_24_P40	BG206642	Similar to	FBG206642	-1.42667	8.38E-09	0.70093	1769.97	1242.395
A_32_P186865	U12390	be THC20046		-1.42667	0.00002	0.70093	826.2625	579.8195
A_23_P52	NM_03236	Homo sapi	ACBD6,MC	-1.42683	5.23E-13	0.70085	10800.38	7551.718
A_23_P37966	DEFM_HU	THC19210		-1.42701	0.00001	0.70077	547.6628	383.7589
A_23_P50	NM_01819	Homo sapi	ZFP64,MG	-1.42716	2.72E-13	0.70069	2309.042	1617.599
A_23_P15	NM_02520	hypothetic	FLJ21945,I	-1.42792	1.05E-07	0.70032	600.7074	420.7086
A_24_P58477	Unknown	A_24_P58		-1.42858	2.91E-12	0.7	18461.84	12938.58
A_23_P17	NM_01765	Homo sapi	DYM,DMC	-1.42858	2.47E-07	0.69999	3338.978	2323.138
A_23_P11	NM_02306	Homo sapi	FOXL2,BP	-1.42869	8.99E-09	0.69994	1239.91	868.1376
A_24_P157720	Unknown	A_24_P15		-1.42885	6.45E-06	0.69987	1650.106	1150.239

A_23_P252125	TRI6_HUM	THC18150	-1.42912	0.00001	0.69973	14941.21	10508.96
A_23_P82	NM_00251	Homo sapi NOV,CCN	-1.42924	0.00009	0.69967	452.0214	316.0035
A_23_P20	NM_01445	Homo sapi PCDH17,P	-1.42935	9.65E-18	0.69962	6238.162	4364.64
A_23_P14	NM_03329	T-cell activ MRFAP1,F	-1.42937	3.69E-17	0.69961	8443.39	5908.21
A_23_P76	NM_00605	reticulon 3 RTN3,HAP	-1.42944	0.00017	0.69957	1582.05	1111.244
A_23_P13	NM_03235	hypothetic MGC11257	-1.43014	2.14E-16	0.69923	2707.385	1893.273
A_23_P12	NM_01633	Selenoprot SEPX1	-1.43054	9.76E-16	0.69903	2542.963	1776.698
A_24_P25	NM_00029	phosphogl PGK1,PGK	-1.43058	1.70E-06	0.69902	3516.895	2470.448
A_23_P33	NM_01546	Homo sapi GEMIN5,D	-1.43081	6.82E-08	0.69891	1695.322	1187.89
A_23_P37	NM_01598	Homo sapi UBE2D4,H	-1.43086	1.72E-09	0.69888	1360.461	949.8873
A_23_P99	AL832695	Homo sapi DKFZp313	-1.43086	1.26E-08	0.69888	781.322	545.9064
A_24_P20	AY033611	Homo sapi AY033611	-1.43093	4.33E-09	0.69885	83543.33	58654.23
A_32_P200120	Unknown	THC19035	-1.43097	0.00037	0.69883	531.3474	371.1122
A_23_P16	NM_01209	adaptor prc APPL,APP	-1.43117	1.85E-06	0.69873	2224.867	1567.948
A_23_P54	BX161435	Thiamine tr THTPA	-1.43128	2.00E-07	0.69868	1394.533	975.5614
A_23_P10	NM_02089	Homo sapi ZBTB4,KAI	-1.43129	8.14E-07	0.69867	2328.757	1630.76
A_24_P65	AY007106	Keratin ass KRTAP4-7	-1.43165	1.07E-14	0.6985	3381.196	2363.817
A_32_P14	NM_00693	SMT3 (sup SUMO2,H	-1.43168	8.68E-10	0.69848	27348.39	19034.6
A_23_P53	NM_01425	Homo sapi TMEM4,M	-1.4318	2.68E-08	0.69842	72257.38	50199.56
A_23_P63	NM_18174	Homo sapi LASS2,L3,	-1.43201	3.35E-11	0.69832	6480.04	4518.855
A_24_P13	NM_00114	autocrine n AMFR,GP7	-1.43275	1.87E-13	0.69796	1738.076	1212.956
A_24_P88	AK094413	MRNA full AK094413	-1.43416	1.52E-15	0.69727	1522.031	1061.274
A_24_P19	AL713633	Nudix (nuc NUDT1	-1.43451	3.19E-15	0.6971	1334.876	930.5641
A_24_P85	AB051538	KIAA1751 KIAA1751	-1.43458	0.00052	0.69707	1804.516	1259.292
A_24_P392742	Unknown	A_24_P39	-1.43472	0.00011	0.697	591.1635	411.9706
A_23_P94	AY358408	Clone DNA LGLL338	-1.43516	1.28E-07	0.69678	494.0433	344.2276
A_24_P50	AK074866	Tripartite rr TRIM11	-1.43574	0.00028	0.69651	859.8388	598.6891
A_23_P39	NM_05285	Homo sapi GADD45G	-1.43592	8.65E-17	0.69642	12659.22	8820.03
A_24_P18	NM_17370	Homo sapi NM_17370	-1.43595	2.64E-08	0.69641	6035.036	4222.728
A_23_P38	NM_00593	MHC class MICB	-1.43615	0.0002	0.69631	1133.038	790.9282
A_23_P119794	Unknown	THC18076	-1.43618	0.00033	0.69629	947.3072	660.1759
A_23_P14	NM_00312	Homo sapi SQLE	-1.43629	0.00012	0.69624	3635.877	2555.716
A_24_P32920	AF386504	THC19247	-1.43665	3.91E-14	0.69606	3882.466	2706.617
A_23_P15	BC017032	Glucosamin GCNT3	-1.43681	3.87E-06	0.69599	1794.598	1249.299
A_32_P55	NM_18262	Homo sapi FAM33A,F	-1.43683	8.60E-06	0.69598	828.9543	578.1202
A_23_P36	AL390079	Hypothetic LOC58489	-1.43691	7.32E-12	0.69594	865.1038	602.1091
A_24_P67494	Unknown	ENST0000	-1.43705	0.00004	0.69587	1533.894	1066.297
A_32_P76	NM_15268	hypothetic RWDD4A,I	-1.43729	9.74E-09	0.69575	778.7754	541.639
A_24_P33	NM_14507	Homo sapi HTRA2,OM	-1.43742	8.87E-16	0.69569	2744.194	1909.874
A_24_P94	AK022820	Family with FAM33A	-1.43745	6.30E-08	0.69567	1110.954	772.7986
A_23_P98	NM_00019	Homo sapi HMBS,UP	-1.43753	4.09E-07	0.69564	24665.9	17108.74
A_32_P125177	AY027936	THC19468	-1.4376	0.00017	0.6956	1974.605	1382.124
A_32_P127209	ALU4_HUM	THC18351	-1.43775	2.18E-07	0.69553	1748.22	1222.977
A_24_P28	NM_03075	makorin, rit MKRN4	-1.43779	0.00002	0.69551	604.8485	420.6823
A_23_P21	NM_01867	smooth mu SMAP-1,FI	-1.43789	1.46E-15	0.69546	4262.78	2963.885
A_23_P32	NM_03316	Homo sapi SURF4,ER	-1.43797	1.14E-07	0.69542	5792.291	4013.085
A_23_P43	NM_03229	hypothetic CAMKK1,C	-1.43825	3.91E-06	0.69529	1592.09	1110.29
A_23_P33	NM_15226	hypothetic FLJ38663	-1.43836	0.00017	0.69524	871.3985	607.051
A_32_P82963	Unknown	A_32_P82	-1.43881	3.94E-14	0.69502	2582.509	1794.397
A_24_P23	NM_00566	voltage-dep VDAC3,HC	-1.43887	8.24E-14	0.69499	6027.673	4178.953
A_23_P79	NM_00681	proteasom PSMF1,PI	-1.43891	2.89E-06	0.69497	2615.533	1831.801
A_23_P54	NM_00480	stomatatin (E STOML1,S	-1.43911	3.88E-16	0.69487	4128.707	2869.275

A_24_P460195	AY173017 THC18890	-1.4393	1.17E-11	0.69478	1605.954	1116.408
A_23_P15	NM_01662 archaemet: AMZ2	-1.43971	9.44E-11	0.69459	11524.15	8034.762
A_23_P46	NM_03263 differential PSRC1,DC	-1.43984	5.37E-07	0.69452	2057.952	1424.375
A_23_P93	NM_01399 Homo sapi DDR1,CAK	-1.44002	1.01E-07	0.69444	763.9948	530.4504
A_24_P204204	Unknown A_24_P20	-1.44022	4.45E-13	0.69434	2215.373	1540.369
A_24_P324084	Unknown A_24_P32	-1.44029	1.20E-07	0.6943	5608.924	3914.5
A_23_P16	NM_15346 Homo sapi IL17RC,IL1	-1.44093	0.00017	0.69399	1027.538	717.9764
A_24_P29	NM_01836 hypothetic: AGPAT5,L	-1.44109	3.47E-06	0.69392	1451.099	1010.134
A_23_P15	NM_00119 Homo sapi BID,MGC1	-1.44115	1.13E-07	0.69389	2081.187	1441.882
A_24_P63	NM_00295 ribophorin RPN1,RBF	-1.44124	3.04E-15	0.69385	1720.194	1194.03
A_24_P587882	Unknown A_24_P58	-1.44135	1.02E-07	0.69379	7237.395	5048.522
A_23_P40	NM_15321 Homo sapi C18orf17,F	-1.44172	8.93E-09	0.69361	682.1207	473.0762
A_24_P40	NM_03121 sec13-like SEH1L,Sel	-1.44186	4.21E-06	0.69355	3633.461	2542.961
A_24_P38	AL831891 Stearoyl-C: SCD4	-1.44197	6.54E-07	0.6935	416.9433	289.1661
A_23_P37	NM_03330 hypothetic: MGC4655	-1.44198	3.53E-10	0.69349	6627.049	4584.629
A_23_P12	NM_15283 Mouse Mar FAM89B,M	-1.44207	0.00014	0.69345	1935.64	1334.25
A_24_P58	NM_01910 beta-1,4 m: ALG1,HMT	-1.44235	6.67E-15	0.69331	2936.307	2035.968
A_24_P10	AF070629 RAB2, mer RAB2	-1.44286	6.84E-10	0.69307	1135.623	786.6175
A_32_P11	NM_19899 N-acyl-pho NAPE-PLD	-1.44302	8.79E-10	0.69299	1243.647	861.609
A_24_P79	NM_00606 aldo-keto r: AKR1A1,A	-1.44304	1.40E-07	0.69298	864.95	600.7135
A_24_P38	NM_00155 insulin-like IGFBP4,BF	-1.44329	3.18E-10	0.69286	1184.228	820.0838
A_24_P91	NM_17855 hypothetic: FLJ37478,I	-1.44342	9.73E-07	0.6928	925.0016	641.3955
A_23_P43	NM_01455 Homo sapi CARD10,B	-1.44397	5.56E-08	0.69253	3408.37	2353.82
A_32_P21	AK074567 Similar to A FLJ90086	-1.44399	3.19E-06	0.69252	592.9736	410.6879
A_32_P80	BC068045 CDNA clon BC068045	-1.44402	5.76E-07	0.69251	448.4246	310.5234
A_24_P24230	Unknown A_24_P24	-1.44411	0.00001	0.69247	711.6005	493.1075
A_24_P94	NM_01786 hypothetic: C1orf56,FL	-1.44435	5.20E-14	0.69235	1791.531	1241.198
A_24_P73	NM_00411 flap structu FEN1,MF1	-1.44436	1.77E-09	0.69235	2293.585	1588.867
A_23_P20	NM_01517 KIAA0717 RHOBTB2	-1.44443	2.44E-11	0.69231	905.7474	627.2408
A_32_P15	BX537645 Hypothetic: LOC40139	-1.44472	1.64E-08	0.69218	2813.094	1943.672
A_23_P15	NM_00373 Homo sapi SSNA1,N1	-1.4449	1.14E-12	0.69209	5209.969	3604.363
A_23_P30	BC013438 Hypothetic: LOC37529	-1.44514	0.00082	0.69198	1329.665	930.2498
A_23_P20	BX648591 Contactin 1CNTN1	-1.44517	1.08E-09	0.69196	10355.3	7153.955
A_23_P20	NM_02301 Homo sapi NADK,FLJ	-1.4452	0.00002	0.69195	2118.205	1465.268
A_23_P36	NM_01220 beta-1,3-gl B3GAT3,G	-1.44524	0.00002	0.69193	495.6892	342.8076
A_23_P10	AK130644 Solute carr SLCO3A1	-1.4454	0.00008	0.69185	377.9396	261.5709
A_24_P39	NM_00525 cyclin G as GAK,MGC	-1.44552	2.24E-11	0.69179	2620.266	1808.007
A_23_P60	NM_00243 mannose p MPI,PMI,P	-1.44557	4.51E-07	0.69177	2067.712	1434.613
A_23_P67	NM_13848 Homo sapi LOC15221	-1.44568	1.60E-08	0.69172	15600.91	10713.64
A_32_P13	BC041080 Homo sapi BC041080	-1.44568	2.89E-06	0.69172	733.1467	507.2872
A_23_P16	AK091324 Hypothetic: LOC21985	-1.44595	1.28E-06	0.69158	3574.8	2489.625
A_32_P11	AK092667 FLJ35348 FLJ35348	-1.44595	3.98E-10	0.69159	4898.33	3387.624
A_32_P16	NM_00506 stearoyl-C: SCD,SCD1	-1.44599	8.91E-07	0.69157	22644.49	15571.88
A_23_P11	NM_01899 Homo sapi FBXW5,Fb	-1.4464	1.44E-07	0.69137	4743.612	3257.198
A_32_P16	NM_20737 hypothetic: LOC38788	-1.44652	0.00006	0.69131	1432.034	992.7882
A_23_P30	AK094772 Homo sapi AK094772	-1.44668	0.00001	0.69124	411.5408	284.5637
A_24_P178523	Unknown ENST0000	-1.44678	2.59E-11	0.69119	706.9323	488.644
A_23_P86	NM_15236 Kelch/anky KARCA1	-1.44711	0.00003	0.69103	2334.13	1609.918
A_23_P14	NM_03214 hypothetic: C16orf48,C	-1.44732	2.15E-11	0.69093	1235.104	854.0155
A_23_P87	NM_00633 C2f protein C2F	-1.44761	3.28E-13	0.69079	29999.08	20681.71
A_23_P52	NM_01402 Homo sapi DCPS,HSL	-1.44768	3.30E-07	0.69076	5392.738	3698.56
A_23_P10	NM_00451 integrin-lin ILK,P59,DI	-1.44768	4.09E-14	0.69076	987.5641	682.1569

A_24_P25: NM_03283	hypothetic	CIRH1A,N	-1.4477	1.00E-11	0.69075	4001.713	2761.358
A_24_P23: AB040958	BTB (POZ)	BTBD7	-1.44781	0.00035	0.6907	1759.502	1229.181
A_23_P41: NM_14526	similar to	RLOC13395	-1.448	1.42E-11	0.69061	2253.955	1554.057
A_32_P10: AK094156	Similar to	AK094156	-1.44801	1.37E-08	0.6906	562.0499	388.1611
A_23_P85: NM_00703	RER1	reter RER1	-1.44806	1.23E-18	0.69058	7066.165	4879.025
A_23_P83: NM_00443	Homo sapi	ENDOG	-1.44812	0.00009	0.69055	5060.916	3517.046
A_24_P31: BC040210	Aldo-keto r	AKR1C1	-1.44814	2.61E-09	0.69054	7204.135	4952.058
A_23_P12: NM_00308	Homo sapi	SNCB	-1.44821	7.40E-06	0.69051	2221.207	1550.03
A_32_P12: NM_01323	px19-like p	PX19,PRE	-1.44827	4.48E-12	0.69048	20875.02	14370.72
A_23_P16: NM_00257	Homo sapi	PAFAH1B3	-1.4483	5.20E-11	0.69047	2005.03	1382.876
A_24_P212764	Unknown	A_24_P21:	-1.44832	2.91E-13	0.69046	9458.034	6546.53
A_32_P16: BX363129	Transcribe	BX363129	-1.44833	2.10E-08	0.69045	600.9294	414.8565
A_23_P11: NM_00211	Homo sapi	HDC	-1.44874	0.0004	0.69025	822.1565	568.5785
A_23_P95: NM_00332	Homo sapi	TUFM,EFT	-1.44884	2.54E-18	0.69021	37652.63	25983.58
A_24_P37: BC016333	Cyclin L2	CCNL2	-1.44884	1.21E-14	0.69021	1001.626	691.3278
A_23_P94: NM_00321	Homo sapi	TEAD4,RT	-1.44885	3.68E-11	0.6902	732.3674	505.4503
A_24_P20: NM_01804	hypothetic	PLEKHJ1,(-1.44894	1.47E-12	0.69016	1903.602	1314.107
A_23_P54: NM_00629	Homo sapi	TYRO3,BY	-1.44908	1.95E-08	0.69009	2530.037	1747.059
A_23_P12: NM_02490	hypothetic	WDR76,FL	-1.44913	0.00073	0.69007	622.9981	429.083
A_23_P84: NM_00493	Homo sapi	CDK5,PSS	-1.44925	0.00008	0.69001	2607.153	1801.727
A_23_P84: NM_01848	Homo sapi	HDAC8,RF	-1.44959	0.00088	0.68985	409.4217	282.0219
A_23_P40: NM_13843	hypothetic	MFSD3	-1.44978	7.17E-06	0.68976	1003.076	693.3563
A_23_P43: NM_01418	HSPC159	HSPC159,	-1.45027	6.53E-07	0.68953	2491.252	1730.916
A_23_P16: NM_00684	Homo sapi	ILVBL,AHA	-1.4506	1.19E-09	0.68937	2672.936	1849.735
A_23_P80: NM_00691	Homo sapi	RNF5,RMA	-1.45072	0.0001	0.68931	1647.94	1138.279
A_23_P11: NM_00476	B-cell CLL/	BCL7C	-1.45104	2.71E-14	0.68916	1373.287	946.3322
A_32_P20: NM_01609	ribosomal	RPL26L1,F	-1.45141	2.27E-12	0.68898	20398.19	14091.16
A_24_P18: NM_01661	hypothetic	PLAC8,C1:	-1.45173	1.75E-06	0.68883	5729.384	3982.289
A_23_P22: NM_00635	Homo sapi	SLC9A6,NI	-1.45266	6.48E-06	0.68839	1789.089	1235.14
A_23_P15: NM_03290	Homo sapi	PPP1R16A	-1.45271	8.43E-13	0.68837	5195.483	3581.544
A_23_P14: NM_14565	Homo sapi	RAD52B,R	-1.45314	7.10E-07	0.68816	747.7611	514.7495
A_23_P94: AB018254	KIAA0711	KIAA0711	-1.4532	3.26E-09	0.68814	2789.092	1921.862
A_23_P11: NM_01609	DNA replic	Pfs2,HSPC	-1.45324	4.07E-12	0.68812	4216.611	2909.419
A_32_P12: AK096908	CDNA FLJ:	AK096908	-1.4534	3.71E-13	0.68804	1532.302	1054.886
A_24_P53: NM_00206	glutamate-	GLUL,GS,(-1.45348	3.54E-09	0.68801	1105.756	759.4134
A_23_P41: NM_03294	putative srr	NID67,MGI	-1.45353	4.61E-07	0.68798	504.1435	346.8881
A_23_P34: NM_02121	Homo sapi	ZNF71,EZI	-1.4537	1.26E-09	0.6879	805.7541	554.3506
A_23_P27: NM_01804	Homo sapi	PLEKHJ1,(-1.45404	1.23E-09	0.68774	10017.99	6865.219
A_23_P20: D43947	KIAA0100	KIAA0100	-1.45457	1.32E-14	0.68749	6739.157	4628.704
A_23_P74: NM_14569	Homo sapi	CDCA1,NL	-1.45465	4.10E-09	0.68745	5195.48	3590.177
A_23_P25: NM_00689	Homo sapi	HOXC9,HC	-1.45475	3.85E-08	0.6874	2680.964	1841.115
A_24_P15: NM_20696	Homo sapi	HRMT1L1,	-1.45529	1.44E-15	0.68715	1411.303	969.9635
A_24_P80: NM_02406	hypothetic	PDCL3,VI/	-1.45536	2.03E-07	0.68712	9546.963	6588.329
A_24_P489399	Unknown	A_24_P48:	-1.45551	1.34E-06	0.68704	1259.861	867.6804
A_23_P17: NM_00657	Homo sapi	EBP,CPX,(-1.4559	6.68E-08	0.68686	31504.25	21483.02
A_23_P16: BC044941	Similar to	LOC44045	-1.45591	3.81E-14	0.68686	1165.359	800.5483
A_24_P19: NM_13039	Homo sapi	WRNIP1,V	-1.45594	5.29E-15	0.68684	1207.328	829.2229
A_24_P17: AY029066	Homo sapi	HN1	-1.45617	9.02E-13	0.68673	483883.6	333580.3
A_23_P21: NM_02032	Homo sapi	RARSL,DA	-1.45635	3.89E-12	0.68665	8128.453	5588.559
A_23_P41: NM_02471	Homo sapi	C5orf14,UI	-1.45639	8.89E-13	0.68663	6926.163	4740.151
A_23_P10: NM_02113	Homo sapi	MRPL23,R	-1.45653	5.27E-07	0.68656	10301.89	7007.008
A_32_P35: NM_03078	phosphatid	PTDSS2,P	-1.45687	1.15E-14	0.6864	7195.803	4945.656

A_23_P36	NM_01566	Homo sapi	GIMAP2,IM	-1.45688	2.89E-13	0.6864	1080.061	741.4114
A_24_P37	NM_01822	chromosom	MOBK1B,M	-1.4571	2.09E-10	0.68629	2227.071	1531.272
A_23_P13	NM_00197	Homo sapi	ENO3,MSE	-1.45726	5.21E-09	0.68622	4763.745	3281.014
A_24_P40	NM_00330	Homo sapi	TRIP6,OIP	-1.45746	3.91E-13	0.68613	14321	9850.182
A_23_P48	NM_02283	mitochondr	MRPS11,H	-1.45768	6.14E-16	0.68602	10827.01	7417.717
A_24_P178333	Unknown	A_24_P17		-1.45769	7.71E-06	0.68602	935.3016	638.9793
A_23_P69	NM_02466	hypothetic	FLJ12436	-1.45781	3.13E-08	0.68596	6472.158	4425.291
A_23_P13	NM_19899	Homo sapi	KCTD1,C1	-1.45828	8.07E-09	0.68574	2888.045	1989.695
A_23_P20	BC022779	V-akt murir	AKT2	-1.45837	3.84E-07	0.68569	1253.382	860.7037
A_23_P11	AB067472	Valyl-tRNA	VARS2L	-1.45838	3.65E-11	0.68569	2746.269	1879.429
A_23_P41	XM_37335	similar to d	LOC39247	-1.45858	2.08E-06	0.6856	1311.228	898.1908
A_24_P39	NM_00166	ADP-ribosy	ARF5	-1.45865	7.03E-07	0.68557	2143.003	1461.278
A_32_P21	AK098678	Interleukin-	IRAK1BP1	-1.45892	0.00024	0.68544	753.2844	516.3185
A_23_P76	NM_00421	Homo sapi	PTTG1,EA	-1.45895	5.18E-11	0.68543	55804.39	38411.23
A_23_P66	NM_00037	Homo sapi	UMPS,OPI	-1.45904	6.90E-20	0.68538	12648.95	8669.671
A_32_P36	XM_05977	Homo sapi	FKBP1C	-1.45906	1.41E-08	0.68537	1465.057	1005.222
A_24_P24	NM_00395	myelin prot	MPZL1,PZ	-1.45927	9.03E-13	0.68528	1262.857	864.6925
A_23_P14	NM_00500	Homo sapi	NDUFAB1,	-1.4594	3.20E-17	0.68521	40022.39	27449.79
A_24_P37	BC020242	Protein tyr	PLIP	-1.45952	1.64E-15	0.68516	7609.563	5205.473
A_24_P53	NM_03227	hypothetic	TRAF7,RF	-1.45971	0.00003	0.68507	6757.385	4641.509
A_24_P290188	Unknown	ENST0000		-1.45983	0.00061	0.68501	1393.104	964.9045
A_23_P17	NM_00726	Homo sapi	XAB1,MB	-1.45995	1.66E-16	0.68496	6929.641	4741.162
A_23_P61	NM_00611	CD2 antige	CD2BP2,F	-1.46005	2.62E-08	0.68491	485.0601	332.2234
A_24_P37	NM_00478	vesicle-ass	VAMP3,CE	-1.46013	4.02E-07	0.68487	592.3428	405.7022
A_23_P79	NM_01323	px19-like p	PX19,PRE	-1.46026	2.05E-13	0.68481	32850.62	22441.46
A_23_P46	NM_01783	hypothetic	PIGV,FLJ2	-1.46053	2.27E-12	0.68468	767.8575	525.7556
A_23_P15	NM_02431	Homo sapi	C7orf23,M	-1.46079	1.46E-07	0.68456	3316.3	2281.292
A_24_P531074	Unknown	A_24_P53		-1.46091	0.00004	0.6845	3135.017	2165.087
A_24_P37	BC008564	Protein phc	PPP1R9B	-1.46111	6.37E-19	0.68441	4368.719	2990.35
A_32_P151191	Unknown	A_32_P15		-1.46141	0.00003	0.68427	426.0825	291.3509
A_24_P34	NM_13849	hypothetic	MGC2164	-1.46141	1.68E-06	0.68427	454.5602	311.071
A_24_P67268	Unknown	A_24_P67		-1.46149	4.71E-14	0.68423	8119.799	5543.305
A_23_P20	NM_03281	hypothetic	FLJ14627	-1.46167	1.36E-08	0.68415	1232.567	845.2161
A_32_P75141	Unknown	A_32_P75		-1.4618	3.13E-10	0.68409	649.3702	444.2454
A_23_P31	NM_00053	Homo sapi	RFXAP	-1.4618	0.0007	0.68409	363.6461	249.0201
A_32_P9816	Unknown	A_32_P98		-1.4618	0.00074	0.68409	271.6296	185.693
A_24_P20	NM_03323	promyeloc	PML,MYL,I	-1.46208	0.00005	0.68396	622.0996	425.4105
A_23_P25	X66610	H.sapiens	enolase	-1.46228	8.14E-06	0.68386	3667.482	2483.644
A_23_P65	NM_02191	Homo sapi	CFL2	-1.46237	5.74E-17	0.68382	3081.709	2107.52
A_23_P10	NM_00350	Homo sapi	ACOX2,BC	-1.46273	3.34E-19	0.68365	6948.569	4749.266
A_32_P55	AK096908	CDNA FLJ:	AK096908	-1.46281	5.20E-11	0.68362	2093.367	1433.703
A_24_P27	NM_01606	CGI-135 pr	TTC11,Fis	-1.46424	3.89E-07	0.68295	2905.519	1999.2
A_23_P254288	Unknown	A_23_P25		-1.46453	4.27E-14	0.68281	11854.18	8105.664
A_24_P36	NM_01339	FtsJ homol	FTSJ2,FJH	-1.46547	1.01E-12	0.68237	1217.62	830.0529
A_32_P19	AF435960	Chromosor	C10orf90	-1.46558	0.00001	0.68232	3324.105	2290.974
A_24_P34	NM_00196	eukaryotic	EIF4E,CBF	-1.46564	6.71E-18	0.6823	1663.659	1135.091
A_23_P96	NM_01773	O-linked m	POMGNT1	-1.46573	1.48E-10	0.68225	1158.582	790.4571
A_23_P80	NM_00270	Homo sapi	PPAT,GPA	-1.46576	2.11E-11	0.68224	1696.078	1158.745
A_24_P29	BC014203	Hypothetic	MGC45871	-1.46591	1.41E-08	0.68217	3212.962	2193.505
A_24_P727868	VAPA_HUI	THC18449		-1.46619	1.90E-19	0.68204	8181.356	5579
A_23_P37	NM_01476	24-dehydr	DHCR24,K	-1.46646	7.50E-12	0.68191	2309.561	1569.06
A_24_P38	NM_00138	dihydropyri	DPYSL2,D	-1.46658	0.00008	0.68186	3067.117	2111.174

A_23_P11	NM_01414 nudix (nucl NUDT5,YS	-1.46672	2.99E-18	0.68179	12209	8318.313
A_23_P18	NM_01658 Chromosom C3orf1	-1.46672	6.72E-08	0.68179	34789.91	23511.34
A_24_P14	NM_01824 hypothetical FLJ10847,l	-1.46697	0.00007	0.68168	545.6123	371.8978
A_23_P50	NM_02110 Homo sapi MRPS12,R	-1.46704	2.72E-08	0.68165	43109.52	29251.09
A_23_P92	NM_03233 hypothetical MGC2408	-1.46706	5.42E-11	0.68163	6304.363	4285.646
A_24_P91	AK001936 Eukaryotic EIF4EBP2	-1.46714	0.00006	0.6816	291.1478	198.434
A_23_P55	NM_02515 hypothetical TARSL1,FI	-1.4672	0.00002	0.68157	4013.842	2754.854
A_23_P13	NM_01620 Vestigial-like FLJ38507	-1.46721	2.55E-08	0.68157	646.1403	440.4551
A_32_P16462	ALU6_HUMAN THC19454	-1.46763	7.86E-06	0.68137	796.5883	543.1932
A_23_P57	AF231919 Chromosom C21orf108	-1.4677	2.95E-07	0.68134	941.8628	642.4489
A_24_P19	XM_11589 similar to HLOC20351	-1.46786	0.0001	0.68126	1599.346	1096.931
A_24_P38	NM_19946 Homo sapi NANOS1	-1.4679	1.21E-08	0.68124	636.3596	433.6083
A_24_P671875	AB015066 THC19206	-1.46811	2.59E-19	0.68115	2591.852	1765.433
A_24_P341106	Unknown A_24_P34	-1.46828	0.00001	0.68107	1274.724	870.9518
A_23_P65	NM_19852 Similar to FHDDC3,MC	-1.46838	3.85E-17	0.68102	4238.622	2884.665
A_23_P14	NM_01545 cofactor of COBRA1,N	-1.46839	3.00E-09	0.68102	3885.228	2642.74
A_23_P61	NM_01820 Homo sapi LRRC20,FI	-1.46866	4.15E-07	0.68089	1272.825	864.6879
A_32_P95223	FPPS_HUMAN ENST0000	-1.46907	5.74E-14	0.6807	4998.539	3397.768
A_24_P37	NM_02191 cofilin 2 (m CFL2	-1.46914	1.69E-11	0.68067	3097.392	2114.164
A_24_P46	BC046178 Chromosom C6orf173	-1.4692	1.48E-13	0.68064	5098.063	3478.285
A_24_P67	AK023329 CDNA FLJ AK023329	-1.46921	2.75E-06	0.68064	745.7362	508.4225
A_24_P95	NM_17823 Homo sapi TUSC3,N3	-1.46934	1.54E-15	0.68058	3983.89	2715.695
A_24_P23	NM_02274 hypothetical MOSC1,FL	-1.46944	8.66E-13	0.68053	2062.148	1402.856
A_24_P29	NM_01814 hypothetical FAM82C,p	-1.46955	1.03E-10	0.68048	652.4422	443.9896
A_24_P28	AK074029 Hypothetical FLJ20255	-1.46988	3.67E-08	0.68033	1452.07	987.6099
A_23_P68	NM_02194 Homo sapi NM_02194	-1.46995	0.00031	0.68029	4613.016	3186.23
A_23_P15	NM_19840 Homo sapi PTPLB	-1.47008	0.0003	0.68024	2925.346	2008.94
A_32_P34	AK026418 FLJ39739 FLJ39739	-1.47009	4.47E-15	0.68023	1541.847	1048.016
A_23_P20	NM_03269 Homo sapi MGC16385	-1.47009	7.39E-12	0.68023	1310.84	890.6489
A_23_P54	NM_01780 hypothetical OSGEP,G	-1.47012	2.95E-15	0.68022	3364.155	2285.89
A_24_P63	AK022223 Hypothetical FLJ22955	-1.4702	1.31E-06	0.68018	864.4218	588.5133
A_32_P8806	ALU4_HUMAN THC19993	-1.4709	2.72E-07	0.67986	1772.49	1205.339
A_24_P37	NM_14517 Homo sapi DHRSX,DI	-1.47153	5.82E-08	0.67956	3912.081	2666.053
A_24_P27	NM_00307 SWI/SNF r SMARCA3	-1.4717	8.51E-09	0.67949	2632.95	1794.964
A_23_P25	NM_00294 ribosomal r RPL15,EC	-1.47182	2.83E-14	0.67943	4584.069	3108.247
A_24_P64	NM_00680 adaptor-rel AP3M2,P4	-1.47185	9.25E-08	0.67942	620.8755	421.764
A_23_P54	AB037759 Eukaryotic EIF2AK4	-1.47195	9.06E-17	0.67937	1939.403	1317.051
A_23_P21	BC036001 Dehydroge DHRS8	-1.47214	3.19E-12	0.67928	5285.796	3578.54
A_23_P92	NM_20584 Homo sapi SPINK6,BL	-1.47215	1.35E-10	0.67928	16693.69	11395.88
A_23_P25	NM_01814 Homo sapi FAIM,FAIM	-1.47273	6.92E-14	0.67901	8009.819	5423.173
A_24_P30	NM_01790 two-pore cl TPCN1,TP	-1.47302	0.00028	0.67888	792.9782	538.7365
A_24_P19	NM_19868 Homo sapi RAB15	-1.47304	1.18E-08	0.67887	876.3259	594.6056
A_24_P55	AK057088 AF034176 AK057088	-1.47367	1.24E-13	0.67858	3178.84	2161.84
A_23_P15	NM_01224 Homo sapi SEPHS1,S	-1.47368	1.57E-14	0.67857	9938.685	6752.917
A_23_P15	NM_00578 Homo sapi THOC4,AL	-1.47391	2.09E-16	0.67847	26469.11	17942.23
A_23_P88	NM_17067 Homo sapi MEIS2,MR	-1.47394	8.44E-09	0.67846	604.8561	410.3771
A_32_P160793	Unknown A_32_P16	-1.47441	0.00002	0.67824	5085.741	3495.214
A_32_P99	BC042161 Trafficking TRAPPC5	-1.47447	4.43E-18	0.67821	4807.529	3257.774
A_24_P30	NM_02272 Hpal1 tiny f HTF9C	-1.47497	3.66E-09	0.67798	658.2954	446.2888
A_23_P20	NM_01328 Homo sapi UHRF1,Np	-1.47503	6.75E-15	0.67795	6141.356	4169.727
A_23_P64	NM_02193 likely ortho RIC8A,RIC	-1.47515	5.58E-18	0.6779	17517.15	11882.88
A_24_P84	NM_00411 flap structu FEN1,MF1	-1.47531	1.77E-18	0.67782	2437.311	1652.36

A_23_P39	NM_01513	Myosin phc M-RIP	-1.47546	8.38E-12	0.67775	2563.68	1740.832
A_24_P15	NM_01231	LIV-1 prote SLC39A6,L	-1.47589	0.00028	0.67756	1451.85	986.1314
A_23_P12	BC003353	Hypothetic MGC5309	-1.47614	1.02E-15	0.67744	2511.292	1702.461
A_24_P10	NM_17844	Homo sapi C9orf140	-1.47642	3.53E-06	0.67731	3735.574	2531.831
A_23_P20	NM_00123	Homo sapi CCNE1,CC	-1.47664	1.90E-08	0.67721	7830.338	5281.895
A_24_P24	NM_02461	NOD9 prot NOD9,FLJ	-1.47725	6.95E-17	0.67693	2338.614	1582.082
A_23_P69	NM_01844	30 kDa pro LOC55831	-1.47768	2.03E-09	0.67674	3157.386	2140.201
A_23_P21	NM_01759	rotavirus X ZC3H7B,R	-1.47803	0.00003	0.67658	1599.59	1085.086
A_23_P28	NM_01825	Homo sapi WDR12,YT	-1.4782	0.00003	0.6765	7624.092	5113.967
A_23_P15	NM_00702	Homo sapi TOPBP1,T	-1.47855	0.00023	0.67634	1185.444	805.7053
A_24_P32	NM_01208	ATP-bindin ABCB10,M	-1.47866	8.94E-08	0.67629	422.2302	285.553
A_24_P49	BC020516	Interferon r IRF2BP2	-1.47872	1.57E-17	0.67626	1477.792	999.4742
A_23_P25	NM_00624	Homo sapi PPP2R5A	-1.47874	3.79E-14	0.67625	6626.158	4470.779
A_23_P43	NM_01799	Homo sapi C9orf40,FL	-1.47884	5.04E-07	0.6762	1202.086	814.5612
A_23_P44	235	AF098788 THC18430	-1.47887	3.22E-11	0.67619	3068.038	2078.889
A_23_P35	NM_02468	likely ortho E2F8,FLJ2	-1.479	9.44E-09	0.67613	1063.073	719.4219
A_24_P29	AF451991	PHD finger PHF10	-1.47937	1.37E-07	0.67596	1078.546	730.2294
A_23_P26	NM_00487	mannose-F MPDU1,SL	-1.48076	2.72E-17	0.67533	3782.823	2552.235
A_23_P12	NM_02408	Homo sapi ASB9,MGC	-1.48086	3.11E-11	0.67529	1996.725	1351.845
A_23_P88	NM_03263	cyclin-depe CINP,MGC	-1.48117	4.54E-06	0.67514	1080.781	727.7182
A_24_P88	NM_00384	peroxisom PEX11B,PI	-1.4813	1.58E-12	0.67508	1614.171	1090.503
A_23_P39	NM_01508	Homo sapi SPG20,SP	-1.48132	6.47E-12	0.67507	1332.979	899.5344
A_23_P69	NM_01844	30 kDa pro LOC55831	-1.48133	3.71E-13	0.67507	20569.91	13919.99
A_23_P42	NM_00131	mitogen-ac MAPK14,R	-1.48151	4.78E-16	0.67499	4068.994	2750.775
A_24_P29	NM_01446	Lsm3 prote LSM3,SMX	-1.48198	2.06E-13	0.67477	30923.71	20935.71
A_23_P57	NM_03307	Homo sapi CECR5	-1.48211	1.21E-06	0.67472	4627.741	3106.715
A_23_P33	NM_15236	Homo sapi ZNF573,FL	-1.48258	0.00002	0.6745	1566.117	1062.199
A_23_P13	NM_00175	Homo sapi CAV1,CAV	-1.48274	1.57E-06	0.67443	9461.163	6416.011
A_23_P40	NM_00591	Homo sapi MAT2A,MA	-1.48291	3.36E-07	0.67435	14026.33	9377.31
A_32_P49	NM_02474	disrupted ir ALG9,DIB	-1.48294	3.16E-08	0.67434	1588.724	1076.395
A_23_P12	NM_13039	Homo sapi TRIM34,IFI	-1.48353	0.00016	0.67407	304.0162	204.8534
A_23_P14	NM_00009	Homo sapi CPT2,CPT	-1.48382	9.22E-13	0.67394	1320.189	890.569
A_23_P83	AK022617	IGF-II mRN IMP-1	-1.48394	1.88E-13	0.67388	2213.235	1485.121
A_23_P13	AL132665	BCL2/aden BNIP3L	-1.48401	1.79E-06	0.67385	850.6025	574.3044
A_23_P11	NM_01640	Huntingtin HYPK,HSF	-1.48428	6.07E-20	0.67373	104340.3	70333.26
A_23_P21	NM_02492	Homo sapi NUP210,G	-1.4843	4.53E-11	0.67372	1540.374	1039.391
A_23_P37	NM_17590	Homo sapi ZADH2,MC	-1.48433	2.00E-16	0.67371	1759.783	1185.274
A_24_P12	NM_03223	Ubiquitin s USP48	-1.48468	1.49E-12	0.67355	1530.943	1031.683
A_23_P37	NM_00279	Homo sapi PSMB5,X,I	-1.48475	4.07E-10	0.67351	55958.13	37439.21
A_23_P54	NM_02393	Homo sapi C16orf24,NA	-1.48501	1.21E-08	0.6734	3906.715	2610.24
A_32_P17	NM_17504	Homo sapi PILRB,FD	-1.48525	2.27E-11	0.67329	1781.645	1204.168
A_24_P92	8765	Unknown A_24_P92	-1.48532	8.27E-13	0.67325	1725.708	1165.071
A_23_P16	NM_02403	hypothetic MGC2803	-1.48579	0.00002	0.67304	4678.793	3155.349
A_23_P36	NM_00550	Homo sapi SCARB1,C	-1.48606	2.10E-08	0.67292	719.6045	484.237
A_23_P12	NM_00367	Homo sapi DEGS1,ML	-1.48614	1.50E-06	0.67288	5153.131	3440.341
A_23_P92	NM_03148	Homo sapi ATG10,AP	-1.48639	0.00002	0.67277	877.4622	591.0267
A_23_P69	NM_01819	Homo sapi LEPREL1,I	-1.48642	2.59E-16	0.67276	3355.388	2258.849
A_24_P74	9042	Unknown A_24_P74	-1.48645	1.44E-19	0.67274	6369.006	4286.245
A_32_P17	NM_01450	ubiquitin ce UBE2S,E2	-1.48667	3.53E-16	0.67264	18765.01	12599.15
A_23_P10	NM_01601	CGI-69 pro CGI-69	-1.48685	1.21E-15	0.67256	4887.821	3292.916
A_23_P16	NM_00334	Homo sapi UBE2L3,E	-1.48702	3.19E-09	0.67248	5023.528	3385.306
A_23_P53	NM_01646	Homo sapi C14orf112,	-1.48755	8.74E-11	0.67225	21390.48	14302.33

A_23_P15:	NM_01208	Homo sapi	PGLS,6PG	-1.48781	4.58E-14	0.67213	4530.891	3052.591
A_32_P16:	AF161398	Chromosom	C6orf115	-1.48804	2.28E-08	0.67202	7123.627	4821.181
A_32_P83520		Unknown	THC18594	-1.48811	0.00035	0.67199	473.1502	317.7637
A_32_P11:	AB058761	Zinc finger	ZNF469	-1.48843	6.09E-13	0.67185	3525.524	2371.475
A_23_P35:	NM_01333	Homo sapi	ALG6	-1.48871	4.14E-16	0.67172	1845.824	1240.313
A_23_P12:	NM_01219	Homo sapi	FXC1,Tim	-1.48893	8.10E-15	0.67162	1415.638	950.1639
A_24_P17:	NM_00664	WAS prote	WASF3,SC	-1.48915	3.42E-09	0.67152	940.9676	631.2161
A_32_P57702		Unknown	A_32_P57	-1.48936	1.34E-07	0.67143	6559.686	4435.797
A_23_P12:	NM_00631	aminopepti	NPEPPS,F	-1.4896	1.54E-14	0.67132	23349.27	15633.84
A_23_P24:	NM_01646	hypothetic	HSPC196	-1.48964	2.91E-09	0.6713	1244.833	837.5857
A_24_P72:	BX538231	BTB (POZ)	BTBD7	-1.48964	3.59E-12	0.6713	1544.668	1038.864
A_23_P39:	NM_08067	Homo sapi	KCNE4,MII	-1.49026	1.31E-12	0.67102	4807.428	3231.741
A_23_P77:	NM_01902	protein arg	PRMT7,FL	-1.49044	4.79E-16	0.67094	7462.941	5013.983
A_23_P52:	NM_02393	Homo sapi	MRPL34,L	-1.49045	1.58E-13	0.67094	15128.88	10193.89
A_32_P15466		Unknown	THC19332	-1.49067	0.00002	0.67084	2658.486	1798.896
A_23_P20:	NM_00347	Homo sapi	CUL5,VAC	-1.49068	3.42E-07	0.67084	835.6025	560.9096
A_24_P94:	D80006	Homo sapi	KIAA0184	-1.49077	4.68E-13	0.67079	1107.746	742.8801
A_23_P15:	NM_00432	BCS1 (yea	BCS1L,BC	-1.49078	4.31E-16	0.67079	3145.567	2106.605
A_23_P57:	NM_02086	Homo sapi	DHX36,DD	-1.49079	1.34E-11	0.67078	5799.269	3902.313
A_23_P10:	NM_01423	upstream b	UBTF,UBF	-1.49081	0.00025	0.67078	525.1883	352.0215
A_23_P10:	NM_00248	Homo sapi	NDN,HsT1	-1.49086	3.04E-08	0.67075	1246.81	834.6835
A_23_P81:	NM_01836	Homo sapi	LMBRD1,C	-1.49103	2.65E-09	0.67068	4679.482	3145.111
A_23_P10:	NM_01613	Homo sapi	INSIG2,MC	-1.49138	0.0002	0.67052	560.6072	375.5989
A_24_P39:	AK095244	Cytochrom	CYB561	-1.49166	2.27E-15	0.6704	8080.53	5408.808
A_23_P13:	NM_02407	Homo sapi	ALG8,MGC	-1.49183	1.18E-07	0.67032	8214.961	5478.984
A_23_P34:	NM_17592	hypothetic	MGC4263C	-1.49207	0.00009	0.67021	351.127	235.4598
A_24_P152566		Unknown	A_24_P15:	-1.49208	0.00003	0.67021	583.9337	391.0963
A_23_P61:	AK098362	Chromosom	C7orf32	-1.4921	0.00068	0.6702	530.7892	355.155
A_24_P68:	NM_02084	Homo sapi	NM_02084	-1.49221	7.34E-13	0.67015	1581.653	1059.222
A_24_P42:	NM_02307	hypothetic	PYCRL,FL	-1.49231	3.30E-11	0.6701	3419.607	2285.543
A_23_P15:	NM_02457	Homo sapi	C16orf33,F	-1.49242	9.77E-16	0.67005	9934.382	6645.851
A_23_P42:	NM_17497	Homo sapi	SEC14L4,1	-1.49246	1.29E-16	0.67003	1327.419	889.588
A_23_P49:	NM_00347	Homo sapi	USP7,TEF	-1.49295	4.19E-13	0.66982	7020.307	4714.793
A_23_P31:	NM_00035	Homo sapi	TCOF1,MF	-1.49311	6.81E-10	0.66974	12174.5	8111.616
A_24_P20:	XM_37799	Homo sapi	XM_37799	-1.49343	1.39E-07	0.6696	15088.18	10148.41
A_23_P14:	NM_01818	hypothetic	FLJ10700	-1.49358	0.00006	0.66953	1277.607	859.8047
A_23_P33:	AJ314645	Hypothetic	FLJ22729	-1.49377	2.11E-07	0.66945	3579.275	2394.244
A_24_P35:	NM_01222	basic trans	GTPBP6,P	-1.49392	1.71E-06	0.66938	501.0162	335.4254
A_32_P11:	BC024301	Sorting nex	SNX12	-1.49394	1.40E-18	0.66937	11444.05	7654.456
A_23_P13:	AF182418	LSM3 hom	LSM3	-1.49405	2.47E-13	0.66932	21264.53	14289.03
A_23_P14:	NM_00259	Homo sapi	PDCD2,RF	-1.49416	1.82E-11	0.66927	4989.833	3358.81
A_32_P58:	NM_15381	Homo sapi	PHF13,PH	-1.49434	6.86E-06	0.66919	2889.978	1944.997
A_23_P49:	NM_01222	nucleotide	NUBP2,CF	-1.49434	7.71E-06	0.66919	2127.581	1420.262
A_23_P21:	NM_00394	Homo sapi	ATP6V0E,I	-1.49435	2.82E-17	0.66919	13365.79	8937.855
A_23_P15233		Unknown	A_23_P15:	-1.49455	2.90E-08	0.6691	3000.169	2013.199
A_23_P38:	NM_01910	DNA methy	DMAP1,DI	-1.49508	0.00063	0.66886	324.4376	216.4885
A_23_P52:	NM_00452	Homo sapi	KIF11,EG5	-1.49521	6.33E-11	0.6688	2613.406	1752.926
A_24_P94:	AK024362	Hypothetic	LOC14765	-1.49525	1.07E-13	0.66878	1111.824	743.6737
A_23_P50:	NM_03238	hypothetic	COG8,DOI	-1.49537	3.31E-13	0.66873	4703.738	3154.181
A_23_P16:	NM_02290	hypothetic	SLC30A5,z	-1.49567	5.58E-08	0.6686	1563.175	1049.609
A_32_P11:	BC040651	Isopenteny	IDI2	-1.49567	4.56E-08	0.6686	2314.602	1552.461
A_23_P10:	NM_00453	Homo sapi	NARS,ASN	-1.49573	2.51E-16	0.66857	12397.33	8307.596

A_23_P20	NM_00300	Homo sapi	SEC14L1,5	-1.49578	1.42E-06	0.66855	578.36	386.5787
A_23_P99	NM_13847	hypothetic	MGC5987	-1.49581	3.70E-07	0.66853	3780.023	2549.809
A_23_P30	NM_00041	Homo sapi	HLCS,HCS	-1.49584	3.07E-10	0.66852	935.6932	625.7748
A_23_P36	NM_00214	Homo sapi	HOXB5,HC	-1.49589	3.87E-09	0.6685	1335.635	893.8314
A_24_P33	NM_00121	carbonic ar	CA12,CAX	-1.49675	7.47E-13	0.66811	2137.558	1429.618
A_24_P22	BC042161	Trafficking	TRAPPC5	-1.49676	0.00003	0.66811	11226.57	7432.718
A_23_P33	BC012766	Endoplasm	KIAA1181	-1.49741	8.61E-10	0.66782	10686.34	7157.467
A_32_P87	NM_00591	methionine	MAT2A,MA	-1.49748	1.28E-17	0.66779	55286.07	36879.52
A_32_P60	NM_00410	coagulator	F2RL2,PAI	-1.49749	1.39E-08	0.66778	2043.874	1353.977
A_23_P36	NM_00640	Homo sapi	DCTN2,RE	-1.49799	1.93E-12	0.66756	6105.606	4061.655
A_32_P21	M83234	Human nuc	NSEP	-1.49815	2.43E-14	0.66749	22657.46	15086.39
A_23_P14	NM_00275	Homo sapi	EIF2AK2,P	-1.49827	7.81E-06	0.66744	4543.934	3048.954
A_32_P14	9011	Unknown	A_32_P14	-1.49833	0.00023	0.66741	776.0054	517.613
A_23_P70	NM_05286	hypothetic	MGC21675	-1.4985	4.50E-07	0.66734	995.8611	664.891
A_23_P42	AF161414	Similar to	MGC88387	-1.49852	2.62E-17	0.66732	3339.508	2229.686
A_23_P21	NM_01542	Homo sapi	WDR51A,L	-1.49868	1.61E-19	0.66725	2766.15	1845.05
A_23_P20	NM_03091	Homo sapi	SNX27,MY	-1.49869	4.85E-08	0.66725	959.5991	641.0808
A_24_P46	7449	ALU5_HUM	THC18362	-1.49891	0.00028	0.66715	303.7861	202.9348
A_24_P91	BC011671	Guanine nt	GNB5	-1.49911	0.00054	0.66706	355.8333	237.9553
A_24_P36	NM_00454	NADH dehy	NDUFC2,E	-1.49914	5.13E-07	0.66705	768.6725	513.1554
A_32_P39	AK054902	CDNA FLJ	AK054902	-1.49943	0.00013	0.66692	351.4855	234.2464
A_24_P62	NM_19428	hypothetic	LOC15219	-1.49956	4.58E-06	0.66686	329.3963	219.6618
A_23_P16	NM_01781	Homo sapi	RG9MTD1	-1.4996	1.15E-20	0.66685	11897.71	7931.076
A_23_P33	NM_02112	polymerase	POLR2L,R	-1.50012	8.02E-13	0.66661	7420.974	4960.874
A_23_P11	NM_18280	Arsenate re	ARS2	-1.50014	0.00099	0.6666	567.481	377.522
A_24_P36	NM_03247	mitochondr	MRPS6,RF	-1.50017	3.91E-14	0.66659	4841.419	3226.636
A_32_P19	5924	Unknown	THC19928	-1.50035	1.39E-10	0.66651	1982.172	1321.727
A_23_P38	NM_02076	raptor	raptor,KIA	-1.50044	0.00002	0.66647	3155.454	2124.649
A_24_P32	NM_05304	Homo sapi	EGLN2,EI	-1.50102	3.72E-12	0.66621	667.1416	444.4453
A_23_P20	NM_01341	adenylate	AK3L1,AK	-1.50135	0.00001	0.66607	491.3852	327.0881
A_32_P67	BC067908	CDNA clon	BC067908	-1.50184	8.40E-11	0.66585	779.9356	519.0844
A_24_P48	AK022066	Unc-51-like	ULK4	-1.50221	0.00006	0.66569	813.4232	541.2316
A_23_P40	NM_15227	Homo sapi	FAM58A,M	-1.50229	2.35E-11	0.66565	12131.17	8029.345
A_24_P27	NM_05294	hypothetic	LRRC42,M	-1.50251	0.00059	0.66555	298.5631	198.3993
A_24_P37	NM_00710	solute carri	SLC22A18	-1.50272	3.82E-18	0.66546	12410.52	8250.128
A_23_P20	NM_00271	Homo sapi	PPP2R5C,	-1.50277	2.81E-16	0.66544	4045.994	2691.979
A_23_P45	NM_01230	Homo sapi	EPB41L3,4	-1.5035	3.36E-14	0.66511	2369.455	1575.948
A_32_P87	32	Unknown	THC18707	-1.50366	5.40E-11	0.66504	995.7799	661.6406
A_23_P34	NM_00669	jumping tra	JTB,PAR,h	-1.50383	1.35E-12	0.66497	36628.76	24270.95
A_23_P21	NM_00472	Ac-like tran	ZBED1,AL	-1.50393	6.55E-08	0.66492	928.5645	617.9867
A_32_P26	956	Unknown	A_32_P26	-1.50403	0.0001	0.66488	1588.434	1052.343
A_24_P17	AB007891	Homo sapi	KIAA0431	-1.50432	0.00032	0.66475	849.199	566.2645
A_23_P78	NM_01452	Homo sapi	MYBBP1A,	-1.50455	7.57E-09	0.66465	2563.753	1703.973
A_23_P20	NM_00191	Homo sapi	CYC1	-1.5053	2.80E-22	0.66432	5293.413	3516.209
A_24_P19	AF333487	Kazal-type	KAZALD1	-1.50536	3.62E-17	0.66429	10212.4	6770.937
A_24_P24	AB028949	Kazrin	KIAA1026	-1.50664	0.00072	0.66373	475.0431	314.8248
A_23_P25	NM_00166	ADP-ribosy	ARF5	-1.5069	7.05E-06	0.66361	1453.736	958.0612
A_23_P15	NM_01563	DKFZP586	EIF2B4,EIF	-1.50712	5.52E-12	0.66352	1242.207	823.3709
A_23_P68	NM_03211	hypothetic	WDR54,FL	-1.50759	5.15E-08	0.66331	3720.276	2485.919
A_23_P95	NM_01646	hypothetic	ANKRD39,	-1.50769	2.26E-14	0.66326	844.3137	560.0452
A_23_P60	NM_00141	Homo sapi	EIF4B,EIF-	-1.50789	4.07E-14	0.66318	1714.518	1135.696
A_24_P30	NM_00349	stannin	SNN	-1.508	0.00004	0.66313	1015.617	679.5656

A_23_P15	NM_03342	Homo sapi	C20orf161,	-1.5081	4.41E-06	0.66309	1630.836	1076.489
A_23_P11	AK129920	Son of sev	SOS2	-1.50817	0.00005	0.66305	504.7969	334.2775
A_24_P12	NM_17070	Homo sapi	ICMT,PCM	-1.50863	2.55E-10	0.66285	2083.824	1383.07
A_23_P15	NM_00710	signal seq	SSR3,TRA	-1.50884	3.03E-08	0.66276	6744.877	4510.657
A_23_P24	NM_00007	Homo sapi	CDK4,CMM	-1.50896	3.67E-17	0.66271	8819.059	5850.523
A_24_P91	AK001134	Hypothetic	PCDH17	-1.50957	3.07E-06	0.66244	499.2452	330.4786
A_23_P41	AF151783	Chromosom	C9orf88	-1.50957	6.23E-18	0.66244	24956.21	16521.43
A_23_P15	NM_00727	Prohibitin	2 REA	-1.50959	4.51E-09	0.66243	65067.76	42700.99
A_23_P11	NM_00693	Homo sapi	SUMO2,H	-1.50974	4.45E-21	0.66237	29852.83	19762.95
A_23_P42	NM_01806	Homo sapi	C6orf166,F	-1.50984	5.69E-13	0.66232	7883.479	5208.969
A_32_P11	NM_17851	hypothetic	FLJ36878	-1.50999	6.77E-07	0.66226	1033.011	685.2963
A_23_P25	AB058721	E1A bindin	EP400	-1.51007	3.05E-08	0.66222	689.4923	456.6191
A_23_P75	NM_00662	Homo sapi	ZMYND11,	-1.51011	5.13E-06	0.6622	423.5128	280.3591
A_23_P16	AK024408	Homo sapi	FLJ14346,I	-1.51043	0.00003	0.66206	701.9845	464.9722
A_24_P37	NM_18279	Homo sapi	TMSL4	-1.51088	4.87E-07	0.66186	227981.9	149013.9
A_23_P27	NM_00484	Homo sapi	IL27RA,CF	-1.51101	0.00001	0.66181	555.5935	367.5317
A_32_P19	AY030238	LQK1 hypc	LQK1	-1.51117	1.01E-16	0.66151	2248.744	1488.673
A_32_P6	2026	Unknown	THC19196	-1.51173	5.04E-08	0.66149	435.4901	288.0635
A_23_P29	NM_00175	carbonyl re	CBR1,CBF	-1.51119	2.37E-16	0.66142	8514.168	5620.749
A_23_P12	NM_01523	likely ortho	CSTF2T,C	-1.51201	0.00005	0.66137	820.5665	540.9009
A_23_P39	NM_02110	thymosin, t	TMSB4X,F	-1.51211	3.18E-23	0.66133	100984.2	66788.42
A_32_P11	7186	Unknown	THC18412	-1.51217	0.00075	0.6613	463.2285	306.0828
A_32_P23	AF132203	Stearoyl-C	SCD	-1.51225	2.32E-21	0.66127	6220.663	4114.58
A_32_P71	NM_00201	FK506-bin	FKBP4,HB	-1.5124	2.59E-10	0.6612	816.7252	539.7991
A_24_P19	XM_17115	similar to r	LOC25388	-1.51247	0.00027	0.66117	491.929	324.6376
A_24_P34	NM_00471	synaptogyr	SYNGR2	-1.51267	2.88E-06	0.66108	2210.227	1467.876
A_32_P18	4092	Unknown	A_32_P18	-1.51272	0.00003	0.66106	656.0686	433.8607
A_23_P42	NM_01446	Homo sapi	LSM3,SM	-1.51272	2.10E-13	0.66106	13010.68	8641.69
A_23_P25	NM_03081	Homo sapi	MRPS26,C	-1.51289	3.51E-16	0.66098	20512.02	13577.22
A_23_P74	NM_01806	Homo sapi	IARS2,FLJ	-1.51324	4.92E-16	0.66083	20231.54	13332.02
A_24_P27	NM_00737	TAR DNA I	TARDBP,T	-1.51326	1.44E-17	0.66082	3935.774	2597.737
A_23_P99	NM_03327	Homo sapi	BTBD6,BD	-1.51329	1.31E-12	0.66081	19112.78	12614.68
A_23_P54	NM_01433	CREBBP/E	CR1,EID1	-1.51331	2.56E-13	0.6608	34450.96	22838.01
A_23_P14	NM_00432	Homo sapi	BMPR1A,A	-1.51361	9.35E-07	0.66067	728.9262	481.9085
A_23_P20	NM_02003	Homo sapi	ACCN2,AS	-1.51369	1.22E-07	0.66064	672.2165	443.9337
A_23_P13	NM_00495	Homo sapi	SLC29A1,E	-1.51391	3.88E-13	0.66054	2533.852	1673.861
A_24_P24	5246	U85245 ph	THC18059	-1.51408	3.83E-21	0.66047	3745.963	2473.821
A_32_P69	NM_02520	FAD-synth	FLAD1,FAI	-1.51416	1.83E-17	0.66043	1191.299	786.8801
A_23_P38	NM_00416	RAB1A, m	RAB1A	-1.51426	2.47E-21	0.66039	7547.694	4982.167
A_24_P29	NM_20735	hypothetic	LOC33944	-1.51435	2.61E-20	0.66035	7295.869	4816.138
A_32_P20	BC040051	U2-associ	SR140	-1.51459	3.35E-14	0.66024	1239.857	817.902
A_23_P46	NM_02236	MMS19 (M	MMS19L,M	-1.51465	7.83E-11	0.66022	3509.887	2331.28
A_23_P11	NM_13924	methionyl-t	MTFMT,FM	-1.51465	8.20E-15	0.66022	1427.341	942.6278
A_24_P26	AJ420430	Full length	AJ420430	-1.51479	1.41E-20	0.66016	2485.643	1641.563
A_23_P88	NM_00008	Homo sapi	CLN3,BTS	-1.51501	1.33E-11	0.66006	3052.673	2011.67
A_24_P12	NM_01399	discoidin d	DDR1,CAK	-1.51506	7.72E-21	0.66004	5958.433	3934.701
A_24_P53	0690	Unknown	A_24_P53	-1.51518	8.99E-13	0.65999	1065.239	703.3169
A_23_P11	NM_01813	Homo sapi	C10orf3,FL	-1.51573	3.71E-11	0.65975	7641.647	5059.703
A_32_P20	NM_18170	Quiescin	QQSCN6L1	-1.51575	1.57E-18	0.65974	1236.37	815.7343
A_24_P21	NM_01632	chemokine	CKLF,C32,	-1.51587	1.88E-09	0.65969	3050.385	2019.587
A_24_P39	XM_37296	Homo sapi	ENST0000	-1.51623	2.85E-22	0.65953	11293.1	7450.387
A_24_P18	AL713644	Neurofilam	NEFL	-1.51625	1.09E-11	0.65952	36963.54	24257.16

A_23_P20	NM_02440	Homo sapi	NOTCH2, h	-1.51635	0.00006	0.65948	5534.666	3614.345
A_32_P13	NM_13927	Homo sapi	AKAP1, AK	-1.51721	8.36E-10	0.6591	10126.64	6700.487
A_32_P10	NM_15279	hypothetic	Ells1, FLJ2	-1.51748	8.81E-06	0.65899	603.1057	397.0363
A_24_P73	BC013276	Homo sapi	BC013276	-1.51775	2.26E-12	0.65887	1004.362	662.5018
A_23_P35	NM_17235	Homo sapi	MCP, TLX, C	-1.51811	5.09E-07	0.65871	616.5984	406.2426
A_24_P26	NM_01560	lamina-ass	TOR1AIP1	-1.51848	1.06E-07	0.65855	502.2726	330.6745
A_23_P72	AK000681	Hypothetic	FLJ20674	-1.51852	2.62E-10	0.65854	646.5678	425.8348
A_23_P65	NM_00240	Homo sapi	MGAT2, G	-1.51853	4.02E-14	0.65853	5987.272	3957.653
A_23_P94	NM_01820	Homo sapi	TBC1D13, I	-1.51873	1.05E-19	0.65844	13219.21	8706.693
A_23_P25	NM_00568	Homo sapi	ABCC5, MF	-1.51885	1.60E-09	0.65839	1469.487	971.1887
A_23_P13	NM_00641	ribonuclea	RPP30, TS	-1.51887	1.36E-06	0.65839	2437.892	1619.909
A_23_P25	NM_01786	hypothetic	TMEM70, F	-1.51953	1.03E-09	0.6581	10730.84	7017.525
A_32_P33	NM_20747	FLJ27365	FLJ27365	-1.52025	1.83E-07	0.65779	910.0346	596.6726
A_23_P16	NM_01821	Chromosor	C20orf31	-1.52044	0.00004	0.6577	427.8596	281.7481
A_23_P25	NM_01504	KIAA0676	KIAA0676	-1.52047	7.75E-10	0.65769	5337.414	3522.493
A_23_P13	NM_02183	Homo sapi	PEO1, PEC	-1.5205	1.14E-15	0.65768	1017.367	668.9349
A_23_P21	NM_01647	scotin	SCOTIN	-1.52096	5.48E-12	0.65748	62086.17	40994.07
A_32_P187458	Unknown	A_32_P18		-1.5212	9.89E-10	0.65738	1174.526	772.3044
A_23_P32426	Unknown	THC19925		-1.52123	0.00019	0.65736	304.6811	200.0885
A_23_P99	AK074057	Hypothetic	FLJ10357	-1.52125	9.31E-09	0.65736	7113.644	4641.846
A_24_P36	NM_01889	protocadhe	PCDHAC1	-1.52127	7.22E-08	0.65735	796.8307	524.6149
A_23_P13	U35612	Homo sapi	SOX22	-1.52127	0.00081	0.65735	317.9077	208.5941
A_23_P34	NM_00455	nuclease s	YBX1, YB1	-1.52146	7.06E-15	0.65726	90082.59	59096.87
A_23_P10	NM_13046	Homo sapi	D4ST1, HD	-1.52174	3.04E-11	0.65714	4418.82	2903.917
A_23_P53	NM_00012	Homo sapi	ERCC3, XF	-1.52225	7.44E-09	0.65692	967.519	634.864
A_24_P66932	Unknown	ENST0000		-1.52257	0.00001	0.65678	2010.182	1330.356
A_23_P71	NM_00615	Neurofilam	NEFL	-1.52266	1.57E-11	0.65674	4712.002	3100.108
A_24_P24	AF060925	Hematolog	HN1	-1.52274	0.00002	0.65671	2009.755	1335.155
A_24_P14	NM_00414	NADH deh	NDUFB7, B	-1.52293	1.48E-09	0.65663	9452.479	6242.475
A_23_P42	NM_15334	hypothetic	FLJ90586	-1.52294	3.09E-07	0.65662	1596.564	1043.368
A_24_P118361	Unknown	A_24_P11		-1.52334	1.56E-16	0.65645	867.001	569.1533
A_24_P14	NM_02471	Chromosor	C9orf86	-1.52339	1.21E-13	0.65643	5404.434	3551.065
A_23_P81	NM_00573	conserved	CTDSP2, C	-1.52351	3.81E-17	0.65638	5912.821	3889.273
A_24_P23	NM_13839	Homo sapi	HNRPLL, S	-1.52362	0.00004	0.65633	2375.83	1571.034
A_23_P65	BX537558	PMS2 post	PMS2	-1.52377	4.84E-08	0.65627	1102.751	725.9367
A_24_P22	NM_00440	DNA fragm	DFFA, DFF	-1.52394	1.68E-09	0.65619	2002.469	1316.58
A_24_P39	NM_20749	Homo sapi	FLJ44896	-1.52406	1.07E-09	0.65614	674.3428	442.295
A_32_P75	NM_15227	Homo sapi	FAM58A, M	-1.52409	1.43E-15	0.65613	7488.698	4927.884
A_24_P680548	Unknown	A_24_P68		-1.52446	9.96E-22	0.65597	55444.16	36384.4
A_32_P10	AK096022	Hypothetic	LOC28602	-1.5249	0.00063	0.65578	242.0475	158.8477
A_23_P69	NM_00288	Homo sapi	RARS, ArgI	-1.52543	1.79E-10	0.65555	63780.59	41444.73
A_24_P18	NM_00271	protein phc	PPP2CA, R	-1.5256	1.39E-17	0.65548	7727.491	5071.638
A_23_P32	NM_01903	Homo sapi	EXOSC4, S	-1.52572	3.36E-14	0.65543	4600.014	3007.886
A_23_P88	NM_01604	CGI-112 pr	C14orf122,	-1.52601	2.23E-10	0.6553	4082.152	2659.486
A_24_P48	NM_00243	N-methylpt	MPG, AAG,	-1.52604	8.14E-09	0.65529	4496.828	2922.394
A_24_P92	NM_02121	transposon	LOC58486	-1.52615	1.39E-07	0.65524	1973.864	1299.74
A_23_P92	AK025789	Homo sapi	RFWD2, C	-1.52617	2.74E-22	0.65523	4802.156	3146.292
A_23_P13	NM_00373	aldo-keto r	AKR1C3, D	-1.52646	1.57E-19	0.65511	152463	99760.73
A_23_P52	NM_01450	Homo sapi	PRPF19, P	-1.5265	2.43E-10	0.65509	733.4563	480.2167
A_24_P35	NM_01642	zinc finger	ZNF219, ZF	-1.52687	8.01E-07	0.65494	3451.399	2235.554
A_24_P19	NM_01494	KIAA0911	CLSTN1, C	-1.5273	5.62E-14	0.65475	1520.324	994.0369
A_23_P14	NM_00396	ubiquitin-cc	UBE2M, UE	-1.52738	5.01E-06	0.65471	2504.033	1626.374

A_23_P21	AL833042 Mitogen-act	MAPKAP1	-1.52746	0.00029	0.65468	248.5306	162.6668
A_23_P42	NM_20346 Homo sapi	MRFAP1L	-1.52758	2.23E-09	0.65463	3722.473	2442.862
A_23_P11	NM_00196 eukaryotic	EIF4E,CBF	-1.528	3.72E-08	0.65445	2804.313	1842.582
A_24_P25	NM_00633 C2f protein	C2F	-1.52819	2.04E-22	0.65437	10208.42	6677.172
A_23_P37	NM_00121 Homo sapi	CA12,CAX	-1.5283	6.55E-11	0.65432	17260.29	11347.33
A_23_P12	NM_19942 Homo sapi	ZFP64,MG	-1.52865	7.02E-17	0.65417	4258.647	2793.089
A_23_P73	NM_00156 Homo sapi	IRAK1,IRA	-1.52868	1.32E-13	0.65416	7113.6	4647.029
A_23_P10	NM_02006 Homo sapi	SLC2A4RC	-1.52875	2.37E-16	0.65413	24318.63	15868.17
A_23_P72	NM_01792 nuclear prc	NPL4,FLJ2	-1.52878	1.64E-16	0.65411	2113.661	1381.81
A_24_P27	AK057015 Heat shock	HSPA4	-1.5288	4.19E-12	0.65411	1072.818	701.8733
A_23_P42	NM_01806 hypothetic	C6orf166,F	-1.52881	3.91E-06	0.6541	1177.057	772.2858
A_23_P40	BC022374 PHD finger	PHF19	-1.52881	2.22E-11	0.6541	7489.905	4926.053
A_23_P11	NM_00116 Homo sapi	BIRC5,API	-1.52964	2.49E-13	0.65375	27458.13	17911.79
A_24_P38	NM_14478 Homo sapi	DEGS1,ML	-1.53018	9.55E-09	0.65352	3200.022	2089.649
A_23_P48	NM_02405 hypothetic	MGC5576	-1.53031	4.33E-17	0.65346	27947.91	18216.97
A_32_P23	BC007205 Homo sapi	BC007205	-1.53048	6.14E-07	0.65339	3629.391	2368.713
A_23_P17	NM_00062 Homo sapi	IFNAR1,A	-1.53092	6.97E-08	0.6532	647.7289	422.7882
A_23_P41	NM_01332 Homo sapi	SNX8	-1.53094	4.85E-11	0.65319	632.9235	413.4034
A_23_P13	BC003515 Chromosom	C6orf129	-1.53096	1.46E-10	0.65319	1848.304	1209.044
A_24_P38	NM_13079 Nucleolar	rNOL6	-1.53132	1.52E-11	0.65303	3277.65	2144.366
A_23_P60	NM_00307 SWI/SNF r	SMARCA2	-1.53137	5.10E-10	0.65301	738.822	482.4507
A_23_P17	NM_00327 Homo sapi	TSPAN6,T	-1.53139	2.53E-15	0.653	3394.011	2222.307
A_23_P10	AK098212 Protein BA	FLJ10359	-1.53139	5.41E-15	0.653	9053.732	5919.746
A_24_P39	NM_01867 smooth mu	SMAP-1,FL	-1.53141	3.13E-11	0.65299	2179.863	1421.106
A_24_P94	AF113007 DKFZP586	DKFZP586	-1.53141	1.35E-19	0.65299	1165.557	761.1012
A_23_P21	NM_01432 Homo sapi	ZNF278,ZS	-1.53153	1.85E-07	0.65294	579.5297	378.324
A_24_P39	NM_00679 MORF-rela	MORF4L1,	-1.53171	5.94E-23	0.65287	57630.02	37639.97
A_23_P14	NM_00586 Homo sapi	OPRS1,SF	-1.53179	5.16E-18	0.65283	9187.665	5988.727
A_32_P70	AK098629 Cyclin M3	CNNM3	-1.53206	1.08E-07	0.65272	979.9879	638.4961
A_23_P10	NM_03247 Homo sapi	MRPS6,RF	-1.53237	5.84E-11	0.65259	30318.09	19667.68
A_24_P27	NM_01871 calreticulin	CDCA7L,R	-1.53312	1.82E-06	0.65227	6397.312	4174.461
A_24_P530977	Unknown	A_24_P530	-1.53345	2.59E-14	0.65212	57460.96	37348.55
A_23_P49	NM_01901 Homo sapi	FAM64A,F	-1.5336	2.18E-21	0.65206	4474.136	2912.662
A_24_P78	AK092572 Hypothetic	LOC14841	-1.53394	0.00031	0.65191	428.6486	279.4935
A_32_P140262	Unknown	A_32_P140	-1.53436	6.33E-12	0.65174	5187.915	3375.946
A_23_P76	NM_00176 Homo sapi	CD9,BA2,F	-1.5346	2.36E-14	0.65164	862.5508	561.9407
A_23_P11	NM_01605 CGI-120 pr	COPZ1,CC	-1.5347	7.95E-06	0.6516	2068.578	1333.058
A_32_P20	AK023412 CDNA FLJ	AK023412	-1.53485	2.20E-06	0.65153	549.1014	357.6417
A_24_P30	NM_00459 spermine s	SMS,SRS,	-1.53486	1.85E-23	0.65153	16536.71	10771.46
A_24_P26	NM_00433 chromosom	C8orf1,hT4	-1.53495	3.10E-06	0.65149	1056.268	691.2404
A_24_P118813	Unknown	A_24_P118	-1.53517	4.42E-10	0.65139	2914.38	1887.015
A_24_P70	BX537688 Homo sapi	DKFZp686	-1.5352	0.00075	0.65138	297.1651	192.9506
A_24_P56	NM_00134 defender a	DAD1	-1.53532	1.57E-11	0.65133	19209.51	12565.07
A_24_P33	NM_02404 hypothetic	METRN,M	-1.53533	1.35E-06	0.65132	5063.382	3267.707
A_23_P15	NM_01795 FLJ20758	FLJ20758	-1.53536	2.87E-11	0.65131	9917.699	6424.755
A_23_P11	NM_02162 Homo sapi	PLEKHA1,	-1.53547	4.75E-06	0.65126	1963.352	1281.436
A_24_P14	NM_02110 thymosin, t	TMSB4X,F	-1.5355	1.29E-21	0.65126	99332.98	64634.73
A_23_P14	NM_00710 Homo sapi	UBL3,HCC	-1.53567	8.34E-07	0.65118	2400.156	1572.136
A_23_P16	NM_00532 Homo sapi	HADHSC,S	-1.5357	1.06E-11	0.65117	11604.71	7594.757
A_23_P10	NM_02452 Homo sapi	TTC13,FLJ	-1.53593	5.40E-10	0.65107	926.395	603.468
A_24_P840868	Unknown	A_24_P840	-1.53609	1.49E-16	0.65101	76332.08	49849.82
A_23_P16	NM_00568 Homo sapi	TAF1B,SL	-1.53612	0.00001	0.65099	1121.886	729.9764

A_23_P14	NM_00717 protein-O-r POMT1,RT	-1.5364	1.24E-10	0.65087	3866.135	2517.3
A_23_P12	NM_01823 Homo sapi NUP133,hf	-1.53661	6.33E-15	0.65078	3795.005	2478.228
A_23_P12	NM_01426 Homo sapi HKE2,KE-2	-1.53695	1.61E-21	0.65064	6579.984	4277.144
A_23_P50	NM_02053 Homo sapi CSRP2BP,	-1.53698	8.39E-13	0.65063	2717.843	1771.356
A_23_P32	NM_05282 Homo sapi WDR5,BIG	-1.53698	1.00E-07	0.65063	1255.37	812.1544
A_24_P71	AK098433 Hypothetic LOC28410	-1.53756	6.76E-16	0.65038	13507.79	8760.078
A_24_P85	2099 Unknown A_24_P85	-1.53808	1.70E-10	0.65016	103042.3	66435.15
A_24_P20	NM_03310 prostein pr SLC45A3,F	-1.53887	1.93E-15	0.64983	1287.174	835.8102
A_23_P33	NM_17851 hypothetic FLJ36878	-1.53887	0.001	0.64983	473.4501	306.7286
A_32_P46	AK090664 Interleukin IL17D	-1.53891	1.09E-12	0.64981	662.4054	430.4016
A_24_P31	NM_00659 API5-like 1 API5,AAC-	-1.53941	1.82E-17	0.6496	1146.205	744.4408
A_32_P11	AK092180 Homo sapi LOC44124	-1.54002	2.33E-16	0.64934	15907.32	10357.09
A_23_P47	NM_00618 Homo sapi OAS3,p10	-1.54004	3.51E-07	0.64933	1351.753	876.5657
A_24_P21	BC001097 Eukaryotic EIF4B	-1.54027	1.11E-14	0.64924	4293.821	2781.374
A_23_P12	NM_00608 tubulin, alp K-ALPHA-	-1.54098	7.19E-18	0.64894	392266.4	255153.4
A_23_P50	NM_00392 methyl-Cp MBD3	-1.5418	2.17E-18	0.64859	3565.011	2314.781
A_23_P63	NM_02403 Homo sapi LRFN4,MC	-1.54183	1.99E-07	0.64858	7531.223	4867.048
A_23_P11	NM_00196 eukaryotic EIF4E,CBF	-1.5422	1.61E-20	0.64843	10537.9	6825.976
A_23_P11	NM_01516 Homo sapi ARL6IP,All	-1.54225	1.20E-22	0.6484	8205.125	5322.198
A_24_P27	XM_29301 similar to F LOC34433	-1.54225	7.62E-12	0.6484	1344.205	872.8895
A_23_P25	NM_00727 Homo sapi TUSC2,PA	-1.54244	1.89E-11	0.64832	6711.751	4345.498
A_23_P74	NM_00313 Homo sapi SRM,PAP1	-1.54248	1.38E-10	0.64831	8933.396	5780.082
A_23_P19	NM_00633 lysophosph LYPLA1,LF	-1.54254	1.97E-06	0.64828	3146.581	2066.419
A_23_P13	NM_00670 Homo sapi TXNL4A,D	-1.54261	2.74E-12	0.64825	29778.71	19195.39
A_23_P21	NM_01812 Homo sapi TMEM33,F	-1.543	1.27E-18	0.64809	7466.842	4829.753
A_24_P41	NM_00578 Not56 (D. r ALG3,CDC	-1.54334	2.13E-09	0.64795	10096.09	6517.857
A_23_P33	NM_01565 DKFZP564 NOC2L,DK	-1.54355	4.63E-12	0.64786	718.7026	465.6644
A_23_P21	AK092776 Hypothetic MGC16471	-1.54381	1.27E-19	0.64775	3958.525	2567.541
A_23_P11	NM_03194 Homo sapi CENTG3,A	-1.54449	2.82E-15	0.64746	9936.881	6417.563
A_32_P23	NM_01208 ATP-bindin ABCB10,M	-1.54454	0.00007	0.64744	622.0363	402.6053
A_23_P61	NM_00040 galactosid GLB1	-1.54481	1.42E-21	0.64733	7231.549	4687.601
A_32_P23	NM_01812 hypothetic ARMC1,Ar	-1.54484	3.88E-13	0.64732	3172.252	2058.386
A_23_P34	NM_02520 FAD-synth FLAD1,FAI	-1.54504	7.38E-08	0.64723	4032.94	2600.333
A_23_P35	NM_01597 Homo sapi SNX7,MGC	-1.54513	2.91E-15	0.6472	3936.473	2556.268
A_32_P31	BC035691 Glutathione GSR	-1.54603	2.00E-08	0.64682	2005.176	1285.961
A_32_P43	NM_02407 hypothetic DDX54,DP	-1.54695	3.88E-15	0.64643	36433.16	23585.32
A_23_P20	NM_05305 Homo sapi CCND1,BC	-1.54703	2.33E-12	0.6464	3131.715	2023.728
A_23_P62	NM_01239 Homo sapi PHLDA3,T	-1.54762	4.48E-10	0.64615	3421.112	2207.809
A_32_P19	7899 Unknown A_32_P19	-1.54791	2.71E-06	0.64603	1388.124	898.4457
A_23_P13	NM_03148 Homo sapi MARVELD	-1.54842	1.61E-11	0.64582	16884.48	10853.36
A_23_P67	NM_02507 Homo sapi UXS1,FLJ2	-1.54877	1.75E-09	0.64567	2781.166	1800.543
A_23_P13	NM_15293 Homo sapi GLT8D1,Al	-1.54893	1.86E-11	0.64561	3920.186	2545.406
A_23_P10	NM_00693 Homo sapi SUMO3,SM	-1.54932	1.43E-19	0.64544	2435.853	1573.493
A_23_P66	NM_00227 Homo sapi KRT19,K19	-1.54961	2.08E-08	0.64532	60792.98	38943.11
A_24_P13	AK090397 Hypothetic MGC15875	-1.54975	1.40E-19	0.64527	1425.176	919.7593
A_23_P14	NM_00364 Homo sapi SLC27A2,\	-1.54993	2.21E-15	0.64519	4790.884	3081.903
A_23_P56	NM_00215 Homo sapi HSPE1,CP	-1.55005	5.44E-15	0.64514	268359.9	172320.8
A_23_P61	NM_01233 Homo sapi MSRA	-1.5501	2.00E-19	0.64512	1548.884	999.167
A_23_P41	NM_00132 C-terminal CTBP1,MC	-1.55066	1.70E-07	0.64489	1596.033	1030.353
A_32_P34	365 Unknown A_32_P34	-1.55111	7.28E-08	0.6447	428.1437	275.951
A_23_P35	NM_15234 hypothetic FLJ33817	-1.55134	7.84E-16	0.6446	2743.605	1768.09
A_23_P34	NM_01612 COP9 (con COPS4,MC	-1.55165	2.94E-24	0.64448	6608.43	4259.79

A_23_P39	NM_15326	hypothetic	FLJ38101,	-1.55165	2.49E-13	0.64447	1321.833	853.3542
A_32_P10	AK130366	Similar to h	AK130366	-1.55223	6.81E-07	0.64423	472.5023	304.2144
A_24_P40	NM_00632	progesterone	PGRMC2,	-1.55238	4.37E-24	0.64417	14103.45	9081.535
A_23_P30	NM_17370	Homo sapi	NM_17370	-1.55244	7.81E-19	0.64415	165552.6	106422.9
A_23_P40	NM_13868	Homo sapi	PIP5K2B,F	-1.55251	0.00059	0.64412	461.651	295.9236
A_23_P32	NM_01546	Homo sapi	C2orf32,D	-1.55255	0.00002	0.6441	541.7922	348.5602
A_24_P89	NM_00088	IMP (inosin	IMPDH1,IM	-1.55273	2.67E-10	0.64403	25395	16407.88
A_24_P18	NM_18129	Homo sapi	WDR20,DM	-1.55276	2.09E-10	0.64401	682.2271	439.4946
A_23_P11	NM_00257	Homo sapi	PRDX1,PA	-1.55322	5.74E-15	0.64382	153498	98374.13
A_23_P20	NM_03057	outer mitoc	CYB5-M,O	-1.55337	3.34E-14	0.64376	1192.361	767.8427
A_24_P21	XM_37661	Homo sapi	ENST0000	-1.55347	4.30E-20	0.64372	12172.41	7847.094
A_32_P67	NM_00416	succinate c	SDHA,FP,	-1.55383	5.12E-10	0.64357	7091.164	4587.141
A_23_P60	NM_02122	Zinc finger	ZNF462	-1.55394	1.13E-09	0.64353	1810.909	1170.667
A_24_P41	NM_01504	Golgi-asso	GGA2,VEA	-1.55396	1.41E-10	0.64352	869.9683	559.5426
A_24_P92	AK056260	LOC44075	AK056260	-1.55401	3.96E-06	0.6435	1248.202	800.6543
A_23_P49	NM_00692	splicing fac	SFRS1,AS	-1.55415	7.41E-08	0.64344	2631.136	1699.63
A_23_P76	NM_00598	Homo sapi	SIX1	-1.5542	2.30E-17	0.64342	4224.662	2719.866
A_23_P15	NM_00588	Homo sapi	DLEU1,BC	-1.55429	7.29E-12	0.64338	9145.692	5877.327
A_23_P20	NM_01208	Homo sapi	ABCB10,M	-1.55454	1.02E-07	0.64328	2830.905	1832.114
A_24_P127362		Unknown	A_24_P12	-1.55465	2.92E-12	0.64323	986.6144	634.1252
A_24_P135933		Unknown	ENST0000	-1.55467	0.00013	0.64323	789.5481	509.1154
A_24_P93	AK021980	Transcripti	TCF4	-1.55476	0.00004	0.64319	1033.347	666.5801
A_24_P33	NM_00411	fibroblast g	FGF12,FH	-1.55536	0.00015	0.64294	343.2061	220.8523
A_23_P88	NM_00252	nth (E.coli	INTHL1,NT	-1.55544	9.82E-13	0.6429	5545.958	3549.21
A_23_P41	NM_00408	Homo sapi	DLG1,hd g	-1.55558	1.25E-07	0.64285	574.1536	369.0342
A_24_P29	XM_37231	Homo sapi	ENST0000	-1.55588	5.23E-12	0.64272	10233.65	6605.845
A_23_P32	NM_01884	Homo sapi	KLHL7,KL	-1.55594	2.07E-12	0.6427	562.7161	361.6596
A_24_P39	NM_17850	NS5ATP13	NS5ATP13	-1.55632	6.13E-08	0.64254	693.7556	445.4661
A_23_P21	NM_13333	Homo sapi	WHSC1,N	-1.55688	7.37E-11	0.64231	2316.15	1488.231
A_24_P227350		GDFB_HU	THC19778	-1.55762	0.00001	0.64201	288.1962	184.9889
A_23_P11	NM_01211	Homo sapi	AHSA1,p3	-1.55777	6.45E-08	0.64194	10969.27	6965.835
A_23_P11	NM_01495	Homo sapi	BAHD1,KI	-1.55819	1.28E-16	0.64177	3970.114	2550.819
A_24_P45	AK024653	Hypothetic	LOC90246	-1.55835	0.00001	0.6417	1285.438	823.6099
A_23_P10	NM_01326	Homo sapi	C11orf2,A	-1.55842	1.12E-12	0.64167	2679.307	1726.423
A_23_P15	NM_03264	likely ortho	SPSB2,SS	-1.55843	3.85E-15	0.64167	2831.479	1816.463
A_23_P15	NM_00087	insulin-like	IGF2R,MPI	-1.56049	0.00004	0.64083	2010.39	1292.667
A_23_P33	NM_15336	Homo sapi	C10orf56,F	-1.56061	5.17E-08	0.64078	1144.229	736.4049
A_24_P610945		Unknown	ENST0000	-1.56074	1.80E-09	0.64072	2085.751	1332.369
A_24_P20	AK054960	Homo sapi	APBB3	-1.56141	1.90E-15	0.64045	977.5592	625.7362
A_24_P11	NM_00642	CD27-bind	SIVA,CD27	-1.56184	1.55E-09	0.64027	41795.87	26486.66
A_24_P67	NM_00583	serine/argi	SRRM1,16	-1.56212	9.18E-08	0.64016	1326.466	852.0892
A_23_P10	NM_00134	Homo sapi	DAD1	-1.56318	9.41E-19	0.63972	40518.75	25972.79
A_23_P74	NM_03275	hypothetic	GLOXD1,M	-1.56357	2.34E-14	0.63956	20321.24	12960.06
A_32_P15	NM_20741	Similar to F	MGC57827	-1.56378	2.00E-23	0.63948	2555.668	1634.667
A_32_P13991		Unknown	THC18823	-1.56482	7.35E-09	0.63905	992.7398	635.3904
A_23_P25	NM_00664	Homo sapi	SDCCAG3	-1.56541	7.76E-20	0.63881	10458.54	6673.252
A_23_P42	NM_14505	Homo sapi	FUK,FLJ3	-1.56569	2.20E-09	0.6387	719.2615	459.6393
A_23_P31	NM_18171	Homo sapi	PRR6,p30,	-1.56577	5.46E-13	0.63866	1031.959	659.4431
A_23_P21	NM_01343	Homo sapi	NFKBIL2,II	-1.56647	1.14E-08	0.63838	552.4336	352.6407
A_24_P409750		Unknown	A_24_P40	-1.56658	3.84E-11	0.63833	1508.018	965.1279
A_23_P94	NM_03142	Homo sapi	C9orf58,IB	-1.56675	2.23E-22	0.63826	3836.137	2447.319
A_24_P92	BC019824	Homo sapi	BC019824	-1.56677	2.55E-10	0.63826	1675.034	1071.815

A_23_P10	NM_01900	hypothetic	FLJ20232,	-1.56717	6.54E-27	0.63809	7385.926	4712.922
A_32_P70	NM_02513	hypothetic	C13orf23,F	-1.56738	3.82E-18	0.63801	2409.96	1540.584
A_24_P923684	AF083108	THC18900		-1.56758	6.90E-07	0.63793	555.1622	353.9605
A_24_P33	NM_00693	SMT3 (sup	SUMO2,H	-1.56788	4.65E-24	0.6378	41297.23	26327.06
A_23_P25	NM_01223	SREBP cle	SCAP	-1.56828	1.07E-10	0.63764	6180.965	3930.766
A_23_P71	NM_00426	Homo sapi	RECQL4,R	-1.56862	6.71E-17	0.6375	2554.333	1629.848
A_23_P16	NM_01553	Homo sapi	GORASP2	-1.5689	2.80E-11	0.63739	3130.272	2006.104
A_23_P35	NM_18261	hypothetic	FLJ34283,	-1.56904	6.26E-25	0.63733	10491.64	6683.331
A_23_P11	NM_00188	Homo sapi	CRYZ	-1.56911	6.40E-06	0.63731	2071.298	1328.271
A_24_P34	BC000147	Malic enzym	ME2	-1.56913	2.91E-08	0.6373	1287.819	822.3301
A_24_P24	NM_00637	vesicle-ass	VT11B,VT1	-1.56919	9.78E-20	0.63727	1993.212	1270.962
A_24_P29	NM_01514	PAS-serine	PASK,STK	-1.5692	0.00001	0.63727	351.2724	224.0758
A_23_P50	NM_00647	Homo sapi	GAS2L1,G	-1.56978	1.15E-14	0.63703	5123.541	3255.708
A_24_P16	NM_14528	similar to h	LOC15119	-1.56993	3.06E-25	0.63697	3898.767	2483.223
A_23_P13	NM_15230	Homo sapi	MRFAP1L	-1.57001	1.27E-08	0.63694	2064.344	1318.474
A_24_P30	NM_00373	Sjogren's s	SSNA1,N1	-1.57025	3.76E-14	0.63684	3423.776	2174.217
A_23_P31	NM_01344	Homo sapi	ST6GALNA	-1.57027	0.00037	0.63683	334.7925	212.6914
A_24_P846810	Unknown	A_24_P84		-1.57041	0.00059	0.63678	544.089	345.4171
A_32_P37	AK096908	CDNA FLJ	AK096908	-1.5706	1.28E-18	0.6367	2273.052	1445.921
A_23_P12	NM_03144	Homo sapi	CCM2,C7c	-1.57123	4.70E-07	0.63644	1690.543	1070.645
A_23_P32	NM_15303	hypothetic	FLJ32065,	-1.57142	7.46E-07	0.63637	748.225	476.1561
A_24_P64	AK124953	Similar to h	AK124953	-1.5716	1.22E-07	0.63629	396.3807	252.1373
A_23_P58	NM_03137	Homo sapi	HNRPD,P3	-1.5717	1.96E-11	0.63625	55532.5	35075.88
A_23_P57	NM_01416	Homo sapi	MKRN2,RN	-1.57182	6.77E-23	0.63621	2290.616	1456.976
A_23_P35	NM_01786	Homo sapi	PIGX,FLJ2	-1.57196	0.00004	0.63615	759.7793	483.8185
A_23_P36	NM_17371	Homo sapi	NM_17371	-1.572	2.88E-16	0.63613	19124.55	12134.59
A_23_P12	NM_01639	p53-inducit	P53CSV,W	-1.57211	6.14E-13	0.63609	4186.525	2646.071
A_23_P13	NM_00359	Homo sapi	TPST2	-1.5722	9.58E-11	0.63605	563.2969	358.2905
A_23_P20	NM_01593	CGI-01 pro	KIAA0859,	-1.57308	1.01E-13	0.6357	1479.704	938.6274
A_23_P44	NM_02075	Homo sapi	CASKIN2,F	-1.57351	1.79E-07	0.63552	378.4934	240.4766
A_24_P324644	Unknown	A_24_P32		-1.57383	5.40E-16	0.63539	1794.608	1141.542
A_23_P15	NM_00276	Homo sapi	PRPSAP1,	-1.57435	5.50E-11	0.63518	3394.779	2169.215
A_23_P21	NM_02067	Homo sapi	ABHD6	-1.57438	4.67E-07	0.63517	3286.941	2084.962
A_23_P53	NM_03057	hypothetic	MGC10993	-1.57438	3.59E-15	0.63517	2574.706	1632.872
A_23_P11	NM_01215	Homo sapi	F8A1,F8A,	-1.57487	7.54E-22	0.63497	9186.043	5830.087
A_23_P55	NM_00067	Homo sapi	ADORA2B	-1.57498	2.97E-13	0.63493	2849.435	1812.885
A_23_P39	NM_00483	Homer, net	HOMER3,F	-1.57566	1.12E-07	0.63465	3240.05	2076.504
A_24_P11	NM_00670	similar to	STXNL4A,D	-1.57593	7.18E-13	0.63455	9483.834	6046.094
A_24_P470809	Unknown	A_24_P47		-1.5763	9.66E-07	0.6344	409.4334	259.5541
A_23_P54	NM_03283	Homo sapi	CIRH1A,N	-1.57631	6.17E-07	0.63439	3434.604	2209.177
A_24_P48	AF086017	Hypothetic	LOC28614	-1.57651	2.86E-08	0.63431	590.9533	374.8252
A_23_P51	NM_01784	tRNA seler	SECP43,P	-1.57656	2.60E-14	0.63429	1600.201	1012.809
A_23_P10	NM_01207	Homo sapi	C16orf35,C	-1.57684	5.80E-19	0.63418	9267.65	5888.554
A_24_P74	NM_00340	zinc finger	ZFP161,ZF	-1.57708	1.28E-06	0.63409	779.8339	494.8652
A_23_P59	NM_00059	Homo sapi	TAP1,APT	-1.57737	8.47E-17	0.63397	6284.218	3978.492
A_24_P548453	QOR_HUM	THC20023		-1.57743	2.24E-07	0.63394	550.8826	349.0497
A_23_P13	NM_00513	Homo sapi	SCO2,SCC	-1.57778	5.06E-21	0.6338	4943.259	3137.455
A_24_P47	NM_00134	defender a	DAD1	-1.5785	1.25E-15	0.63351	36624.66	23312.04
A_23_P12	NM_02015	Homo sapi	STARD7,G	-1.57891	1.45E-15	0.63335	18659.18	11783.6
A_23_P82	NM_00093	Homo sapi	PLAT,TPA,	-1.57898	5.60E-07	0.63332	1236.974	784.8079
A_23_P12	NM_00633	Homo sapi	TIMM17A,	-1.57915	1.70E-21	0.63325	55579.14	35231.75
A_32_P14	NM_00693	SMT3 (sup	SUMO2,H	-1.57924	4.11E-20	0.63322	28058.84	17791.96

A_32_P135489	Unknown	A_32_P135489	-1.5797	0.00063	0.63303	567.9817	358.6906
A_24_P37	NM_18279	Homo sapiens TMSL4	-1.57991	2.17E-20	0.63295	38278.2	24258.8
A_24_P24	NM_01628	adenylate kinase AK3, AK6, F	-1.58106	1.48E-06	0.63249	497.9041	314.5897
A_24_P41	NM_01681	8-oxoguanine OGG1, HML	-1.58142	2.42E-20	0.63234	2515.035	1590.656
A_23_P20	NM_00238	Homo sapiens MCP, TLX, C	-1.58145	1.10E-08	0.63233	1262.833	800.8938
A_24_P19	NM_01812	hypothetical TMEM33, F	-1.58166	4.64E-07	0.63225	580.6303	366.9873
A_23_P32	NM_03091	Homo sapiens C20orf129,	-1.58224	1.80E-13	0.63202	2701.296	1711.655
A_23_P10	NM_00599	Homo sapiens TBCD, KIAA	-1.58263	1.43E-14	0.63186	752.8716	475.76
A_24_P46	AK056550	Syncoilin, isoform 1 SYNCOIL1	-1.5831	1.70E-09	0.63167	1185.999	751.4546
A_32_P58999	VAPA_HUI	THC18449	-1.58318	1.41E-16	0.63164	6006.754	3800.666
A_24_P34	NM_20128	Homo sapiens USP51	-1.58399	9.80E-06	0.63132	347.0684	219.3149
A_24_P50	AK022936	CDNA FLJ AK022936	-1.58443	9.88E-14	0.63114	1700.446	1075.805
A_23_P15	NM_00176	Homo sapiens CCT6A, CC	-1.58457	2.38E-22	0.63109	19398.18	12256.67
A_23_P39	NM_01752	Homo sapiens LY6K, FLJ3	-1.58461	6.73E-17	0.63107	16781.5	10606.38
A_23_P25	NM_00416	RAB1A, member 1 RAB1A	-1.58475	1.10E-25	0.63102	3784.611	2387.288
A_23_P51	BC004501	Hypothetical MGC33867	-1.58526	1.48E-24	0.63081	6827.229	4309.177
A_32_P12	BQ935463	AGENCOL BQ935463	-1.5864	0.00001	0.63036	1149.776	728.2628
A_23_P13	NM_14503	Homo sapiens FAM55C, M	-1.5865	2.19E-26	0.63032	2738.861	1726.257
A_24_P306896	Unknown	A_24_P306896	-1.58696	1.41E-07	0.63014	1012.572	636.7063
A_23_P36	NM_13907	delta-notch DNER	-1.58699	3.91E-14	0.63012	2049.021	1294.25
A_23_P25	NM_00281	Homo sapiens PSMD10, p	-1.58769	1.08E-14	0.62984	10762.7	6758.114
A_24_P32	NM_18270	hypothetical MGC45871	-1.58772	0.00013	0.62983	582.1554	366.1547
A_32_P11	NM_00627	splicing factor SFRS6, B5,	-1.58833	4.83E-09	0.62959	3292.633	2082.954
A_24_P94	AF074331	Homo sapiens PAPSS2	-1.58835	1.94E-08	0.62958	1740.48	1099.175
A_24_P37	NM_01510	Homo sapiens PLXND1, P	-1.58881	4.77E-12	0.6294	1805.417	1137.732
A_24_P25	NM_00416	RAB1A, member 1 RAB1A	-1.58886	3.65E-13	0.62938	3791.912	2396.031
A_23_P42	AL050297	Hypothetical LOC20306	-1.58912	2.22E-18	0.62928	931.1633	585.9016
A_23_P21	NM_02290	Homo sapiens SLC30A5, 2	-1.58983	3.37E-11	0.629	3368.573	2132.006
A_23_P25	NM_01208	Homo sapiens HDHD1A, C	-1.59045	9.46E-06	0.62875	1676.52	1059.637
A_24_P94	D80001	KIAA0179 KIAA0179	-1.59069	7.12E-14	0.62866	2685.863	1690.374
A_24_P30	L40403	YLP motif c YLPM1	-1.59092	7.15E-12	0.62857	779.8997	490.4133
A_23_P30	NM_15226	Homo sapiens RPUSD2, C	-1.59117	1.24E-12	0.62847	7391.569	4647.197
A_23_P11	NM_00655	Homo sapiens PMVK, PMV	-1.59129	4.19E-15	0.62842	5873.029	3676.773
A_32_P57013	Unknown	A_32_P57013	-1.59201	2.78E-12	0.62813	631.7015	396.7558
A_32_P194704	Unknown	A_32_P194704	-1.59296	0.00022	0.62776	1125.963	711.6055
A_23_P12	NM_00467	Homo sapiens ANGPTL1,	-1.59317	0.00042	0.62768	425.0366	265.5614
A_23_P11	NM_01228	Homo sapiens KEAP1, INr	-1.59323	9.03E-22	0.62766	5607.757	3522.917
A_23_P15	NM_01568	B9 protein EPPB9, B9	-1.59331	1.60E-06	0.62762	936.8209	590.0565
A_23_P38	AK092352	Similar to s LOC44207	-1.59348	7.06E-08	0.62756	583.0765	365.8044
A_24_P84112	Unknown	A_24_P84112	-1.5935	3.49E-14	0.62755	25399.13	16001.61
A_23_P20	NM_03286	hypothetical TIGD5, FLJ	-1.59389	2.13E-18	0.6274	2404.989	1506.937
A_23_P16	NM_20738	FLJ43339 FLJ43339, l	-1.59418	8.91E-25	0.62728	3191.427	2000.686
A_24_P64	NM_15263	hypothetical MGC17301	-1.59501	9.75E-07	0.62696	899.1252	563.8505
A_23_P85	NM_00399	CDC-like kinase CLK2, hCLI	-1.59509	1.03E-10	0.62692	671.1211	420.5053
A_24_P367397	Unknown	A_24_P367397	-1.59509	4.68E-11	0.62692	812.7275	509.0264
A_24_P92	NM_01642	cisplatin resistance CROP, LUC	-1.59521	3.26E-06	0.62688	3790.659	2397.028
A_24_P20	U22030	Cytochrome CYP2A7	-1.59578	7.67E-09	0.62665	574.2971	359.8604
A_23_P16	NM_02248	Homo sapiens CENTD3, A	-1.59596	2.73E-15	0.62658	1165.846	730.3115
A_23_P83	NM_02111	Homo sapiens RECK, ST1	-1.59599	2.23E-06	0.62657	585.4509	366.5088
A_24_P82	BC001196	Heparan sulfate HS6ST1	-1.59623	1.04E-16	0.62648	3904.937	2452.807
A_23_P13	NM_00435	CD81 antigen CD81, S5.7	-1.5965	5.55E-12	0.62637	12169.33	7582.528
A_23_P97	NM_02283	hypothetical C1orf80, FL	-1.59657	4.01E-12	0.62634	710.4656	444.7928

A_23_P88	NM_14460 Homo sapi	CKLFSF3,1	-1.59686	2.31E-06	0.62623	4244.543	2694.045
A_24_P837537	Unknown	A_24_P83	-1.59689	6.86E-08	0.62622	921.4216	578.058
A_23_P40	NM_00737 Homo sapi	TARDBP,T	-1.59852	1.21E-12	0.62558	1060.544	662.6863
A_23_P96	NM_00276 Homo sapi	PRPS2,PR	-1.5989	5.23E-06	0.62543	626.036	391.0267
A_23_P15	NM_03193 Homo sapi	RAB34,RA	-1.59973	3.06E-25	0.62511	73939.2	46245.28
A_32_P885123	Unknown	A_32_P88	-1.60052	3.43E-11	0.6248	1626.792	1014.774
A_24_P815062	U66040 D	THC19576	-1.60071	2.16E-06	0.62472	569.5425	355.5062
A_24_P91	NM_05287 transcriptio	BTBD14B,1	-1.6009	1.09E-19	0.62465	4768.912	2986.253
A_24_P30	NM_01433 CREBBP/E	CR11,EID1	-1.6013	1.88E-14	0.62449	23020.07	14418.61
A_23_P17	(NM_00294 Homo sapi	MRPL12,5	-1.60135	2.39E-16	0.62447	6844.21	4257.329
A_23_P42	NM_15245 hypothetic	MGC26690	-1.60183	9.45E-10	0.62428	792.382	494.6613
A_23_P34	NM_03145 selenoprot	SELO	-1.60213	3.61E-11	0.62417	638.4519	398.4625
A_24_P19	NM_00487 mannose-F	MPDU1,SL	-1.60281	2.20E-07	0.6239	2498.927	1546.238
A_24_P40	XM_37336 similar to	c LOC39253	-1.60282	3.70E-14	0.6239	4229.429	2644.605
A_24_P14	NM_18271 Homo sapi	SYPL1,SY	-1.60336	6.87E-12	0.62369	1435.886	900.0892
A_24_P11	NM_00662 S-adenosy	AHCYL1,D	-1.60363	1.57E-21	0.62359	2562.292	1596.343
A_23_P38	NM_00357 VAMP (ves	VAPA,VAP	-1.60368	7.23E-21	0.62356	9154.99	5719.36
A_32_P54	NM_00176 chaperonin	CCT6A,CC	-1.60448	6.72E-25	0.62326	16542.48	10318.58
A_23_P71	(NM_00462 Homo sapi	FANCG,FA	-1.60461	1.12E-27	0.62321	6068.433	3782.845
A_23_P38	NM_00144 Homo sapi	EXTL3,RE	-1.60497	3.29E-24	0.62307	1283.428	799.6687
A_32_P22	NM_00693 SMT3 (sup	SUMO2,H	-1.60518	4.20E-21	0.62298	28555.02	17769.45
A_24_P17	NM_00306 solute carri	SLC22A5,C	-1.60519	1.26E-23	0.62298	2925.849	1823.641
A_32_P92	AK097893 Hypothetic	LOC25739	-1.60526	0.00002	0.62295	650.934	405.8065
A_23_P40	NM_02019 Homo sapi	C2orf33,GI	-1.60541	1.91E-28	0.6229	124756.6	77692.27
A_23_P21	NM_00637 Homo sapi	SYNCRIP,1	-1.60577	1.76E-17	0.62275	7665.402	4782.125
A_23_P25	NM_00135 Homo sapi	AKR1C1,C	-1.60587	2.28E-09	0.62271	239183.6	146768.6
A_23_P21	NM_00569 Homo sapi	ABCF2,AB	-1.60596	3.03E-21	0.62268	1477.311	919.213
A_23_P38	NM_01589 B cell RAG	GALNAC4	-1.60648	1.96E-08	0.62248	1633.125	1017.61
A_24_P38	NM_00578 Homo sapi	THOC4,AL	-1.60666	2.64E-26	0.62241	3408.056	2121.698
A_32_P89	NM_13835 Sterile alph	SAMD1	-1.60683	0.00004	0.62234	584.3496	363.1927
A_23_P16	NM_19896 Homo sapi	AES,GRG,	-1.60712	3.16E-15	0.62223	4824.302	3010.369
A_23_P38	NM_01629 heat shock	TRAP1,HS	-1.60723	1.20E-16	0.62219	2763.714	1715.679
A_24_P11	NM_00583 serine/argi	SRRM1,16	-1.60756	6.37E-09	0.62206	4367.406	2742.407
A_23_P11	NM_17347 Homo sapi	TMEM32	-1.60912	1.31E-08	0.62146	8675.447	5400.216
A_23_P51	NM_01900 ETAA16 pr	ETAA16	-1.6094	2.77E-08	0.62135	826.2062	514.3701
A_23_P31	NM_15267 Homo sapi	SLC10A4,1	-1.60952	2.61E-17	0.6213	806.5225	501.137
A_32_P57002	Unknown	THC19357	-1.60955	0.00002	0.62129	763.6915	474.5781
A_32_P158524	PSA_HUM	THC19423	-1.60969	0.00003	0.62124	1050.13	644.0425
A_32_P101264	AB023583	THC19580	-1.60989	4.23E-06	0.62116	819.8195	510.5498
A_23_P21	(NM_01233 nuclear pre	NARF,FLJ	-1.61069	3.40E-25	0.62085	1800.695	1118.115
A_32_P14	(NM_00693 SMT3 (sup	SUMO2,H	-1.6108	7.45E-19	0.62081	30004.29	18580.2
A_24_P62	NM_01821 Chromosor	C20orf31	-1.61081	6.21E-17	0.62081	2034.346	1264.082
A_23_P30	(NM_00267 promyeloc)	PML,MYL,1	-1.61098	2.51E-06	0.62074	719.8904	446.7932
A_23_P11	(NM_02410 methylosor	WDR77,MI	-1.61165	8.58E-26	0.62048	2422.588	1502.797
A_23_P10	(NM_02006 GLUT4 ent	SLC2A4RC	-1.61187	7.17E-16	0.6204	5096.064	3170.2
A_23_P19	(NM_17814 Solute carr	SLC35B2	-1.61267	5.42E-20	0.62009	6379.949	3948.563
A_24_P28	(NM_02406 hypothetic	MGC2731	-1.61277	1.14E-19	0.62005	1040.613	645.0917
A_32_P205478	Unknown	A_32_P20	-1.61332	2.53E-07	0.61984	857.0112	530.8395
A_24_P41	(NM_00113 amino-terr	AES,GRG,	-1.6137	5.44E-07	0.61969	343.4984	212.8399
A_23_P91	(NM_00473 Homo sapi	VAPB,ALS	-1.61483	1.03E-24	0.61926	1724.413	1067.568
A_24_P33	(NM_19430 Homo sapi	C10orf39,F	-1.61536	1.70E-20	0.61906	1665.457	1031.162
A_23_P91	(NM_13838 hypothetic	MGC1576	-1.61553	2.11E-10	0.61899	2214.971	1365.713

A_23_P14	NM_01514 KIAA0153 KIAA0153	-1.61566	9.92E-17	0.61894	12220.63	7537.282
A_24_P22	(AB028987 Chromosom C19orf7	-1.61575	1.01E-07	0.61891	3531.205	2210.639
A_24_P93	NM_02283 mitochondri MRPS11,H	-1.61587	2.03E-23	0.61886	1734.334	1073.741
A_23_P87	NM_03284 hypothetic FLJ14827	-1.61589	2.20E-11	0.61885	6051.902	3725.529
A_24_P33	NM_02017 Nicalin homolog NCLN	-1.61602	9.27E-16	0.61881	9735.268	6044.011
A_23_P38	NM_15333 Homo sapi KCTD6,MC	-1.61646	1.70E-15	0.61863	1130.767	698.3872
A_23_P16	NM_01567 Homo sapi TRUB2,CL	-1.61674	2.22E-16	0.61853	5832.637	3620.72
A_23_P41	NM_02523 nucleotide COASY,NE	-1.61702	9.73E-19	0.61842	3214.321	1982.455
A_23_P13	NM_03322 Homo sapi PURB,PUF	-1.61709	6.88E-08	0.61839	1043.018	646.7089
A_23_P50	(NM_01421 Homo sapi IMPA2	-1.61785	3.81E-14	0.6181	29326.15	18046.53
A_23_P19	BC008643 Hypothetic MGC3196	-1.61806	9.09E-20	0.61802	4442.82	2740.181
A_23_P20	NM_01901 Homo sapi POLR1B,R	-1.61836	5.70E-12	0.61791	1770.464	1090.297
A_23_P16	NM_00631 aminopepti NPEPPS,F	-1.61846	6.85E-07	0.61787	574.1473	354.5432
A_23_P35	BX247964 Methyltran METTL3	-1.61884	0.00001	0.61773	677.191	417.7775
A_24_P35	NM_01659 DHHC1 precursor ZDHHC3,C	-1.61925	2.17E-16	0.61757	1973.578	1221.786
A_24_P92	L40630 Homo sapi GA11	-1.61927	6.21E-14	0.61756	2897.632	1790.148
A_24_P81	6844 ALU1_HUMAN THC19686	-1.61939	1.01E-14	0.61752	1378.259	852.0033
A_32_P28	(NM_00309 small nucle SNRPA1	-1.61961	1.52E-15	0.61743	14363.91	8830.001
A_23_P25	NM_02453 Homo sapi GPR172A,	-1.61962	4.68E-13	0.61743	6206.929	3820.044
A_24_P39	NM_00429 guanine de GDA,CYPI	-1.61973	3.53E-22	0.61739	1386.894	856.0836
A_24_P21	3284 Unknown A_24_P21	-1.61985	3.73E-19	0.61734	3128.54	1935.174
A_23_P12	(NM_00038 Homo sapi ALDH3A2,	-1.62011	9.36E-17	0.61724	2210.808	1364.886
A_23_P28	(NM_01474 sorting nex SNX17,KIA	-1.62068	0.00005	0.61703	567.1433	349.3878
A_23_P13	(NM_00387 Homo sapi GGH,GH	-1.62083	2.05E-12	0.61697	7441.674	4615.687
A_23_P14	(NM_00572 Homo sapi TSPAN3,T	-1.62097	1.47E-20	0.61691	8584.885	5308.174
A_23_P33	(NM_02456 FLJ21963 FLJ21963	-1.62107	5.46E-18	0.61688	997.1523	615.1901
A_23_P81	(NM_02283 Homo sapi USP46,FL	-1.62108	3.33E-17	0.61687	980.2823	605.0364
A_23_P21	(NM_19445 Homo sapi UBE2J2,N	-1.62126	5.00E-07	0.6168	394.1236	243.2589
A_24_P30	(AK055441 O-sialoglyc OSGEPL1	-1.62221	0.00002	0.61644	847.1793	523.0847
A_24_P91	4922 Unknown A_24_P91	-1.62227	0.00066	0.61642	202.2924	124.6671
A_32_P14	(NM_01595 CGI-27 pro C2orf4,CG	-1.62235	1.11E-20	0.61639	1904.349	1173.124
A_24_P24	806 Unknown A_24_P24	-1.6232	1.69E-08	0.61607	2614.41	1597.55
A_23_P11	(NM_00289 Homo sapi RBBP7,Rb	-1.62334	5.46E-12	0.61601	48959.05	29893.44
A_23_P43	(NM_01612 Homo sapi COPS4,MC	-1.62425	1.46E-18	0.61567	4746.203	2928.959
A_23_P14	(AL834433 EH domain EHBP1L1	-1.62426	3.61E-06	0.61566	685.7227	421.5517
A_23_P11	(NM_00503 Homo sapi PLK1,PLK,	-1.62426	2.33E-09	0.61567	1891.467	1166.057
A_24_P89	(NM_00088 IMP (inosin IMPDH1,IM	-1.62467	0.00005	0.61551	799.7571	489.4012
A_23_P38	(NM_17886 Homo sapi SFXN4,BC	-1.62484	1.58E-08	0.61545	4111.527	2515.387
A_23_P83	(NM_00712 Homo sapi VCP,p97,T	-1.62497	1.62E-16	0.6154	44395.88	27264.5
A_24_P36	(BC029755 Synaptogyl SYNGR2	-1.625	5.57E-10	0.61538	61841.54	37798.21
A_23_P76	(NM_15232 Homo sapi ADSSL1,FI	-1.6253	4.47E-07	0.61527	3469.604	2150.903
A_23_P21	(AF063592 Chromosom C7orf2	-1.62566	1.25E-08	0.61513	1161.413	715.4776
A_23_P68	(AK026838 Protein phc PPM1F	-1.6262	0.00004	0.61493	457.436	280.9482
A_23_P88	(BC047071 BTB (POZ) BTBD7	-1.62641	0.00013	0.61485	297.7245	182.5393
A_24_P91	(BC004338 Uroporphyl UROS	-1.62642	4.17E-13	0.61485	2120.05	1300.503
A_24_P80	(NM_03299 general tra GTF2I,DI	-1.62685	4.71E-12	0.61469	1104.889	680.7562
A_23_P28	(NM_02119 Homo sapi CTDSP1,S	-1.62699	6.17E-20	0.61463	9578.973	5871.479
A_24_P22	(XM_29219 similar to b LOC34165	-1.62702	3.43E-23	0.61462	22093.73	13604.14
A_24_P34	(AK023152 Hypothetic FLJ10099	-1.62714	3.46E-23	0.61458	3894.382	2395.007
A_23_P15	(NM_00253 Homo sapi NUP88,MC	-1.62747	9.48E-14	0.61445	6518.149	4029.082
A_23_P14	(NM_01639 hypothetic HSPC111,	-1.62795	2.44E-21	0.61427	68116.34	41716.03
A_23_P21	(NM_03271 Homo sapi C14orf151,	-1.62802	1.36E-14	0.61424	2028.665	1242.292

A_23_P40	NM_01624	Homo sapi	DHRS8,PA	-1.62803	3.39E-13	0.61424	1919.117	1182.555
A_32_P16	NM_01910	beta-1,4 m	ALG1,HMT	-1.62824	4.53E-09	0.61416	4055.532	2475.852
A_23_P16	NM_00121	Homo sapi	BUB1B,SS	-1.62892	2.21E-11	0.6139	3046.496	1871.288
A_32_P235159	AF117065	THC19025		-1.62902	6.77E-18	0.61387	1121.788	689.1671
A_24_P592060	ALU6_HUM	THC19531		-1.62909	0.00001	0.61384	635.4022	390.0868
A_23_P72	NM_02019	Homo sapi	MRPS22,C	-1.62997	9.55E-25	0.61351	12765.05	7841.488
A_23_P65	NM_03281	hypothetic	FLJ14624,I	-1.63108	2.80E-21	0.61309	1727.718	1058.016
A_23_P31	NM_13857	Bcl-XL-bin	MGC5352,	-1.63208	4.92E-12	0.61271	947.2044	581.0228
A_32_P18	NM_14580	Homo sapi	ZNF511,M	-1.63211	5.44E-18	0.6127	6001.255	3672.192
A_23_P40	D86971	KIAA0217	KIAA0217	-1.63222	1.36E-16	0.61266	4073.925	2505.231
A_23_P12	NM_03284	hypothetic	FLJ14803	-1.63312	5.04E-19	0.61233	5734.599	3509.805
A_23_P30	NM_15238	Homo sapi	KCTD18,FI	-1.63334	8.93E-06	0.61224	367.2288	224.3744
A_23_P21	NM_01398	Neuregulin	NRG2	-1.63382	0.00008	0.61206	515.6882	314.7922
A_24_P761386	Unknown	A_24_P76		-1.63408	4.41E-08	0.61197	1181.679	730.4451
A_23_P38	NM_00052	Homo sapi	HOXA13,H	-1.63437	2.18E-06	0.61186	658.746	403.2828
A_23_P31	NM_01547	DKFZP564	DKFZP564	-1.63496	2.68E-16	0.61164	2314.846	1410.185
A_23_P12	NM_03196	Homo sapi	CCNB1,CC	-1.63497	7.69E-15	0.61163	16889.73	10266.61
A_24_P930597	Unknown	A_24_P93		-1.63513	1.32E-23	0.61157	16213.32	9928.535
A_23_P19	NM_00562	Homo sapi	SGK,SGK1	-1.63594	5.24E-15	0.61127	5999.495	3662.657
A_23_P12	NM_02469	Homo sapi	ECHDC3,F	-1.63644	2.06E-07	0.61108	7029.203	4326.575
A_23_P36	NM_00132	Homo sapi	CSTF1,Cst	-1.63656	9.24E-15	0.61104	4834.992	2963.032
A_23_P43	NM_13835	Sterile alph	SAMD1	-1.63664	7.24E-11	0.61101	632.9265	386.6133
A_23_P39	NM_01531	la related p	LARP1,LAI	-1.63678	1.55E-17	0.61096	1154.047	705.4369
A_23_P85	NM_01423	Homo sapi	GNPAT,DA	-1.63714	1.53E-15	0.61082	6338.361	3883.45
A_24_P10	NM_18169	Homo sapi	PPP2R1B,	-1.63728	5.78E-14	0.61077	1911.75	1172.425
A_24_P17	AK027667	Homo sapi	FLJ14761	-1.63752	4.38E-07	0.61068	1961.681	1206.837
A_24_P80	NM_13848	Homo sapi	LOC15221	-1.63756	1.26E-10	0.61066	1942.125	1187.51
A_23_P16	NM_02393	Homo sapi	MRPS34,M	-1.63768	7.96E-25	0.61062	61548.57	37537.44
A_24_P22	NM_00213	heterogene	HNRPD,P	-1.63785	6.36E-30	0.61056	12509.68	7635.808
A_23_P58	NM_00123	Homo sapi	CCNA2,CC	-1.63889	0	0.61017	2406.156	1468.317
A_24_P31	NM_00735	low density	COG2,LDL	-1.63894	6.84E-21	0.61015	909.8315	555.0992
A_23_P18	NM_00482	Homo sapi	ABCG2,MF	-1.63924	2.84E-20	0.61004	19422.32	11817.55
A_32_P46	NM_15321	hypothetic	FLJ38608	-1.63927	0.00019	0.61003	921.3047	555.7577
A_23_P36	AK001563	Sno, straw	SBNO1	-1.63939	1.81E-11	0.60998	855.9389	521.9153
A_32_P71	NM_01526	KIAA0056	hCAP-D3,I	-1.63943	9.05E-21	0.60997	2451.028	1495.974
A_23_P12	NM_01648	hypothetic	PPHLN1,H	-1.63964	1.80E-17	0.60989	6268.314	3837.05
A_24_P918875	Unknown	THC20024		-1.64013	2.16E-07	0.60971	41366.84	24892.78
A_32_P56415	Unknown	A_32_P56		-1.6406	2.07E-22	0.60953	1347.55	821.7928
A_24_P57	AL831944	Homo sapi	AL831944	-1.64061	3.51E-09	0.60953	2070.896	1269.825
A_24_P12	NM_07846	Homo sapi	BCCIP,TO	-1.64108	3.46E-07	0.60935	620.2048	377.8577
A_24_P25	NM_00509	solute carri	SLC27A4,F	-1.64161	1.59E-06	0.60916	421.4525	256.4858
A_23_P11	NM_00241	Homo sapi	MGST2,G	-1.64242	5.74E-15	0.60886	7161.989	4383.924
A_23_P80	NM_02120	Homo sapi	SRPRB,AF	-1.64254	4.75E-11	0.60881	6564.147	4021.883
A_23_P41	BC010607	Insulin-like	IGF1R	-1.64318	5.05E-13	0.60858	1931.718	1178.141
A_23_P12	NM_02301	Homo sapi	C1orf60,FL	-1.64319	4.39E-08	0.60857	1531.154	935.2924
A_23_P10	NM_03218	hypothetic	MYST1,MC	-1.64428	8.36E-08	0.60817	482.4954	293.1567
A_32_P27	AB033062	Kinesin fan	KIF26A	-1.64432	3.42E-12	0.60816	2281.744	1384.424
A_24_P59	NM_01830	hypothetic	RHOT1,AR	-1.64442	2.81E-06	0.60812	352.3988	213.9975
A_23_P86	NM_00514	Homo sapi	SART1,Ara	-1.64446	1.44E-15	0.6081	1922.347	1169.926
A_24_P508946	Unknown	A_24_P50		-1.64487	1.93E-25	0.60795	1690.999	1027.977
A_23_P39	NM_01223	Homo sapi	PTRF,FKS	-1.64552	5.75E-32	0.60771	30939.02	18802.84
A_24_P20	NM_03056	hypothetic	PRR7,MGC	-1.64691	0.00002	0.6072	287.4075	174.3283

A_23_P98	NM_14564	Homo sapi	SLC15A4,F	-1.64694	5.27E-30	0.60719	3750.299	2276.66
A_32_P15	NM_01910	beta-1,4 m	ALG1,HMT	-1.64715	6.00E-23	0.60711	2095.27	1272.27
A_23_P40	NM_00133	Homo sapi	CTSZ,CTS	-1.64718	0.00076	0.6071	1411.771	846.1129
A_23_P34	NM_13847	Homo sapi	CDAN1,CC	-1.64756	3.08E-07	0.60696	469.2231	285.0596
A_23_P11	NM_02237	Homo sapi	TOR3A,AD	-1.6484	5.71E-15	0.60665	15671.53	9464.254
A_24_P34	NM_17686	Homo sapi	PSME3,Ki,	-1.64855	3.31E-15	0.60659	1282.848	780.5114
A_24_P374973		Unknown	ENST0000	-1.64864	0.00012	0.60656	707.8234	428.6711
A_23_P37	NM_13344	Homo sapi	GPT2,ALT	-1.64887	4.12E-25	0.60648	1371.189	831.5118
A_23_P68	NM_00404	Homo sapi	ATIC,PURI	-1.649	6.89E-31	0.60643	9838.184	5964.223
A_23_P15	NM_02038	Homo sapi	REXO4,XF	-1.6496	1.43E-21	0.60621	1082.674	656.4316
A_23_P10	NM_00464	Chromosor	C21orf33	-1.65036	3.03E-24	0.60593	7319.269	4440.932
A_24_P79	NM_00510	RNA bindir	RBM8A,Y1	-1.65082	4.27E-13	0.60576	2903.328	1762.728
A_32_P167631		Unknown	THC19493	-1.65084	0.00001	0.60575	429.958	259.9341
A_24_P22	NM_00135	aldo-keto r	AKR1C1,C	-1.65112	8.95E-17	0.60565	260753.8	157117.3
A_23_P43	NM_18261	hypothetic	FLJ34283,I	-1.65135	1.37E-12	0.60557	1366.026	824.3355
A_24_P62	XM_37329	similar to a	LOC39234	-1.65197	9.23E-06	0.60534	274.3295	165.9948
A_23_P30	NM_00419	Homo sapi	P4HA2,4-F	-1.65214	3.25E-10	0.60527	5264.156	3199.755
A_23_P39	NM_03142	Homo sapi	C9orf58,IB	-1.65219	1.05E-20	0.60526	1543.582	934.6214
A_23_P21	NM_00229	Homo sapi	LAMB2,LA	-1.65244	3.18E-08	0.60516	4974.756	3030.747
A_24_P303193		Unknown	THC18193	-1.65249	6.58E-11	0.60515	810.7919	490.4849
A_23_P34	NM_00014	Homo sapi	FH,MCL,LF	-1.6526	5.14E-22	0.60511	14309.31	8672.118
A_32_P13	NM_00630	putative hu	ANP32A,L	-1.65309	2.57E-10	0.60493	875.4631	529.7042
A_24_P95	NM_03299	general tra	GTF2I,DIW	-1.65343	3.73E-23	0.6048	1028.541	622.0206
A_23_P13	NM_00360	Homo sapi	STK6,AIK,	-1.6538	4.67E-23	0.60467	5590.74	3374.218
A_24_P19	AK027248	Sin3A assc	SAP30L	-1.65421	0.00014	0.60452	272.2748	164.9987
A_23_P10	NM_00682	Homo sapi	EBNA1BP2	-1.65436	5.41E-13	0.60446	71566.5	42878.32
A_23_P73	NM_03282	hypothetic	C9orf3,AP	-1.65468	2.16E-20	0.60435	3127.202	1893.899
A_23_P43	NM_14457	Homo sapi	C16orf34,L	-1.65514	5.44E-12	0.60418	17688.44	10608.89
A_32_P179005		ALU8_HUM	THC18442	-1.65527	3.39E-15	0.60413	718.9862	434.4325
A_24_P23	AB033004	CD2 antige	CD2BP2	-1.6553	6.72E-08	0.60412	535.0022	323.2349
A_23_P84	NM_03217	hypothetic	TOMM40L	-1.65578	6.20E-09	0.60395	3197.677	1946.52
A_23_P92	NM_17680	Homo sapi	MOCS2,MI	-1.6565	6.39E-10	0.60368	2548.061	1553.867
A_24_P834646		Unknown	A_24_P83	-1.65703	3.78E-18	0.60349	910.1744	549.0225
A_24_P17	AK022463	Homo sapi	FLJ12649,I	-1.65742	9.96E-11	0.60335	1121.835	677.2878
A_24_P36	NM_02045	Homo sapi	THAP11,C	-1.65746	2.59E-13	0.60333	1039.704	628.0939
A_24_P191357		Unknown	THC19455	-1.65796	0.00003	0.60315	547.1714	329.2522
A_23_P14	NM_13844	Homo sapi	C7orf30	-1.65824	6.06E-27	0.60305	7841.203	4734.365
A_32_P112623		BC032035	THC18073	-1.6584	4.42E-20	0.60299	941.8418	568.0272
A_24_P12	NM_01818	hypothetic	SBNO1,Sn	-1.65888	1.23E-06	0.60282	3472.048	2122.859
A_24_P26	NM_01209	CD3-epsilo	CD3EAP,C	-1.6589	1.73E-24	0.60281	2085.836	1257.45
A_23_P11	NM_02237	ATP-deper	TOR3A,AD	-1.65914	2.06E-07	0.60272	38204.09	22717.86
A_32_P83	NM_08062	Homo sapi	BRI3BP,KC	-1.65938	2.00E-14	0.60263	2571.398	1548.334
A_32_P20	NM_13833	Homo sapi	POLR3H,R	-1.65949	0.00059	0.60259	916.3817	555.2297
A_23_P29	NM_00326	Homo sapi	TLR3	-1.65979	1.01E-07	0.60249	508.8738	306.4049
A_32_P45	AF037219	Ubiquitin p	UBE3A	-1.66009	3.18E-08	0.60238	765.9003	461.2455
A_23_P34	AK024475	Pleckstrin	DKFZP434	-1.6605	5.27E-09	0.60223	1932.115	1167.27
A_23_P35	NM_02506	Homo sapi	BXDC5,RF	-1.66061	2.00E-14	0.60219	4206.295	2547.147
A_23_P25	NM_00573	Homo sapi	KIF20A,Mk	-1.66067	7.99E-30	0.60217	2961.892	1785.609
A_23_P35	NM_00566	Homo sapi	C5orf18,DF	-1.66135	6.69E-25	0.60192	9241.814	5554.711
A_23_P18	NM_03280	glycosyltra	AGO61,FL	-1.66154	2.02E-11	0.60185	8913.396	5337.744
A_32_P23	BX108897	BX108897	BX108897	-1.66201	1.51E-10	0.60168	582.4652	350.4307
A_24_P11	NM_02272	DiGeorge ε	DGCR8,Gy	-1.6626	1.63E-13	0.60147	1294.706	779.7498

A_23_P96:	AK074719	FLJ20105 FLJ20105	-1.66268	3.23E-07	0.60144	1088.069	652.6907
A_23_P41:	NM_00669	bladder carcinoma BLCAP, BC	-1.66292	2.89E-19	0.60135	2290.272	1379.125
A_32_P127454	Unknown	A_32_P127454	-1.6634	4.52E-15	0.60118	5079.998	3040.969
A_32_P228254	Unknown	A_32_P228254	-1.66378	0.00018	0.60104	2803.011	1636.071
A_32_P23:	AY007155	Hypothetical LOC43994	-1.6639	3.19E-08	0.601	1220.172	733.8173
A_24_P92:	BC020784	Similar to FMGC14289	-1.66458	3.96E-16	0.60075	3698.24	2229.863
A_24_P41:	NM_03262	hypothetical SSBP4, MGC	-1.66471	3.11E-21	0.60071	2368.061	1423.389
A_24_P10:	NM_02507	hypothetical PTGES2, G	-1.66484	1.49E-10	0.60066	737.1216	442.2913
A_23_P39:	NM_15268	hypothetical RWDD4A, I	-1.66523	2.32E-12	0.60052	1734.913	1045.462
A_24_P26:	AK095775	Pleckstrin domain PLEKHA1	-1.66559	6.78E-17	0.60039	3058.73	1840.996
A_24_P21:	NM_03288	U7 snRNP LSM10, MS	-1.66663	1.01E-13	0.60001	2446.214	1458.854
A_23_P68:	NM_00312	sepiapterin SPR	-1.66672	4.61E-17	0.59998	5647.006	3400.893
A_32_P42:	AA521057	Transcription factor GGA2, VEA	-1.66757	1.06E-15	0.59967	724.0416	434.1465
A_23_P43:	NM_02161	Homo sapiens PRDM12, F	-1.66788	6.02E-06	0.59956	334.139	200.0321
A_23_P39:	AB020684	KIAA0877 KIAA0877	-1.66793	4.10E-09	0.59954	1385.356	834.8175
A_24_P83:	NM_00456	serine (or cysteine) SERPINB6	-1.66978	2.97E-09	0.59888	1392.495	828.5269
A_24_P75:	XR_00018	similar to GLOC34632	-1.6709	3.28E-11	0.59848	1699.103	1018.765
A_23_P20:	NM_01591	Homo sapiens FAM26B	-1.67105	9.18E-13	0.59843	1051.64	628.357
A_23_P12:	NM_00728	protein tyrosine kinase PTK9L, A6r	-1.67106	2.47E-07	0.59842	1232.698	733.4645
A_32_P136427	Unknown	A_32_P136427	-1.67119	1.11E-14	0.59838	1017.27	607.9282
A_23_P10:	NM_00643	Homo sapiens CCT4, SRB	-1.67173	6.29E-33	0.59818	206029.6	123212.8
A_24_P204067	NPL4_HUMAN	THC19917	-1.67174	1.18E-14	0.59818	846.3718	506.4454
A_32_P64820	Unknown	A_32_P64820	-1.67175	0.00011	0.59817	766.3756	453.621
A_23_P19:	NM_03126	Homo sapiens HNRPAB, A	-1.67241	1.65E-08	0.59794	3236.939	1945.139
A_23_P40:	BC029424	Similar to FLOC49386	-1.67304	7.29E-28	0.59771	5696.937	3402.556
A_23_P17:	NM_02515	Nucleotide NUBPL	-1.67333	3.38E-08	0.59761	608.3868	363.3198
A_23_P54:	NM_01610	comparative CGI-37, KD	-1.67355	9.35E-33	0.59753	6210.618	3710.388
A_23_P12:	NM_00557	Homo sapiens LOXL1, LOX	-1.67536	1.41E-28	0.59689	9439.41	5637.942
A_24_P94:	AB023147	Chromosome C22orf9	-1.6761	0.00026	0.59662	208.0893	124.1238
A_23_P15:	NM_03232	Homo sapiens ZDHHC16,	-1.67631	7.02E-20	0.59655	2206.892	1315.235
A_23_P64:	NM_17679	Homo sapiens P2RY6, P2R	-1.67686	6.47E-15	0.59635	542.2422	323.3644
A_23_P12:	NM_01614	PTD009 protein TRAPPC4,	-1.6771	1.22E-14	0.59627	10467.5	6258.297
A_23_P21:	AL831922	DKFZP586DLG1	-1.67711	3.85E-12	0.59626	1464.578	870.7206
A_24_P46:	AK095803	Malate dehydrogenase MDH2	-1.67891	3.58E-19	0.59562	1617.529	964.7528
A_23_P56:	NM_00677	Homo sapiens DDX18, MRL	-1.67896	2.85E-21	0.59561	3261.2	1954.196
A_23_P13:	NM_02033	Homo sapiens RAI17, Zim	-1.6802	6.53E-24	0.59517	8139.831	4854.292
A_23_P12:	NM_02408	Homo sapiens KDELC1, E	-1.68074	1.96E-27	0.59497	1509.678	899.8655
A_32_P62:	NM_00510	RNA binding protein RBM8A, Y1	-1.68104	1.62E-14	0.59487	2013.537	1200.934
A_23_P13:	NM_00616	Homo sapiens NFE2, p45,	-1.68124	6.69E-06	0.5948	775.3591	460.1918
A_24_P29:	NM_01884	SBBI26 protein KLHL7, KLH	-1.68163	9.95E-06	0.59466	241.2581	143.4538
A_24_P40:	NM_17387	Homo sapiens CLCN3, CL	-1.68178	6.68E-06	0.59461	645.8963	383.4603
A_23_P98:	NM_17679	Homo sapiens HRAS, HRAS	-1.68188	6.81E-22	0.59457	2901.303	1721.965
A_23_P13:	NM_19929	Homo sapiens APITD1, M	-1.68267	3.25E-13	0.59429	4045.983	2415.636
A_23_P36:	NM_00176	cyclin D3 CCND3	-1.68301	2.19E-14	0.59417	2777.618	1643.837
A_24_P31:	NM_03194	centaurin, gamma CENTG3, A	-1.68363	2.59E-13	0.59395	611.1805	363.0111
A_23_P74:	NM_01512	Mid-1-related CLCC1, MC	-1.6839	5.35E-16	0.59386	1442.031	857.5291
A_24_P25:	NM_02183	chromosome PEO1, PEC	-1.68429	5.25E-13	0.59372	7947.738	4726.203
A_24_P72:	NM_00640	actin related ARPC1A, A	-1.68446	1.29E-32	0.59366	12273.4	7282.97
A_23_P16:	NM_14564	DEAH (Asp) DHX57	-1.68455	8.72E-15	0.59363	721.8563	428.3921
A_24_P373844	Unknown	THC19679	-1.68469	5.34E-14	0.59358	562.1621	333.6771
A_23_P40:	NM_15335	Homo sapiens FBXL16, Ft	-1.68547	4.29E-20	0.59331	1436.074	852.0975
A_24_P30:	BC002759	Hypothetical FLJ20489	-1.6856	6.85E-18	0.59326	9396.859	5562.128

A_24_P41	(NM_001111 adenylate c	ADCY7,KI	-1.68581	4.29E-14	0.59319	2128.607	1254.328
A_23_P20	(NM_01623 Homo sapi	ANAPC5,A	-1.6868	4.76E-33	0.59284	36544.46	21660.04
A_24_P18	(NM_00331 thyroid tran	TITF1,BC	-1.68713	1.51E-10	0.59272	554.5931	328.6415
A_23_P12	(NM_00598 Homo sapi	SLC25A1,(-1.68734	5.17E-09	0.59265	4492.516	2662.003
A_23_P34	7100 Unknown A_23_P34		-1.68749	3.08E-22	0.5926	45935.58	27259.78
A_32_P10	(AK094772 Homo sapi	AK094772	-1.68868	4.03E-12	0.59218	970.6335	575.0377
A_24_P33	(NM_19848 Homo sapi	RPL7L1,M	-1.68892	2.94E-15	0.59209	2887.435	1716.123
A_24_P31	(NM_00249 NIMA (nev	NEK2,NLK	-1.68941	7.02E-09	0.59192	969.8776	573.958
A_32_P20	(NM_00141 ELAV (emt	ELAVL1,HI	-1.68977	4.62E-08	0.5918	577.7113	341.6354
A_23_P25	(NM_14891 Homo sapi	PSMB8,LV	-1.69092	8.75E-19	0.59139	2552.779	1503.924
A_24_P33	(NM_03092 Homo sapi	NIPA2,MG	-1.69172	1.78E-20	0.59112	1956.917	1154.878
A_24_P32	(NM_00620 phosphodi	PDE4A,PD	-1.6918	2.63E-23	0.59109	1709.82	1010.326
A_24_P55	M28016 Human mit cytochrom		-1.69346	3.25E-11	0.59051	43304.69	25879.59
A_24_P33	AB032966 Tetratricop	TTC7A	-1.6945	1.11E-15	0.59015	1311.608	774.524
A_23_P12	(NM_01208 Homo sapi	FRAT2,MC	-1.69503	5.25E-07	0.58996	2706.24	1588.071
A_23_P19	AB028946 IQ motif co	IQCE	-1.69518	8.87E-14	0.58991	2858.714	1687.771
A_24_P35	(NM_00043 platelet-act	PAFAH1B1	-1.69525	2.74E-30	0.58988	4088.81	2413.04
A_32_P16	(AK056476 Trinucleotic	TNRC8	-1.69528	1.52E-15	0.58987	1535.628	903.3362
A_23_P35	(NM_17234 Homo sapi	SPAG9,HS	-1.69538	4.56E-16	0.58984	736.0813	434.157
A_24_P12	(NM_00513 DiGeorge s	DGCR2,ID	-1.69545	1.04E-15	0.58981	3173.683	1876.515
A_32_P25	(NM_17203 Homo sapi	RDH10	-1.69578	1.31E-18	0.5897	2476.563	1464.728
A_24_P58	(NM_01910 beta-1,4 m	ALG1,HMT	-1.69623	2.35E-11	0.58954	2253.896	1330.774
A_24_P30	(AK021848 Homo sapi	FLJ11786	-1.69629	2.85E-17	0.58952	679.968	400.8265
A_23_P20	(NM_01766 Homo sapi	NDE1,NUC	-1.69636	8.54E-20	0.5895	1900.83	1119.854
A_23_P16	(NM_02206 Homo sapi	C14orf133,	-1.69708	3.91E-22	0.58925	1429.736	842.7019
A_23_P20	(NM_00208 Homo sapi	GPC1	-1.69752	2.26E-16	0.5891	26416.28	15518.98
A_32_P15	(XM_37962 hypothetic	LOC40149	-1.69786	6.06E-22	0.58898	5842.139	3447.589
A_23_P23	(NM_01443 Homo sapi	SLC39A1,2	-1.69837	1.25E-08	0.5888	2600.528	1524.235
A_24_P40	(NM_01466 KIAA0140	FAM53B,K	-1.69966	2.07E-22	0.58835	2621.084	1539.498
A_24_P37	(NM_03247 Homo sapi	MRPL36,R	-1.70045	1.77E-14	0.58808	8054.951	4748.753
A_32_P99	(BM709498 Transcriber	BM709498	-1.7012	1.42E-12	0.58782	891.7142	524.3997
A_23_P16	(NM_00568 phenylalan	FARSLB,F	-1.70183	1.32E-11	0.5876	1972.984	1153.949
A_32_P12	3027 Unknown A_32_P12		-1.70198	0.0001	0.58755	1566.995	895.1842
A_32_P31	(NM_00555 Homo sapi	LAMA1,LA	-1.7025	4.83E-25	0.58737	1464.121	859.3559
A_23_P67	(NM_02457 UDP-N-ac	GALNT14	-1.70278	2.17E-30	0.58727	11457.52	6724.051
A_23_P14	(NM_08065 Homo sapi	ATP6V1E2	-1.7035	3.66E-17	0.58703	917.2247	538.6003
A_23_P36	(BC014522 Protein phc	PPP1R14E	-1.70368	8.70E-18	0.58697	31232.32	18377.48
A_23_P68	(AL137442 Chromosom	C20orf177	-1.7039	1.05E-10	0.58689	3245.676	1914.595
A_23_P25	(NM_00601 Homo sapi	CD164,MC	-1.70453	4.39E-17	0.58667	3032.593	1786.712
A_24_P45	(NM_02044 Hypothetic	DJ462O23	-1.70495	3.59E-21	0.58653	1050.616	616.5858
A_23_P30	(NM_02402 hypothetic	MGC3265	-1.70503	4.53E-11	0.5865	2357.943	1378.875
A_24_P16	(NM_14456 Homo sapi	SLC39A3,2	-1.70602	2.17E-07	0.58616	522.4097	305.7545
A_24_P28	(NM_00280 proteasom	PSMD3,S3	-1.70688	4.17E-06	0.58586	4091.402	2354.978
A_23_P61	(NM_00040 Homo sapi	GLB1	-1.70711	1.08E-09	0.58578	1540.516	908.2465
A_23_P25	(NM_01603 Homo sapi	TTC15,CG	-1.70736	1.19E-15	0.5857	1894.812	1108.096
A_23_P21	(NM_07862 Homo sapi	MSL3L1,DI	-1.70762	5.09E-12	0.58561	1162.036	680.0205
A_24_P93	(AB011092 Adenylate c	ADCY9	-1.70781	3.49E-07	0.58554	1039.532	610.0609
A_23_P25	(NM_00557 Homo sapi	LMNB1,LV	-1.70847	2.59E-19	0.58532	3514.549	2049.607
A_23_P36	(NM_02207 EGL nine	(EGLN3,PH	-1.70856	3.53E-09	0.58529	526.933	308.1873
A_24_P91	(NM_01869 hypothetic	NXT2,P15-	-1.70864	5.42E-26	0.58526	5314.78	3115.357
A_23_P14	(NM_01481 Homo sapi	KIAA0649,	-1.70887	1.11E-17	0.58518	2065.024	1207.384
A_32_P82	(NR_00145 Homo sapi	C14orf62	-1.70906	6.44E-14	0.58512	1939.61	1129.672

A_24_P323916	Unknown	A_24_P32:	-1.70907	3.84E-22	0.58511	1634.121	957.2812	
A_23_P92:	NM_02019	Homo sapi	C2orf33,GI	-1.71002	3.19E-29	0.58479	14149.16	8280.153
A_23_P16:	NM_00032	Homo sapi	QDPR,DHF	-1.71037	2.97E-14	0.58467	3427.079	2013.229
A_23_P25:	NM_02209	transposon	LOC63920	-1.71038	6.89E-06	0.58466	638.9652	373.2386
A_24_P93:	AB020706	Adaptor-rel	AP2A2	-1.71076	1.42E-27	0.58453	1965.749	1149.137
A_24_P12:	M69012	Similar to	CGNAI2	-1.71139	2.46E-06	0.58432	343.4407	200.8941
A_24_P11:	NM_00613	Homo sapi	C11orf11,N	-1.71181	1.92E-06	0.58418	293.3825	171.2797
A_24_P31:	NM_00688	calmodulin	CALM1,CA	-1.712	9.50E-14	0.58411	1427.195	836.6531
A_23_P84:	NM_00631	Homo sapi	NPEPPS,F	-1.71212	2.24E-25	0.58407	2005.66	1172.327
A_23_P16:	NM_03299	Homo sapi	GTF2I,DIW	-1.71266	7.71E-20	0.58389	16815.02	9875.208
A_23_P16:	NM_00299	Homo sapi	SDC1,SDC	-1.71307	1.13E-16	0.58375	8648.686	5066.465
A_24_P13:	NM_19840	Homo sapi	PTPLB	-1.71373	5.42E-15	0.58352	3522.976	2067.302
A_23_P92:	NM_01489	Homo sapi	RHOBTB3	-1.71424	1.18E-16	0.58335	1122.968	655.7974
A_24_P16071	Unknown	A_24_P16:		-1.71447	1.38E-29	0.58327	2303.864	1343.935
A_23_P74:	NM_00285	ATP-bindin	ABCD3,AB	-1.7147	6.73E-10	0.58319	2208.127	1292.04
A_23_P50:	NM_00269	polymerase	POLD1,PC	-1.71554	2.21E-11	0.58291	6093.203	3522.22
A_32_P75:	BC039509	Hypothetic	LOC34010	-1.71629	0.00022	0.58265	298.6617	173.7602
A_23_P23:	NM_00678	Homo sapi	WDR3	-1.71702	1.27E-12	0.5824	8112.097	4754.683
A_24_P56:	AL832817	Dual-specif	DYRK2	-1.71742	2.59E-28	0.58227	1313.255	764.5548
A_24_P75:	BC043374	Homo sapi	LOC34050	-1.71932	3.38E-09	0.58162	657.2575	382.1463
A_23_P37:	NM_00586	STIP1 hom	STUB1,CH	-1.72099	6.68E-17	0.58106	10767.56	6236.114
A_24_P552987	Unknown	A_24_P55:		-1.72127	2.13E-28	0.58097	1398.735	812.8037
A_23_P25:	NM_08062	Homo sapi	BRI3BP,KC	-1.7213	1.03E-34	0.58096	10143.31	5894.559
A_24_P21:	NM_03191	complemer	C1QTNF6,	-1.72158	9.05E-24	0.58086	2553.338	1484.943
A_24_P10:	NM_03204	hypothetic	DKFZP564	-1.72268	8.80E-07	0.58049	382.6575	222.4737
A_23_P34:	NM_01406	Homo sapi	MAGEH1,/	-1.72399	0.0005	0.58005	5100.64	3022.479
A_23_P85:	NM_00349	chromosom	CXorf12,IT	-1.72473	1.02E-21	0.5798	1725.353	1001.476
A_23_P13:	NM_00219	interferon	r IRF2,IRF-2	-1.72496	2.89E-11	0.57972	1310.832	762.3611
A_24_P41:	AB014599	Bicaudal D	BICD2	-1.72531	6.68E-30	0.57961	2621.878	1520.629
A_23_P13:	AF220046	Protein phc	PPP2R2D	-1.72541	4.44E-12	0.57957	2596.668	1509.97
A_23_P86:	NM_00382	Homo sapi	FADD,GIG	-1.72588	2.63E-18	0.57941	9132.123	5275.863
A_23_P28:	NM_00462	Homo sapi	TSN,TRSL	-1.72627	9.26E-33	0.57928	3010.534	1744.609
A_23_P11:	NM_00106	Homo sapi	TOP2A,TO	-1.72649	4.17E-10	0.57921	5618.322	3270.152
A_24_P195454	Unknown	A_24_P19:		-1.72678	1.43E-17	0.57911	1638.129	951.5527
A_24_P38:	NM_01229	Ras-GTPa:	G3BP2	-1.729	1.07E-29	0.57837	11209.12	6488.453
A_32_P146826	Unknown	THC20051		-1.73071	0.00012	0.5778	336.9954	193.9913
A_23_P71:	NM_00521	Homo sapi	DEFB1,BD	-1.73124	1.79E-15	0.57762	2836.188	1634.301
A_24_P92:	U31733	Hypothetic	hppt	-1.73129	1.63E-19	0.5776	4410.994	2542.737
A_23_P12:	NM_00226	Homo sapi	KPNA2,QII	-1.73175	2.59E-27	0.57745	53682.67	30917.1
A_24_P15:	NM_00135	aldo-keto	r AKR1C1,C	-1.73203	8.17E-28	0.57736	16582.95	9603.355
A_23_P15:	NM_00562	Homo sapi	SLC6A8,C	-1.73366	3.55E-13	0.57682	3202.704	1857.875
A_32_P17:	AK026783	CDNA: FL:	AK026783	-1.73515	7.38E-14	0.57632	886.3999	511.0794
A_23_P54:	NM_03287	Homo sapi	JUB,Ajuba	-1.73516	1.17E-21	0.57631	3313.036	1912.431
A_23_P14:	NM_00672	cadherin 1	(CDH10	-1.73579	2.71E-20	0.57611	2303.255	1326.235
A_23_P10:	NM_00523	Homo sapi	NR2F6,EA	-1.73817	2.30E-15	0.57532	1563.892	898.4884
A_23_P41:	NM_01528	KIAA0239	PHF15,JA	-1.73829	5.91E-08	0.57528	498.9901	286.6277
A_23_P40:	NM_00394	Homo sapi	USP13,ISC	-1.7383	5.22E-08	0.57527	1208.909	696.695
A_23_P57:	NM_00377	Homo sapi	HYAL2,LU	-1.73928	1.63E-08	0.57495	2675.225	1529.339
A_24_P21:	NM_01818	hypothetic	DCUN1D2,	-1.74022	3.74E-06	0.57464	379.3563	217.4209
A_24_P93:	AK097197	Hepatocyte	HGS	-1.74076	1.88E-06	0.57446	1291.798	747.6188
A_32_P16625	Unknown	A_32_P16:		-1.741	1.55E-10	0.57438	562.2381	322.9006
A_24_P28:	NM_05285	hypothetic	ADCK2,AA	-1.74124	8.68E-12	0.5743	1669.537	953.2772

A_24_P349756	Unknown	A_24_P349756	-1.74135	3.48E-17	0.57427	18193.01	10412.5
A_23_P25(AK023831)	Hypothetical	FLJ13769	-1.74155	0.00019	0.5742	393.8984	225.2012
A_24_P17(NM_02206)	likely ortho	E2-230K,F	-1.74287	5.00E-16	0.57377	6750.164	3887.804
A_23_P11(NM_02474)	Homo sapi	HPS6,FLJ2	-1.74329	1.56E-36	0.57363	9446.876	5420.437
A_23_P90(NM_00662)	Homo sapi	POP4,RPF	-1.74365	4.59E-35	0.57351	7226.649	4146.601
A_24_P23(NM_00103)	ribonucleot	RRM2,R2,I	-1.74427	4.80E-23	0.57331	6224.17	3580.638
A_23_P49(NM_00345)	Homo sapi	ZNF174,ZS	-1.7453	4.15E-12	0.57297	582.5364	333.7228
A_24_P110591	Unknown	A_24_P110591	-1.74628	1.49E-21	0.57264	15088.74	8649.555
A_24_P341489	Unknown	A_24_P341489	-1.7465	5.24E-16	0.57257	1304.763	749.3091
A_32_P94(BC053353)	Sarcoma a	FLJ10330	-1.74695	1.06E-07	0.57243	356.6429	204.3413
A_24_P36(NM_01594)	CGI-11 pro	ATP6V1H,1	-1.74922	1.98E-10	0.57168	590.5379	337.5283
A_24_P36(NM_02468)	hypothetic	C17orf42,F	-1.74991	4.94E-18	0.57146	3605.103	2069.62
A_23_P21(NM_05402)	Homo sapi	CNOT7,CA	-1.75079	2.32E-15	0.57117	1834.92	1050.956
A_32_P64990	Unknown	A_32_P64990	-1.75083	5.08E-19	0.57116	1283.752	733.0822
A_23_P83(NM_00586)	Homo sapi	NET1,NET	-1.75103	8.24E-22	0.57109	4844.25	2757.011
A_24_P41(AF318364)	Protein phc	PPP1R14E	-1.75136	5.57E-20	0.57099	21490.14	12229.35
A_23_P55(NM_01546)	DKFZP586	NOL11,DK	-1.75177	8.41E-33	0.57085	4678.67	2668.614
A_23_P21(NM_01836)	Homo sapi	AGPAT5,L	-1.75258	7.55E-17	0.57059	751.0172	428.5804
A_23_P61(NM_02526)	hole gene	MGC4659	-1.75274	6.08E-08	0.57053	3936.896	2232.998
A_23_P35(NM_00249)	Homo sapi	NEK2,NLK	-1.75356	8.39E-07	0.57027	1625.52	930.3596
A_23_P50(NM_00122)	Homo sapi	CASP6,MC	-1.75371	1.22E-09	0.57022	2751.453	1579.493
A_23_P21(NM_00504)	Homo sapi	PRKX,PKX	-1.75478	1.96E-16	0.56987	838.0341	477.148
A_23_P34(AK094181)	Tetratricop	TTC15	-1.75485	2.26E-24	0.56985	772.2532	440.068
A_24_P39(AK098354)	BS 3076	AK098354	-1.75534	8.29E-24	0.56969	1125.816	641.2755
A_23_P32(NM_00174)	Homo sapi	CALM2,PH	-1.75543	1.03E-25	0.56966	120142.5	68598.66
A_23_P14(NM_01516)	homolog of	RRS1,KIAA	-1.75634	6.83E-21	0.56936	11359.37	6438.571
A_32_P13(NM_01752)	Homo sapi	LY6K,FLJ3	-1.75638	1.41E-18	0.56935	5558.69	3177.813
A_32_P50(NM_00459)	spermine s	SMS,SRS,	-1.75652	4.63E-19	0.56931	2001.283	1140.693
A_23_P10(NM_01607)	Homo sapi	MRPS23,C	-1.757	2.73E-34	0.56915	7120.904	4055.085
A_24_P31(BX647782)	Nuclear tra	NUTF2	-1.75712	4.93E-25	0.56911	985.9233	560.9743
A_24_P35(NM_01827)	hypothetic	TMEM19,F	-1.75726	1.68E-07	0.56907	1698.745	965.3742
A_32_P38(AB037835)	KIAA1414	KIAA1414	-1.75787	7.88E-12	0.56887	388.0234	220.7495
A_32_P22(AK054569)	Methionine	MAT2A	-1.75834	0.00072	0.56872	2451.68	1343.644
A_23_P50(NM_01828)	Homo sapi	PHF10,XAI	-1.75907	3.28E-36	0.56848	7415.418	4217.008
A_24_P34(NM_02489)	hypothetic	NOL10,FL	-1.7595	3.05E-15	0.56834	620.7053	352.7548
A_23_P50(NM_01231)	Homo sapi	SLC39A6,L	-1.7596	1.82E-11	0.56831	2316.519	1324.62
A_23_P50(NM_02479)	Homo sapi	FAM57A,C	-1.7599	1.20E-06	0.56821	1825.492	1052.665
A_23_P51(NM_15333)	Homo sapi	PUSL1,FL	-1.76027	7.76E-30	0.5681	5445.426	3097.31
A_32_P44(NM_01411)	PRO0149	PRO0149	-1.76036	9.78E-15	0.56806	2837.444	1622.136
A_32_P11(NM_01234)	F-box only	FBX09,FB	-1.76041	4.07E-15	0.56805	1894.724	1074.582
A_32_P186725	Unknown	THC18314	-1.76067	8.01E-11	0.56797	395.2585	224.5472
A_24_P35(NM_00448)	UDP-N-acc	GALNT2,G	-1.76118	4.17E-13	0.5678	3253.999	1848.352
A_24_P18(AL133052)	Chromosor	C1orf37	-1.76125	1.22E-12	0.56778	1539.244	875.8135
A_23_P13(NM_00263)	Homo sapi	PHB	-1.76133	1.11E-34	0.56775	5298.5	3007.047
A_24_P238286	Unknown	A_24_P238286	-1.762	1.88E-08	0.56754	1021.054	579.8789
A_24_P281468	Unknown	ENST0000	-1.76256	2.67E-13	0.56736	3383.448	1926.952
A_24_P94(AB007896)	Prolyl endc	KIAA0436	-1.76307	2.54E-12	0.56719	3986.619	2263.404
A_32_P35486	Unknown	THC18095	-1.76444	4.26E-06	0.56675	410.6939	232.7228
A_23_P10(NM_02475)	Homo sapi	AGMAT,FL	-1.7652	3.84E-14	0.56651	2333.558	1324.682
A_23_P12(NM_01214)	Homo sapi	DTYMK,CI	-1.76585	5.78E-35	0.5663	35685.13	20193.13
A_23_P36(NM_02463)	Sin3A assc	SAP30L,FL	-1.76641	2.25E-30	0.56612	4109.137	2327.218
A_23_P10(NM_17070)	Homo sapi	ICMT,PCM	-1.76716	1.23E-18	0.56588	2843.958	1605.716

A_23_P25	NM_00641	Homo sapi	AGPAT1,G	-1.76752	8.56E-16	0.56576	2811.626	1587.252
A_23_P33	NM_15301	hypothetic	FLJ30596,I	-1.76776	1.26E-12	0.56569	1072.324	607.1244
A_24_P486503		Unknown	A_24_P48	-1.76811	3.66E-09	0.56557	986.1834	558.7034
A_32_P56489		Unknown	A_32_P56	-1.76837	0.00026	0.56549	1896.948	1038.847
A_24_P27	AF284094	BRI3 bindir	BRI3BP	-1.76951	7.97E-36	0.56513	1734.655	980.3065
A_23_P42	NM_02011	hypothetic	ZC3HAV1,	-1.7696	1.58E-17	0.5651	4516.984	2564.125
A_23_P89	NM_00576	tripartite m	TRIM28,K/	-1.7706	4.13E-22	0.56478	6599.111	3737.879
A_23_P15	NM_00611	Homo sapi	PECI,DRS	-1.77063	1.65E-34	0.56477	14842.86	8389.33
A_24_P18	NM_00010	dystrophin	DMD,BMD	-1.77142	0.00009	0.56452	200.5103	113.223
A_24_P12	NM_00678	WD repeat	WDR3	-1.77152	4.83E-12	0.56449	4665.05	2651.961
A_23_P10	NM_00467	3'-phospho	PAPSS2,S	-1.77176	1.27E-09	0.56441	509.4305	287.2949
A_23_P16	NM_00568	Homo sapi	FARSLB,F	-1.77186	7.61E-32	0.56438	27911.77	15739.37
A_24_P14	NM_03199	Williams-B	WBSCR1,E	-1.77359	0.00059	0.56383	491.5413	274.9515
A_32_P226356		Unknown	THC18095	-1.77385	4.51E-07	0.56375	376.4702	211.9933
A_23_P28	NM_00259	Homo sapi	PCNA,MG	-1.77409	2.36E-19	0.56367	43065.28	24141.23
A_23_P10	NM_01595	thioredoxin	TXNDC14,	-1.77415	3.55E-29	0.56365	5072.613	2853.063
A_23_P39	AK021772	Hypothetic	FLJ11710	-1.77489	9.24E-07	0.56342	307.2542	173.2542
A_23_P35	AB046773	KIAA1553	KIAA1553	-1.77552	2.27E-17	0.56322	2560.563	1446.339
A_32_P204205		Unknown	THC19329	-1.77579	1.10E-10	0.56313	496.1301	279.4695
A_23_P41	NM_01426	Homo sapi	MAPRE2,E	-1.77581	1.04E-23	0.56312	835.2102	470.2802
A_24_P29	CD703575	EST20102	CD703575	-1.77838	5.15E-20	0.56231	1354.273	762.5449
A_23_P67	NM_00320	Homo sapi	TCF3,E2A,	-1.77869	1.63E-20	0.56221	4274.293	2398.468
A_23_P33	NM_05716	Homo sapi	GIT2,CAT-	-1.77938	3.50E-24	0.56199	2841.443	1597.704
A_32_P99131		Unknown	A_32_P99	-1.77958	0.0004	0.56193	255.6908	144.6327
A_23_P21	NM_00190	Homo sapi	CTPS	-1.77961	2.57E-24	0.56192	10980.77	6187.793
A_23_P15	AL117509	KIAA1666	KIAA1666	-1.77999	8.44E-11	0.5618	1050.635	591.3191
A_24_P409471		Unknown	A_24_P40	-1.78027	1.03E-34	0.56171	3347.873	1878.912
A_24_P255845		Unknown	A_24_P25	-1.78064	0.00022	0.56159	415.0198	232.1017
A_23_P10	AK000900	Hypothetic	FLJ10038	-1.78068	4.35E-32	0.56158	2489.818	1397.654
A_32_P10	BF911038	SAR1a ger	SARA1	-1.78084	2.77E-22	0.56153	30944.64	17419.38
A_24_P55	NM_00016	gap junctio	GJA1,ODD	-1.78192	1.35E-09	0.56119	2228.309	1257.245
A_24_P221285		Unknown	A_24_P22	-1.7824	1.61E-10	0.56104	1505.438	846.4653
A_23_P16	NM_02407	Homo sapi	DDX54,DP	-1.78245	1.07E-23	0.56103	5643.445	3174.151
A_24_P930337		Unknown	A_24_P93	-1.78289	0.00052	0.56089	226.0912	126.5167
A_23_P25	NM_00539	pyruvate d	PDK3	-1.78545	0.00005	0.56008	667.415	373.1261
A_24_P827096		Unknown	THC18321	-1.78612	1.87E-08	0.55987	2032.041	1145.647
A_32_P49	NM_02192	hypothetic	FLJ13220	-1.78653	0.00099	0.55975	619.3296	343.816
A_23_P16	NM_14471	Homo sapi	KLHL23,M	-1.78705	4.16E-10	0.55958	819.0016	459.5587
A_23_P12	NM_03286	Homo sapi	NUDCD1,C	-1.78717	1.01E-19	0.55954	2330.264	1298.724
A_23_P34	NM_01502	Homo sapi	POP1,KIA/	-1.7873	2.05E-17	0.5595	2658.195	1486.487
A_23_P30	NM_03334	Homo sapi	TRIM7,GN	-1.78765	8.66E-34	0.55939	5215.22	2920.526
A_23_P75	NM_03234	hypothetic	NUDT22,M	-1.78775	1.70E-19	0.55936	3139.121	1747.818
A_23_P28	NM_03190	mitochondr	MRPS5,Mf	-1.78863	2.26E-25	0.55909	3089.396	1723.457
A_32_P15	NM_01425	solute carri	SLC25A13	-1.78978	1.03E-16	0.55873	2310.795	1293.235
A_24_P29	NM_18180	Homo sapi	UBE2C,UE	-1.79119	3.22E-26	0.55829	22410.75	12479.01
A_23_P206598		AY184957	THC19654	-1.79125	4.96E-09	0.55827	3685.246	2070.414
A_24_P13	NM_02237	ATP-deper	TOR3A,AD	-1.79141	1.43E-08	0.55822	2048.743	1136.891
A_23_P21	NM_00163	Homo sapi	AMD1,AMI	-1.79142	5.30E-10	0.55822	10690.78	5986.86
A_23_P12	NM_00252	Homo sapi	NPTX1,NP	-1.7915	3.72E-24	0.55819	956.434	534.0422
A_23_P253701		AF516696	THC20051	-1.79191	2.60E-11	0.55806	580.2112	323.6914
A_23_P38	NM_02429	hypothetic	PHF23,MC	-1.79198	4.76E-23	0.55804	2780.646	1554.855
A_32_P74814		Unknown	THC18592	-1.79218	0.00026	0.55798	254.9264	141.5803

A_23_P165819	Unknown	A_23_P165819	-1.79231	1.96E-31	0.55794	192994.5	107459.9	
A_24_P482017	U44751 py	THC18660	-1.79332	6.05E-08	0.55762	736.8441	409.0946	
A_24_P161	NM_18182	Homo sapi	NF2,ACN,5	-1.79512	0.00075	0.55707	256.9132	141.7871
A_24_P39	NM_00166	aryl hydroc	ARNT,HIF	-1.7953	2.88E-15	0.55701	464.5796	258.7625
A_24_P75994	Unknown	A_24_P75994	-1.79539	9.68E-31	0.55698	1832.076	1019.667	
A_23_P42	NM_19848	Homo sapi	RPL7L1,M	-1.79569	7.18E-23	0.55689	4568.3	2532.617
A_23_P33	NM_00087	Homo sapi	IGF2R,MPI	-1.79613	7.73E-14	0.55675	2627.61	1466.503
A_24_P15	NM_13846	Homo sapi	ZMYND19,	-1.79715	1.02E-18	0.55644	1522.284	843.7708
A_24_P40	NM_01594	CGI-20 pro	C7orf20,C	-1.79769	2.03E-27	0.55627	2344.658	1301.756
A_32_P27327	Unknown	A_32_P27327	-1.79792	1.34E-32	0.5562	7246.269	4028.256	
A_23_P46	NM_00287	RAB13, me	RAB13,GIC	-1.79795	1.18E-07	0.55619	2718.76	1522.927
A_32_P94176	Unknown	THC19089	-1.79864	0.00005	0.55598	235.442	130.6964	
A_24_P178475	Unknown	A_24_P178475	-1.79931	0.00004	0.55577	518.1564	286.6335	
A_23_P16	NM_00599	Homo sapi	CCT3,CCT	-1.79968	5.76E-09	0.55566	112066	61146.26
A_23_P23	NM_00174	Homo sapi	CAPN2,mC	-1.7997	7.29E-43	0.55565	58494.77	32502.43
A_24_P27	NM_03228	hypothetic	MGC3207	-1.80018	2.14E-10	0.5555	931.4149	518.5862
A_23_P15	NM_01621	Homo sapi	C3orf18,G	-1.80213	4.67E-20	0.5549	1232.062	682.5172
A_23_P57	NM_01336	Homo sapi	PCOLCE2,	-1.80237	2.13E-39	0.55482	6497.694	3603.707
A_24_P127462	Unknown	A_24_P127462	-1.80254	1.77E-26	0.55477	13555.96	7501.209	
A_32_P45	NM_00662	FUS-intera	FUSIP1,N	-1.80293	3.42E-26	0.55465	7164.492	3983.972
A_23_P14	NM_00278	Homo sapi	PSMA3,HC	-1.80334	1.15E-33	0.55453	36489.51	20265.93
A_23_P41	(AB040955	KIAA1522	KIAA1522	-1.80426	1.70E-20	0.55424	4927.745	2741.164
A_23_P13	NM_00210	H2A histon	H2AFZ,H2	-1.80609	1.67E-27	0.55368	81546.98	44998.14
A_24_P850336	Unknown	A_24_P850336	-1.80694	1.99E-39	0.55342	38039.95	21064.63	
A_23_P33	NM_18275	hypothetic	FAM79A,F	-1.8076	0.00042	0.55322	317.6386	173.4419
A_24_P91	BX108121	Transcribe	BX108121	-1.8084	2.97E-16	0.55297	663.944	367.0958
A_23_P21	NM_00191	Homo sapi	CYB561	-1.80847	5.00E-31	0.55295	2685.341	1485.708
A_23_P16	NM_03079	EGFR-coal	ECOP,FLJ	-1.80912	4.65E-11	0.55276	548.4099	302.9415
A_23_P20	NM_05284	Homo sapi	WDR34,M	-1.80929	1.14E-14	0.5527	5875.951	3272.183
A_32_P26	NM_00641	1-acylglyce	AGPAT2,B	-1.80943	0.00005	0.55266	318.6866	177.2927
A_23_P15	NM_00708	Homo sapi	NUDT6,gf	-1.80998	1.85E-15	0.55249	2266.212	1255.326
A_24_P89	NM_03282	hypothetic	C9orf3,AP	-1.81088	5.31E-34	0.55222	1401.035	773.7697
A_24_P24	NM_00462	translin	TSN,TRSL	-1.81284	4.25E-19	0.55162	4461.361	2469.403
A_23_P39	BC009229	Hypothetic	LOC90120	-1.81336	8.95E-09	0.55146	2906.168	1588.836
A_24_P47	NM_02409	hypothetic	MGC5508	-1.81387	3.71E-37	0.55131	7906.373	4360.799
A_32_P125568	Unknown	THC18118	-1.81402	1.13E-17	0.55126	539.3076	297.297	
A_23_P31	NM_03338	Homo sapi	FAM78A,C	-1.81417	0.00071	0.55121	288.3491	158.4649
A_32_P23	NM_00737	TAR DNA I	TARDBP,T	-1.81449	8.57E-29	0.55112	44450.19	24541.68
A_23_P20	NM_00648	Homo sapi	DYRK2	-1.81669	6.19E-10	0.55045	2095.318	1149.222
A_23_P15	NM_03271	hypothetic	MGC1133	-1.81682	0.00006	0.55041	659.7014	361.459
A_23_P15	(NM_00684	hepatitis d	DIPA	-1.81752	2.40E-25	0.5502	4326.25	2371.286
A_23_P70	NM_03192	Homo sapi	REPS1,RA	-1.81899	9.32E-14	0.54976	5180.706	2858.857
A_24_P94	AB037811	Family with	FLJ11280	-1.819	1.56E-08	0.54975	494.1214	271.8882
A_24_P41	(NM_00210	H2A histon	H2AFZ,H2	-1.81906	3.65E-16	0.54973	103772.8	56583.44
A_23_P13	NM_01323	putative rib	RNASEN,F	-1.82039	3.73E-36	0.54933	13147.19	7229.006
A_23_P40	NM_00282	Homo sapi	PTPN3,PT	-1.82084	1.76E-13	0.5492	742.2969	407.8268
A_23_P20	NM_00443	Homo sapi	EML1,EMA	-1.82124	2.23E-22	0.54908	1173.25	644.4197
A_24_P38	NM_02403	hypothetic	MGC2803	-1.82134	1.88E-14	0.54905	2104.372	1160.083
A_23_P50	NM_00269	Homo sapi	POLD1,PC	-1.82203	2.97E-32	0.54884	3297.877	1807.775
A_23_P50	BC001528	Guanine nt	GNA11	-1.82226	5.04E-18	0.54877	9820.952	5406.941
A_32_P17	(NM_01335	CCR4-NO	CNOT7,CA	-1.8223	2.05E-13	0.54876	7311.391	3991.589
A_24_P20	U69127	Far upstre	FUBP3	-1.82323	4.76E-34	0.54848	4202.676	2306.929

A_24_P22	NM_02451	hypothetic	C16orf53,1	-1.82328	5.95E-18	0.54846	2329.43	1277.594
A_23_P33	NM_17370	Homo sapi	NM_17370	-1.82377	5.37E-09	0.54832	148430	82861.77
A_23_P41	NM_20350	Ras-GTPa	G3BP2	-1.82411	2.47E-16	0.54821	1083.303	594.7592
A_24_P528213		Unknown	A_24_P52	-1.82438	1.24E-17	0.54813	699.3704	383.2297
A_32_P14	NM_17812	Homo sapi	DLX1	-1.82539	4.67E-09	0.54783	480.907	263.1822
A_24_P203976		Unknown	A_24_P20	-1.82614	2.21E-16	0.5476	784.7671	429.4817
A_24_P57	XM_37490	Homo sapi	XM_37490	-1.82649	3.32E-17	0.5475	578.7745	316.885
A_24_P77	NM_01895	homeo box	HOXA10,P	-1.8269	7.22E-10	0.54738	552.7	302.3434
A_24_P600622		Unknown	A_24_P60	-1.82716	3.50E-33	0.5473	1864.832	1020.559
A_23_P87	NM_00683	ATP/GTP-t	HEAB,CLP	-1.82731	4.49E-12	0.54725	1138.784	624.8268
A_32_P62056		Unknown	A_32_P62	-1.82836	1.86E-06	0.54694	445.3469	243.1139
A_24_P24	NM_00648	dual-specif	DYRK2	-1.82838	2.24E-08	0.54693	540.5255	295.1245
A_24_P35	NM_00210	H2A histon	H2AFZ,H2	-1.8294	4.16E-23	0.54663	150923.2	81971.98
A_23_P10	NM_00142	Homo sapi	EMP2,XMF	-1.82967	7.33E-21	0.54655	9223.984	5072.092
A_23_P36	NM_01984	Homo sapi	SLC10A3,F	-1.83038	6.02E-22	0.54633	3739.286	2037.444
A_23_P60	NM_01617	Ubiquitin a	UBADC1	-1.83156	9.92E-19	0.54598	7657.425	4156.819
A_23_P32	NM_02459	Homo sapi	RHBDL6,R	-1.83226	8.29E-28	0.54577	2715.267	1483.741
A_24_P7380		Unknown	ENST0000	-1.83434	3.49E-28	0.54516	20179.76	10982.59
A_24_P48	NM_00683	ubiquinol-c	UQCR,071	-1.83453	5.13E-27	0.5451	1026.209	559.5992
A_24_P91	M12996	Human glu	G6PD	-1.83488	1.27E-17	0.54499	4783.603	2624.117
A_24_P12	NM_00175	caveolin 1,	CAV1,CAV	-1.83756	5.37E-15	0.5442	4532.823	2484.466
A_23_P73	NM_00031	protein S (PROS1,PS	-1.83933	0.00009	0.54368	487.9702	264.6328
A_23_P28	NM_01474	Homo sapi	SNX17,KIA	-1.83986	4.49E-07	0.54352	1221.256	657.2603
A_23_P38	NM_03282	Homo sapi	ZNF587,ZF	-1.83995	9.62E-12	0.54349	3694.936	2016.597
A_24_P37	AK093577	MKL/myoc	MKL2	-1.84018	0.00013	0.54342	3703.224	2041.471
A_24_P88	BC047753	Postmeioti	POM121	-1.84075	5.27E-06	0.54326	365.8951	197.7527
A_23_P34	NM_00286	RAB2, mer	RAB2,RAB	-1.84377	2.76E-15	0.54237	3864.192	2103.71
A_23_P96	NM_02071	Homo sapi	LRRC47,K	-1.84391	4.87E-18	0.54233	3793.194	2061.675
A_23_P21	NM_00534	Homo sapi	HMGB3,HI	-1.84414	5.24E-26	0.54226	17992.65	9724.479
A_23_P34	NM_01651	putative se	SCLY,SCL	-1.84549	1.67E-18	0.54186	2585.585	1398.079
A_24_P48	AF086257	ST3 beta-g	SIAT4B	-1.84572	3.16E-07	0.54179	1186.586	644.1187
A_23_P58	NM_00265	paired-like	PITX1,BFT	-1.84602	9.42E-17	0.54171	1897.451	1027.97
A_23_P36	BC012476	Hypothetic	LOC11623	-1.84894	5.02E-11	0.54085	530.2435	286.6952
A_24_P19	NM_00721	SEC63 pro	SEC63,ER	-1.84929	3.43E-12	0.54075	2295.583	1248.417
A_23_P14	NM_02407	Homo sapi	ZFYVE21,I	-1.84968	5.31E-25	0.54063	7727.251	4167.872
A_24_P928031		Unknown	A_24_P92	-1.85126	1.11E-15	0.54017	466.9503	252.2156
A_23_P12	NM_00213	Homo sapi	HMOX1,HC	-1.85141	4.31E-07	0.54013	106924.7	55809.11
A_23_P88	NM_02100	Homo sapi	NR2F2,AR	-1.85169	7.70E-40	0.54005	5567.393	3009.027
A_23_P14	NM_01812	hypothetic	DARS2,FL	-1.85183	4.25E-24	0.54001	2831.74	1525.184
A_23_P30	AK096403	Epithelial n	EMP2	-1.85308	1.71E-20	0.53964	1652.763	895.4522
A_24_P63	AK056245	Homo sapi	AK056245	-1.85496	1.43E-08	0.53909	299.4089	161.3942
A_24_P27	NM_00440	DNA fragm	DFFB,CAC	-1.85586	0.00001	0.53883	264.7742	142.6563
A_24_P69	NM_00429	ring finger	RNF14,AR	-1.85596	1.50E-31	0.53881	1659.194	893.1516
A_23_P59	NM_01635	solute carri	SLCO4A1,	-1.85656	4.40E-18	0.53863	2617.828	1417.481
A_23_P63	NM_00658	chromosom	C1orf2,CO	-1.85814	8.48E-36	0.53817	11110.66	5986.654
A_24_P594293		Unknown	A_24_P59	-1.86214	1.54E-14	0.53702	12236.09	6533.282
A_24_P27	NM_03281	hypothetic	FLJ14624,I	-1.86325	0.00021	0.5367	205.3561	110.1659
A_24_P94	BC024302	Homo sapi	BC024302	-1.86407	0.00016	0.53646	354.9595	189.6669
A_23_P15	NM_01333	Homo sapi	ALG5,bA4	-1.86437	4.07E-27	0.53637	14619.93	7866.885
A_23_P21	NM_00721	Homo sapi	SEC63,ER	-1.86447	1.99E-31	0.53635	3046.012	1631.913
A_23_P76	BC013353	Hypothetic	FLJ14166	-1.86504	1.66E-13	0.53618	437.7408	234.7062
A_24_P41	AL117555	SEC6-like	SEC6L1	-1.86559	3.95E-35	0.53602	2221.134	1189.799

A_24_P13:	NM_00596	nucleosom	NAP1L4,N	-1.86834	1.51E-14	0.53523	1173.184	627.091
A_23_P39:	NM_00674	Homo sapi	PSKH1	-1.87049	4.13E-35	0.53462	2117.756	1131.417
A_24_P66:	AL832146	Pre-B-cell	IPBX1	-1.87109	6.65E-27	0.53445	1144.846	611.7993
A_23_P25:	NM_01234	F-box only	FBXO9,FB	-1.87115	6.00E-36	0.53443	9580.347	5116.132
A_23_P10:	NM_01524	Homo sapi	CENTD2,A	-1.87173	5.34E-10	0.53427	640.1021	341.5344
A_23_P12:	NM_01813	Homo sapi	HRMT1L6,	-1.87327	4.68E-27	0.53382	3322.524	1778.851
A_24_P39:	NM_03195	MSTP028	KCTD10,U	-1.87481	4.90E-08	0.53339	557.5599	296.9861
A_32_P46700	Unknown	A_32_P46:		-1.87975	2.72E-12	0.53198	1477.398	780.9035
A_23_P13:	NM_17075	Homo sapi	CDYL,CDY	-1.88046	3.25E-14	0.53178	462.6083	246.0024
A_23_P26:	NM_00280	proteasom	PSMD3,S3	-1.88283	1.24E-39	0.53112	20892.92	11085.91
A_23_P15:	NM_00240	Homo sapi	MEST,PEC	-1.88357	7.05E-24	0.53091	1116.815	593.5382
A_23_P18:	NM_00419	procollager	P4HA2,4-F	-1.88382	5.43E-14	0.53084	388.4828	206.2251
A_32_P15:	AK022268	Transducin	TBL1XR1	-1.88464	5.99E-11	0.53061	759.1531	402.5364
A_24_P40:	NM_00721	SEC63 pro	SEC63,ER	-1.88622	3.29E-09	0.53016	923.1334	490.4825
A_24_P36:	NM_00272	protein phc	PPP6C	-1.88641	1.86E-15	0.53011	1175.933	624.2118
A_24_P69:	AK092810	CDNA FLJ:	AK092810	-1.88718	3.17E-22	0.52989	8017.754	4267.12
A_24_P711050	Unknown	A_24_P71:		-1.88844	0	0.52954	4062.96	2151.634
A_24_P17:	XM_37226	similar to	DLOC38990	-1.8889	3.17E-07	0.52941	498.5679	263.1738
A_24_P85:	BC036885	Nuclear fac	NFIX	-1.89054	1.28E-15	0.52895	772.9211	408.7485
A_24_P26:	BC014029	Ubiquitin p	UBE3C	-1.89138	5.23E-12	0.52871	514.3779	271.7574
A_23_P98:	NM_00132	Homo sapi	CSTF3,CS	-1.89224	3.08E-18	0.52847	5120.078	2718.149
A_23_P16:	NM_00307	Homo sapi	SMARCE1	-1.89349	1.65E-27	0.52812	9174.155	4838.108
A_23_P21:	AK000619	Two pore s	TPCN1	-1.89397	1.24E-30	0.52799	7014.3	3707.177
A_23_P93:	NM_01910	homeo box	HOXA5,HC	-1.89447	2.21E-31	0.52785	1111.016	586.6301
A_23_P13:	NM_03231	Homo sapi	C2orf7,PAI	-1.89524	8.39E-39	0.52764	2219.892	1171.011
A_24_P89:	NM_03316	Homo sapi	SURF4,ER	-1.89573	2.15E-15	0.5275	13760.21	7297.895
A_32_P12:	NM_01508	Homo sapi	SPG20,SP	-1.89623	8.04E-15	0.52736	604.6515	318.8326
A_23_P98:	NM_00027	Homo sapi	OAT,HOG/	-1.89665	3.36E-22	0.52725	6026.324	3194.05
A_32_P60:	NM_00551	homeo box	HLXB9,HB	-1.89695	8.74E-37	0.52716	1471.834	775.7974
A_23_P13:	NM_01214	Homo sapi	DTYMK,C[-1.89712	0	0.52711	7397.59	3899.934
A_23_P11:	NM_01649	Homo sapi	TBC1D7,FI	-1.89812	6.91E-18	0.52684	3042.9	1602.843
A_24_P29:	NM_01221	tripartite m	TRIM32,H[-1.89956	3.51E-42	0.52644	1928.433	1015.161
A_24_P11:	NM_00276	phosphorib	PRPS2,PR	-1.90012	1.79E-21	0.52628	2691.14	1421.052
A_32_P71:	NM_18252	Homo sapi	FAM19A4,	-1.90016	0.00003	0.52627	210.8555	111.0868
A_23_P14:	BC032221	KIAA0391	KIAA0391	-1.90243	2.29E-28	0.52564	1622.506	853.6868
A_23_P55:	NM_00447	flotillin 2	FLOT2,ES,	-1.90275	9.22E-27	0.52555	3454.843	1811.714
A_23_P96:	NM_00213	Homo sapi	RBMX,RNI	-1.90359	1.14E-28	0.52532	12533.55	6595.374
A_23_P33:	NM_01828	mitochondr	IMP3,BRM	-1.90416	2.03E-41	0.52517	12606.76	6629.958
A_23_P13:	NM_01984	Homo sapi	EIF4ENIF1	-1.90434	2.54E-33	0.52512	1307.755	686.3364
A_23_P40:	NM_01239	Homo sapi	PFAS,PUR	-1.90584	2.49E-14	0.5247	6305.804	3332.665
A_32_P112401	Unknown	THC19528		-1.90611	1.20E-10	0.52463	371.3541	194.9507
A_23_P13:	NM_00395	Homo sapi	SLC25A14	-1.90739	1.79E-23	0.52428	2803.189	1461.9
A_23_P34:	NM_03262	hypothetic	RBBP6,MY	-1.90835	1.84E-06	0.52401	642.8257	336.1096
A_24_P63:	NM_00128	adaptor-rel	AP1S1,AP	-1.90956	0.00071	0.52368	400.5645	210.5546
A_24_P40:	NM_01427	Homo sapi	OLFM1,AV	-1.90994	2.31E-06	0.52358	429.4741	224.1008
A_24_P26:	NM_17497	Homo sapi	CCNK	-1.91022	8.92E-08	0.5235	255.2759	133.6719
A_23_P10:	NM_00556	Homo sapi	LAMA5,KI/	-1.91058	4.44E-12	0.5234	2000.164	1040.193
A_24_P35:	NM_01401	HSPC009	HSPC009	-1.91103	2.79E-21	0.52328	8809.26	4628.313
A_23_P20:	NM_02020	Homo sapi	PRTFDC1,	-1.9113	5.94E-24	0.5232	2014.62	1055.439
A_23_P25:	NM_01863	Homo sapi	WSB2,SBA	-1.91257	2.80E-45	0.52286	12533.37	6555.165
A_23_P41:	AF108138	Chromosor	C15orf20	-1.91338	2.92E-17	0.52263	2584.023	1355.021
A_24_P30:	AK024670	GATA zinc	p66alpha	-1.91558	6.79E-16	0.52203	1120.743	586.9389

A_24_P35	NM_00275	mitogen-ac	MAP2K5,M	-1.91678	0.00005	0.52171	242.4248	126.9924
A_23_P20	NM_03232	Homo sapi	C1orf57,FL	-1.91769	2.14E-16	0.52146	3124.423	1635.297
A_23_P14	NM_00206	Homo sapi	GNA11,GN	-1.91771	1.73E-17	0.52146	1531.729	798.1881
A_24_P14	XM_06574	Homo sapi	ENST0000	-1.91939	4.06E-39	0.521	1665.21	867.4628
A_24_P40	AF131831	Full length	CDC42SE2	-1.9195	1.40E-45	0.52097	4055.648	2112.101
A_32_P35	AK092508	Methyl-Cp	MBD3	-1.92071	1.07E-18	0.52064	1029.314	536.6025
A_24_P22	NM_00295	RYK recep	RYK,JTK5,	-1.92091	3.65E-14	0.52059	768.0076	399.8062
A_23_P14	NM_00378	Homo sapi	B4GALT2,I	-1.92126	7.43E-44	0.52049	2244.328	1167.912
A_23_P20	NM_01790	Homo sapi	TMCO3,C1	-1.92359	7.57E-25	0.51986	2848.552	1482.775
A_23_P34	AB011166	SMC5 stru	SMC5L1	-1.92531	2.45E-12	0.5194	2095.511	1090.709
A_32_P22	NM_13928	Homo sapi	CDC26,C9	-1.92562	5.61E-45	0.51931	11176.93	5800.066
A_23_P39	NM_02265	Homo sapi	HOXC8,HC	-1.92672	0.00029	0.51902	282.4984	145.5387
A_23_P76	NM_17384	Homo sapi	GSC	-1.92921	5.96E-16	0.51835	1192.157	616.3743
A_24_P19	NM_02071	Homo sapi	LRRC47,K	-1.92994	4.46E-11	0.51815	314.7104	163.0627
A_24_P94	AK024870	Dual-specif	DYRK2	-1.93029	2.97E-17	0.51806	524.1566	271.5781
A_23_P13	NM_00466	Homo sapi	CDC23,AP	-1.93254	1.61E-25	0.51745	2485.78	1282.367
A_32_P18	NM_02162	guanine nu	GNB4	-1.93379	5.65E-16	0.51712	893.7925	462.7038
A_32_P18	BC033117	Homo sapi	BC033117	-1.93449	3.83E-31	0.51693	1507.613	778.4321
A_24_P84	AK096589	Hypothetic	LOC13224	-1.93599	3.22E-10	0.51653	537.0047	277.329
A_23_P32	NM_02504	hypothetic	C15orf20,F	-1.93604	1.50E-21	0.51652	809.4919	418.2126
A_23_P12	BC029424	Similar to	FLOC49386	-1.93713	0	0.51623	5750.643	2976.103
A_24_P24	NM_00404	ATP synth	ATP5A1,O	-1.93794	5.77E-23	0.51601	40701.38	20860.86
A_23_P36	NM_17812	SEC14 anc	SESTD1	-1.93904	0.00037	0.51572	387.6038	196.4088
A_23_P65	NM_00470	Homo sapi	CCNB2,Hs	-1.93917	3.03E-33	0.51568	18739.21	9677.504
A_23_P11	NM_03328	Homo sapi	TBL1Y,TBI	-1.94002	5.24E-40	0.51546	2003.502	1033.248
A_23_P11	NM_00448	Homo sapi	GCSH,GCI	-1.94021	0	0.51541	18617.09	9593.541
A_23_P87	NM_01240	Homo sapi	ANP32D,P	-1.94108	1.05E-11	0.51518	694.5945	357.7637
A_24_P48	AK021467	Pre-B-cell	IPBX1	-1.9414	4.32E-07	0.51509	344.1803	177.3885
A_24_P375550	Unknown	A_24_P37		-1.94267	2.74E-18	0.51476	458.9377	236.2302
A_23_P50	NM_00410	Homo sapi	EFEMP1,D	-1.94321	3.76E-35	0.51461	1480.362	761.1035
A_24_P921933	ALU5_HUM	THC18696		-1.94429	4.57E-09	0.51433	482.7653	248.4258
A_24_P144149	Unknown	ENST0000		-1.94511	5.20E-26	0.51411	2902.835	1495.042
A_24_P70	BC022928	Homo sapi	BC022928	-1.94544	0	0.51402	20719.98	10644.9
A_23_P30	NM_02101	H3 histone	HIST1H3F,	-1.94807	1.68E-15	0.51333	8309.317	4290.039
A_23_P74	NM_00176	Homo sapi	CD1D	-1.94925	1.15E-10	0.51302	380.9814	195.4375
A_23_P12	BC000665	T-complex	TCP1	-1.95266	8.49E-25	0.51212	67603.2	34802.48
A_23_P14	NM_00660	Homo sapi	TCFL5,CH	-1.95309	4.04E-41	0.51201	2442.907	1250.658
A_24_P31	NM_18251	Spindle pol	Spc24	-1.95412	1.96E-13	0.51174	958.2234	489.501
A_24_P41	AL833329	Similar to	FDKFZp686	-1.95423	0.00007	0.51171	248.5036	127.1446
A_32_P70	BC029038	Ring finger	RNF34	-1.95442	1.89E-21	0.51166	632.3988	323.5773
A_23_P99	NM_00602	Homo sapi	PNMA1,M/	-1.95699	1.66E-39	0.51099	13193.04	6730.531
A_24_P55	AF391805	Homo sapi	unknown	-1.95712	2.57E-30	0.51096	63099.94	32397.64
A_23_P25	NM_15274	Homo sapi	C7orf27,M/	-1.95912	0	0.51043	15936.3	8136.168
A_24_P72	NM_01531	la related	pLARP1,LA	-1.96289	1.47E-13	0.50945	1257.031	642.7763
A_24_P29	NM_19905	Homo sapi	C21orf25,Λ	-1.96643	0.00003	0.50853	286.5275	144.6866
A_23_P13	NM_00289	Homo sapi	RBM4,LAR	-1.97024	2.47E-20	0.50755	2136.575	1088.995
A_23_P46	NM_00168	Homo sapi	ATP5F1,M	-1.97161	2.91E-34	0.5072	22565.48	11492.9
A_24_P79	NM_00637	NS1-assoc	SYNCRIP,	-1.97318	1.84E-24	0.5068	21991.95	11210.89
A_32_P214011	Unknown	A_32_P21		-1.97366	3.07E-11	0.50667	874.1026	444.2609
A_23_P17	NM_00279	Homo sapi	PSMB2,HC	-1.97717	9.75E-40	0.50577	8178.533	4142.888
A_24_P21	NM_03129	hypothetic	CDCA3,GF	-1.97728	9.81E-45	0.50575	3013.352	1523.278
A_32_P130536	Unknown	THC19474		-1.97862	5.46E-06	0.5054	249.0871	125.7242

A_23_P50	NM_13916	Homo sapi	SMCR7,MC	-1.97892	6.64E-24	0.50533	757.7865	382.9778
A_24_P27	NM_02470	hypothetic	ASB13,FL	-1.98277	4.90E-44	0.50435	4694.08	2368.224
A_24_P34	AK092698	CDNA FLJ	AK092698	-1.98287	1.43E-08	0.50432	1698.455	860.3406
A_23_P47	NM_02456	Homo sapi	MPZL1,PZ	-1.98337	2.96E-29	0.50419	5450.35	2755.872
A_32_P139708	Unknown	THC18395		-1.98523	1.28E-07	0.50372	337.5693	169.9982
A_32_P80	AK026192	CDNA: FL	AK026192	-1.98559	1.60E-07	0.50363	960.8893	482.6632
A_23_P88	NM_00390	Homo sapi	CDC16,AP	-1.98751	1.01E-16	0.50314	2563.328	1282.506
A_32_P226700	AF258585	THC18482		-1.99021	9.74E-24	0.50246	709.6539	356.5209
A_24_P34	NM_00484	class I cyto	IL27RA,CF	-1.99316	1.85E-26	0.50172	632.7239	317.4404
A_23_P20	NM_03285	Homo sapi	MUM1,MU	-1.99578	2.36E-15	0.50106	1797.048	900.351
A_32_P22	BX115350	Transcribe	BX115350	-1.99685	0.00037	0.50079	204.9637	103.0459
A_24_P28	NM_00368	calcium/cal	CASK,LIN	-2.00111	1.54E-18	0.49972	1035.692	518.4987
A_23_P10	NM_00510	Homo sapi	RBM8A,Y1	-2.00126	8.90E-26	0.49969	80946.91	40198.16
A_32_P38	NM_18297	Homo sapi	RIMS4,RIM	-2.00203	5.65E-11	0.49949	3168.708	1572.933
A_24_P76	AK024037	Homo sapi	AK024037	-2.00311	1.80E-28	0.49922	6294.285	3153.535
A_23_P36	NM_15323	Homo sapi	CRI2,EID2	-2.0037	3.08E-26	0.49908	4017.713	2012.824
A_23_P16	NM_00670	neuropathy	NTE,sws	-2.00391	0.00023	0.49903	193.4583	96.23705
A_23_P10	BC000907	Vang-like 1	VANGL1	-2.01057	2.85E-19	0.49737	1145.975	569.5579
A_32_P123204	Unknown	A_32_P12		-2.01158	0.00001	0.49712	1821.918	877.467
A_24_P92	BC016950	Phosphata	MGC2610	-2.01317	0.00055	0.49673	400.7838	195.904
A_23_P25	NM_01791	hypothetic	FLJ20628,I	-2.0144	1.01E-16	0.49643	597.9614	296.8037
A_32_P71	AK124776	CDNA FLJ	AK124776	-2.01467	1.11E-09	0.49636	2269.957	1128.086
A_24_P17	BC025792	Splicing fac	SF1	-2.01535	2.01E-06	0.49619	469.1491	231.8155
A_23_P331087	Unknown	A_23_P33		-2.01559	3.23E-16	0.49613	4352.358	2171.916
A_23_P12	NM_00135	DEAD/H (A	DDX3X,DE	-2.01665	1.19E-23	0.49587	9611.649	4791.551
A_23_P25	NM_00588	Homo sapi	SLC25A3,F	-2.01757	1.40E-45	0.49565	129665.2	64149.77
A_23_P30	NM_01532	Homo sapi	KIAA0776,	-2.01773	1.11E-13	0.49561	2193.973	1087.793
A_24_P187154	Unknown	ENST0000		-2.01819	9.61E-23	0.49549	8257.927	4109.557
A_23_P21	NM_00153	HMT1 (hnF	HRMT1L1,	-2.01874	3.51E-12	0.49536	478.8552	237.0534
A_23_P82	NM_01425	solute carri	SLC25A13	-2.01886	0	0.49533	2358.901	1168.598
A_23_P35	NM_00272	Homo sapi	PPP6C	-2.01896	1.41E-33	0.4953	4190.758	2082.86
A_23_P34	NM_00361	Homo sapi	RAE1,MIG	-2.02103	5.08E-21	0.4948	2964.443	1461.395
A_24_P34	NM_00679	mortality fa	MORF4,C9	-2.02434	4.35E-36	0.49399	5924.707	2923.189
A_23_P25	NM_01780	hypothetic	FLJ20397	-2.02616	2.51E-22	0.49354	3409.901	1682.061
A_23_P34	NM_01329	Homo sapi	SAC3D1,S	-2.02892	4.04E-23	0.49287	7557.615	3696.552
A_24_P237896	Unknown	A_24_P23		-2.02954	7.33E-11	0.49272	533.4862	262.8483
A_23_P21	NM_14778	Homo sapi	CTSB,APP	-2.03142	2.40E-17	0.49227	5846.034	2893.778
A_23_P13	NM_00480	Homo sapi	NMT2	-2.03256	1.46E-07	0.49199	400.2418	196.3497
A_32_P138432	Unknown	THC19065		-2.03479	0.00015	0.49145	232.2405	114.5955
A_23_P97	NM_00023	lipase A, ly	LIPA,LAL,C	-2.03512	3.58E-38	0.49137	8413.397	4146.973
A_23_P20	NM_02451	Homo sapi	CYP2R1,M	-2.03632	1.26E-22	0.49108	2129.163	1042.252
A_24_P43	NM_02411	hypothetic	MAPKAP1	-2.03887	2.11E-07	0.49047	952.1256	464.5651
A_23_P24	NM_02467	hypothetic	FLJ23441	-2.03905	2.49E-25	0.49043	1344.585	660.714
A_23_P20	NM_14528	similar to h	LOC15119	-2.03983	2.12E-26	0.49024	4145.189	2039.826
A_23_P10	NM_00297	Homo sapi	SBF1,MTV	-2.04123	2.21E-11	0.4899	499.5004	244.6858
A_23_P47	NM_01655	seven tran	TM7SF3	-2.0416	1.50E-26	0.48981	6227.777	3037.347
A_23_P25	NM_00685	Homo sapi	LIAS,LAS,I	-2.04191	1.70E-12	0.48974	1523.235	756.2648
A_24_P307184	Unknown	ENST0000		-2.04454	6.08E-25	0.48911	14895.37	7337.745
A_23_P15	BC010144	Likely orthc	LARP	-2.04775	1.04E-30	0.48834	27399.04	13325.7
A_24_P714707	Unknown	A_24_P71		-2.04885	2.90E-20	0.48808	583.2824	284.6693
A_24_P12	NM_20350	Ras-GTPa	G3BP2	-2.04909	6.55E-12	0.48802	1420.071	697.3742
A_32_P84	NM_00637	NS1-assoc	SYNCRIP,I	-2.05051	9.88E-10	0.48768	8141.834	4008.655

A_23_P39	NM_15249	Homo sapi	MANEAL,F	-2.0508	3.06E-12	0.48761	904.1445	441.8205
A_23_P50	NM_03239	Homo sapi	MKI67IP,N	-2.05421	2.71E-29	0.48681	4961.552	2404.078
A_32_P41	BC037580	Pseudoger	BC037580	-2.05482	0.00044	0.48666	420.624	205.3556
A_23_P16	NM_02464	Homo sapi	C14orf169,	-2.05488	0	0.48665	6187.865	3011.849
A_23_P42	NM_00028	Homo sapi	PEX6,PAF	-2.05606	2.57E-22	0.48637	1645.547	797.2062
A_23_P21	NM_01428	Homo sapi	FREQ,FLU	-2.05709	0	0.48612	3300.308	1605.175
A_24_P60	AK056073	CDNA FLJ	AK056073	-2.05773	3.27E-06	0.48597	379.4956	184.247
A_24_P11	NM_03204	hypothetic	DKFZP564	-2.05788	2.77E-10	0.48594	1478.305	717.2752
A_32_P13	1342	U12390 be	THC18495	-2.05905	6.02E-08	0.48566	255.0869	123.9724
A_23_P11	NM_00635	Homo sapi	TIMM44,TI	-2.06088	1.24E-09	0.48523	1929.35	938.7679
A_23_P80	NM_03228	Homo sapi	FYTTD1,D	-2.06297	2.20E-17	0.48474	2226.071	1081.447
A_23_P15	NM_00375	eukaryotic	EIF3S9,PR	-2.0635	2.15E-33	0.48461	24263.75	11720.32
A_24_P67	NM_01782	hypothetic	FLJ20436,I	-2.06356	4.63E-17	0.4846	494.2712	239.4385
A_24_P56	NM_01910	beta-1,4 m	ALG1,HMT	-2.0654	2.10E-43	0.48417	1187.094	574.7296
A_23_P14	NM_15294	developme	DRB1	-2.06544	2.73E-43	0.48416	2621.216	1269.88
A_24_P91	BC009775	Multiple su	MuLK	-2.06546	0.00026	0.48415	633.457	314.9742
A_23_P34	NM_01779	Homo sapi	RPP25,FL	-2.06592	3.68E-30	0.48405	7604.986	3669.766
A_24_P37	NM_01774	hypothetic	ZDHC7,Z	-2.06668	5.43E-42	0.48387	3505.634	1692.315
A_24_P22	AF271776	Homo sapi	DC48	-2.06703	3.25E-29	0.48379	8333.636	4011.454
A_23_P10	NM_01401	Homo sapi	HSPC009	-2.06717	0	0.48375	19658.79	9506.447
A_23_P13	NM_00040	Homo sapi	EXT2,SOT	-2.07203	1.86E-39	0.48262	1897.721	916.3586
A_23_P66	NM_01572	gem (nucle	GEMIN4,H	-2.07716	1.12E-31	0.48143	8729.296	4186.105
A_24_P30	NM_05304	Homo sapi	NM_05304	-2.07958	2.03E-09	0.48087	250.1187	120.265
A_23_P37	NM_01623	Homo sapi	ANAPC5,A	-2.0827	3.08E-42	0.48015	5259.553	2521.069
A_32_P17	557	Unknown	A_32_P17	-2.08322	2.76E-09	0.48003	3055.099	1438.14
A_23_P25	NM_00094	Homo sapi	PRIM1,p4	-2.08366	0	0.47992	6447.188	3097.208
A_23_P38	NM_17801	Homo sapi	TUBB,M40	-2.08398	8.77E-17	0.47985	4039.616	1921.155
A_24_P30	6921	Unknown	A_24_P30	-2.08458	2.43E-25	0.47971	4719.514	2278.528
A_24_P12	NM_03137	heterogene	HNRPD,P	-2.08462	2.57E-17	0.4797	1239.034	595.6007
A_23_P15	NM_00719	chromatin-	SUPT16H,	-2.08536	3.36E-44	0.47953	2357.328	1130.672
A_23_P72	NM_03288	hypothetic	MGC1130	-2.08565	4.57E-27	0.47947	4641.396	2212.79
A_23_P99	NM_01639	butyrate-in	HSPC121,I	-2.08577	8.02E-36	0.47944	27722.97	13333.07
A_23_P21	NM_00231	Homo sapi	LIMK1,LIM	-2.09089	2.95E-06	0.47827	267.8467	127.9088
A_23_P66	NM_01572	Homo sapi	GEMIN4,H	-2.09105	5.51E-43	0.47823	6062.785	2902.323
A_32_P18	4746	Unknown	THC19927	-2.09125	4.51E-07	0.47818	404.3386	192.9394
A_32_P10	BX641009	Spermatog	SPATA5	-2.09282	2.17E-06	0.47782	476.4067	226.4159
A_23_P72	NM_03288	hypothetic	MGC1130	-2.09327	3.58E-32	0.47772	1370.328	653.9497
A_24_P45	BC032042	Homo sapi	BC032042	-2.09775	3.89E-15	0.4767	550.0076	261.983
A_23_P15	NM_03279	Homo sapi	ZDHC12,	-2.09892	1.82E-30	0.47643	1009.452	480.5441
A_23_P43	NM_00256	purinergic	P2RX5,P2	-2.09901	0	0.47641	1314.908	626.4182
A_23_P31	NM_01329	transforme	TRA2A,HS	-2.10215	7.64E-18	0.4757	932.9008	444.5
A_24_P38	NM_00284	protein tyrc	PTPRF,LA	-2.10301	2.14E-31	0.47551	7252.495	3434.102
A_32_P45	NM_00262	Phosphogl	PGAM1	-2.10452	2.04E-25	0.47517	22450.45	10591.36
A_23_P33	5190	Unknown	THC20013	-2.10549	4.68E-09	0.47495	438.0792	207.6168
A_24_P33	NM_00517	ATPase, H	ATP6V0A1	-2.11147	3.62E-21	0.4736	3971.48	1878.455
A_32_P16	NM_20683	Homo sapi	MORF4L1,	-2.11751	3.53E-32	0.47225	16121.88	7640.069
A_24_P77	NM_00145	filamin C, g	FLNC,ABP	-2.12588	4.58E-08	0.47039	923.5294	430.2679
A_23_P43	BC029662	Homo sapi	C20orf142,	-2.12758	2.41E-31	0.47002	1628.331	766.432
A_24_P19	NM_00568	ATP-bindin	ABCB6,AB	-2.12952	5.68E-28	0.46959	3101.774	1449.058
A_23_P44	NM_00440	Homo sapi	DFFB,CAD	-2.13049	9.02E-11	0.46937	344.6246	161.5659
A_24_P37	NM_00163	S-adenosy	AMD1,AMI	-2.13058	0	0.46936	21731.39	10199.52
A_32_P19	NM_00213	RNA bindir	RBMX,RNI	-2.13175	0	0.4691	4545.228	2132.611

A_24_P926450	ALU8_HUM	THC18475	-2.13369	7.31E-36	0.46867	1420.778	664.4771
A_23_P12	NM_00225	potassium KCNS3,KV	-2.13429	9.67E-07	0.46854	297.3446	138.6768
A_23_P36	NM_17371	Homo sapi NM_17371	-2.13569	0	0.46823	82458.03	38641.77
A_24_P27	NM_01653	apoptosis-r FLJ39616	-2.1364	5.81E-21	0.46808	1785.252	829.9733
A_23_P20	NM_01830	hypothetic FLJ11029	-2.13757	6.13E-11	0.46782	439.811	205.21
A_23_P25	NM_05817	Homo sapi PSAT1,PS	-2.13804	0	0.46772	26712.41	12475.3
A_24_P11	NM_00456	6-phospho PFKFB3,PI	-2.14212	1.19E-09	0.46683	3559.161	1656.624
A_23_P21	NM_01478	Homo sapi TRIM14,KI	-2.14454	3.94E-14	0.4663	1735.046	813.2651
A_24_P84711	Unknown	A_24_P84	-2.14498	1.41E-22	0.4662	2113.931	989.1417
A_23_P43	NM_00451	interleukin ILF3,DRBF	-2.14502	3.85E-17	0.4662	2300.889	1071.506
A_24_P25	NM_00359	cullin 2 CUL2	-2.14778	6.21E-28	0.4656	763.7575	355.4705
A_24_P12	NM_01866	papillomav ZNF395,PE	-2.14832	3.03E-43	0.46548	2257.688	1049.734
A_23_P33	NM_18337	Homo sapi ARRDC4,F	-2.15272	5.48E-12	0.46453	356.4233	165.7286
A_23_P35	AL832919	1-acylglyce AGPAT3	-2.15294	1.74E-35	0.46448	3143.474	1461.8
A_23_P36	NM_00461	transmemb TSPAN8,C	-2.15567	6.32E-35	0.46389	17438.25	8117.366
A_23_P39	NM_02245	Ring finger RFWD2	-2.1564	0	0.46374	6046.962	2807.501
A_32_P106246	Unknown	THC19718	-2.1571	8.72E-10	0.46359	1705.028	790.1803
A_24_P30	NM_02044	actin-relate ACTR3B,A	-2.15956	1.12E-09	0.46306	294.1664	136.0798
A_23_P32	BC064969	Acidic (leuc ANP32A	-2.16038	7.60E-35	0.46288	4346.309	2004.524
A_24_P40	NM_02483	hypothetic FLJ12443,I	-2.16171	3.62E-36	0.4626	933.4324	431.6643
A_23_P39	NM_00704	Homo sapi BTN3A2,B	-2.16529	1.65E-37	0.46183	1510.76	696.6762
A_24_P29	NM_00712	valosin-cor VCP,p97,T	-2.16701	4.69E-21	0.46147	6119.192	2795.488
A_32_P42	BC018676	Homo sapi BC018676	-2.16763	7.21E-08	0.46133	2214.916	1022.738
A_23_P54	NM_02182	Homo sapi NIF3L1,CA	-2.1711	4.48E-23	0.4606	6352.96	2916.246
A_23_P16	NM_00611	Homo sapi PRAME,M	-2.17232	1.56E-22	0.46034	3058.313	1417.933
A_23_P14	NM_01866	Homo sapi ZNF395,PE	-2.17357	1.80E-35	0.46007	1074.969	494.7086
A_24_P28	NM_00279	proteasom PSMB2,HC	-2.18275	0	0.45814	12102.63	5554.437
A_23_P48	NM_00682	transmemb CKAP4,p6	-2.18517	5.08E-19	0.45763	8320.01	3829.253
A_24_P349580	Unknown	A_24_P34	-2.18581	1.76E-19	0.4575	1439.549	658.2628
A_24_P94	BC032643	Synaptota SYNCRIP	-2.18738	1.26E-06	0.45717	444.6506	201.9302
A_32_P50	NM_03075	t-complex TCP1,CCT	-2.18978	0	0.45667	3885.232	1772.823
A_23_P12	BC030623	Homo sapi BC030623	-2.19317	1.63E-39	0.45596	1479.148	672.9546
A_24_P25	AB051504	SET domai SET7	-2.19802	4.20E-45	0.45495	1261.573	574.1476
A_24_P19	NM_13895	Homo sapi VANGL1,L	-2.20055	8.34E-21	0.45443	3308.96	1506.412
A_24_P701814	Unknown	A_24_P70	-2.20085	7.90E-06	0.45437	258.01	116.4555
A_24_P612890	Unknown	A_24_P61	-2.206	2.56E-06	0.45331	269.8066	121.7742
A_23_P39	NM_01335	CCR4-NO CNOT7,CA	-2.21034	7.86E-43	0.45242	4120.912	1858.895
A_24_P25	U90143	Butyrophili BTN3A2	-2.2145	6.06E-12	0.45157	939.192	425.7188
A_24_P69	NM_03078	mitochondr SLC25A32	-2.21465	2.77E-17	0.45154	2521.365	1137.656
A_32_P13	AK024240	Chromosor C13orf25	-2.21516	0	0.45143	1773.613	800.4854
A_23_P64	NM_01434	Homo sapi MTCH2,HS	-2.21556	0	0.45135	20838.04	9397.472
A_23_P15	NM_03310	Homo sapi PNPT1,OL	-2.22078	3.14E-31	0.45029	7553.75	3418.87
A_23_P25	NM_01641	Homo sapi NF2,ACN,S	-2.22155	1.46E-41	0.45014	6617.903	2986.855
A_24_P14	XM_37167	similar to p LOC38917	-2.22167	1.83E-17	0.45011	574.3898	258.4428
A_23_P65	NM_13337	hypothetic MGC4562,	-2.22734	1.02E-29	0.44897	3310.158	1480.931
A_24_P90	NM_01849	G protein-c LGR4,GPR	-2.22857	0	0.44872	6792.264	3052.364
A_23_P12	NM_00642	Homo sapi MRPL28,p	-2.22868	5.01E-18	0.4487	9770.707	4345.323
A_23_P13	NM_01450	malonyl-Cc MT,MCT,M	-2.23112	1.48E-30	0.4482	1464.667	654.2673
A_24_P37	NM_00458	RAB27A, n RAB27A,G	-2.23125	1.79E-17	0.44818	391.4812	175.4506
A_23_P42	NM_03277	hypothetic FLJ14397	-2.23354	1.75E-12	0.44772	660.2221	295.0667
A_23_P92	NM_02526	Homo sapi TSEN2,SE	-2.23382	3.37E-16	0.44766	5561.826	2470.324
A_24_P59	BC013074	Mitochondr MCART1	-2.23569	1.12E-15	0.44729	638.3827	285.3729

A_23_P80	NM_02487	hypothetic	FLJ12748	-2.23632	0	0.44716	1178.392	526.7952
A_23_P59	NM_01404	Homo sapi	MRPS18B,	-2.23754	0	0.44692	24243.73	10833.03
A_32_P17	NM_00290	reticulocal	RCN2,E6B	-2.2444	2.67E-29	0.44555	12865.07	5755.896
A_23_P30	NM_14458	Homo sapi	C1orf59,FL	-2.24704	8.63E-11	0.44503	2053.796	912.9913
A_23_P15	NM_02499	Homo sapi	GFM1,EFC	-2.24749	1.40E-45	0.44494	3472.5	1546.74
A_24_P33	NM_01220	heterogene	HNRPH3,2	-2.24866	1.30E-35	0.44471	4300.628	1918.054
A_23_P12	NM_00121	Homo sapi	CALU	-2.24973	1.09E-24	0.4445	1984.188	885.3013
A_23_P78	NM_01608	CGI-150 pr	C17orf25,t	-2.25082	2.91E-28	0.44428	7698.071	3438.5
A_24_P22	BC064492	Ubiquitin-c	UBE2I	-2.25273	5.43E-12	0.44391	1487.924	659.2163
A_24_P25040	Unknown	A_24_P25		-2.25298	1.96E-12	0.44386	13850.8	6071.303
A_24_P12	NM_00135	DEAD/H (A	DDX3X,DE	-2.25768	0	0.44293	3059.917	1355.372
A_24_P19	NM_03321	tripartite m	TRIM14,KI	-2.26096	4.77E-14	0.44229	1341.311	594.7585
A_32_P161755	Unknown	A_32_P16		-2.2633	1.51E-06	0.44183	284.2322	125.9608
A_23_P13	NM_05283	Homo sapi	C6orf192,d	-2.26492	1.31E-30	0.44152	5860.64	2576.516
A_23_P77	NM_02451	Homo sapi	C16orf53,Λ	-2.26581	0	0.44134	27001.97	11897.27
A_23_P74	NM_01432	Homo sapi	OPN3,ECF	-2.26738	3.85E-10	0.44104	1588.422	702.9977
A_24_P82	AK092875	CDNA FLJ	AK092875	-2.27361	4.69E-14	0.43983	375.2612	165.1708
A_23_P10	NM_14516	Homo sapi	PIGM,MGC	-2.27407	2.10E-09	0.43974	1124.495	498.332
A_24_P38	NM_00350	frizzled (Dr	FZD1	-2.27515	1.15E-29	0.43953	953.5396	418.6635
A_32_P13	AB007950	Transmem	TMCC2	-2.27532	1.07E-08	0.4395	1385.653	606.386
A_23_P31	AB028999	KIAA1076	KIAA1076	-2.27766	0.00036	0.43905	321.238	138.3159
A_24_P14	NM_01401	dexametha	DEXI,MYLI	-2.28758	1.92E-27	0.43714	8507.424	3723.022
A_24_P221485	Unknown	A_24_P22		-2.28771	1.40E-44	0.43712	1183.43	517.3111
A_24_P20	NM_03056	hypothetic	C6orf148,Λ	-2.28857	2.73E-17	0.43695	436.2451	190.506
A_23_P31	NM_01864	Homo sapi	CHST12,C	-2.29271	4.85E-26	0.43617	1186.656	518.1835
A_23_P88	BC013943	Hypothetic	MGC24381	-2.29333	0	0.43605	4793.585	2089.978
A_23_P73	NM_02409	Homo sapi	ELOVL6,F	-2.29363	2.20E-35	0.43599	2271.075	989.3865
A_23_P25	NM_01234	Homo sapi	FBXO9,FB	-2.29916	8.01E-33	0.43494	5937.092	2571.087
A_23_P94	NM_01450	Homo sapi	TOR1B,DC	-2.30073	5.65E-26	0.43464	829.1321	360.7357
A_23_P33	NM_00307	SWI/SNF r	SMARCE1	-2.30208	8.02E-31	0.43439	1239.582	539.1132
A_32_P22	BG436185	602508593	BG436185	-2.30676	2.66E-44	0.43351	10307.92	4481.661
A_23_P96	NM_02071	Homo sapi	LRRC47,K	-2.31321	9.35E-42	0.4323	2606.909	1129.43
A_23_P87	BC004815	Hypothetic	MGC5139	-2.31396	1.76E-23	0.43216	2465.656	1060.885
A_32_P22	NM_00213	RNA bindir	RBMX,RNI	-2.31469	1.12E-32	0.43202	4657.151	2019.212
A_32_P45	NM_03075	t-complex	TCP1,CCT	-2.31669	0	0.43165	8954.625	3861.011
A_24_P32	BC015689	Collagen, t	COL9A3	-2.31876	5.81E-15	0.43126	2467.299	1063.316
A_23_P28	NM_01620	cleavage a	CPSF3,CP	-2.32469	0	0.43016	3826.411	1641.677
A_23_P50	NM_02450	Homo sapi	LRFN3,MG	-2.32561	4.20E-45	0.43	2104.288	902.5368
A_23_P50	NM_00704	E1B-55kDε	HNRPUL1,	-2.32926	1.50E-31	0.42932	3980.03	1722.875
A_32_P119233	Unknown	A_32_P11		-2.33073	7.01E-07	0.42905	2037.328	846.5637
A_24_P20	AK074041	DnaJ (Hsp	DNAJC5	-2.33756	2.42E-24	0.4278	1105.131	473.3045
A_23_P67	NM_03273	hypothetic	LMNB2,LM	-2.33952	0	0.42744	5382.181	2304.623
A_23_P16	NM_02208	Homo sapi	TXNDC5,E	-2.34414	5.75E-44	0.4266	8324.371	3536.146
A_24_P40	NM_00645	synaptoner	SC65,NOL	-2.34595	2.76E-23	0.42627	1658.922	705.1071
A_23_P54	NM_00568	ATP-bindin	ABCB6,AB	-2.34772	6.98E-41	0.42594	8582.859	3641.875
A_23_P13	NM_05304	hypothetic	MGC14327	-2.36376	0	0.42305	4836.148	2045.767
A_23_P11	NM_01536	Homo sapi	SKIV2L2,K	-2.3638	0	0.42305	9177.499	3873.341
A_32_P18	NM_00082	gamma-glu	GGCX,VK	-2.36868	9.95E-21	0.42218	967.4735	407.5985
A_32_P92	NM_00681	stress-indu	STIP1,HOF	-2.37027	0	0.42189	2095.855	883.2996
A_32_P133038	Unknown	THC18645		-2.37495	9.89E-33	0.42106	660.2041	277.9622
A_23_P63	NM_01629	Homo sapi	HSPA14,H	-2.37602	0	0.42087	7556.185	3189.803
A_24_P41	NM_00180	centromere	CENPA	-2.38151	2.12E-37	0.4199	1514.818	637.1782

A_23_P69	NM_00177	Homo sapi	CD47,IAP,	-2.3818	0	0.41985	7315.958	3073.443
A_24_P36	NM_00320	Homo sapi	TCF3,E2A,	-2.38411	0	0.41944	5969.789	2502.671
A_24_P20	NM_01503	Formin bin	FNBP1	-2.38717	7.22E-16	0.41891	1024.337	429.6919
A_23_P39	NM_00184	Homo sapi	CNTN1,F3	-2.39665	5.55E-09	0.41725	423.7026	176.2679
A_23_P93	NM_01602	Homo sapi	TFB1M,CC	-2.40137	0	0.41643	1697.818	707.278
A_23_P38	NM_05824	Homo sapi	DNAJB6,M	-2.40182	1.00E-16	0.41635	661.0651	275.0477
A_32_P47285	ALU7_HUM	THC18713		-2.40609	9.34E-31	0.41561	698.8428	290.4211
A_23_P15	NM_02077	Homo sapi	TBC1D14,I	-2.40666	0	0.41551	2153.263	894.9534
A_23_P11	NM_01220	Homo sapi	HNRPH3,2	-2.40937	5.61E-45	0.41505	5061.349	2106.535
A_23_P25	NM_02162	Homo sapi	PRDM13,F	-2.41375	1.34E-16	0.41429	2567.459	1058.566
A_32_P28	AF074986	Full length	AF074986	-2.41377	4.88E-43	0.41429	758.2539	314.1398
A_23_P64	NM_01741	Homo sapi	HOXC13,H	-2.41514	3.48E-13	0.41405	538.9367	222.8421
A_32_P77977	A40201	art	THC19995	-2.41618	1.18E-42	0.41388	807.6481	334.2885
A_23_P23	NM_00368	Homo sapi	EXO1,HEX	-2.41765	0	0.41363	5798.22	2409.538
A_23_P13	NM_03131	carboxypep	CPVL,HVL	-2.41775	3.65E-20	0.41361	1436.286	592.6003
A_24_P12	NM_00457	RAB4A, m	RAB4A	-2.4258	4.23E-37	0.41224	2576.293	1066.098
A_23_P37	NM_00679	MORF-rela	MORF4L1,	-2.43249	3.32E-33	0.4111	7183.032	2959.004
A_23_P21	NM_00459	Homo sapi	SFRS10,TI	-2.4369	0	0.41036	27086.09	11133.77
A_24_P37	NM_02283	hypothetic	USP46,FL	-2.44438	4.52E-09	0.4091	277.1823	113.0764
A_24_P37	NM_01646	hypothetic	ORMDL1,L	-2.44557	2.13E-06	0.4089	232.2916	94.46548
A_24_P616082	Unknown	A_24_P61		-2.44976	2.82E-23	0.4082	1103.343	450.5841
A_24_P568645	Unknown	A_24_P56		-2.45218	3.24E-14	0.4078	2625.615	1072.149
A_23_P99	NM_01653	apoptosis-r	FLJ39616	-2.45352	0	0.40758	1234.74	503.1235
A_24_P13	AL831922	DKFZP586	DLG1	-2.45738	0	0.40694	2610.09	1064.109
A_24_P30	NM_00359	cullin 1	CUL1	-2.46173	0	0.40622	3386.556	1378.287
A_23_P76	NM_00500	Homo sapi	NDUFA9,N	-2.46871	0	0.40507	7599.358	3080.958
A_24_P35	NM_13083	Homo sapi	OPA1,NPC	-2.47057	5.08E-21	0.40476	1385.406	560.9412
A_23_P13	NM_00717	Homo sapi	STRAP,MA	-2.47425	0	0.40416	50742.66	20500.07
A_23_P40	D86973	GCN1 gen	GCN1L1	-2.48113	0	0.40304	19242.11	7754.462
A_24_P24	NM_01560	Homo sapi	WDR21A,V	-2.48351	1.35E-42	0.40266	2932.311	1180.018
A_32_P55414	ALU6_HUM	THC19988		-2.49636	2.23E-07	0.40058	1395.313	548.9785
A_23_P21	NM_00339	Homo sapi	WNT5A,hV	-2.4964	4.64E-21	0.40058	3302.104	1328.399
A_24_P33	NM_00717	unr-interac	STRAP,MA	-2.49985	4.14E-22	0.40002	9703.706	3918.575
A_23_P80	NM_00462	Homo sapi	WRB,CHD	-2.50239	8.52E-11	0.39962	2704.024	1074.813
A_23_P15	NM_02216	Homo sapi	XYLT2,XT	-2.50733	0	0.39883	4272.969	1708.19
A_32_P12	BU581060	KIAA0664	KIAA0664	-2.51224	2.33E-16	0.39805	312.2306	124.2885
A_23_P12	NM_02044	Homo sapi	ACTR3B,A	-2.51988	2.64E-24	0.39684	4460.758	1774.948
A_23_P70	NM_00689	Homo sapi	HOXA7,AN	-2.52261	8.01E-21	0.39641	2498.585	993.8191
A_24_P21	AF318321	Casein kin	CSNK1D	-2.52411	0	0.39618	1218.784	482.7451
A_24_P599496	Unknown	THC20035		-2.52801	0	0.39557	1670.854	661.1328
A_23_P10	NM_00643	Homo sapi	CCT2,CCT	-2.52957	0	0.39532	106271.7	41762.85
A_23_P11	NM_00496	Homo sapi	HDAC1,HC	-2.53521	3.16E-22	0.39445	10209.49	4029.731
A_23_P20	NM_00229	Homo sapi	LBR,PHA,L	-2.53711	4.49E-38	0.39415	46989.31	18632.62
A_23_P57	NM_03224	Plexin A1	PLXNA1	-2.53906	0	0.39385	4381.428	1725.228
A_32_P192853	Unknown	A_32_P19		-2.54	3.74E-09	0.3937	437.6613	173.198
A_23_P77	NM_00390	Homo sapi	APPBP1,H	-2.54201	0	0.39339	39454.45	15447.13
A_24_P92	S81524	RC1=NADI	RC1	-2.54544	1.54E-19	0.39286	12236.33	4736.013
A_23_P32	NM_01516	Peptidase	PMPCA	-2.54686	2.02E-31	0.39264	7836.645	3063.677
A_23_P10	NM_01490	Homo sapi	TMEM15,K	-2.54782	0	0.39249	2542.519	997.2194
A_24_P15	NM_00719	chromatin-	SUPT16H,	-2.54932	1.94E-26	0.39226	4135.313	1627.785
A_24_P61	BC000206	Homo sapi	BC000206	-2.5515	1.52E-12	0.39193	13968.75	5243.385
A_23_P16	NM_01844	PC326 pro	IQWD1,NR	-2.56034	8.09E-12	0.39057	1194.351	467.9712

A_23_P205584	AF212245	THC18895	-2.56136	1.20E-06	0.39042	353.5834	138.8067
A_23_P92	NM_02453	hypothetic FLJ12684	-2.5684	2.55E-06	0.38935	222.358	87.01894
A_23_P25	NM_00662	Homo sapi HBS1L,ER	-2.56846	2.09E-12	0.38934	4017.81	1564.287
A_24_P87	AK022110	Similar to FLOC49386	-2.57445	0	0.38843	1511.351	587.3909
A_32_P10	AK127572	Full-length AK127572	-2.57599	1.26E-19	0.3882	1535.286	592.5703
A_24_P108281	Unknown	A_24_P10	-2.58506	4.20E-44	0.38684	752.6724	291.1789
A_23_P59	NM_00359	Homo sapi CUL1	-2.59363	0	0.38556	4632.901	1782.933
A_23_P21	NM_00454	Homo sapi NDUFC2,B	-2.5979	0	0.38493	54118.49	20916.86
A_23_P25	NM_03213	Homo sapi KBTBD7,D	-2.59985	1.17E-18	0.38464	462.3368	177.9551
A_23_P35	AK095583	Similar to FAK095583	-2.60477	3.29E-08	0.38391	582.2373	221.8662
A_24_P36	NM_00500	NADH deH NDUFA9,N	-2.60605	0	0.38372	5533.051	2127.342
A_23_P20	NM_03234	Homo sapi C9orf125,M	-2.60686	0	0.3836	6085.48	2338.743
A_24_P50	AW511222	hd44d11.x AW511222	-2.60802	4.41E-14	0.38343	537.9198	205.9543
A_24_P11	AF289610	Clone pp9 unknown	-2.61574	8.72E-24	0.3823	512.4827	195.8133
A_32_P18	AY007110	Acidic (leuc ANP32A	-2.619	0	0.38183	5444.704	2081.979
A_24_P90	D28589	KIAA0114 KIAA0114	-2.62145	1.20E-32	0.38147	4410.819	1666.547
A_23_P38	NM_01647	Homo sapi C14orf100,	-2.6264	0	0.38075	6794.616	2582.706
A_32_P33213	Unknown	THC18645	-2.64161	1.53E-31	0.37856	992.7858	375.2035
A_23_P99	NM_01649	Homo sapi MRPL51,C	-2.65344	0	0.37687	32840.3	12363.5
A_23_P15	BC035645	G protein-c GPR125	-2.65435	0.00003	0.37674	159.8442	60.33765
A_23_P14	NM_00146	Homo sapi FZD2	-2.65697	1.31E-19	0.37637	991.9603	374.1923
A_23_P40	NM_03285	Homo sapi MUM1,MU	-2.67224	6.96E-08	0.37422	390.2853	143.4117
A_24_P28	NM_00238	membrane MCP,TLX,(-2.67887	1.74E-14	0.37329	1901.248	716.6133
A_24_P32	NM_00681	stress-indu STIP1,HOF	-2.68815	2.06E-17	0.372	637.8642	236.9048
A_23_P36	NM_03210	Splicing fac SRP46	-2.69768	0	0.37069	896.1233	332.1373
A_23_P15	NM_15269	Homo sapi C1GALT1C	-2.69938	0	0.37046	6198.898	2300.607
A_24_P59	AK094159	CDNA FLJ: AK094159	-2.69965	7.42E-28	0.37042	851.9638	315.9639
A_23_P69	NM_18276	Homo sapi SUMF1,FC	-2.70396	4.56E-14	0.36983	3374.348	1234.6
A_23_P33	NM_00364	Homo sapi HAT1	-2.70482	2.56E-30	0.36971	12387.36	4626.677
A_23_P36	NM_00457	RAB4A, m RAB4A	-2.72484	0	0.36699	3269.279	1199.767
A_23_P42	NM_00295	Homo sapi RXRA,NR2	-2.72742	1.48E-24	0.36665	6205.543	2288.591
A_23_P15	NM_00002	adenylosuc ADSL	-2.72896	0	0.36644	4673.406	1707.912
A_24_P31	NM_00503	polo (Dros PLK1,PLK,	-2.73322	0	0.36587	21616.97	7905.23
A_23_P32	NM_17382	mitochondr FLJ38991	-2.73366	0	0.36581	1611.685	590.2645
A_23_P10	NM_01458	Homo sapi SLC40A1,F	-2.73834	4.31E-19	0.36519	343.908	125.5558
A_23_P15	NM_01814	hypothetic C8orf35,FL	-2.74273	9.82E-42	0.3646	5056.57	1852.566
A_24_P10	NM_00210	H3 histone H3F3A,H3I	-2.75231	2.59E-35	0.36333	182067.7	65253.49
A_32_P10	AK057196	CDNA FLJ: AK057196	-2.75329	0.00018	0.3632	186.2682	67.37752
A_23_P12	NM_01234	Homo sapi GTPBP4,N	-2.76305	0	0.36192	15641.35	5650.881
A_23_P63	NM_01329	Homo sapi GPSM2,LC	-2.78828	6.40E-13	0.35864	621.0959	221.9133
A_32_P14	BC062473	Microsoma MGST1	-2.78892	1.44E-43	0.35856	7405.474	2660.026
A_23_P15	NM_01824	Homo sapi NEIL3,FPC	-2.81187	0	0.35564	1038.228	369.351
A_23_P40	NM_00434	carbamoyl- CAD	-2.81629	0	0.35508	2239.37	795.3547
A_24_P245108	Unknown	THC19435	-2.81773	6.54E-16	0.3549	504.3817	178.6432
A_23_P76	AK096242	Dynactin 2 DCTN2	-2.81878	0	0.35476	2892.325	1028.404
A_23_P15	NM_01434	Homo sapi FJX1,FLJ2	-2.82332	0	0.35419	2796.402	989.6598
A_23_P70	NM_17689	Homo sapi PPAP2A,P	-2.82801	0	0.35361	3487.499	1231.905
A_23_P37	NM_17497	Homo sapi CCNK	-2.85945	9.85E-29	0.34972	1834.219	644.6788
A_23_P38	NM_00503	peptidylpro PPID,CYPI	-2.89264	2.63E-33	0.34571	8664.331	3018.566
A_23_P28	NM_02462	hypothetic FLJ21901,I	-2.89367	0	0.34558	2963.718	1021.643
A_23_P12	NM_02453	hypothetic FLJ12118,I	-2.929	2.66E-33	0.34141	22237.7	7502.289
A_32_P198179	Unknown	A_32_P19	-2.92948	0	0.34136	3208.457	1092.914

A_23_P12	NM_01233 acyl-Coenz ACATE2,C	-2.93945	0	0.3402	6256.671	2135.513
A_23_P44	NM_02290 hypothetic FLJ12442	-2.95224	3.24E-30	0.33873	613.5308	207.6989
A_24_P32	NM_01808 hypothetic POLR3B,R	-2.96467	4.86E-35	0.33731	1600.823	540.3835
A_32_P17	NM_14526 similar to C LOC13739	-2.98266	2.27E-20	0.33527	358.9429	120.2684
A_23_P21	NM_01323 Homo sapi PDCD6,AL	-2.99362	0	0.33404	38194.59	12730.85
A_24_P13	XM_29302 similar to U LOC34438	-3.00861	0	0.33238	3516.904	1171.162
A_23_P25	NM_02409 defective ir DCC1,MG	-3.00954	0	0.33228	1129.085	373.7029
A_23_P39	NM_14460 hypothetic FLJ31153	-3.01178	0	0.33203	18413.2	6132.748
A_24_P29	NM_02053 Homo sapi C20orf3,BS	-3.01278	0	0.33192	757.577	251.4243
A_23_P11	NM_02453 hypothetic FLJ12118,I	-3.04274	1.81E-29	0.32865	540.5226	177.5562
A_32_P12	BE877371 Clone pp9 BE877371	-3.05645	3.38E-17	0.32718	372.2598	121.6033
A_23_P14	NM_00093 Homo sapi POLR2B,R	-3.06924	1.02E-23	0.32581	4388.935	1412.71
A_32_P14	2334 ALU1_HUI THC19337	-3.08716	9.40E-10	0.32392	222.157	72.09042
A_23_P11	NM_00379 Homo sapi C19orf2,RI	-3.09789	0	0.3228	2858.594	923.6879
A_23_P16	NM_01431 Trans-pren TPRT	-3.11003	1.09E-32	0.32154	2491.255	794.9889
A_24_P9	26993 unnamed p NP181386	-3.11671	0	0.32085	994.5159	319.0936
A_23_P10	NM_15271 Homo sapi ITM1,TMC	-3.11714	0	0.32081	5887.779	1891.374
A_23_P37	AK128205 DKFZP434 DKFZP434	-3.12026	0	0.32049	2419.524	773.868
A_23_P38	NM_00141 Homo sapi ELAVL1,HI	-3.14696	1.35E-22	0.31777	795.1875	252.553
A_23_P15	M37712 Homo sapi p58/GTA p	-3.15515	0	0.31694	4244.667	1350.698
A_23_P37	NM_01560 Homo sapi WDR21A,V	-3.16847	0	0.31561	3343.793	1057.434
A_24_P33	NM_00710 ubiquitin-lik UBL3,HCG	-3.17297	0	0.31516	2639.34	832.7036
A_24_P40	AK098759 Homo sapi LOC55420	-3.20506	5.22E-12	0.31201	585.4348	181.5747
A_32_P20	2913 Unknown A_32_P20	-3.21554	7.15E-09	0.31099	3348.042	967.0488
A_24_P76	AB007954 KIAA0485 KIAA0485	-3.22005	0	0.31055	1110.127	344.5294
A_23_P18	NM_00588 macrophag MAEA,EMI	-3.23036	0	0.30956	837.8378	259.3926
A_23_P12	NM_00152 Histone de HDAC2	-3.25027	0	0.30767	11025.27	3390.994
A_23_P56	NM_02477 Homo sapi GEMIN6,FI	-3.25109	0	0.30759	20060.62	6199.621
A_23_P83	NM_00204 Homo sapi GAS1	-3.25927	2.80E-22	0.30682	4251.248	1287.838
A_32_P10	NM_00294 ATP-bindin ABCE1,RL	-3.26959	3.72E-30	0.30585	3082.188	937.4766
A_23_P93	NM_00353 Homo sapi HIST1H3J,	-3.28724	0	0.30421	30619.13	9311.637
A_23_P13	NM_19913 hypothetic LOC13414	-3.31711	2.77E-10	0.30147	783.8302	233.7979
A_24_P26	NM_00553 inositol pol INPP5A,5F	-3.33297	1.84E-26	0.30003	630.7437	189.0579
A_23_P25	NM_14498 hypothetic ALG14,MG	-3.33365	0	0.29997	1604.033	481.8415
A_23_P25	NM_00248 Homo sapi NCBP1,NC	-3.34961	0	0.29854	5027.427	1498.217
A_24_P6	71842 PPOL_HUI THC19431	-3.38428	1.50E-39	0.29548	6245.355	1834.256
A_23_P87	NM_01808 hypothetic POLR3B,R	-3.43652	7.24E-37	0.29099	2228.204	651.6686
A_23_P20	NM_01333 hypoxia-inc HIG2	-3.45789	0	0.28919	4981.944	1435.914
A_24_P17	4063 Unknown A_24_P17	-3.5233	4.96E-27	0.28383	1248.617	354.4034
A_23_P18	NM_00291 Homo sapi RFC4,A1,F	-3.57143	0	0.28	20024.36	5568.8
A_23_P31	NM_17371 Homo sapi NM_17371	-3.57755	0	0.27952	6778.245	1903.491
A_23_P39	NM_01646 Homo sapi CXXC5,HS	-3.58658	0	0.27882	10419.48	2907.085
A_32_P23	3049 Unknown A_32_P23	-3.58759	3.04E-20	0.27874	853.0683	235.5847
A_24_P9	2 AK096975 Protein tyrc PTPN3	-3.61226	0.00001	0.27684	418.7156	110.9998
A_23_P42	AK095295 Homo sapi SFXN4,BC	-3.6184	1.07E-20	0.27637	311.3366	86.0016
A_32_P12	NM_00281 proteasom PSMD12,p	-3.63526	0	0.27508	8821.196	2430.185
A_24_P21	NM_00480 transmemb TM9SF2,P	-3.67076	0	0.27242	6334.175	1715.303
A_24_P38	NM_01642 cisplatin re CROP,LUC	-3.67877	0	0.27183	6652.747	1815.731
A_23_P94	NM_00209 general tra GTF2E2,FI	-3.69642	0	0.27053	3272.314	880.7581
A_32_P10	NM_14526 similar to C LOC13739	-3.7269	2.39E-31	0.26832	1159.851	312.3046
A_23_P55	NM_03274 hypothetic SFT2D3,M	-3.74441	1.55E-16	0.26706	302.38	80.63084
A_23_P69	NM_00135 Homo sapi DHX15,DB	-3.77301	0	0.26504	9205.891	2432.664

A_23_P10	NM_02053 Homo sapi	C20orf3,B	-3.788	0	0.26399	21503.11	5661.152
A_24_P92	NM_19913 hypothetic	LOC13414	-3.79754	2.80E-45	0.26333	615.0771	161.9784
A_23_P77	NM_00281 Homo sapi	PSMD12,p	-3.89609	0	0.25667	12704.94	3242.766
A_24_P33	NM_00588 macrophag	MAEA,EMI	-3.93771	0	0.25395	8759.037	2221.171
A_23_P20	NM_00141 ELAV (emt	ELAVL1,HI	-4.05168	0	0.24681	3548.696	877.8359
A_32_P62	NM_01849 PDZ-bindin	PBK,SPK,1	-4.06847	0	0.24579	9524.673	2340.163
A_23_P16	NM_00640 Homo sapi	ARL6IP5,J	-4.09228	0	0.24436	7064.572	1729.13
A_23_P41	NM_00294 Homo sapi	ABCE1,RL	-4.12427	0	0.24247	2569.803	625.0526
A_23_P25	NM_01595 2-deoxyrib	DERA	-4.20029	0	0.23808	12275	2919.993
A_24_P79	XR_00018: Homo sapi	XR_00018:	-4.5183	1.08E-39	0.22132	994.9663	219.8424
A_23_P10	NM_02307 Chromosor	C16orf28	-4.56117	0	0.21924	2568.176	563.397
A_23_P29	NM_01538 CGI-51 pro	SAMM50,C	-4.62341	0	0.21629	10147.3	2192.762
A_23_P51	NM_02067 Homo sapi	SPBC25,A	-4.68019	0	0.21367	7577.211	1622.858
A_24_P521544	Unknown A_24_P52		-4.6844	0	0.21347	1976.542	421.9834
A_23_P11	NM_01603 comparativ	UTP11L,C	-4.69639	0	0.21293	6913.769	1465.964
A_23_P45	NM_02236 Homo sapi	TFB2M,Hk	-4.71295	0	0.21218	11885.91	2513.712
A_24_P524562	Unknown A_24_P52		-4.73234	0	0.21131	21531.96	4552.867
A_23_P10	NM_00452 Homo sapi	MEOX1,M	-4.97961	2.17E-12	0.20082	794.249	169.1716
A_32_P56143	HSHU33 h THC19668		-5.03948	3.77E-11	0.19843	989.6415	187.4557
A_23_P70	NM_00353 Homo sapi	HIST1H3E	-5.19681	0	0.19243	25989.24	5041.698
A_23_P11	NM_00352 Homo sapi	HIST1H3A	-5.2376	0	0.19093	32811.48	6270.298
A_32_P20	NM_00210 H3 histone	H3F3A,H3I	-5.29533	0	0.18885	106689.6	20209.66
A_23_P11	NM_00161 Homo sapi	PARP1,PA	-5.30234	0	0.1886	28104.1	5321.275
A_32_P82179	Unknown THC18098		-5.30451	0	0.18852	2282.315	429.4548
A_32_P180958	ALU8_HUM THC19992		-5.34334	0	0.18715	4778.99	887.491
A_24_P135801	Unknown A_24_P13		-5.42598	0	0.1843	30456.71	5618.697
A_24_P30	NM_00602 D123 gene	C10orf7,D1	-5.43217	0	0.18409	5973.407	1098.97
A_32_P10	AW135423 UI-H-BI1-a	AW135423	-5.50098	1.39E-07	0.18179	409.635	72.86716
A_24_P25	NM_00640 vitamin A r	ARL6IP5,J	-5.5057	0	0.18163	10291.18	1865.988
A_24_P229756	Unknown ENST0000		-5.52656	0	0.18094	59244.81	10761.8
A_23_P23	NM_00210 Homo sapi	H3F3A,H3I	-5.66455	0	0.17654	145174.5	25759.2
A_24_P32	NM_00210 H3 histone	H3F3A,H3I	-5.67759	0	0.17613	139485.8	24556.55
A_32_P10856	Unknown THC18928		-5.68916	0	0.17577	5239.636	926.3623
A_23_P56	NM_01629 prenylcyste	PCYOX1,F	-5.81165	6.43E-10	0.17207	306.9498	52.331
A_23_P29	NM_00588 Homo sapi	MAEA,EMI	-5.89238	0	0.16971	16516.93	2805.651
A_23_P12	NM_03273 Homo sapi	RTN4IP1,N	-5.95733	0	0.16786	4937.315	826.0753
A_32_P15	NM_20737 FLJ45187	FLJ45187,	-5.96196	6.41E-22	0.16773	439.5295	73.2805
A_23_P49	NM_00692 Homo sapi	SFRS1,AS	-5.96484	0	0.16765	18675.62	3098.824
A_23_P21	NM_01542 Gene mod	DKFZP564	-6.06466	0	0.16489	11039.94	1826.267
A_32_P22	BX350880 Transcribe	BX350880	-6.15443	1.81E-15	0.16248	14957.59	2183.518
A_23_P11	NM_00281 proteasom	PSMD12,p	-6.30102	0	0.1587	1864.652	295.5582
A_24_P59	NM_00210 H3 histone	H3F3A,H3I	-6.41973	0	0.15577	62748.96	9748.145
A_23_P78	BC000892 Chromosor	C18orf55	-6.84034	4.27E-43	0.14619	10672.38	1535.31
A_23_P97	NM_00277 Homo sapi	HTRA1,L5	-7.03517	0	0.14214	8346.594	1180.969
A_24_P20	AK094466 TRNA splic	SEN54L	-7.05336	7.76E-16	0.14178	602.3548	84.2941
A_23_P10	NM_00294 Homo sapi	RPA1,HSS	-7.07319	0	0.14138	24009.68	3376.187
A_23_P50	NM_15363 Homo sapi	HOXA3,HC	-7.08814	0	0.14108	5329.335	752.6847
A_24_P94	NM_01629 prenylcyste	PCYOX1,F	-7.22241	0	0.13846	1435.588	196.7744
A_23_P86	NM_00687 UDP-GlcN/	B3GNT6,iC	-8.26239	0	0.12103	1123.573	134.9406
A_23_P15	NM_00480 Homo sapi	TM9SF2,P	-9.0991	0	0.1099	9498.295	1035.576
A_24_P74	NM_01417 HSPC154	C18orf55,	-13.03302	3.28E-20	0.07673	485.272	36.88374