

Protein Alignment Summary Table (Vpr)

Name(s)	Accession No.	Region	First Author	Reference
AGMTYO	X07805	complete genome	Fukasawa, M.	<i>Nature</i> 333 , 457-461 (1988)
AGM155	M29975	complete genome	Johnson, P.R.	<i>J. Virol.</i> 64 , 1086-92 (1990)
AGM3	M30931	complete genome	Baier, M.	<i>Virology</i> 176 , 216-221 (1990)
SIVREV	L40990	complete genome	Hirsch, V.M.	<i>J. Virol.</i> 69 , 955-967 (1995)
AGM677	M66437	complete genome	Fonsgaard, A.	<i>Virology</i> 182 , 397-402 (1991)
SABIC	U04005	complete genome	Jin, M.J.	<i>EMBO J.</i> 13 , 2935-2947 (1994)
U20965	U20965	vpr	Jubier-Maurin, V.	<i>J. Virol.</i> 69 , 7349-7353 (1995)
U20967	U20967	vpr	Jubier-Maurin, V.	<i>J. Virol.</i> 69 , 7349-7353 (1995)
TANI	U58991	complete genome	Soares, M.A.	<i>Virology</i> 228 , 394-399 (1997)

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VERVET.CON	MASG.....	RdPRRARRPGe?EIMDLsREPMdWLRDMLed?NgeAKLHFGRILLfGVWNYCOEE	GER?gtcPm?RRAVkyYrLYOKALFVHFRCGCRRRRQPFEPY...	EERRdGGGGGR	106
AGMTYO	110
AGM155	110
AGM3	110
SIVREV	110
GRIVET.CON	109
AGM677	109
SABAEUS.CON	127
SABIC	130
U20965	128
U20967	130
CONSENSUS.TAN	109
TANI	109
VERVET.CON	a?R...VPPGLd\$	114
AGMTYO	-N.....E	119
AGM155	PG.....E	119
AGM3	-G.....	119
SIVREV	-N.....A	119
GRIVET.CON	...APP-----	117
AGM677	...APP-----	117
SABAEUS.CON	GG.AaaAa---\$	137
SABIC	GG.AASAA---	140
U20965	GG.AAAA---\$	138
U20967	GG.ASAAA---\$	140
CONSENSUS.TAN	DG..REP-----A-	119
TANI	DG..REP-----A-	119