

hum pol gamma aa seq
 Pan troglodytes (chimpanzee)
 Macaca mulatta (rhesus monkey) pol gamma
 mus pol gamma
 rat pol gamma
 Xenopus pol gamma
 Drosophila pol gamma
 scyeast pol gamma mip1
 S. pombe pol gamma
 Neurospora pol gamma
 Consensus

hum pol gamma aa seq
 Pan troglodytes (chimpanzee)
 Macaca mulatta (rhesus monkey) pol gamma
 mus pol gamma
 rat pol gamma
 Xenopus pol gamma
 Drosophila pol gamma
 scyeast pol gamma mip1
 S. pombe pol gamma
 Neurospora pol gamma
 Consensus

hum pol gamma aa seq
 Pan troglodytes (chimpanzee)
 Macaca mulatta (rhesus monkey) pol gamma
 mus pol gamma
 rat pol gamma
 Xenopus pol gamma
 Drosophila pol gamma
 scyeast pol gamma mip1
 S. pombe pol gamma
 Neurospora pol gamma
 Consensus

1 * * * * 55
 (1) -MSRLWWRKVAGATVGP GPVPA PGRWVSSSVP ASDPSDG QRRRQQQQQQQQQQQQQ
 (1) -MSRLWWRKVAGATVGP GPVPA PGRWVSSSVP ASDPSDG QRRRQQQQQQQQQQQQQ
 (1) -MSRLWWRKLAGATVGP GPVPA PGRWVSSSVP ASDPSDG RRRQQQQLQQPQVP--
 (1) -MSRLWKKVAGAKVASGPVPA TARWVSSSVLDPVPSDG--RPPSQMP-----
 (1) -MSRLWKKVAGAKVASGPVPA TGRWVSSSVLDPVPSDG--QPQSQMP-----
 (1) ---MNRLLQKGTSLVPSWRTRGCRYRRCSYAPQLHAKPLEMETSQR-----
 (1) -MQFHLLRKYASKVSREHYASSSVKIFRRVKP PQKVNPKPKPENVENG-----
 (1) ---MTKLMVRSECMLRMVRRRPLRVQFCARWFSTKKN-----
 (1) ---MFYKACPSTLTC SKWIHSIIKTKKFYCRHYSSKS-----
 (1) MLTPVRCRTV PNATVATAARVLRRANLF SRYR PQLGHLRW DSTIAQ VLERKG---
 (1) MSRLWWRKVAGA V GPVPA RWVSSSVP PSDG Q

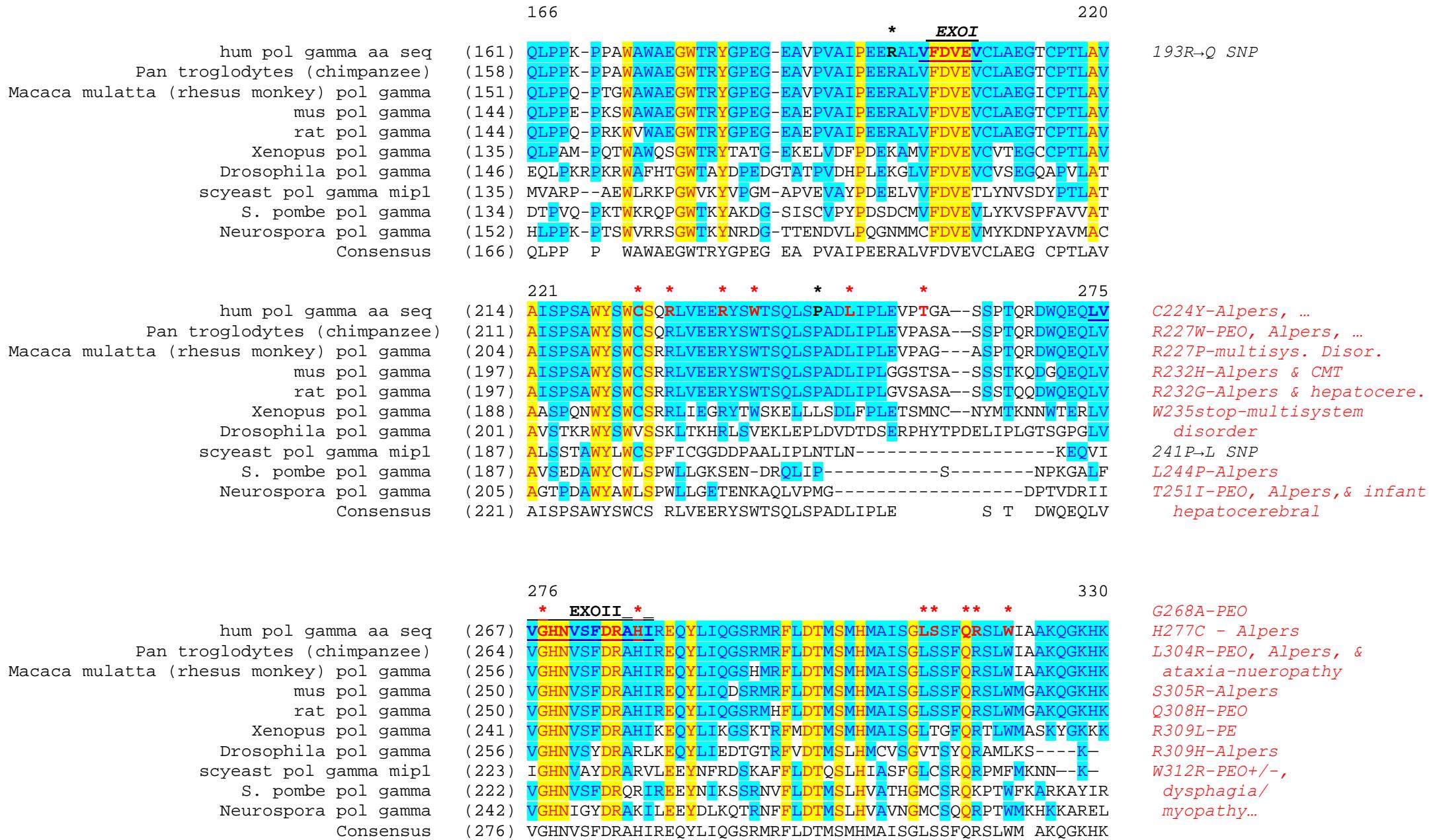
56 * * * 110
 (55) QPQQPQVLSSEGGLRHNPLDIQMLSRGLHEQIFGQGGEMPDE---AAVRSVE
 (53) -PQQPQVLSSEGGLQPRHNPLDIQMLSRGLHEQIFGQGGEMPDE---AAVRSVE
 (53) -----SSEGGLRHNPLHIQMLSKGLHEQIFGQGGEIPGE---AAVRSVE
 (46) -----SSEN GQLRLNPLLIQMLSRGLHEQIFGCGGEMPD E---AAVRSVE
 (46) -----SSEN GQLRLNPLHIQMLSRGLHEQIFGCGGDVPD E---AAVRSIE
 (45) -----MNPNIQMLSKGFHEQIFRGKQVQHAE---EDVQRSIT
 (48) -----PTEYAE NLVKVQMI SRNLHAQLF P QAPRSISEQQ-VASAKVYKD
 (35) -----TAEAPRINPVGIQYLGESLQRQVFGSCCGKDEVEQSDKLME LSKK
 (36) -----FIDNAPLRLINPVGVQYLSPALQNQVFPQ QNTQISQ---LHLDLAKF
 (53) -----LGVPSTARHNEIGVQQLSEHLYKQLF PRGNTDPAP---ELIELAKD
 (56) SSE GQLR NPL IQMLSRGLHEQIFGQGG P E AAV RS E

111 * * 165
 (106) HLQKHGLWGQPAVPLPDVELRLPPL YGDNL DQHF RLLAQKQSLPYLEAANLLIQA
 (103) HLQKHGLWGQPAVPLPDVELRLPPL YGDNL DQHF RLLAQKQSLPYLEAANLLIQA
 (96) HLQKHGLWGQPAAPLPDVELRLPPL YGDNL DQHF RLLAQKQSLPYLEAANSLLIQA
 (89) HLQKHGLWGQPAATPLPDVELRLPRLFGGNLDQHF RLLAQKQSLPYLEAAASLSEA
 (89) HLRKHGLWGQPTTPLPDVQLRLPRLFGGNLDQHF RLLAQKQSLPYLEAAASLSEA
 (80) HLNKHELGQETSTVPDVELQLPKMYGN NIEHFQI LAQKQSLPYLEAANDLLNC
 (91) ELRRHGVDISSA PVSDVQLKLPP LRGANIEEHFHNIAKEQVQFYEELLLPLVQC
 (80) SLKD HGLWGKKLITDPISFP LPPIQGRSLDEHFQKIGRFNSEPYKSF CEDKFT
 (79) HLA KHO LNLKETIKLPSFNFRLPLQOKTISEHFYNIGLEFAEPHILSKAIKFSKI
 (97) HLARHDLLGKTTDKTPPIAFQ LPA LVGDTLDEHFHKLGVDAAEPLTHAKQFADA
 (111) HL KHGLWGQPT PLPDVELRLPPL G NLD HFRLLAQKQSLPYLEAA LL A

R3P-PEO
 G11D-Alpers
 18P→S SNP
 Q43R-idiopathic
 sporadic Parkinsons
 49Q→E SNP
 ΔGlu 43-55 -male
 infertility

64S→L SNP
 Q68X - Alpers
 L83P - Alpers

H110Y-hearing loss, resp.
 failure, & failure to
 thrive
 A143V-Alpers, respir.
 enzyme chain def.



hum pol gamma aa seq
 Pan troglodytes (chimpanzee)
 Macaca mulatta (rhesus monkey) pol gamma
 mus pol gamma
 rat pol gamma
 Xenopus pol gamma
 Drosophila pol gamma
 scyeast pol gamma mip1
 S. pombe pol gamma
 Neurospora pol gamma
 Consensus

331 * * * * 385
 (322) VQ---PP**T**QGQKSQRKARRG--PAISSLWD**WLDI**SSVNSLAE**VHRLYVGGPPLKE**
 (319) VQ---PST**K**QGQKSQRKARRG--PAISSLWD**WLDI**SSVNSLAE**VHRLYVGGPPLKE**
 (311) VQ---GST**K**QGQKSQRKARRG--PAISSLWD**WLDI**SSVNSLAE**VHRLYVGGPPLKE**
 (305) TQ---QS**T**KRGQKS**PRK**-ANG--PAISSLWD**WMDI**S SANNLAD**VHNLYVGGPPLKE**
 (305) TQ---HP**T**KRGQKS**QKN**-ANG--PAISSLWD**WMDI**S SANNLAD**VHNLYVGGPPLKE**
 (296) GL---QE**V**KQH**I**KK**T**RSNFSG--SP**I**SSWD**WVN**I S INNLAD**VHALYVGGPPLKE**
 (305) -----KE-----PA**A**EDLG**W**LE**Q**SP**L**N**S**VE**V**HRLY**C**GGDT**L**IS**K**E
 (274) -----K**KE**AEVESEV**H**P**E**ISIEDYDDP**W**LN**V**SAL**N****S****I****K**D**V**AKFHCK-ID**L**D**K**T
 (277) S**Q**SETSETSEDDDS**S**FDDDYQ--NYL**K**QEP**W**LA**H****S**V**N****S****I****K**D**V**AKFH**C**N-IT**L**D**K**S
 (297) REKA**E**HESASVEL**Q**EVL**Q**GG**S**LT**A**EE**A**DL**W**V**D**K**S****I****N****S****L**R**D**V**A**Q**F**HLN-VK**I**D**K**D
 (331) Q TK GQKSQR G PAISSLWD**WLDI**SS NSLAD**VH** LYVGGPPLKE

hum pol gamma aa seq
 Pan troglodytes (chimpanzee)
 Macaca mulatta (rhesus monkey) pol gamma
 mus pol gamma
 rat pol gamma
 Xenopus pol gamma
 Drosophila pol gamma
 scyeast pol gamma mip1
 S. pombe pol gamma
 Neurospora pol gamma
 Consensus

386 * * * * 440
 (373) PRELFVK**G**TM**K**DIRENFQ**DLM**Q**YCA**Q**D**VWA**T**HEVFQQQLP**L**F**L**ER**C**PHP**V**T**L**AGM
 (370) PRELFVK**G**TM**K**DIRENFQ**DLM**Q**YCA**Q**D**VWA**T**HEVFQQQLP**L**F**L**ER**C**PHP**V**T**L**AGM
 (362) PRELFVK**G**TM**K**DIRENFQ**DLM**Q**YCA**Q**D**VWA**T**HEVFQQQLP**L**F**L**ER**C**PHP**V**T**L**AGM
 (355) PRELFVK**G**MSMRDIRENFQ**DLM**Q**YCA**Q**D**VWA**T**HEVFQQQLP**L**F**L**ER**C**PHP**V**T**L**AGM
 (355) PRELFVK**G**MSMRDIRENFQ**DLM**MEYC**A**RD**V**WA**T**HEVFQQQLP**L**F**L**ER**C**PHP**V**T**L**AGM
 (347) ARELFVK**G**MSDIRTEF**QEL**M**R**Y**C**AL**D**V**Q**A**T**HEVF**Q**E**Q**F**P**LF**M**ER**C**PHP**V**T**L**SGM
 (340) PRNIE**V**EG**T**GLE**QVR**QS**F**QSLTN**YCA**S**D**VE**A**THR**I**RL**V**LY**P**LY**A**ER**F**PHP**A**SL**L**AGM
 (321) DRDFF**A**STDK**S**T**I**EN**F**Q**K**L**V**N**Y**C**A**D**V**T**A**T**S**Q**V**F**D**E**I**F**P**V**F**L**K**K**C**PHP**V**S**F**AGL
 (329) KR**D**D**F**AS**L**E**K**E**P**L**Q**KL**N**E**L**ITY**Y**CA**H**D**T**Y**S**TH**Q**V**F**K**K**V**F**P**Q**F**L**E**V**C**P**PHP**A**T**F**S**A**M
 (351) IRDV**F**A**E**T**D**R**N**V**I**LN**Q**LD**D**L**T**Y**C**AA**D**V**Q**V**T**H**Q**V**Y**Q**V**V**F**P**N**F**L**G**V**C**P**HP**V**S**F**AL
 (386) PRELFVK**G** M DIRENFQ**DLM** Y**C**A DVWAT**H**E**V**F**Q**QQ**Q**L**P**L**F**ER**C**PHP**V**T**L**AGM

hum pol gamma aa seq
 Pan troglodytes (chimpanzee)
 Macaca mulatta (rhesus monkey) pol gamma
 mus pol gamma
 rat pol gamma
 Xenopus pol gamma
 Drosophila pol gamma
 scyeast pol gamma mip1
 S. pombe pol gamma
 Neurospora pol gamma
 Consensus

441 * * * * 495
 (428) LEMGV**S**YL**P**V**N**Q-NW**E**RY**I**LA**E**AQ**G**T**Y**E**E**EL**Q**REM**K**K**S****L****M****D****A**ND**A**QC**Q**LL**S**GERY**K**E
 (425) LEMGV**S**YL**P**V**N**Q-NW**E**RY**I**LA**E**AQ**G**T**Y**E**E**EL**Q**REM**K**K**S****L****M****D****A**ND**A**QC**Q**LL**S**GERY**K**D
 (417) LEMGV**S**YL**P**V**N**Q-NW**E**RY**I**LA**E**AQ**G**T**Y**E**E**EL**Q**REM**K**K**S****L****M****D****A**ND**A**QC**Q**LL**S**GERY**K**E
 (410) LEMGV**S**YL**P**V**N**Q-NW**E**RY**I**LT**E**AQ**Q**NT**Y**E**E**EL**Q**REM**K**K**S****L****M****D****A**ND**A**QC**Q**LL**S**GERY**K**E
 (410) LEMGV**S**YL**P**V**N**Q-NW**E**RY**I**LT**E**AQ**Q**ST**Y**E**E**EL**Q**REM**K**K**S****L****M****E****A**ND**A**QC**Q**LL**S**GERY**K**E
 (402) LEMGV**S**YL**P**V**N**Q-NW**E**RY**I**LT**E**QA**Q**TS**Y**E**E**EL**Q**REM**K**K**S****L****M****K****L****A**ND**A**QC**Q**LL**T**K**D**A**Y****K**E
 (395) LEMGSAY**L**P**V**N**S**-NW**E**RY**I**RE**A**QL**T**Y**E**EL**S****I****E**AK**Y**H**I**LG**R**RA**E****E****A**C**S****L****L****D****D****Q****Y****R****Q**
 (376) KSLSKCI**L**P**T**KL**N**D**W**ND**Y****I**NS**S****E****S****L****Y****Q**QSKV**Q**IES**K**IV**Q**I**I****K**D**I****V****L****L****K**D**P****D****F****Y****L**
 (384) LSLGSVF**L**P**V**N**H**-SW**T**RY**I**NG**V**EE**Q****Y****Q**QM**I**QL**V**D**Q**K**L****S****Q****Y****A****E****K****A****K****D****L****I****N****T****K**D**T****V****L**
 (406) RHLASVI**L**P**V**N**K**-TW**D**TY**I**ET**A**EA**T****Y****L**Q**M**L**H**GV**Q**ER**L**F**T****I****M****E****R****T****L****D****Y****K****A****D****P****E****K****Y****L**
 (441) LEMGV**S**YL**P**V**N**Q NW**E**RY**I** EA**Q** TY**E**EL**Q**REM**K****S****L** LAND**A**QC**Q**LL**S**GERY**K**E

324P→S SNP
 T326fsX387-Alpers
 W347_L356del-Alpers

R374X -Alpers
 G380D-PEO
 392L→V SNP
 L392V-muscle weakness,
 optic atrophy
 R417T - Alpers
 C418R - Alpers
 L424(ct)del452stop-PEO
 Alpers

L428P-Alpers,...
 M430L-PEO
 G431V-PEO
 S433C-PEO, ataxia, ...
 T452stop-PEO
 L463F-PEO
 A467T-PEO, SANDO,
 MERRF, & Alpers....
 N468D-PEO

hum pol gamma aa seq Pan troglodytes (chimpanzee) Macaca mulatta (rhesus monkey) pol gamma mus pol gamma rat pol gamma Xenopus pol gamma Drosophila pol gamma scyeast pol gamma mip1 S. pombe pol gamma Neurospora pol gamma Consensus	496 (482) DPWLWDLQEFKQKKAKKVKK EPAT ASKLPIEGAG AP GDPMDQED LGPC SEE (479) DPWLWDLQEFKQKKAKKVKK RVKKE EPAT ASKLPIEGAG AP GDPMDQED LGPC SEE (471) DPWLWDLQEFKQKKAKKVKK EPAT ASKLPIEGAG AP GDPMDQED LGPC SEE (464) DPWLWDLQEFKQKKAKKVKK -PAS ASKLPIEGAG PF GDPMDQED PGPP SEE (464) DPWLWDLQEFKQKKAKKVKK -TAS ASKLPIEGAG PF GDPMDQED PGPP SEE (456) DPWLWDLWDI QES KQKKTK ISKK --QKKANEAA ESVGNKLVEDHN ED PGPTEK (449) NL WLWDEDWSVQELKLKQPPKRKP ----- LPTVEIKDS (431) KDPWLSQLDWTTKPLRLTKKG VPAK ----- (438) KDPWLRLQLDWTPCNLYRK LKKATQE ----- (460) SDPWLSQLDWSGQEIKMAKP KKKG ----- (496) DPWLWDLQEFKQKKAKKVKK A ASKLPIEGAG GDPMDQED GP SEE	550 <i>Q497H-Ataxia-Neuropathy</i> <i>S511N-PEO</i> <i>G517V- Ataxia-Neuropathy</i>
hum pol gamma aa seq Pan troglodytes (chimpanzee) Macaca mulatta (rhesus monkey) pol gamma mus pol gamma pol gamma (518) Xenopus pol gamma Drosophila pol gamma scyeast pol gamma mip1 S. pombe pol gamma Neurospora pol gamma Consensus	551 (537) EEFQ QDVMARAC LQKLKGTT E LLPKR PQHLPGH PGWY RKLCPRL ----- (534) EEFQ QDVMARAC LQKLKGTT E LLPKR PQHLPGH PGWY RKLCPRL ----- (526) EEFQ QDVMARAC LQKLKGTT E LLPKR PQHLPGH PGWY RKLCPRL ----- (518) EELQ RSVTA HNR LQQLRSTTD LLPKR PQHLPGH PGWY RKLCPRL ----- (518) EELQ QNIMA HTRL LQQLKS TTD LLPKR PQHLPGH PGWY RKLCPRL ----- (509) EESRPSMGKLYLEDLKLK TLP LLPKR NQHLPGH PGWY RKLCPKL ----- (482) GNTPEERRLQAKF QHL YDQQALLPARRPLPGY PLWY RKLCKRKP PAKREDEILED (456) ----- CQLPGF PEWY RQLFPSK ----- (463) ----- VP ----- VVPK WYKKAYCKT ----- (484) ----- DVERPALNQKLPGY PQWY KDL FVKVPKELSGLDEPD (551) EE Q A LQ L TT LLPKR PQHLPGH PGWY RKLCPRL	605 <i>546R→C SNP</i> <i>K561M- dysmorphia, hypotonia, liver insufficiencies, rat</i> <i>R562Q-PEO</i> <i>H569Q - Alpers</i> <i>R574W-PEO+myopathy dysphagia & Alpers</i> <i>R579W-PEO</i>
hum pol gamma aa seq Pan troglodytes (chimpanzee) Macaca mulatta (rhesus monkey) pol gamma mus pol gamma rat pol gamma Xenopus pol gamma Drosophila pol gamma scyeast pol gamma mip1 S. pombe pol gamma Neurospora pol gamma Consensus	606 (581) --- DDPAWT PGP SLLS LQM RVT P KLM AL TWDGF P LHYS ER H GWGYL VPG R RDNL A (578) --- DDPAWT PGP SLLS LQM RVT P KLM AL TWDGF P LHYS ER H GWGYL VPG R RDNL A (570) --- DDPAWT PGP SLLS LQM RVT P KLM AL TWDGF P LHYS ER H GWGYL VPG R RDNL A (562) --- DDPAWA PGP SLLS LQM RVT P KLM AL TWDGF P LHYS DSH GWGYL VPG R RDNL T (562) --- DDPAWT PGP SLLS LQM RVT P KLM AL TWDGF P LHYS DSH GWGYL VPG R RDNL T (553) --- EDPDWL PGP EGL I SLQM RL TP KLM RL TWDGF P LHYS EKH GWGYL VPG R KNK NL (537) --- DEEP WSPG ASE I STGM QIA PK LLS LC WEG YPL HY EREQ GWGF L VPFR SDS -- (474) ----- DTVEPKITIKS RII PI LFKL SWENS PVI WS KES GWCFN V PHEQVET -- (478) ----- EKRAVITAKS RLA P II LR LKWK KH P LAWS DTY GWVFS VERTSKD -- (520) KEQENRKARHEFIN LTVRS RIA P L LK LSW EG YPL FW S DQF GWT FQV PR -- EK -- (606) DDPAW PGP SLLS LQM RVT P KLM AL TWDGF P LHYS HGWGYL VPG R RDNL	660 <i>P587L-PEO, Alpers, & inf. hepatocerebral</i> <i>P589L - Alpers</i> <i>R597W - PEO</i> <i>M603L-PEO w/ ptosis</i> <i>L605R - Alpers</i> <i>R617C-muscle weakness, exercise intolerance, hearing loss, arrhythmia</i> <i>R627W-PEO, Alpers, & SANDO</i> <i>R627Q-ataxia-myopathy, Alpers, PEO, ataxia-neuropathy, & MELAS</i>

hum pol gamma aa seq
 Pan troglodytes (chimpanzee)
 Macaca mulatta (rhesus monkey) pol gamma
 mus pol gamma
 rat pol gamma
 Xenopus pol gamma
 Drosophila pol gamma
 scyeast pol gamma mip1
 S. pombe pol gamma
 Neurospora pol gamma
 Consensus

661 * * 715
 (633) KLPTGTTLESAGVCPYRAIESLYRKHCLEQGKQQLMPOEAGLAEEFLLTDNSAI
 (630) KLPTGTTLESAGVCPYRAIESLYRKHCLEQGKQQLMPOEAGLAEEFLLTDNSAI
 (622) KLPTGTTLESAGVCPYRAIESLYRKHCLEQGKQQLTPQEAGLAEEFLLTDNSAI
 (614) EPPVSPTVESAAVTCPYRAIESLYRKHCLEQGKQQLEPQEVDLAEEFLLTDSSAM
 (614) ELFVSPTEESAAVTCPYRAIESLYRRHCLDQGKQQLETQETDLAEEFLLTDs-AM
 (605) NNEEEE-E--IIPCPYRAIEDIYA**EYSKNKT**DGCLSQHSTIPEEFMLTDDNSM
 (587) -----EGVDRLPMEQLLARCPVPEFARLSASKAESDMAFDMLP
 (520) -----YKAKN-----YVLADSV
 (522) -----EIEMLLD-QGLV
 (571) -----AETFIQRQMTP
 (661) P T ESA V CPYRAIESLYR HCL QGKQQL QE LAEEFLLTD A

hum pol gamma aa seq
 Pan troglodytes (chimpanzee)
 Macaca mulatta (rhesus monkey) pol gamma
 mus pol gamma
 rat pol gamma
 Xenopus pol gamma
 Drosophila pol gamma
 scyeast pol gamma mip1
 S. pombe pol gamma
 Neurospora pol gamma
 Consensus

716 * * * * 770
 (688) WQTVEELDYLEVEAEA**KMENLR**--AAVPGQPLALTARGGPKDTQFSYHHGNGPYN
 (685) WQTVEELDYLEVEAEA**KMENLR**--AAVPGQPLALTARGGPKDTQFSYHHGNGPYN
 (677) WQTVEELDYLEVEAEA**KMENLR**--AAVPGQPLALTAPGGPKDSQPNYHHGNGPYN
 (669) WQTVEELGCLDVEAEA**KMEN**---SGLSQPLVPLAACAPKSSQPTYHHGNGPYN
 (668) WQTVEELGCLDVEAEA**TVES**---SGLSQPLVPPTACAPKTSQPTYHHGNGPYN
 (656) WQKVEELSRTEMDLSSSEVPATAKKRNNNSSEHPVKLEMEDDSLFDNHHGNSPCCG
 (625) GQVEQHLGKREHYKKLSQKQ-----QR-----LETQYQGSGVWCN
 (532) SQEEEE-----IRTHN
 (533) PCSREE-----DT
 (582) VQFEDP-----DVDDR
 (716) WQTVEEL LEVEAEA E QPL PK QP YHHGNGPYN

hum pol gamma aa seq
 Pan troglodytes (chimpanzee)
 Macaca mulatta (rhesus monkey) pol gamma
 mus pol gamma
 rat pol gamma
 Xenopus pol gamma
 Drosophila pol gamma
 S. pombe pol gamma
 Neurospora pol gamma
 Consensus

771 * * * * 825
 (741) DV DIPGCWFFKLPHKDGNSCNVGSPFAKDFLPMEDGTLQA GPGGASGPRALEI
 (738) DV DIPGCWFFKLPHKDGNSCNVGSPFAKDFLPMEDGTLQA GPGGASGPRALEI
 (730) DV DIPGCWFFKLPHKDGNSCNVGSPFAKDFLPMEDGTLQA GPGGASGPRALEI
 (719) DVNIPGCWFFKLPHKDGNNYVGSPFAKDFLPMEDGTLQA GPGGASGPRALEI
 (718) DV DIPGCWFFKLPHKDGNNYVGSPFAKDFLPMEDGTLQA GPGGARGPRALEI
 (711) DV NVSGCWFYKLPHKDGNANVGSPFAKDFLPMEDGTLQA STGDSSATRALEI
 (660) KV LDDDCFFFLKLPHKNGPSFRVGNELSKDFLNKFAENVILSSGDPSCQAAARVIDI
 (541) KLDYNNYIFFKVPHKDGEARCVRHSPNHITHTSKRVFYNO---IMKWLLKALEM
 (593) LRMDVDHKYFKLPHKDGPNARCVNPMAKGYLPYFEKGILSS---EYPYAKEALEM
 (771) DV DIPGCWFFKLPHKDGN NVGSPFAKDFLPMEDGTLQA GPGGASGPRALEI

P648R-PEO+MYOPATHY &
 SANDO
 662E→K SNP

R709stop-PEO
 Q715X-Alpers
 c.2157+5_+6-Alpers
 722R→H SNP
 G737R- Alpers, CMT, &
 Parkinsonism

G746S-PEO
 W748S- Alpers & ataxia
 nueropathy
 F749S-Alpers
 G763R-PEO & SANDO
 A767D-Alpers
 2354ins(g)G785stop-
 PEO

Sequence alignment of Pol Gamma aa seq across various species. The alignment highlights mutations in three regions: 826-880, 881-935, and 936-990. Amino acid changes are color-coded (e.g., red for R807P). Consensus sequences are shown at the bottom of each region.

Region 1: 826 - 880

	826	*	**	*	880
hum pol gamma aa seq	(795) NKMI SFWRNAHKRISSQMVVWLPRSA LPRAVI RHPDYDE ---	EGLYGAILPQVVT			R807P-PEO
Pan troglodytes (chimpanzee)	(792) NKMI SFWRNAHKRISSQMVVWLPRSA LPRAVI RHPDYDE ---	EGLYGAILPQVVT			R807C- SANDO
Macaca mulatta (rhesus monkey) pol gamma	(784) NKMI SFWRNAHKRISSQMVVWLPRSA LPRAVI RHPDYDE ---	EGLYGAILPQVVT			c.2480+1-Alpers
mus pol gamma	(773) NKMI SFWRNAHKRISSQMVVWLPRSA LPRAVI RHPDYDE ---	EGLYGAILPQVVT			IVS15-9 c.2485del- Alpers
rat pol gamma	(772) NKMI SFWRNAHKRISSQMVVWLPRSA LPRAVI RHPDYDE ---	EGLYGAILPQVVT			
Xenopus pol gamma	(765) NKMI SFWRNAHKRISSQMVVWMKKNELHRTITRDP EFE-----	ENKYGAIILAQVVS			
Drosophila pol gamma	(715) ARMM SYWRNNRDRIMGQLVVWLDSQQLP NEFTG--EKCQ---PIA YGAICPQVVA				
scyeast pol gamma mip1	(595) NSSG SYWMSARERIQSQFVVPSCKFPNEFQSLSAKSSLNNEKTNDLAI IIPKIVP				
S. pombe pol gamma	(593) SASC SYWSSARDRIRSQMVVWDKDAEL GVPSSVD--G-----FGIILPCIIIP				
Neurospora pol gamma	(645) NASC SYWISARERIKNQMVVYEDQLPPSQRFVNKDADSN --TPIGGFVLPOQVIP				
Consensus	(826) NKM ISFWRNAHKRISSQMVVWLPRSA LPR V RHP DE E YGAILPQVVT				

Region 2: 881 - 935

	881		<u>PolA</u>	935	
hum pol gamma aa seq	*** *** *	* *	*	** **	
Pan troglodytes (chimpanzee)	(847) AGT ITRRAVEPTWL TASNARPDRV GSELKAMVQAPPGYTLV GADVDS QELWI AAV				G848S-PEO & Alpers, & MELAS
Macaca mulatta (rhesus monkey) pol gamma	(844) AGT ITRRAVEPTWL TASNARPDRV GSELKAMVQAPPGYTLV GADVDS QELWI AAV				T849H-Alpers
mus pol gamma	(836) AGT ITRRAVEPTWL TASNARPDRV GSELKAMVQAPPGYTLV GADVDS QELWI AAV				T851A- Alpers
rat pol gamma	(825) AGT ITRRAVEPTWL TASNARPDRV GSELKAMVQAPPGYVLV GADVDS QELWI AAV				R852C-Alpers
Xenopus pol gamma	(824) AGT ITRRAVEPTWL TASNARPDRV GSELKAMVQAPPGYVLV GADVDS QELWI AAV				R853Q-Myocerebrohepat.
Drosophila pol gamma	(817) AGT ITRRAVEPTWL TASNARADRV GSELKAMVQVPPGYHLI GADVDS QELWI AAI				R853W-PEO w/ptosis & Parkinsonism
scyeast pol gamma mip1	(765) CG TLTRRAVEPTWM TASNRPDRL GSELRS MVQAPPGYRLV GADVDS QELWI ASV				V855A-muscle weakness
S. pombe pol gamma	(650) MGT ITRRAVENAWLTASN AKANRI GSELKTQVKAPPGYCFV GADVDS EELWI ASL				A862T-Ataxia & PEO
Neurospora pol gamma	(638) MGT ITRRAVENTWL TASN SKKNRL GSELKAMIRAPDGTYTFV GADVDS EELWI VAL				N864S-PEO
Consensus	(697) MGT ITRRAVERTWL TASN AKKNR V GSELKAMVRAPPGYVFV GADVDS EELWI ASV				E873stop- Alpers
	(881) AGT ITRRAVEPTWL TASNARPDRV GSELKAMVQAPPGY LVGADVDS QELWIAAV				Q879H-Alpers

Region 3: 936 - 990

	936		<u>PolB</u>	990	
hum pol gamma aa seq	* * * * *		*	* *	
Pan troglodytes (chimpanzee)	(902) LGDAHFAGM HGCTA F GWMTLQCRKSRGTDLHSKTAT TVGISR EHA KIFNYGRIYG				T914P-PEO, Alpers...
Macaca mulatta (rhesus monkey) pol gamma	(899) LGDAHFAGM HGCTA F GWMTLQCRKSRGTDLHSKTAT TVGISR EHA KIFNYGRIYG				W918R - PEO
mus pol gamma	(891) LGDAHFAGM HGCTA F GWMTLQCRKSRGTDLHSKTAT TVGISR EHA KVFNYGRIYG				G923D-PEO
rat pol gamma	(880) LGDAHFAGM HGCTA F GWMTLQCRKSRGTDLHSKTAA TVGISR EHA KIFNYGRIYG				H932Y-PEO, & SANDO
Xenopus pol gamma	(879) LGDAHFAGM HGCTA F GWMTLQCRKSRGTDLHSKTAA TVGISR EHA KVFNYGRIYG				R943C-Myocerebrohep
Drosophila pol gamma	(872) LGEAHFAGI HGCTA F GWMTLQCRKSSGTDLHSKTAST TVGISR EHA KVFNYGRIYG				R943H-PEO
scyeast pol gamma mip1	(820) LGDAYACGE HGATPL GWMTLSGSKSNGSDMHSITAKAV VGISR D HA KVINYARIYG				R953C-PEO
S. pombe pol gamma	(705) VCD SIFN-VHGTTA IC GMCL ECTKNE GTDLHTKTA QIL GCSP N EAKI FNYGRIYG				Y955C-PEO, & Parkinsonism...
Neurospora pol gamma	(693) MC DSQFR-LHGATA L GMML LEGKK SEGT DL HSKTA AIL GV SRDS AKV FNYGRLYG				
Consensus	(752) VCD ATFK-LHGNA IC GMFL ECTK QSQT DL HSRTA SIL GITR ND AKV FNYGRIYG				
	(936) LGDAHFAGMH GCTA F GWMTLQCRKSRGTDLHSKTAT TVGISR EHA KVFNYGRIYG				

hum pol gamma aa seq Pan troglodytes (chimpanzee) Macaca mulatta (rhesus monkey) pol gamma mus pol gamma rat pol gamma Xenopus pol gamma Drosophila pol gamma scyeast pol gamma mip1 S. pombe pol gamma Neurospora pol gamma Consensus	991 * * *** (957) AGQPFAERLLMQFNHRLTQEAAEKAQQMYAATKGLRWYRLSDEGEWLVRELNLP (954) AGQPFAERLLMQFNHRLTQEAAEKAQQMYAATKGLRWYRLSDEGEWLVRELNLP (946) AGQPFAERLLMQFNHRLTQEAAEKAQQMYAATKGLRWYRLSDEGEWLVRELHLP (935) AGQSFAERLLMQFNHRLTRQEAAEKAQQMYAVTKGLRRLYRLSADGEWLVKQLNLP (934) AGQSFAERLLMQFNHRLSRQEAADKAQQMYAVTKGLRRLYRLSDDGEWLVKQLNVP (927) AGQPFAERLLMQFNHRLTQEAAEKAQKQMYAVTKGIRRYILSKEGEWLVEELGIS (875) AGQLFAETLLRQFNPTFSASEAKAKAMKMSFIATKGRKVYRLREEFHDELEDRAYS (759) AGAKFASQLLKRFPNSLTDEETKKIAANKLYENTKGKTK-RSK----- (747) AGLKHTTLLLMQMNPTLKTAEAKELAKKLYASATKGVSKMSKRLQEMGLPKLT-- (806) AGLKFAASQLLRFQVPSLTEAETTAIATKLYDATKGAKTNRKS----- (991) AGQ FAERLLMQFNHRLT QEAAEKAQQMYA TKGLR YRLS EGEWLV L P	1045
hum pol gamma aa seq Pan troglodytes (chimpanzee) Macaca mulatta (rhesus monkey) pol gamma mus pol gamma rat pol gamma Xenopus pol gamma Drosophila pol gamma scyeast pol gamma mip1 S. pombe pol gamma Neurospora pol gamma Consensus	1046 * * * 1100 (1012) VDRTEGGWI SLQDLRKVQREARKS QWKK-WEVVAERA WKGGTESEMFnKLESIA (1009) VDRTEGGWI SLQDLRKVQREARKS QWKK-WEVVAERA WKGGTESEMFnKLESIA (1001) VDRTEGGWI SLQDLRKVQREARKSHRKK-WEVIAERA WKGGTESEMFnKLESIA (990) VDRTEDGWVSLQDLRMRREASRKS RWKK-WEVASERA WTGGTESEMFnKLESIA (989) VDRTEDGWVSLQDLRKIRREASRKS RWKK-WEVVTERA WTGGTESEMFnKLESIA (982) VERGEENSVNLQDLRKIQKDATKRSRRK-WNLVSRRIWTGGTESQMFnKLETIA (930) SYEASRLAIQ-----RNRTLAEVFHRPNWQGGTESAMFnRLEEA (800) -----LFKKFWYGGSESILFnKLESIA (800) -----FWSQGTEESFVFFnKLEAMA (848) -----LYKRPFWRGTEESFVFNMLEEFA (1046) VDRTE GW SLQDLRK RE RKS KK WEV ERAW GGTESEMFnKLESIA	1101 * * ** *** 1155
hum pol gamma aa seq Pan troglodytes (chimpanzee) Macaca mulatta (rhesus monkey) pol gamma mus pol gamma rat pol gamma Xenopus pol gamma Drosophila pol gamma scyeast pol gamma mip1 S. pombe pol gamma Neurospora pol gamma Consensus	(1066) TSDIPRTPVLGC CISRALEPSA--VQEEFMTSRVNWVVQSSAVDYLHMLVLAMK (1063) TSDIPRTPVLGC CISRALEPSA--VQEEFMTSRVNWVVQSSAVDYLHMLVLAMK (1055) TSDIPRTPVLGC CISRALEPSA--VQGEFMTSRVNWVVQSSAVDYLHMLVLAMK (1044) MSDTPRTPVLGC CISRALEPSV--VQGEFITSRVNWVVQSSAVDYLHMLVLAMK (1043) MSDTPRTPVLGC CISRALEPSV--VQGEFMTSRVNWVVQSSAVDYLHMLVLAMK (1035) MSPSPKTPVLGC RISRALEPTA--VKGEFITSRVNWVVQSSAVDYLHMLVLAMK (970) TGSQPRTPFLGGRLSRALEADTGPEQEQRFLPTRI NWVVQSGAVDFLHMLVLAMR (822) EQETPKTPVLGC GITYSLMKKN--LRANSFLPSRINWAIQS SGVDYLHLLCSCM (818) QLPSPRTPVL DAGI QQALSSKN--LSKNSFMTSRVNWAIQS SAVDYLHLLLVSMN (871) EQERPRTPVLGAGITEALMSRWVS--KGFFLTSRINWAIQS SGVDYLHLLIAMD (1101) SD PRTPVLGC ISRALEPS VQGEFMTSRVNWVVQSSAVDYLHMLVLAMK	G1076V-PEO I1079L-adPEO S1095R-adPEO R1096C-PEO & Alpers R1096H-Alpers S1104C-PEO A1105T-PEO V1106I-PEO

		<u>Pol C</u>												
hum pol gamma aa seq		1156	*	*	**	*				* **		1210		
Pan troglodytes (chimpanzee)	(1118)	WLFEFAIDGRFCISI	HDE	VRYL	VREEDRYRAA	ALALQITNLL	T	RCM	FAYKL	G	LND			
Macaca mulatta (rhesus monkey) pol gamma	(1115)	WLFEFAIDGRFCISI	HDE	VRYL	VREEDRYRAA	ALALQITNLL	T	RCM	FAYKL	G	LND			
mus pol gamma	(1107)	WLFEFAIDGRFCISI	HDE	VRYL	VREEDRYRAA	ALALQITNLL	T	RCM	FAYKL	G	LND			
rat pol gamma	(1096)	WLFEFAIDGRFCISI	HDE	VRYL	VREEDRYRAA	ALALQITNLL	T	RCM	FAYKL	G	LND			
Xenopus pol gamma	(1095)	WLFEFAIDGRFCISI	HDE	VRYL	VREEDRYRAA	ALALQITNLL	T	RCM	FAYKL	G	LND			
Drosophila pol gamma	(1087)	WLFEAYDIDGRFCISI	HDE	VRYL	VREEDRYRAA	ALALQITNLL	T	RCM	FAYKL	G	LND			
scyeast pol gamma mip1	(1025)	WLMG---	SHV	R	FCLSF	HDEL	RYL	VKEEL	SPKA	ALA	MHITNLMTR	SFCVSRI	G	
S. pombe pol gamma	(875)	YIIKKYNLEAR	R	L	CISI	HDE	I	RFL	VSEK	DKYRAA	AMALQI	SN	IWTRAMF	
Neurospora pol gamma	(871)	HLIKKYYLEAR	R	L	SLSTV	HDE	V	RYL	SSDK	DKYRVAF	ALQVANLW	TRAFF	CQRIC	
Consensus	(924)	YLTRRFNLAC	R	LA	ITV	HDE	I	RYL	AEP	DKYRVAM	ALQIANLW	TRVMFA	QQVG	
	(1156)	WLFEFAIDGRFCISI	HDE	VRYL	VREEDRYRAA	ALALQITNLL	T	RCM	FAYKL	G	LND			
hum pol gamma aa seq		1211										1265		
Pan troglodytes (chimpanzee)	(1173)	LPQSVAFFSAVDI	D	R	CLR	K	EVT	MDCK	TPSN	PTGMERRY	Y	GIPQGEAL	LDIYQIII	
Macaca mulatta (rhesus monkey) pol gamma	(1170)	LPQSVAFFSAVDI	D	R	CLR	K	EVT	MDCK	TPSN	PTGMERRY	GIPQGEAL	LDIYQIII	ELT	
mus pol gamma	(1162)	LPQSVAFFSAVDI	D	R	CLR	K	EVT	MDCK	TPSN	PTGMERRY	GIPQGEAL	LDIYQIII	ELT	
rat pol gamma	(1151)	LPQSVAFFSAVDI	D	Q	CLR	K	EVT	MDCK	TPSN	PTGMERRY	GIPQGEAL	LDIYQIII	ELT	
Xenopus pol gamma	(1150)	LPQSVAFFSAVDI	D	Q	CLR	K	EVT	MDCK	TPSN	PTGMER	KY	GIPQGEAL	LDIYQIII	
Drosophila pol gamma	(1142)	V	PQSVAFFSAVDI	D	K	CLR	EVT	MDCK	TPSN	PNGMEK	RYGIPQGEAL	LDIYQI	ILKVT	
scyeast pol gamma mip1	(1077)	LPMSVAFFSS	V	E	D	T	V	L	R	K	TMDCK	TPSN	PHGLRIGY	
S. pombe pol gamma	(930)	LPQNC	AFFSQVDI	D	S	V	I	R	E	V	N	MDCI	TPSN	K
Neurospora pol gamma	(926)	LPQSVAFFSS	V	D	I	D	H	V	L	R	K	D	TPSN	KVPIP
Consensus	(979)	LPQSC	AFFSAVDI	D	I	D	H	V	L	R	K	E	TPSN	P
	(1211)	LPQSVAFFSAVDID	C	L	R	K	E	V	T	M	D	C	TPSN	PTGMERRY
hum pol gamma aa seq		1266	*	*									1320	
Pan troglodytes (chimpanzee)	(1228)	KGSLEKRSQPGP	-	-	-	-	-	-	-	-	-	-		
Macaca mulatta (rhesus monkey) pol gamma	(1225)	KGSLEKRSQPGP	-	-	-	-	-	-	-	-	-	-		
mus pol gamma	(1217)	KGSLEKRSQPGP	-	-	-	-	-	-	-	-	-	-		
rat pol gamma	(1206)	KGSLEKRSQPGP	-	-	-	-	-	-	-	-	-	-		
Xenopus pol gamma	(1205)	KGSLEKRSQPGP	-	-	-	-	-	-	-	-	-	-		
Drosophila pol gamma	(1197)	KGVL	-	-	-	-	-	-	-	-	-	-		
scyeast pol gamma mip1	(1132)	GCNDVSQLWDWIKKS	-	-	-	-	-	-	-	-	-	-		
S. pombe pol gamma	(985)	PNLDIDSKVSYAYNY	R	E	P	V	F	E	Y	N	K	S	T	
Neurospora pol gamma	(981)	LEPLEQIQCFVDVKATT	S	A	E	I	T	E	E	D	KKNIAYL	K	QAFY	-
Consensus	(1034)	DSIVPQSQYAPRL	E	N	I	P	Y	T	P	R	V	P	V	-
	(1266)	KGSLEKRSQPGP	-	-	-	-	-	-	-	-	-	-	-	

R1128H-Micorcephaly

R1138C-PEO

1142R→W SNP

1143E→G SNP, PEO, ...

1146R→C SNP

3482+2TX21splice-
Alpers

M1163R-Alpers

F1164I-PEO

L1173fs stop- Alpers

S1176L-PEO

D1184N-PEO

R1187W-?

K1191N-Alpers

K1191R-Myocerebrohep.

D1196N-Myopathy

G1205A-hearing loss,
failure to thrive,&
ETC complex defic.

Y1210fs-1216stop-
Alpers

1230S→F SNP

1236Q→H SNP, PEO

Alpers

		1321		1375
hum pol gamma aa seq	(1240)	-----		
Pan troglodytes (chimpanzee)	(1237)	-----		
Macaca mulatta (rhesus monkey) pol gamma	(1229)	-----		
mus pol gamma	(1218)	-----		
rat pol gamma	(1217)	-----		
Xenopus pol gamma	(1201)	-----		
Drosophila pol gamma	(1146)	-----		
scyeast pol gamma mip1	(1040)	LRECTSKEYARDGNTAEYSLLDYIKDVEKGKRTKVRIMGSNFLDGTKNAKADQRI		
S. pombe pol gamma	(1019)	-----		
Neurospora pol gamma	(1089)	KRIIAETRYSDPYGAFSLASNGRVSGNPHQRHAAVHASTKTAAPSKPSIASRFD		
Consensus	(1321)			
		1376		1430
hum pol gamma aa seq	(1240)	-----		
Pan troglodytes (chimpanzee)	(1237)	-----		
Macaca mulatta (rhesus monkey) pol gamma	(1229)	-----		
mus pol gamma	(1218)	-----		
rat pol gamma	(1217)	-----		
Xenopus pol gamma	(1201)	-----		
Drosophila pol gamma	(1146)	-----		
scyeast pol gamma mip1	(1095)	RLPVNMPDYPTLHKIANDSAIPEKQLLENRRKKENRIDDENKKLTRKNTTPME		
S. pombe pol gamma	(1019)	-----		
Neurospora pol gamma	(1144)	SVSQASRIKSVAAGSDEPTIRATKAQGKAMAKASGTKLAASTKDVTLNVTIKKKV		
Consensus	(1376)			
		1431		1485
hum pol gamma aa seq	(1240)	-----		
Pan troglodytes (chimpanzee)	(1237)	-----		
Macaca mulatta (rhesus monkey) pol gamma	(1229)	-----		
mus pol gamma	(1218)	-----		
rat pol gamma	(1217)	-----		
Xenopus pol gamma	(1201)	-----		
Drosophila pol gamma	(1146)	-----		
scyeast pol gamma mip1	(1150)	RKYKRVYGGRKAFEAFYECANKPLDYTLETEKQFFNIPIDGVIDDVLNDKSNYKK		
S. pombe pol gamma	(1019)	-----		
Neurospora pol gamma	(1199)	AAPEMAAVPSTSSESKSASKATTSTTTENATASPSSSNVDAKTTSKTPTHK		
Consensus	(1431)			

	1486	1537
hum pol gamma aa seq	(1240)	-----
Pan troglodytes (chimpanzee)	(1237)	-----
Macaca mulatta (rhesus monkey) pol gamma	(1229)	-----
mus pol gamma	(1218)	-----
rat pol gamma	(1217)	-----
Xenopus pol gamma	(1201)	-----
Drosophila pol gamma	(1146)	-----
scyeast pol gamma mip1	(1205)	KPSQARTASSSPIRKTAKVHSKKLPARKSSTTNRNLVELERDITISREY--
S. pombe pol gamma	(1019)	-----
Neurospora pol gamma	(1254)	KETEGEPFPSLDDPVIAARLEAVSKTSPGTRASVAAKLDALASFCHASCCGC
Consensus	(1486)	