

[illegible]

NTDB id	1192	PAKAF	RS23870	WP	024947914.1	EDATKKVEVP	GVAAADANKLGTIALKPD PADG...TADITLTf.....T.MGG..AGPKNK	GKI	T	122	
NTDB id	1122866	EL216	RS08560	WP	085390949.1	NAKSK.....DEKATKYVADM	TIICCGNG...N.GA.PGAADG...VGIVITVTL.....QDGNTAGGASFP	TDVQ	GKTI	133	
NTDB id	1188	NELON	RS01755	WP	003772458.1	NKTAA.....TTIAACGGAG...G.AAAAGTQHASKYVKSVKVTEASPWDITVAISATEANGIPTGLDGN	TI	NTI	132		
NTDB id	1424	RS	RS02795	WP	011000517.1	GSSVF...TPTKNVAN	TIACCTC.....TVPGQITVTY.....T.TAAGGCTL	TA	AGGCTI	113	
NTDB id	1002	ACIAD	RS15100	WP	004923779.1	SSCVGVQEISASNATTNNVATA..TCGA.....SSAQGITVTM.....D.....TTKAKGANI	NTI	117			
NTDB id	1274	LPP	RS09530	WP	011214168.1	V.....SPAATPNVASI..AIGA.....NGVITYTY.....TAAAGGCTI	NTI	104			
NTDB id	1258	GC085	RS09745	WP	014844212.1	V.....SPAATPNVQSIL..AIGA.....NGVITYTY.....TAAAGGCTI	NTI	104			
consensus						* * *	* * *	*	* * *	!	* * *

NTDB id	1192 PAKAF RS23870 WP	O24947914.1	TLTR.....TAADGLWKCTSDQDEQ.....FIPKGC ^S K	150
NTDB id	1122866 EL216 RS08560 WP	O85390949.1	LTPN ^V KKAL..AAGAQQATDWACTTTTGENAKNRG.LTATLNGTLPAKYAPSECR.	187
NTDB id	1188 NELON RS01755 WP	O03772458.1	RLEPLV ^N KAVP..AATSTGAIDWACASAAKETADKRFGTGSGVAVG.TLPAYAPSEC ^K .	188
NTDB id	1424 RS RS02795 WP	O11000517.1	ALVPTAAGTALPVSSASPIMWTTCYQAQNKAQAASSV...APSGTMSLAAKYAPAECR.	168
NTDB id	1002 ACIAD RS15100 WP	O04923779.1	LTPT.....YASGAVTWKCTIT.S.DKKYPVEPSCR ^G	147
NTDB id	1274 LPP RS09530 WP	O11214168.1	VMPT.....QANGDVTWTCTGG.....TLLAKYRPASC ^R P	136
NTDB id	1258 GC085 RS09745 WP	O14844212.1	IMPT.....QANGDVTVTCTGG.....TLLAKYRPASC ^R P	136
consensus			* * * * *	! ! *

- X non conserved
- X similar
- X $\geq 50\%$ conserved