

[illegible]

NTDB	id	1006	ACIAD	RS15005	WP	004923843.1	SGQTLAKFDSTKQIVQLPLDTSNTKYTLAVVDTTNNPLLTSAANAIGQGWAIKAMSA...DNNNYSLL...LTNTGF	147
NTDB	id	1002	ACIAD	RS15100	WP	004923779.1	AGSSCVGVQHEISASNATTNVATATCGASS.....AGQIIVTMDT..TKAKGANITLTPTYASCVAWT	130
NTDB	id	1184	GTFT7	RS02335	WP	000649326.1	AAGFTRLGTVEDMG.DGKIVIAPT..ASGAL.....GGTIKYTFDA..GVVSSSKIQLAR...DANGLW	131
NTDB	id	1167	A1552VC	RS11065	WP	000649326.1	AAGFTRLGTVEDMG.DGKIVIAPT..ASGAL.....GGTIKYTFDA..GVVSSSKIQLAR...DANGLW	131
NTDB	id	1160	ABDM36	RS02330	WP	000649326.1	AAGFTRLGTVEDMG.DGKIVIAPT..ASGAL.....GGTIKYTFDA..GVVSSSKIQLAR...DANGLW	131
NTDB	id	23891	PBPR	RS16270	WP	011219760.1	TTPLTDLGKTAKDANPLGVISVS...ADN.....NLQFKFGD.NSASKDLLVTFD...RSA.TGW	117
NTDB	id	1292	VP	RS12235	WP	005479693.1	ATLAQLCVDAANDLGALDSALAGCSP.....TLTFTFDG.DSSIDGGILTFTF...DAA.TGW	124
NTDB	id	1192	PAKAF	RS23870	WP	024947914.1	DATKKEVPLGVAAADANKLGTIALKPDPADGTA.....DITLTFTFMGCAGPKNKGKIITLTR...TAADGLW	133
NTDB	id	1197	PSJM300	03940	AFN	76866.1	EGYLGITDS.TSYCDVLDL.TAADGHTL.....ECTAKGCNAGKFDGKTLTLNR...TADGEW	123
consensus							* * * * *	

NTDB id 1006	ACIAD RS15005 WP 004923843.1	S C K N K T W A N I S S S S	C G T G S E S W	169
NTDB id 1002	ACIAD RS15100 WP 004923779.1	K C T T T S D K K Y V P S E C R G		147
NTDB id 1184	GTFF74 RS02335 WP 000649326.1	T C S T T V T S E I A P K G C T	A G A T A T I N	153
NTDB id 1167	A1552VC RS11065 WP 000649326.1	T C S T T V T S E I A P K G C T	A G A T A T I N	153
NTDB id 1160	ABDM36 RS02330 WP 000649326.1	T C S T T V T S E I A P K G C T	A G A T A T I N	153
NTDB id 23891	PBPR RS16270 WP 011219760.1	A C T Q T T G . A E. I D S C P A P K S ..		135
NTDB id 1292	VP RS12235 WP 005479693.1	S C L P T A P A I A. V D G C S		139
NTDB id 1192	PAKAF RS23870 WP 024947914.1	K C T S D Q D E Q F I P K G C S K		150
NTDB id 1197	PSJM300 03940 AFN76866.1	S C A S T L D A K Y K P G K C S.....		139
consensus		* ! * * *	* ! * * *	

- X non conserved
- X similar
- X >50% conserved