

logo

NTDB id 1274 LPP RS09530 WP 011214168.1	MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAAT.....	69
NTDB id 1258 GC085 RS09745 WP 014844212.1	MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAAT.....	69
NTDB id 1063296 ACG4IV RS08390 WP 000649326.1	MKAYKNKQKQKGFTLIELMIVVAVIGVLAIAAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGT.....	75
NTDB id 1184 GTF74 RS02335 WP 000649326.1	MKAYKNKQKQKGFTLIELMIVVAVIGVLAIAAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGT.....	75
NTDB id 1167 A1552VC RS11065 WP 000649326.1	MKAYKNKQKQKGFTLIELMIVVAVIGVLAIAAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGT.....	75
NTDB id 1160 ABDM36 RS02330 WP 000649326.1	MKAYKNKQKQKGFTLIELMIVVAVIGVLAIAAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGT.....	75
NTDB id 1292 VP RS12235 WP 005479693.1	MKHSKQKQKQKGFTLIELMIVVAIIGVLAIAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGI.AAAA.....	71
NTDB id 1038 HON27 RS01510 WP 000993725.1	MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSSEISSIKTGYENAVNEGAV.PTKLSD.....	68
NTDB id 1192 PAKAF RS23870 WP 024947914.1	MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWS.VKSGTGTEDATK	74
NTDB id 1197 PSJM300 03940 AFN76866.1	...MKAQMKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGKT.PSLTSTD.....	71
consensus	** *** !*!!!!!!!!!!!!!!*!*!!!!*!*!!!!* *	

logo

NTDB id 1274 LPP RS09530 WP 011214168.1	..GYVSPAAT.PNVASIA.....IGANGVITII.....TYTAAAGGGTIV.MPTTL.QANGDVTWTCGGTLLA	127
NTDB id 1258 GC085 RS09745 WP 014844212.1	..GYVSPAAT.PNVQSI.....IGANGVITII.....TYTAAAGGGTII.MPTTL.QANGDVTWTCGGTLLA	127
NTDB id 1063296 ACG4IV RS08390 WP 000649326.1	AAGFTRLGTVED.....MGDGKIVIIAPTASGALGGTIKYT.F.DAGVSSSKIQLAR.DANGLWTCST.TVTS	139
NTDB id 1184 GTF74 RS02335 WP 000649326.1	AAGFTRLGTVED.....MGDGKIVIIAPTASGALGGTIKYT.F.DAGVSSSKIQLAR.DANGLWTCST.TVTS	139
NTDB id 1167 A1552VC RS11065 WP 000649326.1	AAGFTRLGTVED.....MGDGKIVIIAPTASGALGGTIKYT.F.DAGVSSSKIQLAR.DANGLWTCST.TVTS	139
NTDB id 1160 ABDM36 RS02330 WP 000649326.1	AAGFTRLGTVED.....MGDGKIVIIAPTASGALGGTIKYT.F.DAGVSSSKIQLAR.DANGLWTCST.TVTS	139
NTDB id 1292 VP RS12235 WP 005479693.1	TLAQ LGVDVAAND LGAIDSAL...AGGSPTL.....TFT.FDADSSIDGGILTFTR.DAATGWSCLP.TAPA	132
NTDB id 1038 HON27 RS01510 WP 000993725.1	...VGFTTAATSSVCATYSIDAFTETDGAGKI.....TCTLK..GNPKIATKIVELTR.STDGAWTCVS.DVPT	129
NTDB id 1192 PAKAF RS23870 WP 024947914.1	KEVPLGVAADANKLGTIA.LKP.DPADGTADIT.....LTFTMGGAGPKNKGKIITLTRTAADGLWKCTS.DQDE	141
NTDB id 1197 PSJM300 03940 AFN76866.1	.EGYIGITDS.TSYCDVD.LD..TAADGHI.....ECTAKGGNACKFDGKTIITLNR.TADGEWSCAS.TLDA	131
consensus	* *** ***** **! *	

logo

NTDB id 1274 LPP RS09530 WP 011214168.1	KYRPASCRP.....	136
NTDB id 1258 GC085 RS09745 WP 014844212.1	KYRPASCRP.....	136
NTDB id 1063296 ACG4IV RS08390 WP 000649326.1	EIAPKGCTAGATIN.....	153
NTDB id 1184 GTF74 RS02335 WP 000649326.1	EIAPKGCTAGATIN.....	153
NTDB id 1167 A1552VC RS11065 WP 000649326.1	EIAPKGCTAGATIN.....	153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1	EIAPKGCTAGATIN.....	153
NTDB id 1292 VP RS12235 WP 005479693.1	IA.VDGCS.....	139
NTDB id 1038 HON27 RS01510 WP 000993725.1	EFLPKGCTAGTPTEGAITSL	149
NTDB id 1192 PAKAF RS23870 WP 024947914.1	QFIPKGCSK.....	150
NTDB id 1197 PSJM300 03940 AFN76866.1	KYKPKGCS.....	139
consensus	** * * * ! * * * *	

- [X] non conserved
- [X] similar
- [X] ≥ 50% conserved