

logo

NTDB id 1191 AAC28468.1 484..945()	...MLKKVQKGFLLIELMIVIAIIGILAAIALPAYQDYTARAQATEGFKATAGLQTDLGAWRADRGSGFPNAAALVAGNVI	77
NTDB id 1184 GTF74 RS02335 WP 000649326.1	MKAYKKNKQKGFLLIELMIVVAVIGVLAIAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAA..G	78
NTDB id 1167 A1552VC RS11065 WP 000649326.1	MKAYKKNKQKGFLLIELMIVVAVIGVLAIAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAA..G	78
NTDB id 1160 ABDM36 RS02330 WP 000649326.1	MKAYKKNKQKGFLLIELMIVVAVIGVLAIAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAA..G	78
NTDB id 407281 GPY31 RS10845 WP 069545806.1	MKHSKQKQKQKGFLLIELMIVVAIIGVLSAIAVPAYQNYVAKSEAAATLGLSLRALVTPAELKLQQDGELSGVV.....	72
NTDB id 1292 VP RS12235 WP 005479693.1	MKHSKQKQKQKGFLLIELMIVVAIIGVLAAAIAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGIAAAAT.....	72
NTDB id 1192 PAKAF RS23870 WP 024947914.1	MKAQKGFLLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWSVKSGTGTEDATKK	75
NTDB id 1204 PSJM300 03935 AFN76865.1	.MKPVKSSCRGFSLLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISEGHTPSLVSSLP.....	74
NTDB id 1197 PSJM300 03940 AFN76866.1	...MKAQMKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGKTPSLTSTDE.....	72
consensus	** *** **!!!!!!*!*!*!*!*!*!*!*!*!*!* ** ** ** *	

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NTDB id 1191 AAC28468.1 484..945()	F.....DSANGLDGYFAPGGVTVPDGTGAINIAFDAGAN...AGKTMITPTVIAASGQISTWTCAPGAANGIETR	147
NTDB id 1184 GTF74 RS02335 WP 000649326.1	FTRLGTVEDMG.DGKIVIAPTASGALGG..TIKYTFDAGV...VSSSKIQLAR.....D.ANGLWTCSTTVTS....EI	141
NTDB id 1167 A1552VC RS11065 WP 000649326.1	FTRLGTVEDMG.DGKIVIAPTASGALGG..TIKYTFDAGV...VSSSKIQLAR.....D.ANGLWTCSTTVTS....EI	141
NTDB id 1160 ABDM36 RS02330 WP 000649326.1	FTRLGTVEDMG.DGKIVIAPTASGALGG..TIKYTFDAGV...VSSSKIQLAR.....D.ANGLWTCSTTVTS....EI	141
NTDB id 407281 GPY31 RS10845 WP 069545806.1	.ADLGG.S.ASHALGAITTS....GANISAAATLTFTFN...TGS.LNGDAITL.T....K.TSSGWTCTDGTTV.....	129
NTDB id 1292 VP RS12235 WP 005479693.1	LAQLGVDVAANDLGAIDSA....LAGGSPTLTFTFD...ADSSIDGGILTFTFTR.....D.AATGWSCLPTAPA....I.	133
NTDB id 1192 PAKAF RS23870 WP 024947914.1	EVPLGVAADANKLGTIAIKP...DPADGTADITLTFTMGAGPKNGKGIITLTR.....TAADGLWKCTSDQDE....QF	143
NTDB id 1204 PSJM300 03935 AFN76865.1	.GFIGIGASTN.YCTVTLI....ATATGS..IT.CTTQNGMPDRFNGKTIITLAR.....T.TDGLWDCTSDLDS....RY	135
NTDB id 1197 PSJM300 03940 AFN76866.1	.GYIGITDSTS.YCDVDLD...TAADGH..IE.CTAKGGNAGKFDGKTIITLNR.....T.ADGEWSCASTLDA....KY	133
consensus	** *	

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NTDB id 1191 AAC28468.1 484..945()	LPTSCQ*.....	153
NTDB id 1184 GTF74 RS02335 WP 000649326.1	APKGCTAGATIN	153
NTDB id 1167 A1552VC RS11065 WP 000649326.1	APKGCTAGATIN	153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1	APKGCTAGATIN	153
NTDB id 407281 GPY31 RS10845 WP 069545806.1	.LDNCN.....	134
NTDB id 1292 VP RS12235 WP 005479693.1	AVDGC.....	139
NTDB id 1192 PAKAF RS23870 WP 024947914.1	IPKGCSK.....	150
NTDB id 1204 PSJM300 03935 AFN76865.1	KPGKCL.....	141
NTDB id 1197 PSJM300 03940 AFN76866.1	KPGKCS.....	139
consensus	* ! *	

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved