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NTDB id 1038 HON27 RS01510 WP 000993725.1	MNAQKGFLLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGA.VPTK.....	65
NTDB id 1192 PAKAF RS23870 WP 024947914.1	MKAQKGFLLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWSVKSGTGTEDATKK	75
NTDB id 1292 VP RS12235 WP 005479693.1	MKHSKQKKQKGFLLIELMIVVAIIGVLAIAAIPAYQYQNYVTRSEVTSGLATVKKALITPAELHYQENGIAAAAT.....	72
NTDB id 408091 GPY47 RS15625 WP 047008977.1	MKNSKQKKQKGFLLIELMIVVAIIGVLAIAAIPQYQYQYVAKSEVASALATLTGVKTNVEAYAVENGAFPDGSTANQT...	77
NTDB id 1184 GTF74 RS02335 WP 000649326.1	MKAYKKNKQKGFLLIELMIVVAIIGVLAIAAIPQYQYVKKSAIGVGLANITALKTNIEDYIATEGSFPAATTAGTAA..G	78
NTDB id 1167 A1552VC RS11065 WP 000649326.1	MKAYKKNKQKGFLLIELMIVVAIIGVLAIAAIPQYQYVKKSAIGVGLANITALKTNIEDYIATEGSFPAATTAGTAA..G	78
NTDB id 1160 ABDM36 RS02330 WP 000649326.1	MKAYKKNKQKGFLLIELMIVVAIIGVLAIAAIPQYQYVKKSAIGVGLANITALKTNIEDYIATEGSFPAATTAGTAA..G	78
NTDB id 1119 NGFG RS11465 WP 017147169.1	MNTLQKGFLLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAECQKSAVTEYYLNHCKWPENNTSA.....	71
NTDB id 1258 GC085 RS09745 WP 014844212.1	MRQKGFLLIELMIVVAIIGILAAIAAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAAT.....	69
consensus	** *** !*!!!!!!!!!!*!*!*!*!*!*!*!*!*!*!* * * * * * * * * * * * * * * * * * * *	

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NTDB id 1038 HON27 RS01510 WP 000993725.1	LSDVGFTAATSSVCATYSIDAFETDAGGKITCTLKG...NPKIATKIVELTRS...TDGAWTCVSDV.....	127
NTDB id 1192 PAKAF RS23870 WP 024947914.1	EVPLGVAADANK..LGTIALKPD PADGTADITLTFTMGAGPKNKGKIITLTRT...AADGLWKCTSDQ.....	139
NTDB id 1292 VP RS12235 WP 005479693.1	LAQLGVDVAAND...LGAIDSALAGGSPTLTFTFDADS...SIDGGILTFTRD...AATGWSCLPTA.....	130
NTDB id 408091 GPY47 RS15625 WP 047008977.1	EADLGV PSTIPSGV...IAFSTASSAG..GTIKFTFNASGVSNLITSKNFELVRA...SDGTWTCQGSSASP.....	141
NTDB id 1184 GTF74 RS02335 WP 000649326.1	FTRLGTVEDMGDGGKIVIAPTASGALG..GTIKYTFDAGV...VSSSKIQLARD...ANGLWTCSTT.....	136
NTDB id 1167 A1552VC RS11065 WP 000649326.1	FTRLGTVEDMGDGGKIVIAPTASGALG..GTIKYTFDAGV...VSSSKIQLARD...ANGLWTCSTT.....	136
NTDB id 1160 ABDM36 RS02330 WP 000649326.1	FTRLGTVEDMGDGGKIVIAPTASGALG..GTIKYTFDAGV...VSSSKIQLARD...ANGLWTCSTT.....	136
NTDB id 1119 NGFG RS11465 WP 017147169.1	GVASPPTDIKGKY...VKEVEVKNGVVTATMLSSGVNNEIKGKKLSLWGRRENGSVKWF CGQPVTTRADDDTVADAK	144
NTDB id 1258 GC085 RS09745 WP 014844212.1	GYVSPAATPNVQS...I..AIGANGVITITITVTAAA....GGGTIIMTPTTLQANGDVTWTCTGGTL.....	125
consensus	*! *	

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NTDB id 1038 HON27 RS01510 WP 000993725.1	PTEFLPKGCTAGTPTEGAITSL	149
NTDB id 1192 PAKAF RS23870 WP 024947914.1	DEQFIPKGC SK.....	150
NTDB id 1292 VP RS12235 WP 005479693.1	PAIAVDGCS.....	139
NTDB id 408091 GPY47 RS15625 WP 047008977.1	VTDDLFPKNCR.....	152
NTDB id 1184 GTF74 RS02335 WP 000649326.1	VTSEIAPKGCTAGATIN.....	153
NTDB id 1167 A1552VC RS11065 WP 000649326.1	VTSEIAPKGCTAGATIN.....	153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1	VTSEIAPKGCTAGATIN.....	153
NTDB id 1119 NGFG RS11465 WP 017147169.1	DGKEIDIKHLPSTCRDKASDAK.....	166
NTDB id 1258 GC085 RS09745 WP 014844212.1	LAKYRPASCRP.....	136
consensus	* *** *!*!* *	

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