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NTDB id 1204 PSJM300 03935 AFN76865.1	MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQ.AISEGHTPSLVSSLPG	75
NTDB id 1191 AAC28468.1 484..945()	MLKKVQKGFTLIELMIVIAIIGILAAIALPAYQDYTARAQATEGFKATAGLQTDLGAWRADRGSFNPAAALV.A	73
NTDB id 1109 OK783 RS10670 WP 157147416.1	MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGSIWPKDNTSA.G	72
NTDB id 1119 NGFG RS11465 WP 017147169.1	MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGKWPENNTSA.G	72
NTDB id 1188 NELON RS01755 WP 003772458.1	MKAMQKGFTLIELMIVIAIIGILAAIALPAYQDYTVRSKVSEALIAASSPKSALSEAFQTDGITGMTAAAK.A	72
NTDB id 360537 FAZ30 RS07100 WP 124645119.1	MRGNNNGFTLIELMIVVAIIGILAAIALPAYQAYTIRAKVAEGLLAADAIKSAVWDSYTSNNSWPSSNNSA.G	71
NTDB id 1424 RS RS02795 WP 011000517.1	MKSMRHLNKRQKGFTLIELMIVVAIVGILAAIALPAYQDYTVRARVTEGLSLAAQAKALVSENAANAQSDLSV.....	74
NTDB id 1002 ACIAD RS15100 WP 004923779.1	MNAQKGFTLIELMIVIAIIGILAAIALPAYTDYTVRARVSEGLTAASSMKTTVSENILNAGALVAGTPTST.A	71
NTDB id 1258 GC085 RS09745 WP 014844212.1	MRQKGFTLIELMIVVAIIGILAAIALPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAAT.G	70
NTDB id 1274 LPP RS09530 WP 011214168.1	MRQKGFTLIELMIVVAIIGILAAIALPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAAT.G	70
consensus	* **!*!!!!!!!!!!!!!!*!***** ** * * * *	

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NTDB id 1204 PSJM300 03935 AFN76865.1	FI.....GIGA STNYCTV.TLI..ATATGS...ITCTTQNGMPDRFNGKTITITART.....	120
NTDB id 1191 AAC28468.1 484..945()	GNVIFDS.....ANGLDGKYFAPGGVTVT.PDTGAINIAFDA.....GANAGKMTITPTVI.....AASG	128
NTDB id 1109 OK783 RS10670 WP 157147416.1	VA.....STIKGKYVQKVEVAKG....VVT AQMA...STGVNKEIQDKKLSLWA.....KRQD	120
NTDB id 1119 NGFG RS11465 WP 017147169.1	VA.....SPPIDIKGKYVKEVEVKNG....VVTATML...SSGVNNEIKGKKLSLWG.....RREN	121
NTDB id 1188 NELON RS01755 WP 003772458.1	FNKTAAGGGGAGGAAAAGTQHASKYVKSVKVTEA..SPWDITVAISATEANGIPTGLDGNTIRLFLPNKA.VP.AATST	148
NTDB id 360537 FAZ30 RS07100 WP 124645119.1	ILSA.....STYATKYIDSINIVAASS.VSNIVVNFNA.....STVLSGLTVTMP.....QLSS	120
NTDB id 1424 RS RS02795 WP 011000517.1	GSSVF...TPTKNVANLTIAGTGTVPQG IIVTYTT.....AAGGGTLALVPTAAGTALPVSSAPS	131
NTDB id 1002 ACIAD RS15100 WP 004923779.1	GSSCVGVQEISASNATTNVAT..ATCGASSAGQIIVTMDT.....TKAKGANITLTP.....TYAS	125
NTDB id 1258 GC085 RS09745 WP 014844212.1	YV.....SPAATPNVQS..IAIG..ANGVITITYTA.....AAGGGTIIMTPT.....LQAN	113
NTDB id 1274 LPP RS09530 WP 011214168.1	YV.....SPAATPNVAS..IAIG..ANGVITITYTA.....AAGGGTIVMTPT.....LQAN	113
consensus	* ***** ** * * * * * * * *	

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NTDB id 1204 PSJM300 03935 AFN76865.1	TDGLWDCTSD.....LDSRYKPGKCL.....	141
NTDB id 1191 AAC28468.1 484..945()	QISTWTCAPG.....AANGIETRRLPTSCQ*.....	153
NTDB id 1109 OK783 RS10670 WP 157147416.1	GSVKWFCGQPVTRTGDNDDTVA..DANNAIDTKHLPSTCRDESSAT.	164
NTDB id 1119 NGFG RS11465 WP 017147169.1	GSVKWFCGQPVTRADDDTVADA..KDGKEIDTKHLPSTCRDKASDAK	166
NTDB id 1188 NELON RS01755 WP 003772458.1	GAIDWACASAAKETADKRFGTSGSVAVGTLPAKYAPSECK.....	188
NTDB id 360537 FAZ30 RS07100 WP 124645119.1	GSIMWTCQAS.....ATFSQYLPSSCR.....	142
NTDB id 1424 RS RS02795 WP 011000517.1	GPIMWTCYAQNKAQAASSVAPS...GTMSLAAKYAPAECR.....	168
NTDB id 1002 ACIAD RS15100 WP 004923779.1	GAVTWKCTTT.....S.DKKYVPSECRG.....	147
NTDB id 1258 GC085 RS09745 WP 014844212.1	GDVTWTCGG.....TLLAKYRPASCRP.....	136
NTDB id 1274 LPP RS09530 WP 011214168.1	GDVTWTCGG.....TLLAKYRPASCRP.....	136
consensus	* * !! *****!***	

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