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NTDB id 113 BSU 40900 NP 391970.1	...M.LNRRVVLVGRLLTKDPELRYTPNGAAVATFTLAVNRTEFTNQS.GER..EADDFINCVTWRRQAENVANFLKKGSLAGV	73
NTDB id 1131 NGFG RS05740 WP 003695064.1	...MSLNKVVILIGRLGRDPEVRYPMPNGEAVCNFSVATSETWNRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL	76
NTDB id 1103 NMB RS07590 WP 002212976.1	...MSLNKVVILIGRLGRDPEVRYPMPNGEAVCNFSVATSETWNRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL	76
NTDB id 1166 A1552VC RS00795 WP 000168289.1	MASRGVNKVVILIGNLGQDPEVRYPMPNGGAVANIITATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI	80
NTDB id 1065170 ACHWS3 RS03360 WP 012340854.1	..MAGINKVIVGNLGNDPEMRTMPNGEAVANISVATSESWTDKNTGERREVTEWHRIVFYRRYAEICGQYLRKGSKTYI	78
NTDB id 1390 A4U84 RS00055 WP 021115958.1	..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFRRQAEVAGQYLRKGSKVYV	78
consensus	*!*!*****!*!*!*****!*!*!*****!*!*!*****!*!*!*****!*!*!*****!*!*!*****!*!*!*****	

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NTDB id 113 BSU 40900 NP 391970.1	EGRLQTRNYENQQGQRVFVTEVQA....ESVQFLEPKNGGGSGSGG...YNEGNSGGGQYFGGGQNDNPFGGNQNN....	142
NTDB id 1131 NGFG RS05740 WP 003695064.1	EGRIQSRKYQGGKDGIERAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQS.....QEAYQRPAAQSRQPAPDA	143
NTDB id 1103 NMB RS07590 WP 002212976.1	EGRIQSRKYQGGKDGIERAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQS.....QEAYQRPAAQSRQPASDA	143
NTDB id 1166 A1552VC RS00795 WP 000168289.1	EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGM...NVPA....QQGSWGQPQQPAKQHQP..	150
NTDB id 1065170 ACHWS3 RS03360 WP 012340854.1	EGKLRTRKWQDQNGQERYTTEIQG....ELLQMLDSRNATNS.VDGGQSN.....YQA.....	126
NTDB id 1390 A4U84 RS00055 WP 021115958.1	EGKLRTRKWQDQNGQRYTTEIQG....DVLQMLDSRSSGGD.FGQNGSGWNQAPAQTNYNQQGYSDNYAQNNNFNGGN	153
consensus	*!******!*!*!*********!*!*!********	

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NTDB id 113 BSU 40900 NP 391970.1	.QRRNQGNSFNDDPFANDGKPIDISDDDLPF	172
NTDB id 1131 NGFG RS05740 WP 003695064.1	PSHPQEAPAAPRRQPVPAAPVEDIDDDIPF	174
NTDB id 1103 NMB RS07590 WP 002212976.1	PSHPQEAPAAPRRQPVPAAPVEDIDDDIPF	174
NTDB id 1166 A1552VC RS00795 WP 000168289.1	MQQ...SAPQQYSQPQYNEPPMDFDDDI PF	177
NTDB id 1065170 ACHWS3 RS03360 WP 012340854.1	MSQSPKATHPQDLMDNFD DDIPF	150
NTDB id 1390 A4U84 RS00055 WP 021115958.1	ATR...PQPAKPAAQAEPMDNFD DDIPF	180
consensus	*!*!*!!!!!!	

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