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NTDB id 1323 RAOC RS07335 WP 004916218.1	MSLAIVYSRASIGVQAPLVTTIEVHLSNGKPGFTLVGLPETVKESQDRVRSALINSQFEFPARRITVNLAPADLPKEGGR	78
NTDB id 1271 LPP RS03220 WP 011213198.1	MSLAFTTKTRSTIGIVAQPVSVEVHLSNGLPSFTMVGLAETAVKESKDRVRSALINSQFEFPCKRITVNLGPANLPKGTSG	80
NTDB id 1248 GC085 RS03150 WP 011213198.1	MSLAFTTKTRSTIGIVAQPVSVEVHLSNGLPSFTMVGLAETAVKESKDRVRSALINSQFEFPCKRITVNLGPANLPKGTSG	80
NTDB id 1043447 ACEN2W RS23790 WP 014618079.1	MSLAVLRSRALAGIDAPAVSVEVHLANGLPSFTIVGLPDTEVKESKDRVRAAIQNSRFEFPTRRITVNLAPADLPKESGR	80
NTDB id 1153 A1552VC RS00115 WP 000521788.1	MGLAIIHSRASIGVQAPPVTVVEVHISNGMPGFTLVGLPETTVKESRDRVRSALINSRFEFPAKRITVNLAPADLPKEGGR	80
NTDB id 1397 DSB67 RS15690 WP 005535577.1	MGLAIIHSRASVGVQAPSVSVEVHISNGMPGFTLVGLPETTVKESKDRVRSALVNSNFQFPAKRITVNLAPADLPKEGGR	80
NTDB id 1387 A4U84 RS06900 WP 020457569.1	MSLAIIYSRASIGVEAPLVTTIEVHISGGSPGLTTLVGLPEKSVKEAQDRVRSALMNFQFPAKRITVNLAPADLPKEGGR	80
NTDB id 1349 HI 1117 AAC22771.1	MSLAIVYSRASMGVQAPLVTTIEVHLSNGKPGFTLVGLPEKTVKEAQDRVRSALMNAQFKYPAKRITVNLAPADLPKEGGR	80
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NTDB id 1271 LPP RS03220 WP 011213198.1	FDLPIALGILAAAEQIPLTNLANHEFIGELALSGELRGVSAIIPAVLAH.....EKGFGIILPKQNTREAAIVN	150
NTDB id 1248 GC085 RS03150 WP 011213198.1	FDLPIALGILAAAEQIPLTNLANHEFIGELALSGELRGVSAIIPAVLAH.....KDNQHLLIIANANAAEASLTG	150
NTDB id 1043447 ACEN2W RS23790 WP 014618079.1	FDLPIALGILAAARQIPQTHLDTHEFAGELSLSGDLRPIRGALAMAYALSCHPGSPDGDAPLARAFLVPDENAAEAALVS	160
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NTDB id 1387 A4U84 RS06900 WP 020457569.1	FDLPIALGILAAAGQIDSDKLRKRIELLGELSLTGHLRGVHGAIPAVLAAS.....KAKRELIIVPQQNANEVSLVS	150
NTDB id 1349 HI 1117 AAC22771.1	FDLPIALGILAAADQIDASHLKQFEFVAELALTGQLRGVHGVIPAILAAQ.....KSKRELIITAKQNANEASLVS	150
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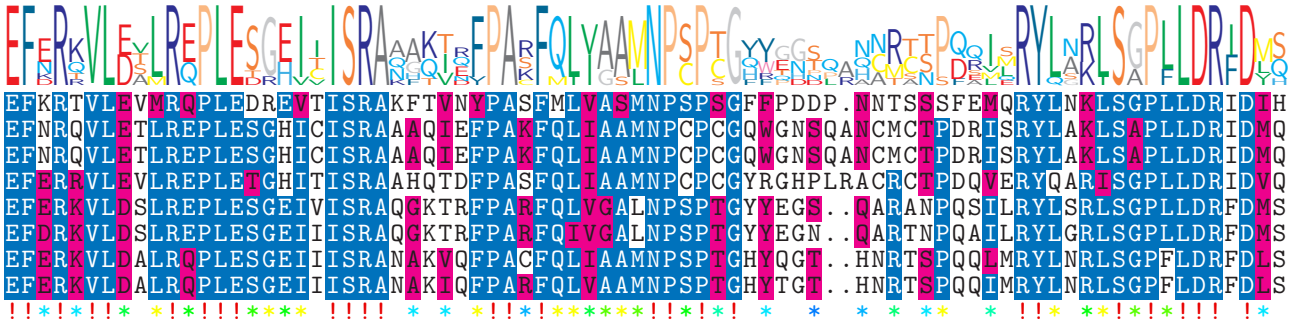
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NTDB id 1271 LPP RS03220 WP 011213198.1	HQKVFTANNLREVCDYLCQGTSLQSLPPKPD....LLNNYELDWSIDKGGQGHAKNAMVIAACGGHSILLSGAPGSGKTM	226
NTDB id 1248 GC085 RS03150 WP 011213198.1	HQKVFTANNLREVCDYLCQGTSLQSLPPKPD....LLNNYELDWSIDKGGQGHAKNAMVIAACGGHSILLSGAPGSGKTM	226
NTDB id 1043447 ACEN2W RS23790 WP 014618079.1	GTRVLPARTLIEVCDHLSAPDRKLAPAAPA....PVGTTGTRYPDADVIRGQPPQARRALEVAAAGGHSMLMIGPPPGTGKSM	236
NTDB id 1153 A1552VC RS00115 WP 000521788.1	VERHKSAAQSLLEVCAELCGQQTLSLFQSSPSV....QQVSQTRDLQDIIGQQQGHAKRALEIAAAGNHNLLFLGPPPGTGKTM	226
NTDB id 1397 DSB67 RS15690 WP 005535577.1	KEQHKSAAQSLLEVCAELCGQHQLNLYQTPKKK....ELKAHGRDLQDIIGQQQGHAKRALEIAAAGNHNLLFLGPPPGTGKTM	226
NTDB id 1387 A4U84 RS06900 WP 020457569.1	NTQTYCASTLLQVVNFLLNNRDQLPIAQQLSQNIAEN.RPLVSRDLTDIIGQQHAKRALIIAASGQHNNLLFLGPPPGTGKTM	229
NTDB id 1349 HI 1117 AAC22771.1	DQNTYFAQTLLDVVQFLNGQEKLPLATEIVKESAVNFSGKNTLDLTDIIGQQHAKRALIIAAGQHNNLLFLGPPPGTGKTM	230
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NTDB id 1323 RAOC RS07335 WP 004916218.1	LAKRIPSLPPLTLKEALETTHSVAGKMGAEETSLMTIRPFRSPHHTISDVALVGGGSYPQPGEISLAHNGVFLDDEMP	309
NTDB id 1271 LPP RS03220 WP 011213198.1	MAKRFTSLPELSETQALECAAINSIRGKLP.DFREWRLLPPFRAPHHTASPVVALVGGGNPPKPGGEISLAHHGVFLDDELP	305
NTDB id 1248 GC085 RS03150 WP 011213198.1	MAKRFTSLPELSETQALECAAINSIRGKLP.DFREWRLLPPFRAPHHTASPVVALVGGGNPPKPGGEISLAHHGVFLDDELP	305
NTDB id 1043447 ACEN2W RS23790 WP 014618079.1	LAQRFPGLLPDMDDEALSAAAVMSLTTRGF.DAQRWKKVRPFRAPHHTASAVAMVGGGNTPRPGEISLAHQGVFLDDELP	315
NTDB id 1153 A1552VC RS00115 WP 000521788.1	LASRLCDLLPEMSDEEAMETASIASLTQSEI.NQHNWKKLRPFRAPHHSSSMAALVGGGTIPRPGEISLAHNGLLFLDDEMP	305
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NTDB id 1387 A4U84 RS06900 WP 020457569.1	LASRLADLPEMEDDEAIETASVTSLVQNEL.NFQNWKKRPPFRAPHHSSSMAALVGGGSIPKPGGEISLAHNGVFLDDELP	308
NTDB id 1349 HI 1117 AAC22771.1	LASRLTGLLPMTDLEAIETASVTSLVQNEL.NFHNWKKRPPFRAPHHSSAMPALVGGGTIPKPGGEISLATNGVFLDDELP	309
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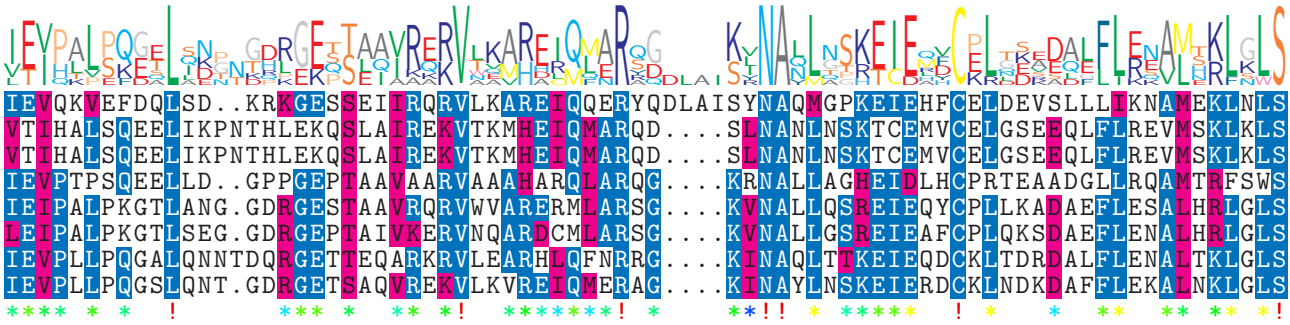
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NTDB id 1387 A4U84 RS06900 WP 020457569.1
NTDB id 1349 HI 1117 AAC22771.1
consensus



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☒ non conserved
☒ similar
☒ ≥50% conserved