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NTDB id 1323 RAOC RS07335 WP 004916218.1	MSLAIVYSRASIGVAPLVTVEVHLSNGKPGFTLVGLPETVKESDRVRSALINSFEFPAKRITVNLAPADLPKEGGR	78
NTDB id 1271 LPP RS03220 WP 011213198.1	.MLVKKVYGSAIHGVSAQTTITIEVNVDQGV.GYHLVGLPDNAIKESSHRISAALKNVGYKLPKGGKITINMAPADLRKEGSA	80
NTDB id 1248 GC085 RS03150 WP 011213198.1	MSLAFTTKTRSTIGIVAQPVSVEVHLSNGLPSFTMVGLAETAVKESKDRVRSALINSQFEFPCRKITVNLGPANLPKGTGSG	80
NTDB id 1162934 ABNR97 RS09195 WP 003689469.1	MSLAFTTKTRSTIGIVAQPVSVEVHLSNGLPSFTMVGLAETAVKESKDRVRSALINSQFEFPCRKITVNLGPANLPKGTGSG	80
NTDB id 1153 A1552VC RS00115 WP 000521788.1	MSLALVYSRALSGMNAPLVEVEAHLANGLPHFNIVGLPDTEVKESRDRVRAAIIQSGFEFPAKKITVNLAPADLPKESGR	80
NTDB id 1397 DSB67 RS15690 WP 005535577.1	MGLAIIHSRASIGVQAPPVTIVEVHISNGMPGFTLVGLPETTVKESRDRVRSALINSRFEFPAKRITVNLAPADLPKEGGR	80
NTDB id 1349 HI 1117 AAC22771.1	MGLAIIHSRASVGVQAPSVSVEVHISNGMPGFTLVGLPETTVKESKDRVRSALVNSNFQFPAKRITVNLAPADLPKEGGR	80
NTDB id 1387 A4U84 RS06900 WP 020457569.1	MSLAIVYSRASMGVQAPLVTTIEVHLSNGKPGFTLVGLPEKTVKEAQDRVRSALMNAQFKYPAKRITVNLAPADLPKEGGR	80
consensus	MSLAIIYSRASIGVEAPLVTTIEVHISGGSPGLTTLVGLPEKS VKEAQDRVRSALMNANFDYPARRITINLAPADLPKEGGR	80

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NTDB id 1323 RAOC RS07335 WP 004916218.1	FDLPIALGILAAASQIAKLNKYEEFELALSGLRGVGYLPAALAAK	158
NTDB id 1271 LPP RS03220 WP 011213198.1	YDLSTALGILAASGQIIAPETIERYLIMGELSLDGGQLPIKGVLPPIAIRAREEGFGKTIILPKQNTREAAIVNDLEVYGVEN	159
NTDB id 1248 GC085 RS03150 WP 011213198.1	FDLPIALGILAASEQIPLTNLANHEFIGELALSGELRGVSAIIPAVLAAHK.DNQHLIIANANAAEASLTGHQKVFTANN	159
NTDB id 1162934 ABNR97 RS09195 WP 003689469.1	FDLPIALGILAASEQIPLTNLANHEFIGELALSGELRGVSAIIPAVLAAHK.DNQHLIIANANAAEASLTGHQKVFTANN	159
NTDB id 1153 A1552VC RS00115 WP 000521788.1	FDLPIALGILAASGQVAPEKLAIEYEFAGELALSGLLRPVRGALAMAWQGMQ.AKRAFVLPEENAGQAAVMRGITVYGARS	159
NTDB id 1397 DSB67 RS15690 WP 005535577.1	FDLPIALGILAASDQIARNKLESYEFIGELALSGETIRGVKGVLPAALAAANQ.VERCLVVPHSNGDQAALVGVERHKSQAS	159
NTDB id 1349 HI 1117 AAC22771.1	FDLPIALGILAASEQIATDKLNKYEFVGLALSGGLRTVKGVLPAALAAANK.KQRHLVVPHANGDQAALVGKEQHKSAQS	159
NTDB id 1387 A4U84 RS06900 WP 020457569.1	FDLPIALGILAASDQLDASHLKQFEFVAELALTGQLRGVHGVIPAILAAQK.SKRELIIAKQNANEASLVSDQNTYFAQT	159
consensus	FDLPIALGILAASGQIDSDKLKRIFELGELSLTGHLRGVHGAIPAVLAASK.AKRETIIVPQQNANEVSLVSNQTQYCAST	159

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NTDB id 1323 RAOC RS07335 WP 004916218.1	LLEVCDYLCQGTSLQSLPPKPDLL...LLNNYELDWS	238
NTDB id 1271 LPP RS03220 WP 011213198.1	LLEVCDYLCQGTSLQSLPPKPDLL...LLNNYELDWS	235
NTDB id 1248 GC085 RS03150 WP 011213198.1	LLEVCDYLCQGTSLQSLPPKPDLL...LLNNYELDWS	235
NTDB id 1162934 ABNR97 RS09195 WP 003689469.1	LGEVAAHLNGIEPLAQTECSVPQ..MPSEHGGQPDLC	237
NTDB id 1153 A1552VC RS00115 WP 000521788.1	LLEVCAELCGQHQLNLYQTPKKK...ELKAHGRDL	235
NTDB id 1397 DSB67 RS15690 WP 005535577.1	LLEVCAELCGQHQLNLYQTPKKK...ELKAHGRDL	235
NTDB id 1349 HI 1117 AAC22771.1	LLDVVQFLNGQEKPLATEIVKESAVNFSGKNTLL	239
NTDB id 1387 A4U84 RS06900 WP 020457569.1	LLQVVNFLNNRDQLPIAQQLSQNIAEN.RPLVSR	238
consensus	LLEVCDYLCQGTSLQSLPPKPDLL...LLNNYELDWS	238

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NTDB id 1323 RAOC RS07335 WP 004916218.1	PPLTLKEALETTKIHSAVGKMGAEISLMTIRPFRSPHHTISDV	318
NTDB id 1271 LPP RS03220 WP 011213198.1	PELSETQALECAAINSIRGKLPD.FREWRLPPFRAPHHTASPV	314
NTDB id 1248 GC085 RS03150 WP 011213198.1	PELSETQALECAAINSIRGKLPD.FREWRLPPFRAPHHTASPV	314
NTDB id 1162934 ABNR97 RS09195 WP 003689469.1	PPLAEDELVEVWALRSLLPNHQQ..QLDSNRPFRRSPHHSASAA	315
NTDB id 1153 A1552VC RS00115 WP 000521788.1	PEMSDEEAMETASIASLTQKEIN.QHNWKLRPFRAPHHSSSMA	314
NTDB id 1397 DSB67 RS15690 WP 005535577.1	PEMSDEEAMETASVASLTQSEIN.EHNWKTTPFRAPHHSSSMA	314
NTDB id 1349 HI 1117 AAC22771.1	PEMTDLEAIIETASVTSVLQNELN.FHNWKRPFRRAPHHSASMP	318
NTDB id 1387 A4U84 RS06900 WP 020457569.1	PEMTDLEAIIETASVTSVLQNELN.FQNWKKRPFRSPHHSASMV	317
consensus	PEMSDEEAMETASIASLTQKEIN.QHNWKLRPFRAPHHSSSMA	318

