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NTDB id 1169 A1552VC RS11075 WP 000648511.1	MAKNTSGGFSLFITKGERYAFKFDRSSKILKISYFNLIEFKSADRIHLSGGVEERUNFIERLSRRITIEAL	72
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NTDB id 1402 DSB67 RS12670 WP 010643257.1	.MKTT.....TPQLKNFRWKGVNSSGKK.TSCQTLAMSEIEVRERLDAQHIKIKKLKKGSSISFLT.LSHRVKKGKDIT	71
NTDB id 1112 NGFG RS09215 WP 003689811.1	.MKTI.....APQLKNYRMKGINSSGKK.TSCNVLAMTEIEVRERLDAQHIKIKKLKKGSSISFITK.ISHRVKGKDITI	74
NTDB id 1113 AAA85695.1 219..1451()	.MAKNGGFSLFIAKKEKRFIFEGRHSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	74
NTDB id 1252 GC085 RS07730 WP 011213805.1	.MAKNGGFSLFIAKKEKRFIFEGRHSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	71
NTDB id 1198 PSJM300 03950 AFN76868.1	.MDKN.....SPPILTFHYQCINKAGQK.MBCDIQARSLATAKADLRKQGIVTNKVVKRRKP.FDR.KNKKITQADITV	70
NTDB id 1016 ACIAD RS01680 WP 004920476.1	.MAQK.....AIKNSVFTWEGLDRQCAK.IKGELSCVSPALVKAQLRKQGINPQKVKKSISLFS..AGKKIKPMDIAL	73
NTDB id 1059 ABD1 RS01610 WP 000279215.1	MAAKK.....TQVMPVFAYEGVDRKGCIK.LKGELPARNMALAKVTLRKQGITITIREKRKNIFEGLLKKVKVPLDIAI	73
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NTDB id 1402 DSB67 RS12670 WP 010643257.1	FTRQIATMTMTGVPIVQALKLVSENHKKAEKSLMSVTRAVEACTPMSKAMRTASTHFDPLYTDLIATGEQSGNSQVF	151
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NTDB id 1252 GC085 RS07730 WP 011213805.1	FTRQLATMTIESGIPLVQAFDIVAKCQSNKRLKDLIETIKHDIETGLTLAESLIKHPLYFNELFCNLVDAGEKSGSLDML	151
NTDB id 1198 PSJM300 03950 AFN76868.1	FTRQMATMMKAGVPLIQSFDIIGEGFDNPNMKLVDDLKQEVAAACNSFASSIRKKPQYFDDLYCNLVDSGEQSGSLETLL	150
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NTDB id 1059 ABD1 RS01610 WP 000279215.1	FTRQLATMMKAGVPLVQCFEIVAEGLNPAMREVLIGKGEVCGSIFASALRKYPPHFDNLFCSLVESGEQSGALETML	153
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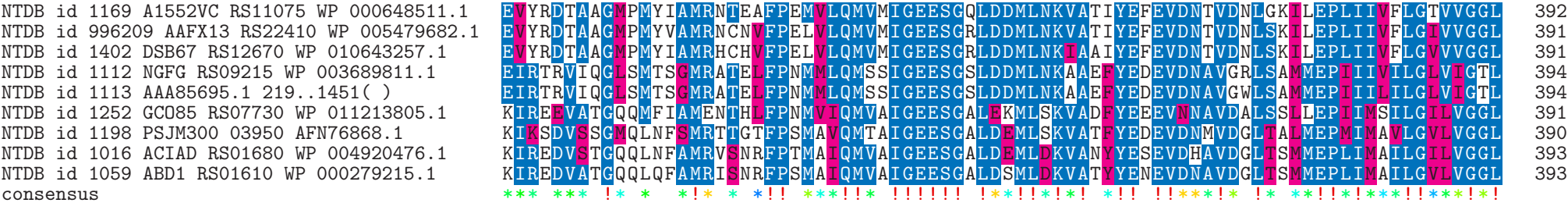
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NTDB id 1252 GC085 RS07730 WP 011213805.1	DKVATYKEKTEIKKKTKKALTYPPIAVMVVALLVTAGLLIYVVPQFESLFKGFADLPAMTRAVITMSEFMQAYWYIIFG	231
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NTDB id 1016 ACIAD RS01680 WP 004920476.1	DRVAVYKEKSEVLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFKDLFTSFAGADLPAFTQLVNVMSNWMQDYWFILIL	233
NTDB id 1059 ABD1 RS01610 WP 000279215.1	DRVATYKEKSELKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFKDLFASFAGADLPAFTQMVNVMSKWMQDYWFIMII	233
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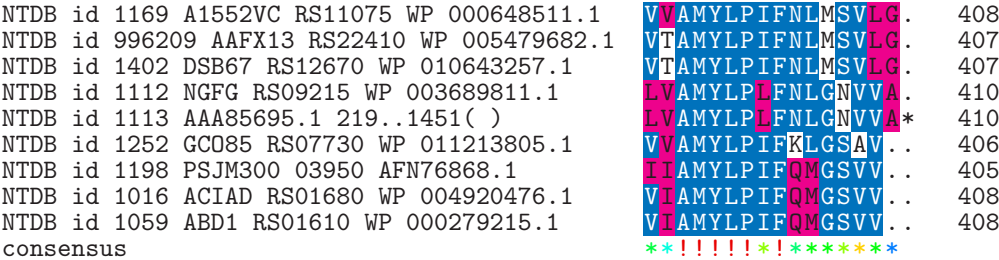
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NTDB id 1252 GC085 RS07730 WP 011213805.1	ALGAAIYSLFLKKRSKFRDRLLKPLIFGDLVKAARSRRLATIFAAGVPLVDALKSTAGATGNLYEAT	311
NTDB id 1198 PSJM300 03950 AFN76868.1	ALGAAIYSLFLKKRSKFRDRLLKPLIFGDLVKAARSRRLATIFAAGVPLVDALKSTAGATGNLYEAT	310
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NTDB id 1059 ABD1 RS01610 WP 000279215.1	ALGAAIYSLFLKKRSKFRDRLLKPLIFGDLVKAARSRRLATIFAAGVPLVDALKSTAGATGNLYEAT	313
consensus	ALGAAIYSLFLKKRSKFRDRLLKPLIFGDLVKAARSRRLATIFAAGVPLVDALKSTAGATGNLYEAT	313

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☐ non conserved
☒ similar
☒ ≥50% conserved