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NTDB id 1020 ACIAD RS04210 WP 011182164.1	MDITELLAFSVQNRASDLHLSAGIPPMIRVDGEVRRINLPAMDHSEVHNLIYDIMNDKQRRDYEEKLE	68
NTDB id 1283 ACE17W RS02630 WP 010947729.1	MDIAELLAFSVKNKSSDLHISAGMPPMIRVDGDLRKINLPPLLEHKDVIKIYDIMNDRQRKEYEEFFFE	68
NTDB id 1259 GC085 RS10305 WP 010947729.1	MDIAELLAFSVKNKSSDLHISAGMPPMIRVDGDLRKINLPPLLEHKDVIKIYDIMNDRQRKEYEEFFFE	68
NTDB id 1177 A1552VC RS01165 WP 000422572.1	MELNQYLDGMLTHKASDLYITVGAPILYRVDGELRAQGEALSVAD.VTALLHAMMDARQAEFKQTRE	67
NTDB id 238197 CFH90 RS07725 WP 126036367.1	MYSAEELLEEARMMMF.HMLSKVVEYGGSDLFISADFPPIKHKQGLMKPLGQQNLPCDQTKLFAYSIMNEKQRLFEFETELE	79
NTDB id 1203 PSJM300 01605 AFN76401.1	MEFEKLLRLMVEKGGSDLFITAGVPPSMKVNGKILPVNKTTPMSPEMTRETVHGVMNEQQRRREFTENHE	68
NTDB id 1019 ACIAD RS04205 WP 004922051.1	MDFNDLLNLMKHKASDLFITAGVEPSLKINGEITPVSKTKLDGDMIAQLIDSITTQKQRQEFIEETHE	68
consensus	! ** * * ! * * * * ! ! * * * * * * * * * * ! * * * * * * * * * * !	

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NTDB id 1020 ACIAD RS04210 WP 011182164.1	TDFSFEPN.LARFRVNVFNQNRGAGAVFRTIPSQVLTLEELGLGKIFRDI CDYPRGLVLVLTGPTGSGKSTTLAAMDYI	147
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TDFSFETIAN.LARFRVNAFNQSRGAAAVFRTIPSEILSMEDLGLPPIFKEMASFSRGLVLVLTGPTGSGKSTTLAAVIDYI	147
NTDB id 1259 GC085 RS10305 WP 010947729.1	TDFSFETIAN.LARFRVNAFNQSRGAAAVFRTIPSEILSMEDLGLPPIFKEMASFSRGLVLVLTGPTGSGKSTTLAAVIDYI	147
NTDB id 1177 A1552VC RS01165 WP 000422572.1	ANFAVVRD..SGRFRVSFAFFQRELPGAVIRRIETRIPTFEELKLPEVLQNLATAKRGVLVLVVGATGSGKSTTMAAMTGyr	145
NTDB id 238197 CFH90 RS07725 WP 126036367.1	CNFAISVPN.VSRFRVNVFQQQLHVGMVIRITITAEIPNFTKLQLPTSCLKDIIMEKRGVLVLVVGATGSGKSTSLAAMIDHR	158
NTDB id 1203 PSJM300 01605 AFN76401.1	CNFAISARG.IGRFRVSFAFYQRNLAGMVLRRITETNIPTIEDLKLDPVLKKLALTKRGVLVLVVGATGTGKSTSLAAMIGYR	147
NTDB id 1019 ACIAD RS04205 WP 004922051.1	CNFAILNREKNERFRVSFAFQQRDMPGMVIIRRIETKIPSPDDLHLPPILKELAMSKRGITIFSGATGTGKSTSLASMIQHR	148
consensus	* ! * * * * * * * * * * ! * * * * * * * * * * ! * * * * * * * * * * ! * * * * * * * * * *	

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NTDB id 1020 ACIAD RS04210 WP 011182164.1	NNHRYDHILTVEDPIEFVHEPKKCLINQREVHRDTLGFNEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGT	227
NTDB id 1283 ACE17W RS02630 WP 010947729.1	NSSRYEHILTVEDPIEFVHQSKKCLVNQREVHKDTLSFNAALRSALREDPDIILVGELRDLETIRLAMTAAETGHLVFGT	227
NTDB id 1259 GC085 RS10305 WP 010947729.1	NSSRYEHILTVEDPIEFVHQSKKCLVNQREVHKDTLSFNAALRSALREDPDIILVGELRDLETIRLAMTAAETGHLVFGT	227
NTDB id 1177 A1552VC RS01165 WP 000422572.1	NQHRTGHILTVEDPIEFVHEHKRCIVTQREVGLDTESYEVALKNSLRQAPDMILIGEIRSRRETMEYAMTFAETGHLCMAT	225
NTDB id 238197 CFH90 RS07725 WP 126036367.1	NENSAGHIITVEDPVEYVHKHKKSMITHREVGVDCSHWNALKNLTRQAPDVLIGEIRDTEHAIATAETGHLCLGT	238
NTDB id 1203 PSJM300 01605 AFN76401.1	NKNSSGHIISIEDPIEFITHQHQCIVTQREVGITDTSFEVALKNTLRQAPDVLIGEVRTRETMDHAVATAETGHLCLAT	227
NTDB id 1019 ACIAD RS04205 WP 004922051.1	NQHSKGHIITIEDPIEFITHEHAGCITITQREVGITDTSFEIALKNLTRQAPDVLIGEIRSRREVMYALATAETGHLVFAT	228
consensus	! * * ! * * * * * ! * * * * * * * * * * ! * * * * * * * * * * ! * * * * * * * * * *	

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NTDB id 1020 ACIAD RS04210 WP 011182164.1	LHTTSAAKTIDRIIDVFPAEEKDMVRTMLSESLSQAVISQTLLKK.EGGGRVAAHEIMTIGIPAIRNLIREGKVAQMYSSIQ	306
NTDB id 1283 ACE17W RS02630 WP 010947729.1	LHTNSATKTIINRIIDVFPAEEKSMVRSMLSESLSQAVVAQSLLKK.NKGGRVAALEIMMCTGAIRNLIREDKIAQMYSSIQ	306
NTDB id 1259 GC085 RS10305 WP 010947729.1	LHTNSATKTIINRIIDVFPAEEKSMVRSMLSESLSQAVVAQSLLKK.NKGGRVAALEIMMCTGAIRNLIREDKIAQMYSSIQ	306
NTDB id 1177 A1552VC RS01165 WP 000422572.1	LHANNANQALERILHLPKEQREQFLDLDSLNLKGVIQAQLLRDKNKGKGRHGVFEVLLNSPRIADLIRRGELHELKATMA	305
NTDB id 238197 CFH90 RS07725 WP 126036367.1	LHANNANQALDRIINFFPDERRLMDLSSNMKATISQRLVRTEGRCRRATVEIMLNTPLMSDLILKGNFHELKEVMS	318
NTDB id 1203 PSJM300 01605 AFN76401.1	LHANNANQALDRIINFFPADRNQVQVMDLSLNLKAIQAQLVPTPDGKRRRAVIEVLINTPLAADLIRKGEVHELKGLMK	307
NTDB id 1019 ACIAD RS04205 WP 004922051.1	LHANNANQTLERIIHFFDADRHQQIFMDLSLNLKAVISQQLIPTHDQGSRRAAVEVLINTPLLSDYIRQGEIYKIKELMA	308
consensus	! * * * * * ! * * * * * * * * * * ! * * * * * ! * * * * * * * * * * ! * * * * * * * * * *	

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