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NTDB id 1177 A1552VC RS01165 WP 000422572.1	MEINQYLDGMLTHKASDLYITVVGAPILYRVDGELRAQGE.ALSVADVTTALLHAMMDARQAEEFKQTREAN	69
NTDB id 189996 A6L28 RS08585 WP 002221204.1	MFTDENMTAKEELFAWLHRHMNQNGSDFVTTTHFPAMKLDGKITRITDEPLTAEKCMETAFSIMSAKQAEESSTNECN	80
NTDB id 1203 PSJM300 01605 AFN76401.1	MEFEKLLRLMVEKGGSDLFITAGVPPSMKVNGKILPVNKTPTSPEMTRETIVHGVNMNEQQRREFTENHECN	70
NTDB id 1019 ACIAD RS04205 WP 004922051.1	MDFNLLNLMAKHKASDLFITAGVEPSLKINGEITPVSKTKLDGDMIAQLIDSITTQKQRQEFIIETHECN	70
NTDB id 1307 DR RS10055 WP 010888596.1	MTSGIDITDILRLAADKGASDITLAAGLPPQFKLSGDYDAQGFEP LSASDTRKLMYSMMNERQQRIFEEREKRELD	74
NTDB id 1086 NMV RS00245 WP 002218487.1	MQITDLLAFGAKNKASDLHLSSGISPMIRVHGDMMRRINLPESASAEVGNMVTSMVNDHQRKIYQQNLEVD	70
NTDB id 1283 ACE17W RS02630 WP 010947729.1	MDIAELLAFSVKKNSSDLHISAGMPPMIRVDGDLRKINLPPLLEHKDVIKIITYDIMNDRQRKEYEEFFETD	70
NTDB id 1259 GC085 RS10305 WP 010947729.1	MDIAELLAFSVKKNSSDLHISAGMPPMIRVDGDLRKINLPPLLEHKDVIKIITYDIMNDRQRKEYEEFFETD	70
NTDB id 1194 PAKAF RS01995 WP 003084552.1	MDITELLAFSAKQGASDLHLSAGLPPMIRVDGDVRRINLPPLLEHKQVHALIYDIMNDKQRKDFEEFLETD	70
NTDB id 1202 PSJM300 01600 AFN76400.1	MDITELLAFSAKQGASDLHLSAGLPPMIRVDGDVRRINLPAMDHKQVHALIYDIMNDKQRKDFEEFLETD	70
NTDB id 1020 ACIAD RS04210 WP 011182164.1	MDITELLAFSVQNRASDLHLSAGIPPMIRVDGEVRRINLPAMDHSEVHNLIYDIMNDKQRRDYEEKLETD	70
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	MDITELLAFSVKNGASDLHLSAGMPPMIRVDGEVRRINLPALPHKDVHRLVYDIMNDKQRRDYEEKLETD	70
NTDB id 1060 ABD1 RS04380 WP 000355489.1	MDITELLAFSVKNGASDLHLSAGMPPMIRVDGEVRRINLPALPHKDVHRLVYDIMNDKQRRDYEEKLETD	70
NTDB id 1040 HON27 RS13360 WP 000355489.1	MDITELLAFSVKNGASDLHLSAGMPPMIRVDGEVRRINLPALPHKDVHRLVYDIMNDKQRRDYEEKLETD	70
consensus	*****!***** *!!*****!*****!*****!*****!*****!*****!*****!*****!*****	

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	FAVVRD..SGRFRVSAFFQRELPGAVIRRIETRIPTFEELKLPVLQNLAIAKRGLVLVVGATGSGKSTTMAAMTGYRNQ	147
NTDB id 189996 A6L28 RS08585 WP 002221204.1	FAISLPD.TSRFRVNAMIQRGATALVFRAITSKIPKFESLNLPPVLKDVALKKRGLVIFVGGTSGSKSTSLASLIDYRNE	159
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NTDB id 1019 ACIAD RS04205 WP 004922051.1	FAILNREKNERFRVSAFQQORDMPGMIIRRIETKIPSPDDLHLPPILKLELAMSKRGIIIFSGATGTGKSTSLASMIQHRNQ	150
NTDB id 1307 DR RS10055 WP 010888596.1	FSFALGD.KARFRVNAFMQRGVNGVMRLIPTTVRTTAEMGLPQTVIDIANAPRGLVLVTGPTGSGKSTTLASMIDHINT	153
NTDB id 1086 NMV RS00245 WP 002218487.1	FSFELPN.VARFRVNAFNIGRGAFAVFRITIPSTVLSLEELKAPSIFQKIAESPRGMVLVTGPTGSGKSTTLAAMINYINE	149
NTDB id 1283 ACE17W RS02630 WP 010947729.1	FSFEIAN.LARFRVNAFNQSRGAFAVFRITIPSEILSMEDLGLPPIFKEMASFSRGLVLVTGPTGSGKSTTLAAVIDYINS	149
NTDB id 1259 GC085 RS10305 WP 010947729.1	FSFEIAN.LARFRVNAFNQSRGAFAVFRITIPSEILSMEDLGLPPIFKEMASFSRGLVLVTGPTGSGKSTTLAAVIDYINS	149
NTDB id 1194 PAKAF RS01995 WP 003084552.1	FSFEVPG.VARFRVNAFNQNRGAGAVFRITIPSKVLTMEELGMGEVFKRVS DVPRLVLVTGPTGSGKSTTLAAMDYLN	149
NTDB id 1202 PSJM300 01600 AFN76400.1	FSFEVPG.VARFRVNAFNQNRGAGAVFRITIPSKVLTMEELGMGEVFKRITDVPRLVLVTGPTGSGKSTTLAAMDYLN	149
NTDB id 1020 ACIAD RS04210 WP 011182164.1	FSFEVPN.LARFRVNQNRGAGAVFRITIPSKVLTLEELGLGKIFRDCDYPRGLVLVTGPTGSGKSTTLAAMDYINN	149
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	FSFEVPN.VARFRVNAFNQNRGAGAVFRITIPSKVLTMEDLGLGQIFKDCDYPRGIVLVTGPTGSGKSTTLAAMDYINE	149
NTDB id 1060 ABD1 RS04380 WP 000355489.1	FSFEVPN.VARFRVNAFNQNRGAGAVFRITIPSKVLTMEDLGLGQIFKDCDYPRGIVLVTGPTGSGKSTTLAAMDYINE	149
NTDB id 1040 HON27 RS13360 WP 000355489.1	FSFEVPN.VARFRVNAFNQNRGAGAVFRITIPSKVLTMEDLGLGQIFKDCDYPRGIVLVTGPTGSGKSTTLAAMDYINE	149
consensus	!***** *!!!!!!***** *****!*****!*****!*****!*****!*****!*****!*****!*****!	

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	HRTGHILTVEDPIEFVHEHKRCIVTQREVGLDTESYEVALKNSLRQAPDMILIGEIRSRETMEYAMTFAETGHLCMATLH	227
NTDB id 189996 A6L28 RS08585 WP 002221204.1	NSFGHIITIEDPIEFVHEHKNCIITQREVGVDTENWMAALKNTLRQAPDVLIGEIRDRETM DYAI AFAETGHLCMATLH	239
NTDB id 1203 PSJM300 01605 AFN76401.1	NSSGHIISIEDPIEFIHQHQCIVTQREVGIDTDSFEVALKNTLRQAPDVLIGEVRTRETM DHA V AFAETGHLCLATLH	229
NTDB id 1019 ACIAD RS04205 WP 004922051.1	HSKGHIITIEDPIEFIEHAGCIITQREVGIDTDSFEIALKNTLRQAPDVLIGEIRSREVM DYAL AFAETGHLVATLH	230
NTDB id 1307 DR RS10055 WP 010888596.1	TKKLHILTIEDPIEFMHPNKASIIINQREVGDSTLSFDDALRALRQAPDVLIVGEMRDYETIKAAVTA AETGHLVMTLH	233
NTDB id 1086 NMV RS00245 WP 002218487.1	TQPAHILTIEDPIEFVHQSKKSLINQRELHQHTLSFANALRSALREDPDVILVGE MRDPETIGLALTA AETGHLVFGTLH	229
NTDB id 1283 ACE17W RS02630 WP 010947729.1	SRYEHIITVEDPIEFVHQSKKCLVNQREVHKDTLSFNAALRSALREDPDIILVGE LRDLETIRLAMTAAETGHLVFGTLH	229
NTDB id 1259 GC085 RS10305 WP 010947729.1	SRYEHIITVEDPIEFVHQSKKCLVNQREVHKDTLSFNAALRSALREDPDIILVGE LRDLETIRLAMTAAETGHLVFGTLH	229
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TKYHHILTIEDPIEFVHESKKCLVNQREVHRDTLGFSEALRSALREDPDIILVGE MRDLETIRLALTA AETGHLVFGTLH	229
NTDB id 1202 PSJM300 01600 AFN76400.1	TKYHHILTIEDPIEFVHESKKCLVNQREVHRDTLGFSEALRSALREDPDIILVGE MRDLETIRLALTA AETGHLVFGTLH	229
NTDB id 1020 ACIAD RS04210 WP 011182164.1	HRYDHILTVEDPIEFVHEPKKCLINQREVHRDTLGFNEALRSALREDPDIILVGE MRDLETIRLALTA AETGHLVFGTLH	229
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	NRYDHILTVEDPIEFVHQSKKCLINQREVHRDTLGFNEALRSALREDPDIILVGE MRDLETIRLALTA AETGHLVFGTLH	229
NTDB id 1060 ABD1 RS04380 WP 000355489.1	NRYDHILTVEDPIEFVHQSKKCLINQREVHRDTLGFNEALRSALREDPDIILVGE MRDLETIRLALTA AETGHLVFGTLH	229
NTDB id 1040 HON27 RS13360 WP 000355489.1	NRYDHILTVEDPIEFVHQSKKCLINQREVHRDTLGFNEALRSALREDPDIILVGE MRDLETIRLALTA AETGHLVFGTLH	229
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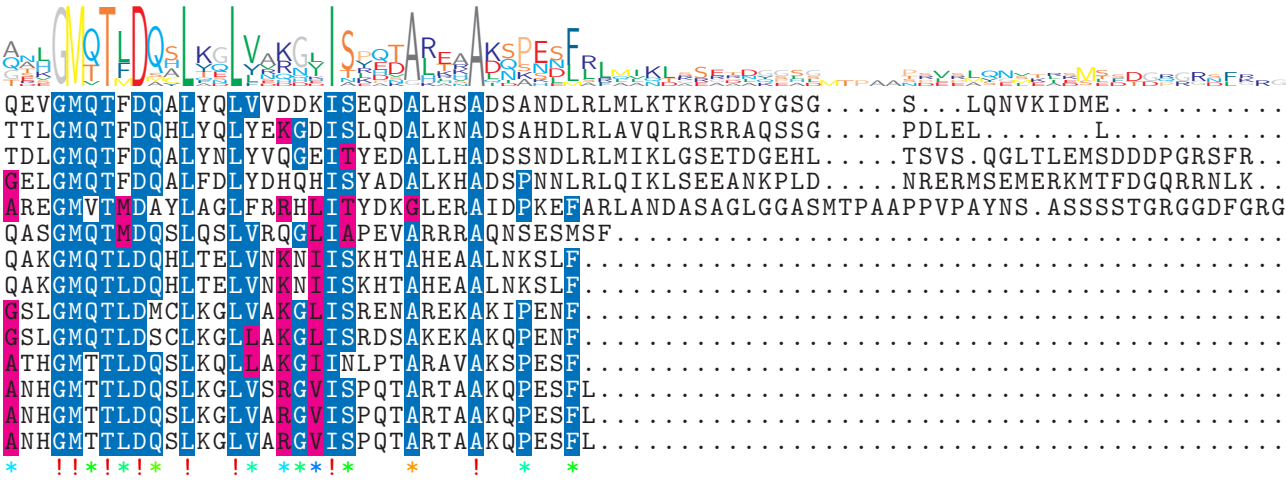
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NTDB	id	1203	PSJM300	01605	AFN76401.1	
NTDB	id	1019	ACIAD	RS04205	WP	004922051.1
NTDB	id	1307	DR	RS10055	WP	010888596.1
NTDB	id	1086	NMV	RS00245	WP	002218487.1
NTDB	id	1283	ACE17W	RS02630	WP	010947729.1
NTDB	id	1259	GC085	RS10305	WP	010947729.1
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NTDB	id	1081	FDQ49	RS05335	WP	006581387.1
NTDB	id	1060	ABD1	RS04380	WP	000355489.1
NTDB	id	1040	HON27	RS13360	WP	000355489.1
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NTDB	id	1040	HON27	RS13360	WP	000355489.1
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NTDB	id	1040	HON27	RS13360	WP	000355489.1
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 non conserved
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 $\geq 50\%$ conserved