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NTDB id 1168	A1552VC	RS11070	WP 000957200.1	KQLEALRPQCMILMTGPTGSGKTVSLYTGLNINTEELNISTAE DPVEINLPGINQVQVQPKIGFGFAEALRSFLRQ	385
NTDB id 1293	VP	RS12240	WP 005479695.1	KQLYLEALRRPQGMILMTGPTGSGKTVSLYTGLNINLKPETINISTAE DPVEINLPGINQVQVQPKIGFGFAEALRSFLRQ	385
NTDB id 1401	DSB67	RS12665	WP 010643259.1	KQLYLDALRRPQGMILMTGPTGSGKTVSLYTGLSILNKPETINISTAE DPVEIYLPGINQVQVQPKIGFGFAEALRSFLRQ	385
NTDB id 1110	NGFG	RS09235	WP 003689817.1	KKLLLEATHRPYCMVLVTGPTGSGKTVSLYTCLNINLNTESVNIAAE DPAEINLPGINQVNVNDKQGTFAAALRSFLRQ	376
NTDB id 1251	GCO85	RS07725	WP 011946523.1	RTNFLKAIQRPQGMILVTGPTGSGKTVTLYTALNINLNTIEVNISTAE DPVEIKVPGINQVNI NPKAGETFSGALRSFLRQ	394
NTDB id 999859	ABFU36	RS05545	WP 011038212.1	QKFLDIAIHKPYCMVLVTGPTGSGKTVSLYTALGILNDETRNISTAE DPVEIRLPGVNVQVQVQVNRGCMTFAAALRSFLRQ	398
NTDB id 1058	ABD1	RS01615	WP 001274986.1	KALFMEALDKPQGMILLTGPTGSGKTVSLYTGLNINLNTESNISTAE DPVEINLEGINQVNVNPKVGLTFAAALRSFLRQ	389
NTDB id 1018	ACIAD	RS01685	WP 004920473.1	KALFMEALNKPQGMILLTGPTGSGKTVSLYTGLNINLNTAHANISTAE DPVEINLEGVNVQVNVNPKVGLTFAAALRSFLRQ	389
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logo

NTDB id 1168	A1552VC	RS11070	WP 000957200.1	DPDWMNGEIRDLETAETAKAAQTGHLVSLHTNSAAETIRLNMGV EPNLASSLSLIAQRLARRLCCKHCKIAV.	464
NTDB id 1293	VP	RS12240	WP 005479695.1	DPDVVMVGEIRDLDTAETAIKASQTGHLVSLHTNSAAETIRLSNMGV ESFNLASSLSLIAQRLARKLCPYCKQPQ.	464
NTDB id 1401	DSB67	RS12665	WP 010643259.1	DPDVVMVGEIRDLDTAETAIKASQTGHLVSLHTNSAAETIRLSNMGV ESFNLASSLSLIAQRLARKLCPYCKQPQ.	464
NTDB id 1110	NGFG	RS09235	WP 003689817.1	DPDIIMVGEIRDLETAETAKAAQTGHLVSLHTNSAAETIRLNMGV APFNIASSVSLIMAQRLARRLCSSCKQEVE	456
NTDB id 1251	GCO85	RS07725	WP 011946523.1	DPDIIMVGEIRDLETAETAIKAAQTGHLVSLHTNSAAETIRLNMGV IPTFNIASSVSLIAQRLARKLCNQCKAVRD	474
NTDB id 999859	ABFU36	RS05545	WP 011038212.1	DPDIIMVGEIRDLETAETAIKAAQTGHLVSLHTNSAAETIRLNMGV IAPYNI TSSVTLVIAQRLARRLCNNCKRK.S	477
NTDB id 1058	ABD1	RS01615	WP 001274986.1	DPDIIMVGEIRDLETAETAIKAAQTGHLVSLHTNSAAETIRLNMGV PSFNIATSVNLVIAQRLARRLCSCCKIP.A	468
NTDB id 1018	ACIAD	RS01685	WP 004920473.1	DPDIIMVGEIRDLETAETAIKAAQTGHLVSLHTNSAAETIRLNMGV ASFNIATSVNLVIAQRLARRLCSCCKRP.I	468
consensus					

logo

NTDB id 1168	A1552VC	RS11070	WP 000957200.1	RPSALLQSQFAHQPNEL...IYEANAAGCNECT.GGYSGRVGIYEVMAFNTEIAEALMQRASIHQTERIAKANGMQTLQ	539
NTDB id 1293	VP	RS12240	WP 005479695.1	EHTVQL.QHIGIQTDDN...IFRANPDGCNECT.HGYSGRGTGIYEVMRFDESISEALIKGASVHELEKLAIANGMSTLQ	538
NTDB id 1401	DSB67	RS12665	WP 010643259.1	EPNSQL.QHIGIANTEBQ...IFQANPDGCNECT.HGYSGRGTGIYEVMMKFDESISEALIKGASVHELEKLAIANGMQTLQ	538
NTDB id 1110	NGFG	RS09235	WP 003689817.1	RPSASALKEVGFTDEDLAK.DWKLYRAVGCDCRGQCYKGRACVYEVMPISEEMQRVIMNNGTEVGLILDVAYKEGMVDLR	535
NTDB id 1251	GCO85	RS07725	WP 011946523.1	DFTNQGLIELGFKEADLVN..LKLYKAVGCDECT.SGYRGRVGLFEVLPMTKELGQLIMSGCNSLDILKLAQSEGMLTF	551
NTDB id 999859	ABFU36	RS05545	WP 011038212.1	TLPEHALLABGFTPEQVAN..IELYEAVGCDECT.EGYKGRGTGIYQVMPMTDEIGATVLEGGNAMQIAEAAQKIGTRDLR	554
NTDB id 1058	ABD1	RS01615	WP 001274986.1	DIPKQSLLEMGFTEQDLAHPDFRVQPVGCLECR.EGYKGRVGIYEVMMKVTPEISKIIMEDCNALEIAAASEKLCFNCLR	547
NTDB id 1018	ACIAD	RS01685	WP 004920473.1	QVPERSLLEMGFTEPDLAQPEFQIFEPVGCHDCR.EGYKGRVGIYEVMMKVTPEISKIIMEDCNALEIAATAETLCFNCLR	547
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logo

NTDB id 1168	A1552VC	RS11070	WP 000957200.1	ESGLEKIREGITSFAELQRVLYF	562
NTDB id 1293	VP	RS12240	WP 005479695.1	MSGIEKIKQGITSFSELQRVLYF	561
NTDB id 1401	DSB67	RS12665	WP 010643259.1	MSGIEKIKQGITSFRELQRVLYF	561
NTDB id 1110	NGFG	RS09235	WP 003689817.1	RAGILKIMQGITSLEEV TANTND	558
NTDB id 1251	GCO85	RS07725	WP 011946523.1	QSGIEKVKEGITTIEEVNRVTVD	574
NTDB id 999859	ABFU36	RS05545	WP 011038212.1	QSALMKASHGVTSIAEINRVTKD	577
NTDB id 1058	ABD1	RS01615	WP 001274986.1	RSCLKKVMQCVTSLQEVNRVTSE	570
NTDB id 1018	ACIAD	RS01685	WP 004920473.1	RSGLKKVMQCVTSLQEINRVTSE	570
consensus					

 non conserved
 similar
 $\geq 50\%$ conserved