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NTDB id 1251 GCO85 RS07725 WP 011946523.1	MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEEKISLLQYIVKNKILSAEQIALTAAQNFQVPLMDINCIDVGTI	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1MLTNLVAILRQAEELISATQEQAVVTQVSASGTSVPEALLELSIFHAQELTEQLSHIFGLPETDLSRYDYANL	72
NTDB id 1086 NMV RS00245 WP 002218487.1	0
NTDB id 1116 NGFG RS10615 WP 003690200.1	0
NTDB id 1185 GTF74 RS11940 WP 000350195.1	0
NTDB id 1176 A1552VC RS01160 WP 000350195.1	0
NTDB id 1283 ACE17W RS02630 WP 010947729.1	0
NTDB id 1259 GCO85 RS10305 WP 010947729.1	0
NTDB id 1202 PSJM300 01600 AFN76400.1	0
NTDB id 1194 PAKAF RS01995 WP 003084552.1	0
NTDB id 1020 ACIAD RS04210 WP 011182164.1	0
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	0
NTDB id 1060 ABD1 RS04380 WP 000355489.1	0
NTDB id 1040 HON27 RS13360 WP 000355489.1	0
NTDB id 1145336 DXE33 RS03965 WP 331851814.1	0
NTDB id 1307 DR RS10055 WP 010888596.1	0
NTDB id 1177 A1552VC RS01165 WP 000422572.1	0
NTDB id 1203 PSJM300 01605 AFN76401.1	0
NTDB id 1019 ACIAD RS04205 WP 004922051.1	0
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NTDB id 1251 GCO85 RS07725 WP 011946523.1	PVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTGLNTHAIVVETDKLSALIDNLLTAKESQGLSEYVE	160
NTDB id 1168 A1552VC RS11070 WP 000957200.1	CQQLGLRELITRYDALPIAKQGNLLLLAVSDPTLLQAEEEFRFATGLQVELALADHRALQAAIRRLYGRS.IQGAANQGK	151
NTDB id 1086 NMV RS00245 WP 002218487.1	0
NTDB id 1116 NGFG RS10615 WP 003690200.1	0
NTDB id 1185 GTF74 RS11940 WP 000350195.1	0
NTDB id 1176 A1552VC RS01160 WP 000350195.1	0
NTDB id 1283 ACE17W RS02630 WP 010947729.1	0
NTDB id 1259 GCO85 RS10305 WP 010947729.1	0
NTDB id 1202 PSJM300 01600 AFN76400.1	0
NTDB id 1194 PAKAF RS01995 WP 003084552.1	0
NTDB id 1020 ACIAD RS04210 WP 011182164.1	0
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	0
NTDB id 1060 ABD1 RS04380 WP 000355489.1	0
NTDB id 1040 HON27 RS13360 WP 000355489.1	0
NTDB id 1145336 DXE33 RS03965 WP 331851814.1	0
NTDB id 1307 DR RS10055 WP 010888596.1	0
NTDB id 1177 A1552VC RS01165 WP 000422572.1	0
NTDB id 1203 PSJM300 01605 AFN76401.1	0
NTDB id 1019 ACIAD RS04205 WP 004922051.1	0
consensus		

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NTDB id 1251	GCO85	RS07725	WP 011946523.1	QCKAVRDDFTNQGLIELGFKEADLVMLKLYKAVGCEQCTSGYRGRVGLFVLPMTKEIGQLIMSGGN.SLDLTKLAQSE	545
NTDB id 1168	A1552VC	RS11070	WP 000957200.1	HCKIAVRP...SALLQSQFAFQPNEILYEANAAGCNECTGGYSGRVGIYEVMAFNTEIAEAIMGRAS.IHQIERLAKAN	533
NTDB id 1086	NMV	RS00245	WP 002218487.1	312
NTDB id 1116	NGFG	RS10615	WP 003690200.1HDCNGRVASHEILLIANPAVRNLIRENKITQINSVLQTGQAS	312
NTDB id 1185	GTF74	RS11940	WP 000350195.1VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAH	311
NTDB id 1176	A1552VC	RS01160	WP 000350195.1VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAH	311
NTDB id 1283	ACE17W	RS02630	WP 010947729.1NKGGRVAALIEIMCTGAIRNLIREDKIAQMYSIIQTGQAK	311
NTDB id 1259	GCO85	RS10305	WP 010947729.1NKGGRVAALIEIMCTGAIRNLIREDKIAQMYSIIQTGQAK	311
NTDB id 1202	PSJM300	01600	AFN76400.1VGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSIIQTGQSL	311
NTDB id 1194	PAKAF	RS01995	WP 003084552.1IGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSIIQTGQSL	311
NTDB id 1020	ACIAD	RS04210	WP 011182164.1EGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSIIQTGATH	311
NTDB id 1081	FDQ49	RS05335	WP 006581387.1NCGGRVAAHEIMIGTPAIRNLIRENKVAQMYSIIQTGANH	311
NTDB id 1060	ABD1	RS04380	WP 000355489.1NCGGRVAAHEIMIGTPAIRNLIRENKVAQMYSIIQTGANH	311
NTDB id 1040	HON27	RS13360	WP 000355489.1NCGGRVAAHEIMIGTPAIRNLIRENKVAQMYSIIQTGANH	311
NTDB id 1145336	DXE33	RS03965	WP 331851814.1KDAFGRIGAFVMTCPNPAVRNLIRENKVFCIPSVMQMARGE	214
NTDB id 1307	DR	RS10055	WP 010888596.1ADCGGRVLAIEILLANPAVRALIREGKTFQITSVMQTGARE	316
NTDB id 1177	A1552VC	RS01165	WP 000422572.1KNCKGRHCVFVLLNSPRIADLIRGETHELKATMARSQEV	310
NTDB id 1203	PSJM300	01605	AFN76401.1PDCGRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDL	312
NTDB id 1019	ACIAD	RS04205	WP 004922051.1HDCQSRRAAVEVLINTPLISDYIRGETYKIKELMARSCEL	313
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NTDB id 1251	GCO85	RS07725	WP 011946523.1	GMLTIFQSGIEKVKEGIIIEEVNRVTVD.....	574
NTDB id 1168	A1552VC	RS11070	WP 000957200.1	GMQTLQESGLEKIREGITSFAELQRVLYF.....	562
NTDB id 1086	NMV	RS00245	WP 002218487.1	GMQTMDDSLQSLVRQGLIAPPEARARRAQNSEMSF.....	347
NTDB id 1116	NGFG	RS10615	WP 003690200.1	GMQTMDDSLQSLVRQGLIAPPEARARRAQNSEMSF.....	347
NTDB id 1185	GTF74	RS11940	WP 000350195.1	GMQTMENAKQLIARCVVDAQEVQSKIELDLKAF.....	345
NTDB id 1176	A1552VC	RS01160	WP 000350195.1	GMQTMENAKQLIARCVVDAQEVQSKIELDLKAF.....	345
NTDB id 1283	ACE17W	RS02630	WP 010947729.1	GMQTLDDQHLTELNVNKNIIISKHTAHEAALNK.SLF.....	344
NTDB id 1259	GCO85	RS10305	WP 010947729.1	GMQTLDDQHLTELNVNKNIIISKHTAHEAALNK.SLF.....	344
NTDB id 1202	PSJM300	01600	AFN76400.1	GMQTLDSCLKGLIAKGLISRDSAKEKAKQP.ENF.....	344
NTDB id 1194	PAKAF	RS01995	WP 003084552.1	GMQTLDMCLKGLVAKGLISRRENAREKAKIP.ENF.....	344
NTDB id 1020	ACIAD	RS04210	WP 011182164.1	GMTTLDQSLKQLIAKGLINLPTARAVAKSP.ESF.....	344
NTDB id 1081	FDQ49	RS05335	WP 006581387.1	GMTTLDQSLKGLVSRGVISPQTARTAAKQP.ESFL.....	345
NTDB id 1060	ABD1	RS04380	WP 000355489.1	GMTTLDQSLKGLVSRGVISPQTARTAAKQP.ESFL.....	345
NTDB id 1040	HON27	RS13360	WP 000355489.1	GMTTLDQSLKGLVSRGVISPQTARTAAKQP.ESFL.....	345
NTDB id 1145336	DXE33	RS03965	WP 331851814.1	GMMTMEASLQQLVQSGQIEAT.....	235
NTDB id 1307	DR	RS10055	WP 010888596.1	GMVTMDAYLAGLFRHIIITDYDKLERALDPKEFARLANDASAGLGG.....ASMTPAAPPVPAYNSASSSSTGRGGDFGR	391
NTDB id 1177	A1552VC	RS01165	WP 000422572.1	GMQTFDQALYQLVDDKISEQDALHSADSANDLRLMLKTK.RGDDY...GSGSLQNVKI.....DME.....	368
NTDB id 1203	PSJM300	01605	AFN76401.1	GMQTFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLG.SETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR.	381
NTDB id 1019	ACIAD	RS04205	WP 004922051.1	GMQTFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLS.EEANKPLDNREREMSEMER.....KMTFDGQRRNLK.	383
consensus				!!!*** * ** ***** * *	

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NTDB id 1251 GCO85 RS07725 WP 011946523.1	574
NTDB id 1168 A1552VC RS11070 WP 000957200.1	562
NTDB id 1086 NMV RS00245 WP 002218487.1	347
NTDB id 1116 NGFG RS10615 WP 003690200.1	347
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NTDB id 1040 HON27 RS13360 WP 000355489.1	345
NTDB id 1145336 DXE33 RS03965 WP 331851814.1	235
NTDB id 1307 DR RS10055 WP 010888596.1	GAAAPATPAAPAAGRSEPGRAGGFGRR	418
NTDB id 1177 A1552VC RS01165 WP 000422572.1	368
NTDB id 1203 PSJM300 01605 AFN76401.1	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	383
consensus		

⊠ non conserved
⊠ similar
⊠ ≥ 50% conserved