

logo

NTDB id 1251 GCO85 RS07725 WP 011946523.1	MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVKNKILSAEQIALTAAGNFQVPMLEINCLDVETI	80
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
NTDB id 101170 AABJ78 RS04005 WP 001295381.1		0
NTDB id 1307 DR RS10055 WP 010888596.1		0
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
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NTDB id 1251 GCO85 RS07725 WP 011946523.1	PVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTGLNTHAIVVETDKLSALIDNLLTAKESQGLSEYVE	160
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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
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consensus		

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NTDB id 1251	GC085	RS07725	WP 011946523.1	DSGDLEGLEISADDEDQDIDTATSVTDDAPIVKFV	NDITELLAFSVKNGASDLHL	SA	240
NTDB id 1177	A1552VC	RS01165	WP 000422572.1				44
NTDB id 1203	PSJM300	01605	AFN76401.1				45
NTDB id 1019	ACIAD	RS04205	WP 004922051.1				45
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NTDB id 1307	DR	RS10055	WP 010888596.1				49
NTDB id 1086	NMV	RS00245	WP 002218487.1				45
NTDB id 1116	NGFG	RS10615	WP 003690200.1				45
NTDB id 1185	GTF74	RS11940	WP 000350195.1				45
NTDB id 1176	A1552VC	RS01160	WP 000350195.1				45
NTDB id 1283	ACE17W	RS02630	WP 010947729.1				45
NTDB id 1259	GC085	RS10305	WP 010947729.1				45
NTDB id 1202	PSJM300	01600	AFN76400.1				45
NTDB id 1194	PAKAF	RS01995	WP 003084552.1				45
NTDB id 1020	ACIAD	RS04210	WP 011182164.1				45
NTDB id 1081	FDQ49	RS05335	WP 006581387.1				45
NTDB id 1060	ABD1	RS04380	WP 000355489.1				45
NTDB id 1040	HON27	RS13360	WP 000355489.1				45
consensus				***	*****	!!*****	*****!*****

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NTDB id 1251	GC085	RS07725	WP 011946523.1	SRITARIKVMSNLDISERRIPQDGGFKMKISKSR	AVFRVNAFNGRCAGAVRTIP	SV	319
NTDB id 1177	A1552VC	RS01165	WP 000422572.1				114
NTDB id 1203	PSJM300	01605	AFN76401.1				116
NTDB id 1019	ACIAD	RS04205	WP 004922051.1				117
NTDB id 101170	AABJ78	RS04005	WP 001295381.1				113
NTDB id 1307	DR	RS10055	WP 010888596.1				120
NTDB id 1086	NMV	RS00245	WP 002218487.1				116
NTDB id 1116	NGFG	RS10615	WP 003690200.1				116
NTDB id 1185	GTF74	RS11940	WP 000350195.1				116
NTDB id 1176	A1552VC	RS01160	WP 000350195.1				116
NTDB id 1283	ACE17W	RS02630	WP 010947729.1				116
NTDB id 1259	GC085	RS10305	WP 010947729.1				116
NTDB id 1202	PSJM300	01600	AFN76400.1				116
NTDB id 1194	PAKAF	RS01995	WP 003084552.1				116
NTDB id 1020	ACIAD	RS04210	WP 011182164.1				116
NTDB id 1081	FDQ49	RS05335	WP 006581387.1				116
NTDB id 1060	ABD1	RS04380	WP 000355489.1				116
NTDB id 1040	HON27	RS13360	WP 000355489.1				116
consensus				**	*****	*****	*****

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NTDB id 1251 GCO85 RS07725 WP 011946523.1
NTDB id 1177 A1552VC RS01165 WP 000422572.1
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NTDB id 1019 ACIAD RS04205 WP 004922051.1
NTDB id 101170 AABJ78 RS04005 WP 001295381.1
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NTDB id 1116 NGFG RS10615 WP 003690200.1
NTDB id 1185 GTF74 RS11940 WP 000350195.1
NTDB id 1176 A1552VC RS01160 WP 000350195.1
NTDB id 1283 ACE17W RS02630 WP 010947729.1
NTDB id 1259 GCO85 RS10305 WP 010947729.1
NTDB id 1202 PSJM300 01600 AFN76400.1
NTDB id 1194 PAKAF RS01995 WP 003084552.1
NTDB id 1020 ACIAD RS04210 WP 011182164.1
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 1251	GC085 RS07725 WP 011946523.1	KAVRDDFTNQGLIELGFKEADLVNLKLYKAVGCEQCTSGYRGRVCLFVLPMTKEIGQLIMSGGNS..LDIILKLAQSEGM	547
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NTDB id 1203	PSJM300 01605 AFN76401.1	TPDGKGRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLGM	314
NTDB id 1019	ACIAD RS04205 WP 004922051.1	THDCQSRRAAVEVLINTPLISDYIRGEIYKIKELMARSCELGM	315
NTDB id 101170	AABJ78 RS04005 WP 001295381.1	D.KQEGRVALFELLINTPAVGNLIREGKTHQLPHVIQTGQQVGM	310
NTDB id 1307	DR RS10055 WP 010888596.1	HADGQGRVLAYEILLANPAVRALIREGKTFQITSVMTGAREGM	318
NTDB id 1086	NMV RS00245 WP 002218487.1	THDCNGRVASHEILLIANPAVRNLIRENKITQINSVLQTGQASGM	314
NTDB id 1116	NGFG RS10615 WP 003690200.1	THDCNGRVASHEILLIANPAVRNLIRENKITQINSVLQTGQASGM	314
NTDB id 1185	GTF74 RS11940 WP 000350195.1	R.VGGGRVACHEIMLATPAIRNLIREDKVAQMYSTIQTGAHGM	313
NTDB id 1176	A1552VC RS01160 WP 000350195.1	R.VGGGRVACHEIMLATPAIRNLIREDKVAQMYSTIQTGAHGM	313
NTDB id 1283	ACE17W RS02630 WP 010947729.1	K.NKGGRVAALEIMMCTGAIRNLIREDKIAQMYSSIQTGQAKGM	313
NTDB id 1259	GC085 RS10305 WP 010947729.1	K.NKGGRVAALEIMMCTGAIRNLIREDKIAQMYSSIQTGQAKGM	313
NTDB id 1202	PSJM300 01600 AFN76400.1	K.VGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSSIQTGCSLGM	313
NTDB id 1194	PAKAF RS01995 WP 003084552.1	K.IGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSAIQTGCSLGM	313
NTDB id 1020	ACIAD RS04210 WP 011182164.1	K.EGGGRVAAHEIMIGTPAIRNLIREGKVAQMYSSIQTGATHGM	313
NTDB id 1081	FDQ49 RS05335 WP 006581387.1	K.NGGGRVAAHEIMIGTPAIRNLIRENKVAQMYSAIQTGANHGM	313
NTDB id 1060	ABD1 RS04380 WP 000355489.1	K.NGGGRVAAHEIMIGTPAIRNLIRENKVAQMYSAIQTGANHGM	313
NTDB id 1040	HON27 RS13360 WP 000355489.1	K.NGGGRVAAHEIMIGTPAIRNLIRENKVAQMYSAIQTGANHGM	313
consensus		* **!*!!!!!!*****!!** *!!!!!! ***** !!	

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NTDB id 1251	GC085 RS07725 WP 011946523.1	LITFQSGIEKVKEGITTIEEVNRVTVD.....	574
NTDB id 1177	A1552VC RS01165 WP 000422572.1	QTFDQALYQLVDDKISEQDALHSADSANDLRLMLKTKR.GDDY...GSGSLQNVKI.....DME.....	368
NTDB id 1203	PSJM300 01605 AFN76401.1	QTFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGS.ETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR...	381
NTDB id 1019	ACIAD RS04205 WP 004922051.1	QTFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLSE.EANKPLDNRERMSEMER.....KMTFDGQRRNLK...	383
NTDB id 101170	AABJ78 RS04005 WP 001295381.1	ITFQQSYQHRVGEGR.....	326
NTDB id 1307	DR RS10055 WP 010888596.1	VTMDAYLAGLFRHILITYDKCLERAIDPKEFARLANDASAGLGG....ASMTPAAPPVPAYNSASSSSTGRGGDFGRGA	393
NTDB id 1086	NMV RS00245 WP 002218487.1	QTMDDQLQSLVRQGLIAPEVARRRAQNSESMSF.....	347
NTDB id 1116	NGFG RS10615 WP 003690200.1	QTMDDQLQSLVRQGLIAPEAARRRAQNSESMSF.....	347
NTDB id 1185	GTF74 RS11940 WP 000350195.1	QTIMEQNAKQLIARGVVDAAQEVQSKIELDLKAF.....	345
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NTDB id 1283	ACE17W RS02630 WP 010947729.1	QTLDDQHLTELNVKNNTISKHTAHEAALNK.SLF.....	344
NTDB id 1259	GC085 RS10305 WP 010947729.1	QTLDDQHLTELNVKNNTISKHTAHEAALNK.SLF.....	344
NTDB id 1202	PSJM300 01600 AFN76400.1	QTLDSCLKGLIAKGLISRDSAKEKAKQP.ENF.....	344
NTDB id 1194	PAKAF RS01995 WP 003084552.1	QTLDMCLKGLIAKGLISRRENAKEKAKIP.ENF.....	344
NTDB id 1020	ACIAD RS04210 WP 011182164.1	TTLDDQLKQLIAKGLINLPTARAVAKSP.ESF.....	344
NTDB id 1081	FDQ49 RS05335 WP 006581387.1	TTLDDQLKGLVSRGVISPQTARTAAKQP.ESFL.....	345
NTDB id 1060	ABD1 RS04380 WP 000355489.1	TTLDDQLKGLVARGVISPQTARTAAKQP.ESFL.....	345
NTDB id 1040	HON27 RS13360 WP 000355489.1	TTLDDQLKGLVARGVISPQTARTAAKQP.ESFL.....	345
consensus		*!*** * ** **** * *	

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NTDB id 1251	GCO85	RS07725	WP 011946523.1		574
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
NTDB id 101170	AABJ78	RS04005	WP 001295381.1		326
NTDB id 1307	DR	RS10055	WP 010888596.1	AAPATPAAPAAGRSEPGRAGGFGRR	418
NTDB id 1086	NMV	RS00245	WP 002218487.1		347
NTDB id 1116	NGFG	RS10615	WP 003690200.1		347
NTDB id 1185	GTF74	RS11940	WP 000350195.1		345
NTDB id 1176	A1552VC	RS01160	WP 000350195.1		345
NTDB id 1283	ACE17W	RS02630	WP 010947729.1		344
NTDB id 1259	GCO85	RS10305	WP 010947729.1		344
NTDB id 1202	PSJM300	01600	AFN76400.1		344
NTDB id 1194	PAKAF	RS01995	WP 003084552.1		344
NTDB id 1020	ACIAD	RS04210	WP 011182164.1		344
NTDB id 1081	FDQ49	RS05335	WP 006581387.1		345
NTDB id 1060	ABD1	RS04380	WP 000355489.1		345
NTDB id 1040	HON27	RS13360	WP 000355489.1		345
consensus					

-  non conserved
-  similar
-  ≥ 50% conserved