

logo

NTDB	id	1159	ABDM36	RS04765	WP	001166105.1		0
NTDB	id	1150	A1552VC	RS08580	WP	001166105.1		0
NTDB	id	1395	DSB67	RS04445	WP	010649155.1		0
NTDB	id	1362	HI	1008	AAC22669.1		0	
NTDB	id	1382	A4U84	RS07375	WP	062924234.1		0
NTDB	id	595	KW2	RS08725	WP	021037687.1	MEKILDKLKEYWKIM.LLVVICALIAGGIFYIFTNSPKPAETLSVEN....LSSS..STKSSVSKFNSSSSSEK	65
NTDB	id	521	SMSK321	RS04960	WP	000443804.1	MESIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPHTPVKETNLQAE...VVAISKDSSTEKDVKKEEKEEPV	69
NTDB	id	493	SM12261	RS04300	WP	000387351.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAVSKDSVSEKEVKKEEKEEPL	69
NTDB	id	263	KZH43	RS04195	WP	000387330.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VAAVSKDLVSEKEVNKEEKEEPL	69
NTDB	id	222	SPD	RS04535	WP	000387330.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VAAVSKDLVSEKEVNKEEKEEPL	69
NTDB	id	188	SPR	RS04295	WP	000387330.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VAAVSKDLVSEKEVNKEEKEEPL	69
NTDB	id	153	SP	RS04730	WP	000387344.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VAAVSKDSSTEKEVKKEEKEEPL	69
NTDB	id	1069	ABD1	RS03060	WP	043960841.1		0
NTDB	id	1041	HON27	RS14740	WP	005124713.1		0
NTDB	id	105	BSU	25590	NP	390437.1	MNWLNQHKKAIILAASAAVFTAIMIFLATG....KNKE....PVKQAVPT....ETENTVVVKQEANNDE	57
NTDB	id	36	MW	RS08230	WP	001793467.1	MVLLYQFLLRYKDFLTQWKLYIISAVVLMVLIGFIFWRQDDYTSRNFENKDNALKQSTSENSSLSKVEDVQVKDGDNSK	80
NTDB	id	20	SA	RS08005	WP	001802022.1	MVLLYQFLLRYKDFLTQWKLYIISAVVLMVLIGFIFWRQDDYTSRNFENKDTALKQSTSENNSLSKLEDVQVKDGDNSK	80
NTDB	id	1014	ACIAD	RS13855	WP	004924483.1		0
NTDB	id	196660	FORC46	RS00050	WP	002853093.1		0
NTDB	id	1242	Cj0011c	YP	002343483.1			0
NTDB	id	1127	NGFG	RS10560	WP	003703428.1		0
NTDB	id	1126	NGFG	RS09375	WP	003703428.1		0
NTDB	id	1125	NGFG	RS07215	WP	003703428.1		0
NTDB	id	1108	NGFG	RS06525	WP	003703428.1		0
consensus								

logo

NTDB	id	1159	ABDM36	RS04765	WP	001166105.1	MQIKTK..IVTLFISLCLPT.....LPLLANAETAPAAQ.....	33
NTDB	id	1150	A1552VC	RS08580	WP	001166105.1	MQIKTK..IVTLFISLCLPT.....LPLLANAETAPAAQ.....	33
NTDB	id	1395	DSB67	RS04445	WP	010649155.1	MKWMLTLCLLT.....LAPMSWAE..TKTKAD.....	25
NTDB	id	1362	HI	1008	AAC22669.1	MKTL...FTSVVLCGAL.....VVSSSFAEEKATXQTAQSVVTT	36	
NTDB	id	1382	A4U84	RS07375	WP	062924234.1	MKKL...SRYTFSLLFG..L.....FSATTFABEKTVTSTEHPTVVM	37
NTDB	id	595	KW2	RS08725	WP	021037687.1	NKNEIMVDLKGAVAKPNVYQISSDE.RIVDLIRQAGGFTDQADQKSINLSAKIKDEEVIYVPKVCSSSSSESTDSP TG..	142
NTDB	id	521	SMSK321	RS04960	WP	000443804.1	EQDLITVDVKGAVKSPGIYDLVSVGS.RVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP..	146
NTDB	id	493	SM12261	RS04300	WP	000387351.1	EQDLITVDVKGAVKSPGIYDLPVGS.RVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEAASQQAGPGTA..	146
NTDB	id	263	KZH43	RS04195	WP	000387330.1	EQDLITVDVKGAVKSPGIYDLPVGS.RVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEAVSQQTGLGTA..	146
NTDB	id	222	SPD	RS04535	WP	000387330.1	EQDLITVDVKGAVKSPGIYDLPVGS.RVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEAASQQTGLGTA..	146
NTDB	id	188	SPR	RS04295	WP	000387330.1	EQDLITVDVKGAVKSPGIYDLPVGS.RVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEAASQQTGLGTA..	146
NTDB	id	153	SP	RS04730	WP	000387344.1	EQDLITVDVKGAVKSPGIYDLPVGS.RVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEAASQQTGS GTA..	146
NTDB	id	1069	ABD1	RS03060	WP	043960841.1	MQYSMRFWKNKYFVLLIMLC LGV.....ISPSVCAQSFDQNFQEWKVKQMYDQ	48
NTDB	id	1041	HON27	RS14740	WP	005124713.1	MQYSMRFWKNKYFLLIMLC LGV.....ISPSVCAQSFDQNFQEWKAKQMYDQ	48
NTDB	id	105	BSU	25590	NP	390437.1	SNETIVVDIKGAVQHHPGVYEMRTGD.RVSAIEKAGGTSEQADEAQVNLA EILQDGTIVVYIPKKCEETAV.....Q	127
NTDB	id	36	MW	RS08230	WP	001793467.1	NKGPVYVDVKGAVKHPNVYKMTSKD.RVVDLLDKAQ.LLEDADV SQINLSEKLTQDKMIFIPH.....K.GQKNVEP	149
NTDB	id	20	SA	RS08005	WP	001802022.1	NKGPVYVDVKGAVKHPNVYKMTSKD.RVVDLLDKAQ.LLDDADV SRINLSEKLTQDKMIFIPH.....K.GQKNVEP	149
NTDB	id	1014	ACIAD	RS13855	WP	004924483.1	MSIR.HFF...T....FFWAT.....LL..TLFSMQLVFAAQGFDQ QYDAWK TQQKAHDT	44
NTDB	id	196660	FORC46	RS00050	WP	002853093.1	MKKLLFLF...F.....	9
NTDB	id	1242	Cj0011c	YP	002343483.1	MKKLLFLF...F.....		9
NTDB	id	1127	NGFG	RS10560	WP	003703428.1	MSRGGA....TPYRFLLIR.HIV...ARCG..LLFAT.....LKGKTMKKMFVLF CML.....	43
NTDB	id	1126	NGFG	RS09375	WP	003703428.1	MSRGGA....TPYRFLLIR.HIV...ARCG..LLFAT.....LKGKTMKKMFVLF CML.....	43
NTDB	id	1125	NGFG	RS07215	WP	003703428.1	MSRGGA....TPYRFLLIR.HIV...ARCG..LLFAT.....LKGKTMKKMFVLF CML.....	43
NTDB	id	1108	NGFG	RS06525	WP	003703428.1	MSRGGA....TPYRFLLIR.HIV...ARCG..LLFAT.....LKGKTMKKMFVLF CML.....	43
consensus							* ** *	

* ** * * * * *

logo

NTDB id 1159	ABDM36	RS04765	WP 001166105.1	VEEGIVITVNINTASABELATLLKIGIGLKKAAQAVDYREANGPFTHIDDLTNVKGIGEATV	94
NTDB id 1150	A1552VC	RS08580	WP 001166105.1	VEEGIVITVNINTASABELATLLKIGIGLKKAAQAVDYREANGPFTHIDDLTNVKGIGEATV	94
NTDB id 1395	DSB67	RS04445	WP 010649155.1	KYEGIEITVMVNSATAQETATLLIGIGBKKAQDIVYRNEHGPFKTAADLTQVKGIGEATV	86
NTDB id 1362	HI 1008	AAC22669.1		QAEA.....QVAPAVVSDKLNINTATASEIQKSLTGICAKKABAVQYREKHCNFXNAEQLLLEVQGIGKATL	103
NTDB id 1382	A4U84	RS07375	WP 062924234.1	QQ.V.....AATQKVNANVVNINTATASEIQDKLVGIGAKKAAQAVYREKNGKFLNVEQLTEVSGIGKATL	103
NTDB id 595	KW2	RS08725	WP 021037687.1	S.....S..VSNQISTTSGPKVNINKADITELQK.LTGIGQKKAAQDIDDERMKNCGDFKSIDDLGKVSQFGDKTL	208
NTDB id 521	SMSK321	RS04960	WP 000443804.1	SSTSCKDKKLNINKASLEELKQ.VKCLGGKRAQDIDHREANGKFKSVDLKKVSGIGAKTI	206
NTDB id 493	SM12261	RS04300	WP 000387351.1	SSTSKEKKVNINKASLEELKQ.VKCLGGKRAQDIDHREANGKFKSVDLKKVSGIGAKTI	206
NTDB id 263	KZH43	RS04195	WP 000387330.1	SSISKEKKVNINKASLEELKQ.VKCLGGKRAQDIDHREANGKFKSVDLKKVSGIGGKTI	206
NTDB id 222	SPD	RS04535	WP 000387330.1	SSISKEKKVNINKASLEELKQ.VKCLGGKRAQDIDHREANGKFKSVDLKKVSGIGGKTI	206
NTDB id 188	SPR	RS04295	WP 000387330.1	SSISKEKKVNINKASLEELKQ.VKCLGGKRAQDIDHREANGKFKSVDLKKVSGIGGKTI	206
NTDB id 153	SP	RS04730	WP 000387344.1	SSTSKEKKVNINKASLEELKQ.VKCLGGKRAQDIDHREANGKFKSVDLKKVSGIGGKTI	206
NTDB id 1069	ABD1	RS03060	WP 043960841.1	KLNISKSSSHSNSSKISQTKNFNDSTGQIHLNQANVNEFQQ.LKGIGEKKAQAVYRQKNCSFKNIDETKNVKGIGPAIF	127
NTDB id 1041	HON27	RS14740	WP 005124713.1	KLNITKSSSHSNGSKISQTKNFNDSTGQIHLNQANVNEFQQ.LKGVGKKKAQAVYRQKNCSFKNIDETKNVKGIGPAIF	127
NTDB id 105	BSU	25590	NP 390437.1	QGGG.....GSV....QSDGGKGALVNINTATLEELQG.LSCVGPSKABATAYREENGCRFQITIEDITKVSGIGESF	195
NTDB id 36	MW	RS08230	WP 001793467.1	QIGVNSV.....HV....KNGNTNNTKVNINTASVSELSMS.VPCVQQAQANAIVEYRNQCGAFQETDDLKKVKGFCSKTF	219
NTDB id 20	SA	RS08005	WP 001802022.1	QIEVNSV.....HV....KNGNTNNTKVNINTASVSELSMS.VPCVQQAQANAIVEYRNQCGAFQETDDLKKVKGFCSKTF	219
NTDB id 1014	ACIAD	RS13855	WP 004924483.1	RLNSSASHHSLSKP....TQSGTGKVNINSATAKELQQ.LHCVGEKKALAIYRQQYCKFASVDDLQKVKGIGPKFI	119
NTDB id 196660	FORC46	RS00050	WP 002853093.1	ALTAFLFGAVNINTATLKEKLS.LNGICEAKAKAILEYRKE.ANFTSIDDLKKVKGIGDKLF	69
NTDB id 1242	Cj0011c	YP 002343483.1		ALTAFLFGAVNINTATLKEKLS.LNGICEAKAKAILEYRKE.ANFTSIDDLKKVKGIGDKLF	69
NTDB id 1127	NGFG	RS10560	WP 003703428.1	FSCAFSLAAVNINAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDLTKVKGIGPAVL	104
NTDB id 1126	NGFG	RS09375	WP 003703428.1	FSCAFSLAAVNINAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDLTKVKGIGPAVL	104
NTDB id 1125	NGFG	RS07215	WP 003703428.1	FSCAFSLAAVNINAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDLTKVKGIGPAVL	104
NTDB id 1108	NGFG	RS06525	WP 003703428.1	FSCAFSLAAVNINAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDLTKVKGIGPAVL	104
consensus				***! !** !* * !*! **!*!!***! ** !***** *!*!*! ***	

logo

NTDB id 1159	ABDM36	RS04765	WP 001166105.1	RNAARILL.....	103
NTDB id 1150	A1552VC	RS08580	WP 001166105.1	RNAARILL.....	103
NTDB id 1395	DSB67	RS04445	WP 010649155.1	RKNEDRILL.....	95
NTDB id 1362	HI 1008	AAC22669.1		EKNRDRILF.....	112
NTDB id 1382	A4U84	RS07375	WP 062924234.1	EKNRDCIVIE.....	113
NTDB id 595	KW2	RS08725	WP 021037687.1	EKLKDEISID.....	218
NTDB id 521	SMSK321	RS04960	WP 000443804.1	EKLKDYVTVD.....	216
NTDB id 493	SM12261	RS04300	WP 000387351.1	EKLKDYVTVD.....	216
NTDB id 263	KZH43	RS04195	WP 000387330.1	EKLKDYVTVD.....	216
NTDB id 222	SPD	RS04535	WP 000387330.1	EKLKDYVTVD.....	216
NTDB id 188	SPR	RS04295	WP 000387330.1	EKLKDYVTVD.....	216
NTDB id 153	SP	RS04730	WP 000387344.1	EKLKDYVTVD.....	216
NTDB id 1069	ABD1	RS03060	WP 043960841.1	EKNKSRIAL.....	136
NTDB id 1041	HON27	RS14740	WP 005124713.1	EKNKSRIAL.....	136
NTDB id 105	BSU	25590	NP 390437.1	EKLKSSITVK.....	205
NTDB id 36	MW	RS08230	WP 001793467.1	DKLKSIFTI.....	228
NTDB id 20	SA	RS08005	WP 001802022.1	DKLKSIFTI.....	228
NTDB id 1014	ACIAD	RS13855	WP 004924483.1	EKNRDRILAI.....	128
NTDB id 196660	FORC46	RS00050	WP 002853093.1	EKLKNDITIE.....	79
NTDB id 1242	Cj0011c	YP 002343483.1		EKLKNDITIE.....	79
NTDB id 1127	NGFG	RS10560	WP 003703428.1	AKLKDQASVGAPAPKGPAPVLPVAVKK	131
NTDB id 1126	NGFG	RS09375	WP 003703428.1	AKLKDQASVGAPAPKGPAPVLPVAVKK	131
NTDB id 1125	NGFG	RS07215	WP 003703428.1	AKLKDQASVGAPAPKGPAPVLPVAVKK	131
NTDB id 1108	NGFG	RS06525	WP 003703428.1	AKLKDQASVGAPAPKGPAPVLPVAVKK	131
consensus				***** ***	

 non conserved
 similar
 $\geq 50\%$ conserved