

logo

NTDB id 1159	ABDM36	RS04765	WP 001166105.1	MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAAPTEYKEINLSAEVVAVSKDLYEKEVSKKEEKKEEPL	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..STKSSSVSKFNSSSSSEK	65
NTDB id 521	SMSK321	RS04960	WP 000443804.1	MESIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPHTPVKETNLQAE...VVAISKDSSTEKDVKKEEKEEPV	69
NTDB id 493	SM12261	RS04300	WP 000387351.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAVSKDSVSEKEVNKKEEKKEEPL	69
NTDB id 263	KZH43	RS04195	WP 000387330.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAVSKDLVSEKEVNKKEEKKEEPL	69
NTDB id 222	SPD	RS04535	WP 000387330.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAVSKDLVSEKEVNKKEEKKEEPL	69
NTDB id 188	SPR	RS04295	WP 000387330.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAVSKDLVSEKEVNKKEEKKEEPL	69
NTDB id 153	SP	RS04730	WP 000387344.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAVSKDSSTEKEVNKKEEKKEEPL	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	MVLLYQFLLRYKDFTLTQWKLYIISAVVLMVLI GFIFWRQDDYTSRNFENKDNALKQSTSENSSLSKVEDVQVKDGDNSK	80
NTDB id 20	SA	RS08005	WP 001802022.1	MVLLYQFLLRYKDFTLTQWKLYIISAVVLMVLI GFIFWRQDDYTSRNFENKDTALKQSTSENNSLSKLEDVQVKDGDNSK	80
NTDB id 1014	ACIAD	RS13855	WP 004924483.1		0
NTDB id 1158117	QOR69	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	EQDLITVDVKCAVKSPGIYDLVSVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKCEESASQHTGSGAP..	33
NTDB id 1150	A1552VC	RS08580	WP 001166105.1	MQIKTK..IVTLFISLCLPT.....LPLLANAETAPAAQ.....	33
NTDB id 1395	DSB67	RS04445	WP 010649155.1	MQIKTK..IVTLFISLCLPT.....LPLLANAETAPAAQ.....	33
NTDB id 1362	HI 1008	AAC22669.1		MKWMITLCLL.....LAPMSWAE..TKTKAD.....	25
NTDB id 1382	A4U84	RS07375	WP 062924234.1	MKTL.....FTSVLCCAL.....VSSSFAEEKATXQTAQSVVTT	36
NTDB id 595	KW2	RS08725	WP 021037687.1	MKKL...SRYTFSLLFC..L.....FSATTFABEKVTTSTEHTPVVM	37
NTDB id 521	SMSK321	RS04960	WP 000443804.1	NKNEIMVDLKGAVAKPNVYQISSDE.RIVDLIRQAGGFTDQADQKSINLSAKLKDEEVIYVPKVCSSSSSESTDSP TG..	142
NTDB id 493	SM12261	RS04300	WP 000387351.1	EQDLITVDVKCAVKSPGIYDLVSVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKCEESASQHTGSGAP..	146
NTDB id 263	KZH43	RS04195	WP 000387330.1	EQDLITVDVKCAVKSPGIYDLVSVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKCEEASQQAGPGTA..	146
NTDB id 222	SPD	RS04535	WP 000387330.1	EQDLITVDVKCAVKSPGIYDLVSVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKCEEAVSQQTGLGTA..	146
NTDB id 188	SPR	RS04295	WP 000387330.1	EQDLITVDVKCAVKSPGIYDLVSVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKCEEAVSQQTGLGTA..	146
NTDB id 153	SP	RS04730	WP 000387344.1	EQDLITVDVKCAVKSPGIYDLVSVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKCEEAVSQQTGLGTA..	146
NTDB id 1069	ABD1	RS03060	WP 043960841.1	MQYSMRFWKNKYFVLLIMLCLGV.....ISPSVCAQSFDQNFQEWKVKQMYDQ	48
NTDB id 1041	HON27	RS14740	WP 005124713.1	MQYSMRFWKNKYFVLLIMLCLGV.....ISPSVCAQSFDQNFQEWKAKQMYDQ	48
NTDB id 105	BSU 25590	NP 390437.1		SNETIVVDIKGAVQHHPGVYEMRTGD.RVSAIEKAGGTSEQADEAQVNLAIEIQDGTVVYIPKKCEETAV.....Q	127
NTDB id 36	MW	RS08230	WP 001793467.1	NKGPVYVDVKCAVKHPNVYKMTSKD.RVVDLIDKAQ..LLEDADV SQINLSEKLTQDKMIFIPH.....K.GQKNVEP	149
NTDB id 20	SA	RS08005	WP 001802022.1	NKGPVYVDVKCAVKHPNVYKMTSKD.RVVDLIDKAQ..LLEDADVSRINLSEKLTQDKMIFIPH.....K.GQKNVEP	149
NTDB id 1014	ACIAD	RS13855	WP 004924483.1	MSIR.HFF...T....FFWAT.....LL..TLFSMQLVFAAQGFDQYDAWK TQQKAHDT	44
NTDB id 1158117	QOR69	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....TPYRFLLR.HIV...ARCG..LLFAT.....LKGKTMKKMFVLF CML.....	43
NTDB id 1126	NGFG	RS09375	WP 003703428.1	MSRGGA....TPYRFLLR.HIV...ARCG..LLFAT.....LKGKTMKKMFVLF CML.....	43
NTDB id 1125	NGFG	RS07215	WP 003703428.1	MSRGGA....TPYRFLLR.HIV...ARCG..LLFAT.....LKGKTMKKMFVLF CML.....	43
NTDB id 1108	NGFG	RS06525	WP 003703428.1	MSRGGA....TPYRFLLR.HIV...ARCG..LLFAT.....LKGKTMKKMFVLF CML.....	43
consensus					

* ** * * * * * * * *

logo

NTDB id 1159	ABDM36	RS04765	WP 001166105.1	VEEGIVITVNINTASABELATLLKIGIGLKKAAQAIVDYREANGPFTHIDDLTNVKGIGEATV	94
NTDB id 1150	A1552VC	RS08580	WP 001166105.1	VEEGIVITVNINTASABELATLLKIGIGLKKAAQAIVDYREANGPFTHIDDLTNVKGIGEATV	94
NTDB id 1395	DSB67	RS04445	WP 010649155.1	KYEGIEITVMVNSATAQETATLLIGIGBKKAQDIVYRNEHGPFKTAADLTQVKGIGEATV	86
NTDB id 1362	HI 1008	AAC22669.1		QAEA.....QVAPAVVSDKLNINTATASEIQKSLTGICAKKABAIQYREKHCNFXNAEQLLLEVQGIGKATL	103
NTDB id 1382	A4U84	RS07375	WP 062924234.1	QQ.V.....AATQKVNANVVNINTATASEIQDKLVGIGAKKAQAIVEYREKNGKFLNVEQLTEVSGIGKATL	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.LKGIGEKKAQAIVEYRQKNCSFKNIDETKNVKGIGPAIF	127
NTDB id 1041	HON27	RS14740	WP 005124713.1	KLNITKSSSHSNGSKISQTKNFNDSTGQIHLNQANVNEFQQ.LKGIGEKKAQAIVEYRQKNCSFKNIDETKNVKGIGPAIF	127
NTDB id 105	BSU	25590	NP 390437.1	QGGG.....GSV....QSDGGKGALVNINTATLEELQG.LSCVGPSSKABAIAYREENGCRFQITIEDITKVSGIGESF	195
NTDB id 36	MW	RS08230	WP 001793467.1	QIGVNSV.....HV....KNGNTNNTKVNINTASVSELSMS.VPCVQQAQANAIVEYRNQQGAFQETDDLKKVKGFCSKTF	219
NTDB id 20	SA	RS08005	WP 001802022.1	QIEVNSV.....HV....KNGNTNNTKVNINTASVSELSMS.VPCVQQAQANAIVEYRNQQGAFQETDDLKKVKGFCSKTF	219
NTDB id 1014	ACIAD	RS13855	WP 004924483.1	RLNSSASHHSLSKP....TQSGTGKVNINSATAKELQQ.LHCVGEKKALAIVEYRQQYCKFASVDDLQKVKGIGPKFI	119
NTDB id 1158117	QOR69	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 1158117	QOR69	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