

logo

NTDB id 1159	ABDM36	RS04765	WP 001166105.1	MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKDVKKEEKEEPV.....	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...STKSSVSKFNSSSSEK	65
NTDB id 521	SMSK321	RS04960	WP 000443804.1	MESIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPHTPVKETNLQAE...VVAISKDSSTEKDVKKEEKEEPV	69
NTDB id 493	SM12261	RS04300	WP 000387351.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSVSEKEVNKKEEKEEPL	69
NTDB id 263	KZH43	RS04195	WP 000387330.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDLVSEKEVNKKEEKEEPL	69
NTDB id 222	SPD	RS04535	WP 000387330.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDLVSEKEVNKKEEKEEPL	69
NTDB id 188	SPR	RS04295	WP 000387330.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDLVSEKEVNKKEEKEEPL	69
NTDB id 153	SP	RS04730	WP 000387344.1	MEAIIEKIKEYKIIIV.ICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKEVNKKEEKEEPL	69
NTDB id 1069	ABD1	RS03060	WP 043960841.1		0
NTDB id 1041	HON27	RS14740	WP 005124713.1		0
NTDB id 36	MW	RS08230	WP 001793467.1	MVLLYQFLLRYKDFLTQWKLYIISAVVLMVLI GFIFWRQDDYTSRNFENKDNALKQSTSENSSLSKVEDVQVKDGDNSK	80
NTDB id 20	SA	RS08005	WP 001802022.1	MVLLYQFLLRYKDFLTQWKLYIISAVVLMVLI GFIFWRQDDYTSRNFENKDTALKQSTSENSSLSKLEDVQVKDGDNSK	80
NTDB id 105	BSU 25590	NP 390437.1		MNWLNQHKKAIILAASAAVFTAIMIFLATG...KNKE...PVKQAVPTE...TENTVVVKQEANNDE	57
NTDB id 1014	ACIAD	RS13855	WP 004924483.1		0
NTDB id 1127	NGFG	RS10560	WP 003703428.1	MSRGG	5
NTDB id 1126	NGFG	RS09375	WP 003703428.1	MSRGG	5
NTDB id 1125	NGFG	RS07215	WP 003703428.1	MSRGG	5
NTDB id 1108	NGFG	RS06525	WP 003703428.1	MSRGG	5
NTDB id 152956	AEI04	RS08195	WP 002855574.1		0
NTDB id 1242	Cj0011c	YP 002343483.1			0
consensus					0

logo

NTDB id 1159	ABDM36	RS04765	WP 001166105.1	EQDLITVDYKGA VKSPG Y L L T Y S R P K V K L T L C M L E L A S K A G G L T E A D S K S L N L A Q K V S D E A L V Y V P T K G E E A V	33
NTDB id 1150	A1552VC	RS08580	WP 001166105.1	MQIKT.KIVTFLFLSLCLPTLPLLANAEETAPAAQ.....	33
NTDB id 1395	DSB67	RS04445	WP 010649155.1	MKWMLTLCLLILAPMSWAE.TKTKAD.....	25
NTDB id 1362	HI 1008	AAC22669.1		MKT.L.FTSVVLGALVSSSF AEEKATXQTAQSVV.....TTQA.....	38
NTDB id 1382	A4U84	RS07375	WP 062924234.1	MKK.LSRYTFSLLFG.LFSATTFAEEKVTTSTEHTPV.....VMQQ.....	39
NTDB id 595	KW2	RS08725	WP 021037687.1	NKNEIMVDLKGAVAKPNVYQISSDERLVDL.....RQAGGFTDQADQKSINLSAKLKDEEVIYVPKVGESS	133
NTDB id 521	SMSK321	RS04960	WP 000443804.1	EQDLITVDVKGAVKSPGIYDLVSGSRVNDV.....QKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEA	137
NTDB id 493	SM12261	RS04300	WP 000387351.1	EQDLITVDVKGAVKSPGIYDLVSGSRVNDV.....QKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEA	137
NTDB id 263	KZH43	RS04195	WP 000387330.1	EQDLITVDVKGAVKSPGIYDLVSGSRVNDV.....QKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEA	137
NTDB id 222	SPD	RS04535	WP 000387330.1	EQDLITVDVKGAVKSPGIYDLVSGSRVNDV.....QKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEA	137
NTDB id 188	SPR	RS04295	WP 000387330.1	EQDLITVDVKGAVKSPGIYDLVSGSRVNDV.....QKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEA	137
NTDB id 153	SP	RS04730	WP 000387344.1	EQDLITVDVKGAVKSPGIYDLVSGSRVNDV.....QKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEA	137
NTDB id 1069	ABD1	RS03060	WP 043960841.1	MQYSMRFWKN...K.YFVLLIMLC LGVISPSVCAQSFDQNFQEWKVKQQM..YDQK.....	49
NTDB id 1041	HON27	RS14740	WP 005124713.1	MQYSMRFWKN...K.YFLLIMLC LGVISPSVCAQSFDQNFQEWKAKQQM..YDQK.....	49
NTDB id 36	MW	RS08230	WP 001793467.1	NKGPVYVDVKGAVKHPNVYKMTSKDRVVD.LL.....DKAQLLEDADVSRINLSEKLTQDKMIFIPHKGQKNV	147
NTDB id 20	SA	RS08005	WP 001802022.1	NKGPVYVDVKGAVKHPNVYKMTSKDRVVD.LL.....DKAQLLEDADVSRINLSEKLTQDKMIFIPHKGQKNV	147
NTDB id 105	BSU 25590	NP 390437.1		SNETIVIDIKGAVQHPGVYEM.RTGDRVS.....QAI...EKAGGTSEQADEAQVNLAELIQDGTVVYIPKKGEETA	125
NTDB id 1014	ACIAD	RS13855	WP 004924483.1	MSIRHFFT...FFWAT.....LLTLFSMQI...VFAQQGFDQYDQAW.....KTQQKAHDTR	45
NTDB id 1127	NGFG	RS10560	WP 003703428.1	ATPYRFLIRHIVARCGLLFATLKGKTMKKMFVLF CMLF.....	44
NTDB id 1126	NGFG	RS09375	WP 003703428.1	ATPYRFLIRHIVARCGLLFATLKGKTMKKMFVLF CMLF.....	44
NTDB id 1125	NGFG	RS07215	WP 003703428.1	ATPYRFLIRHIVARCGLLFATLKGKTMKKMFVLF CMLF.....	44
NTDB id 1108	NGFG	RS06525	WP 003703428.1	ATPYRFLIRHIVARCGLLFATLKGKTMKKMFVLF CMLF.....	44
NTDB id 152956	AEI04	RS08195	WP 002855574.1	MKKL LFLF..FA.....	10
NTDB id 1242	Cj0011c	YP 002343483.1		MKKL LFLF..FA.....	10
consensus					10

logo

NTDB id 1159	ABDM36	RS04765	WP 001166105.1	VEEGIVITVNINTASAEELATLLKGI	GLKKAQAI	VDYREANGP	FTH	IDDLTNVKGIG	EATVR	95	
NTDB id 1150	A1552VC	RS08580	WP 001166105.1	VEEGIVITVNINTASAEELATLLKGI	GLKKAQAI	VDYREANGP	FTH	IDDLTNVKGIG	EATVR	95	
NTDB id 1395	DSB67	RS04445	WP 010649155.1	KYEGIEITVNVNSATAQEIATLLIGI	GEKKAQD	IVEYRNEHG	PFKTAADL	TKVKGIG	EATVR	87	
NTDB id 1362	HI 1008	AAC22669.1	EAQVAPAVVSDKLNINTATASEIQKS	LTGIGAKKA	EAI	VQYREKHGN	FXNAE	QQLLEVQ	GIGKATLE	104	
NTDB id 1382	A4U84	RS07375	WP 062924234.1	VAATQKVNANV	VNINTATASEIQDK	LVGIGAKKAQAI	IVEYREKNGK	FLNVEQL	TEVSGIGKATLE	104	
NTDB id 595	KW2	RS08725	WP 021037687.1	SESTDSPTG...SSVSNQISTTSGPKVNINKADL	TELQK	LTGIGQKKAQD	IIDFRMKN	GDFKSI	EDLQKVS	GFGDKTLE	209
NTDB id 521	SMSK321	RS04960	WP 000443804.1	SQHTGSGAP.....SSTS	SKDKKINLN	NKASLEELKQ	VKGLGGKRAQD	IIDHREANGK	FKSVDELKKVSGIGAKTIE	207	
NTDB id 493	SM12261	RS04300	WP 000387351.1	SQQAGPGTA.....SSTS	KEKKVNL	NKASLEELKQ	VKGLGGKRAQD	IIDHREANGK	FKSVDELKKVSGIGAKTIE	207	
NTDB id 263	KZH43	RS04195	WP 000387330.1	SQQTGLGTA.....SSISKEKKVNL	NKASLEELKQ	VKGLGGKRAQD	IIDHREANGK	FKSVDELKKVSGIGGKTIE	207		
NTDB id 222	SPD	RS04535	WP 000387330.1	SQQTGLGTA.....SSISKEKKVNL	NKASLEELKQ	VKGLGGKRAQD	IIDHREANGK	FKSVDELKKVSGIGGKTIE	207		
NTDB id 188	SPR	RS04295	WP 000387330.1	SQQTGLGTA.....SSISKEKKVNL	NKASLEELKQ	VKGLGGKRAQD	IIDHREANGK	FKSVDELKKVSGIGGKTIE	207		
NTDB id 153	SP	RS04730	WP 000387344.1	SQQTGSGTA.....SSTS	KEKKVNL	NKASLEELKQ	VKGLGGKRAQD	IIDHREANGK	FKSVDELKKVSGIGGKTIE	207	
NTDB id 1069	ABD1	RS03060	WP 043960841.1	LNISKSSHSNSSKISQTKNFNDSTGQ	IHLNQANVNEFQQ	LKGIGEKKAQAI	IVEYRQKNGS	FKNIDEI	KNVKGIGPAIFE	128	
NTDB id 1041	HON27	RS14740	WP 005124713.1	LNITKSSHSNGSKISQTKNFNDSTGQ	IHLNQANVNEFQQ	LKGIGEKKAQAI	IVEYRQKNGS	FKNIDEI	KNVKGIGPAIFE	128	
NTDB id 36	MW	RS08230	WP 001793467.1	EP....QIGVNS..VHVKNGNTNNTK	VNLNTASVSELMS	VPGVGQAKANA	IVEYRNQQGAFQE	IDDLK	KKVKGFGSKTFD	220	
NTDB id 20	SA	RS08005	WP 001802022.1	EP....QIEVNS..VHVKNGNTNNTK	VNLNTASVSELMS	VPGVGQAKANA	IVEYRNQQGAFQE	IDDLK	KKVKGFGSKTFD	220	
NTDB id 105	BSU 25590	NP 390437.1	V....QQGGGGS..V..QSDGGKGALVNINTATLEELQG	ISGVGPSKAEAI	IAYREENGRFQTI	IEDIT	TKVSGIG	EKSFE	196		
NTDB id 1014	ACIAD	RS13855	WP 004924483.1	LNSSASHHSLSK..P..TQQSGTG	VKNINSATAKELQQ	LHGVGEKKALAI	IVEYRQYQK	FASVDDL	QKVKGIGPKFIE	120	
NTDB id 1127	NGFG	RS10560	WP 003703428.1	SCAFSLAAVNINAASQQELEA	LPGIGPAKAKAIAEYRAQNGAFKSVDDL	IKVKGIGPAVLA	105				
NTDB id 1126	NGFG	RS09375	WP 003703428.1	SCAFSLAAVNINAASQQELEA	LPGIGPAKAKAIAEYRAQNGAFKSVDDL	IKVKGIGPAVLA	105				
NTDB id 1125	NGFG	RS07215	WP 003703428.1	SCAFSLAAVNINAASQQELEA	LPGIGPAKAKAIAEYRAQNGAFKSVDDL	IKVKGIGPAVLA	105				
NTDB id 1108	NGFG	RS06525	WP 003703428.1	SCAFSLAAVNINAASQQELEA	LPGIGPAKAKAIAEYRAQNGAFKSVDDL	IKVKGIGPAVLA	105				
NTDB id 152956	AEI04	RS08195	WP 002855574.1	LTAFLFGA	VNINTATL	KLKLS	LNGIGEAKAKAILEYRKE	ANFTS	IDDLK	KKVKGIGDKLFE	70
NTDB id 1242	Cj0011c	YP 002343483.1	LTAFLFGA	VNINTATL	KLKLS	LNGIGEAKAKAILEYRKE	ANFTS	IDDLK	KKVKGIGDKLFE	70	
consensus				***! !** !* * !*! ***!!*!!*! * * !***** *!***! ****							

logo

NTDB id 1159	ABDM36	RS04765	WP 001166105.1	NNAARILL	103	
NTDB id 1150	A1552VC	RS08580	WP 001166105.1	NNAARILL	103	
NTDB id 1395	DSB67	RS04445	WP 010649155.1	KNEDRILL	95	
NTDB id 1362	HI 1008	AAC22669.1	KNRDRIIF	112		
NTDB id 1382	A4U84	RS07375	WP 062924234.1	KNRDCIVIE	113	
NTDB id 595	KW2	RS08725	WP 021037687.1	KLKDEISID	218	
NTDB id 521	SMSK321	RS04960	WP 000443804.1	KLKDYVTVD	216	
NTDB id 493	SM12261	RS04300	WP 000387351.1	KLKDYVTVD	216	
NTDB id 263	KZH43	RS04195	WP 000387330.1	KLKDYVTVD	216	
NTDB id 222	SPD	RS04535	WP 000387330.1	KLKDYVTVD	216	
NTDB id 188	SPR	RS04295	WP 000387330.1	KLKDYVTVD	216	
NTDB id 153	SP	RS04730	WP 000387344.1	KLKDYVTVD	216	
NTDB id 1069	ABD1	RS03060	WP 043960841.1	KNKSRLIAL	136	
NTDB id 1041	HON27	RS14740	WP 005124713.1	KNKSRLIAL	136	
NTDB id 36	MW	RS08230	WP 001793467.1	KLKSYFTI	228	
NTDB id 20	SA	RS08005	WP 001802022.1	KLKSYFTI	228	
NTDB id 105	BSU 25590	NP 390437.1	KIKSSIIVK	205		
NTDB id 1014	ACIAD	RS13855	WP 004924483.1	KNRDRLAI	128	
NTDB id 1127	NGFG	RS10560	WP 003703428.1	KLKDQASVGAPAPKGP	PAKPVLPVAVKK	131
NTDB id 1126	NGFG	RS09375	WP 003703428.1	KLKDQASVGAPAPKGP	PAKPVLPVAVKK	131
NTDB id 1125	NGFG	RS07215	WP 003703428.1	KLKDQASVGAPAPKGP	PAKPVLPVAVKK	131
NTDB id 1108	NGFG	RS06525	WP 003703428.1	KLKDQASVGAPAPKGP	PAKPVLPVAVKK	131
NTDB id 152956	AEI04	RS08195	WP 002855574.1	KIKNDITVE	79	
NTDB id 1242	Cj0011c	YP 002343483.1	KIKNDITIE	79		
consensus				*****		

 non conserved
 similar
 $\geq 50\%$ conserved