

logo

NTDB	id	1029	TT	RS02230	WP	011228203.1	MPVYQYKARDRQGR.L	VEATIEAEDLRTAARLRDRGLFVAETKEPGKG	LQAEVRIPALER	GPGLK	65								
NTDB	id	1402	DSB67	RS12670	WP	010643257.1	.MKT.....IAPQLKNYRWKGINS	SGKK.TSGNVLAMTEIEVRERLDAQH	IKKLKKSISFITK	ISHRVK	KGK	67							
NTDB	id	1169	A1552VC	RS11075	WP	000648511.1	MKAT.....QTLPLKNYRWKGINS	NGKK.VSQMLAISEIEVRDKLKDQHI	QIKKLKKSIVS	LLAR.....LTHR	VKSK	68							
NTDB	id	1113	AAA85695.1	219..1451(	)		.MAKNGGFSLFAKKEKRFIFEGRHS	ASDKLVNGEVS	AFTEEEARKKLAKRGIRPLQITRV	K.....T....SSK	RKITQE	70							
NTDB	id	1112	NGCF	RS09215	WP	003689811.1	.MAKNGGFSLFAKKEKRFIFEGRHS	ASDKLVNGEVS	AFTEEEARKKLAKRGIRPLQITRV	K.....T....SSK	RKITQE	70							
NTDB	id	1252	GC085	RS07730	WP	011213805.1	.MDK.....NSPPLTLTFHYQCIN	KAQK.MEGDIQARSLATAKADLRKQ	GIVTNKVVKKRKP	LFDR.....KNKKI	TQA	67							
NTDB	id	1016	ACIAD	RS01680	WP	004920476.1	MAAK.....KTQVMPVFAYEGVDR	KGIK.LKGELPARNMALAKVTLRKQ	GITRTIREKRKN	IFEG....LLK	KKV	KPL	69						
NTDB	id	1059	ABD1	RS01610	WP	000279215.1	MAVK.....KAQMMPTFAYEGVDR	KGVK.IKGELPAKNMALAKVTLRKQ	GVTVRNIREKRKN	ILEG....LFK	KKV	TTL	69						
NTDB	id	164112	AU380	RS01350	WP	003107299.1	.MAD.....KALKTSVFWEGTDK	KGKT.VKGELSSQNPTLVKAQLRK	QGITPVKVRKKG	SISLLG.....AG	KKIK	PM	66						
NTDB	id	1198	PSJM300	03950	AFN	76868.1	.MAQ.....KAIKNSVFTWGLDR	QGAQ.IKGELSCVSPALVKAQLRKQ	GINPVKVRKKS	SISLFS.....AG	KKIK	PM	66						
consensus							***	*	*****	*	*****	*	*****	!	*****	*****	*	*****	

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[illegible]

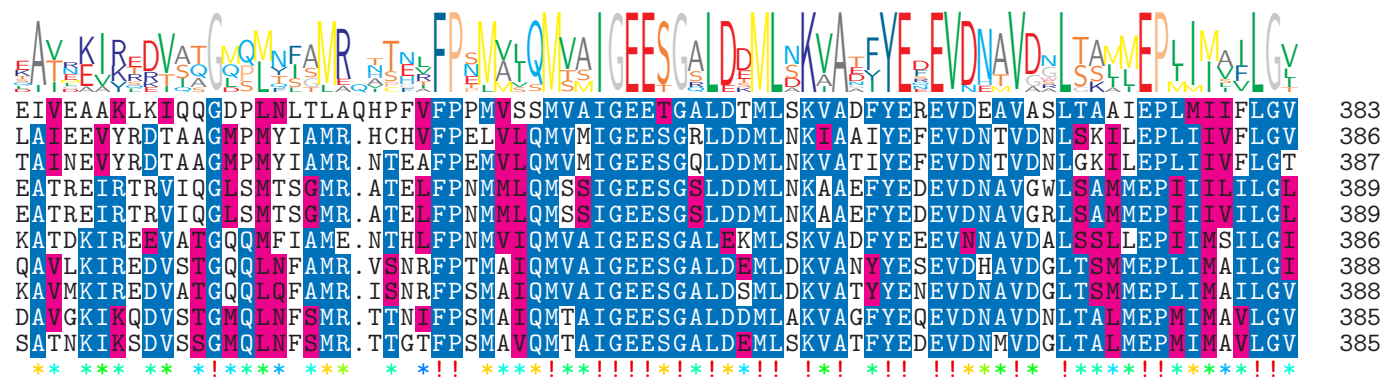
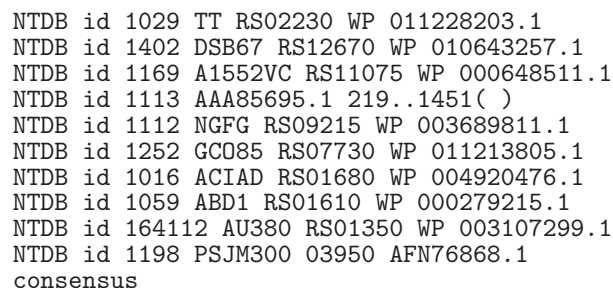
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NTDB id 1029	TT RS02230 WP 011228203.1	DLLDRLRSLFLEKEELRGKIRSASTYPIVVFVAVGVAYFLLTGIVPQFAQLILTDLGSELPLLTRFLIAVSDDLRAATL	224
NTDB id 1402	DSB67 RS12670 WP 010643257.1	SQVFERLATYREKSEQLRAKVIKALITYPAMVVVLVALGVSYLMLTRVIPEFEKMFTGFAGDLPAFTQMVLNLSAWTQNWGP	227
NTDB id 1169	A1552VC RS11075 WP 000648511.1	PEVFERLATYREKSEQLRAKVIKALIYPSMVVLVALGVSYLMVTMVIPEFESMFKGFGAELPWFTQQVLKLSHWVQAYSLS	228
NTDB id 1113	AAA85695.1 219..1451()	ESELLDKLATYKEKTQAIRKKVKLTALTYPVSVIAVAIGLVFVMIMIFVLPAFKEVYANMGGAELPALTQTVMDSDFVSYGW	230
NTDB id 1112	NGFG RS09215 WP 003689811.1	ESELLDKLATYKEKTQAIRKKVKLTALTYPVSVIAVAIGLVFMMIFVLPAFKEVYANMGGAELPLTQTVMDSDFVSYGW	230
NTDB id 1252	GCO85 RS07730 WP 011213805.1	DIMLDKLVATYKEKTEILKKKIKKALTYPJAVMVVALVLTAGLLIYVVPQFESLFKGGFADLPAMTRAVITMSSEFMQAYWY	227
NTDB id 1016	ACIAD RS01680 WP 004920476.1	EIMLDRVAVYKEKSEVLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFKDLFTSFGADLPFTQLVNNMSNWMQDYWF	229
NTDB id 1059	ABD1 RS01610 WP 000279215.1	ETMLDRVAVYKEKSELLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFDLFFASFGADLPFTQMNVNMSKWMDQEYWF	229
NTDB id 164112	AU380 RS01350 WP 003107299.1	ETLDRVATYKEKTEALKAKIKKAMTYPIAVIVVAIVVSAILLIKVVPQFQSVEFGFGAELPAFTMMVINISNVLQEWL	226
NTDB id 1198	PSJM300 03950 AFN76868.1	ETLDRVATYKEKTEALKAKIKKANPYIAVVLVAVIVTAILLIKVVPQFDVFANFGAELPAFTLMVIGLSEALQAWY	226
consensus		* * * * *	

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NTDB id 1029 TT RS02230 WP 011228203.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
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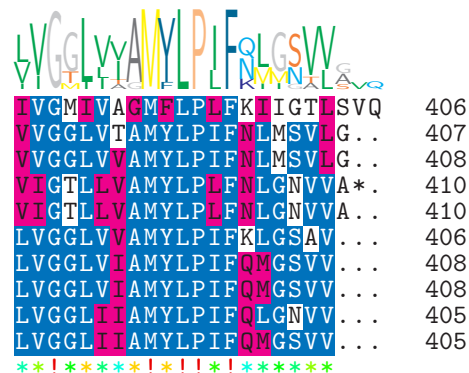
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NTDB id 1029 TT RS02230 WP 011228203.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
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NTDB id 1113 AAA85695.1 219..1451( )
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NTDB id 164112 AU380 RS01350 WP 003107299.1
NTDB id 1198 PSJM300 03950 AFN76868.1
consensus

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- X non conserved
- X similar
- X > 50% conserved