

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGRLVEATIEAEDLRTAARL	LRDRGLFVAETKEPGKGQLQA	EVRIPALERGPGLK	65						
NTDB id 1402	DSB67 RS12670 WP 010643257.1	.MKT.....IAPQLKNYRWKGINSSGKK.TSGNVLAMTEIEVRERLDAQHIIKIKKLKKG	SISFITK....ISHRVKGGK	67							
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKAT.....QTLP LKNYRWKGINSN GKK.VSGQMLAI SEIEVRDKLKDQHIQIKKLKKG	SVSLLLAR.....LTHR VKSK	68							
NTDB id 1113	AAA5695.1 219..450()	.MAKNGGFSLFAKKEKRIFEGRHASADKLNVGEVS	AFTEEEARKKLAKRGIRPLQITRVK....T....SSK RKITQE	70							
NTDB id 1112	NFGF RS09215 WP 003689811.1	.MAKNGGFSLFAKKEKRIFEGRHASADKLNVGEVS	AFTEEEARKKLAKRGIRPLQITRVK....T....SSK RKITQE	70							
NTDB id 1252	GCO85 RS07730 WP 011213805.1	.MDK.....NSPPLLTFHYQGINKAGQK.MEGDIQARSLATAKADLRKQGIVTNKVVK	KRKPLFDR....KNKKITQA	67							
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAK.....KTQVM PVFAYEGVDRKG IK.LKGELPARNMALAKVT LRKQGITIRTIREKRKN I	FEG...LLKKKVKPL	69							
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVK.....KAQM MP TFAYEGVDRKG VIK.KGELPAKNMALAKVT LRKQGVTVRNIREKRKN I	LEG...LFKKKV TTL	69							
NTDB id 95363	AOY09 RS05730 WP 003161763.1	.MAD.....KALKTSVFVWEGTDKKGT K.VKGELSQQNP TLVKAQLRKQGIPV KV RKKGIS	LLG.....AGKKIKPM	66							
NTDB id 1198	PSJM300 03950 AFN76868.1	.MAQ.....KAIKNSVFTWEGLDRQGA K.IKGELSGVSPALVKAQLRKQGIN PQKVRKK SIS	LFS.....AGKKIKPM	66							
consensus		***	*	* * *	*****	*	****	!*****	*****	*	*****

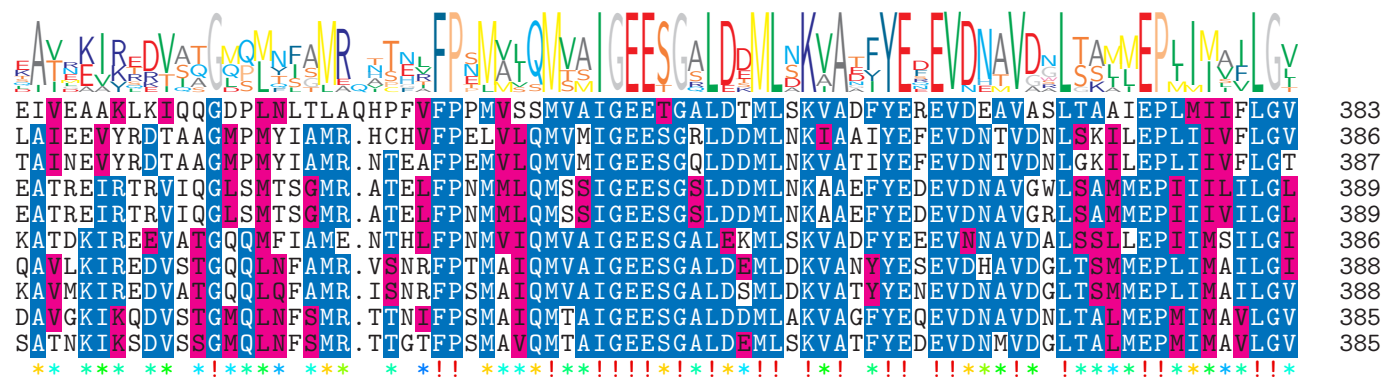
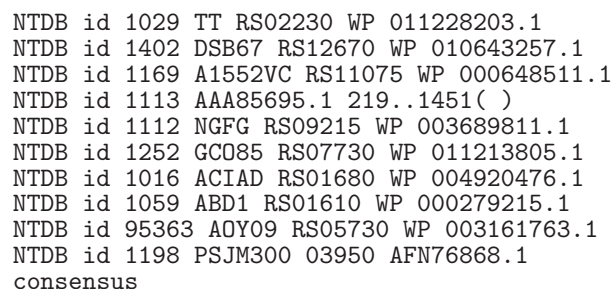
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NTDB id 1402	DSB67 RS12670 WP 010643257.1	DITIFTRQIIATMLMTGVPIVQALKLVSENHKKAEKMSILMSVTRAVEAGTPMSKAMRTASTHFDPLYTDLIATGEQSGNL	147
NTDB id 1169	A1552VC RS11075 WP 000648511.1	DITI L TRQLATMLTGTGPVIVQALKLVGDNHRKAEMKSILAQITKSVEAGTPLSKAMRTASAHFDTLYVDLVETGEMSGNL	148
NTDB id 1113	AAA85695.1 219..1451()	DITVFTRQLSTMINAGLPLMQAFEIVARGHGNSPMTEMLMEIRGQVEQGSSLSRAFSNHPKYFDRFYCNLVAAGETGGVL	150
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NTDB id 1016	ACIAD RS01680 WP 004920476.1	DIAIFTRQLATMMKAGVPLVQAFEIVAEGLENPAMRDVVVLGLKN E VEGGNTFAGALRKYPQYFDNFLC S SLIESGEQSGSL	149
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NTDB id 1198	PSJM300 03950 AFN76868.1	DIALFTRQMATTMMKAGVPLLQSFDIIGEGFDNPNNMRK L VDDLKQE V AAGNSFASSLRKKPKQYFDDL C NLVDSEGEQSGSL	146
consensus		!*****!!**!* *!*****!*****!***** ***** !*****!*****!*****!*****!	

[illegible]

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consensus

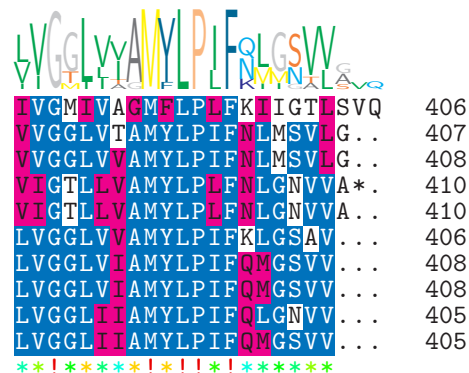
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NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 95363 AOY09 RS05730 WP 003161763.1
NTDB id 1198 PSJM300 03950 AFN76868.1
consensus

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