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NTDB id 1029 TT RS02230 WP 011228203.1	MAK	65
NTDB id 1402 DSB67 RS12670 WP 010643257.1	MPVYQYKARDRQGRLLVEATTEAEDLRTAARLLRDRGLFVAEIKKEPGKGLQAEVRIPALERGPGGLK	67
NTDB id 1169 A1552VC RS11075 WP 000648511.1	.MKT.....IAPQLKNYRWKGINSSGKK.TSGNVLAMTEIEVRERLDAQHIKIKKLLKKSISFITK.....ISHRVKGK	68
NTDB id 1113 AAA85695.1 219..1451()	MKAT.....QTLPLKNYRWKGINSSGKK.VSQMLAISEIEVRDKLKDQHIQIKKLLKKSVSLLAR.....LTHRVKSK	70
NTDB id 1112 NGFG RS09215 WP 003689811.1	.MAKNGGFSLFAKKEKRFFIEGRHSASDKLVNGEVSATETEEEARKKLAKRGIRPLQITRVK.....TS....SKRKITQE	70
NTDB id 1252 GC085 RS07730 WP 011213805.1	.MAKNGGFSLFAKKEKRFFIEGRHSASDKLVNGEVSATETEEEARKKLAKRGIRPLQITRVK.....TS....SKRKITQE	67
NTDB id 1016 ACIAD RS01680 WP 004920476.1	.MDK.....NSPPLILT FHYQGINKAGQK.MEGDIQARSLAIADLRKQGI VTNKVVKRKPLFDR.....KNKKITQA	69
NTDB id 1059 ABD1 RS01610 WP 000279215.1	MAAK.....KTQVMPVFAYEGVDRKGKIK.LKGELPARNMALAKVTLRKQGITRTIREKRKNIFEGL....LKKKVKPL	69
NTDB id 415462 GVI47 RS23390 WP 160288285.1	MAVK.....KAQMMPTFAYEGVDRKGKIK.LKGELPAKNMALAKVTLRKQGVTVRNIREKRKNILEGL....FKKKVTTL	66
NTDB id 1198 PSJM300 03950 AFN76868.1	.MAE.....KALKTSVFTFEGTDRKGGK.IKGELSGTNTALVKAQLRKQGINPTKVRKKGMSLLS.....KGKKIKPM	66
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NTDB id 1402 DSB67 RS12670 WP 010643257.1	DLAIFSRQLATMLGAGLTLLQALATILERQTEENRKFREILKQVRTDVEGGMAFSEALSCHKH.IFSRLYVNLVRAGETSGGL	147
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NTDB id 1112 NGFG RS09215 WP 003689811.1	DITVFTRQLSTMINAGLPLMQAFEIVARGHGNPSMTEMLMEIRGQVEQGSSLSRAFSNHPKYFDRFYCNLVAAGETGGVL	150
NTDB id 1252 GC085 RS07730 WP 011213805.1	DITVFTRQLSTMIKAGLPLMQAFEIVARGHGNPSMTEMLMEIRGQVEQGSSLSRAFSNHPKYFDRFYCNLVAAGETGGVL	147
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NTDB id 415462 GVI47 RS23390 WP 160288285.1	DITIFTRQLATMMKAGVPLVQAFEIVAEGLENPAMREVLGLKGEVEGGSTFASALRKYPQHFDNLFCSLIESGEQSGAL	146
NTDB id 1198 PSJM300 03950 AFN76868.1	DIALFTRQMATMMQSGVPLQLSFDIIGEGFDNPNMKKLVVEIKQEVAAGNSLANSLRKKKPLYFDDLYCNLVDAGEQSGAL	146
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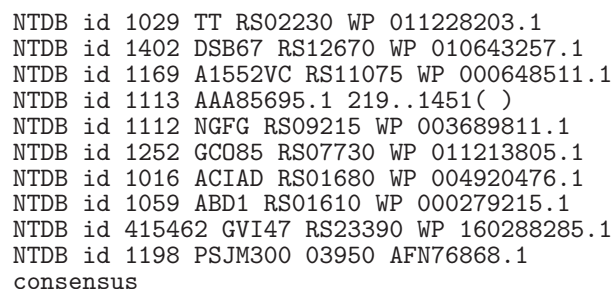
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NTDB id 415462 GVI47 RS23390 WP 160288285.1	ETMLDRVAIYKEKSELLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFFQDLFASFGADLPFTQMVVNMSKWMQYWF	226
NTDB id 1198 PSJM300 03950 AFN76868.1	ETLLDRVATYKEKTESLKGKIKKAMTYPIAVVVVAIVVSAILLIKVVPQFQSVFANFGAELPAFTQMVINLSEILQKWWL	226
consensus	ETLLDRVATYKEKTEALKAKIKKAMNYPIAVVVLVAVIVTAILLIKVVPQFQDVFANFGAELPAFTLMVIGLSEALQAWWY	226

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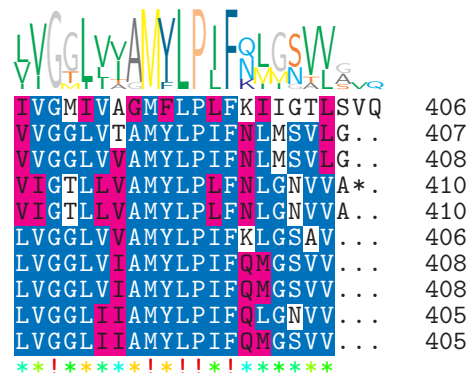
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NTDB id 415462 GVI47 RS23390 WP 160288285.1
NTDB id 1198 PSJM300 03950 AFN76868.1
consensus

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