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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.TSGNVLAMTEIEVRRERLDAQHIKIKKLLKKGSIKFITK.....ISHRVKGK	68
NTDB id 1113 AAA85695.1 219..1451()	MKAT.....QTLPLKNYRWKGINSSGKK.VSQQLAISEIEVRDKLKDQHIQIKKLLKKGSIKLAR.....LTHRVKSK	70
NTDB id 1112 NGFG RS09215 WP 003689811.1	.MAKNGGFSLF AKKEKR FIFEGRHSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRVK.....T.....SSKRKITQE	70
NTDB id 1252 GC085 RS07730 WP 011213805.1	.MAKNGGFSLF AKKEKR FIFEGRHSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRVK.....T.....SSKRKITQE	67
NTDB id 1016 ACIAD RS01680 WP 004920476.1	.MDK.....NSPPLILT FHYQG INKAGQK.MEGDIQARS LA IAKADLRKQGI V TNKKVVKRKPFLFDR.....KNKKKITQA	69
NTDB id 1059 ABD1 RS01610 WP 000279215.1	MAAK.....KTQVMPV FAYEG VDRKG IK .LKGELPARNMALAKVTLRKKGIT IRT IREKRKNIFEG....LLKKKV K P	69
NTDB id 124901 HW07 RS24160 WP 003107299.1	MAVK.....KAQMMPT FAYEG VDRKG VK .IKGELPAKNMALAKVTLRKKGV T VRNIREKRKNILEG....L F KKKV T TL	66
NTDB id 1198 PSJM300 03950 AFN76868.1	.MAD.....KALKTSV FV WEGTDKKG TK .VKGELSSQNPTLVKAQLRKKGIT P VKVRKKGISLLG.....AGKKIKPM	66
consensus	.MAQ.....KAIKNSV FT WEG LD RQGA K .IKGELSGVSPALVKAQLRKKGINPQKVRKKSSISLFS.....AGKKIKPM	66

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NTDB id 1402 DSB67 RS12670 WP 010643257.1	DLAIFSRQLATMLGAGLTLLQALAITLERQTE NR KFREILKQV RT DVEGGMAFSEALS SKH .KIFSRLYVNLVRAGETSGGL	147
NTDB id 1169 A1552VC RS11075 WP 000648511.1	DITIFTRQ IA TMLMTGVPIVQALKLVSENHKKAE MS ILMSVTRAVEAGTPMSKAMRTASTHFDPLYTDLIATGEQSGNL	148
NTDB id 1113 AAA85695.1 219..1451()	DITILTRQLATMLTTGVPIVQALKLVGDNHRKAEMKSILAQITKSVEAGTPLSKAMRTASAHFDTLYVDLIVETGEMSGNL	150
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NTDB id 1252 GC085 RS07730 WP 011213805.1	DITVFTRQLSTM IK AGLPLMQAFEIVARGHG NPS MT EM LMEIRGQVEQGSSLSRAFSNHPKYFDRFYCNLVAAGETGGVL	147
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NTDB id 124901 HW07 RS24160 WP 003107299.1	DITIFTRQLATMMKAGVPLVQAFEIVAEGLENPAMREVVLGIKGEVEGGSTFASALRKYPQHFDNLFCSLIVESGEQSGAL	146
NTDB id 1198 PSJM300 03950 AFN76868.1	DIALFTRQ MS TMMAGVPL LQ SFDI I SEGFDNPNMRKLV EE IKQEVAGGNSLANSLRKKPKQYFDSLYCNLVDAGEQSGAL	146
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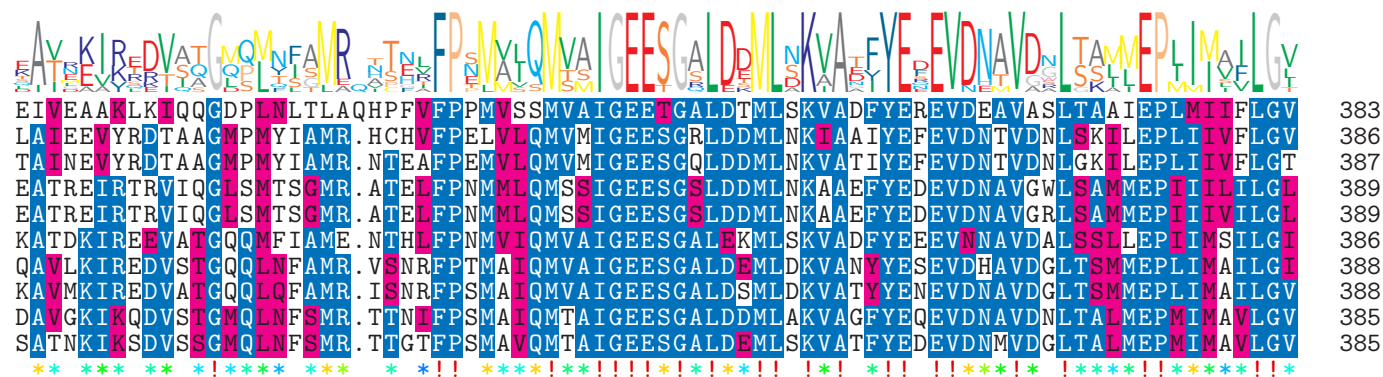
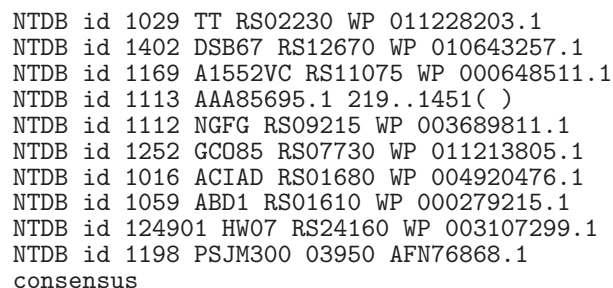
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NTDB id 1029 TT RS02230 WP 011228203.1	EILLDRYAYK EK IELR K IKK AM YPV V VAIVIGTYLLI W PPF E ELFAGELPAETQV IN MS Q YWL	224
NTDB id 1402 DSB67 RS12670 WP 010643257.1	DLILDRLASFLEKELELRGKIRSAMTYPVIVFVFAVGVA Y LLTGIVPQFAQILTDLGSSELPLLTRFLIAVSDLLRAATL	227
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NTDB id 1059 ABD1 RS01610 WP 000279215.1	EIMLDRVAIVYKEKSEVLKQKIKKAMKYPATVIVVAIVVTI LM VKVVPVFKDLFTSFGADLP FT QLVVNMSNWMQDYWF	229
NTDB id 124901 HW07 RS24160 WP 003107299.1	ETMLDRVAIVYKEKSELLKQKIKKAMKYPATVIVVAIVVTI LM VKVVPV FQ DLFASFGADLP FT QMVVNMSKWMQ EY WF	226
NTDB id 1198 PSJM300 03950 AFN76868.1	ETLLDRVATYKEKTEALKAKIKKAMTYPIAVIVVAIV VS AILLIKVVPQFQSVFEGFAGELPAFTMMVINISNVLQEWL	226
consensus	ETLLDRVATYKEKTEALKAKIKKAMNYPIAVVVLVAVIVTAILLIKVVPQFQDV F ANFAGELPAFTLMVIGLSEALQAWWY	226

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consensus

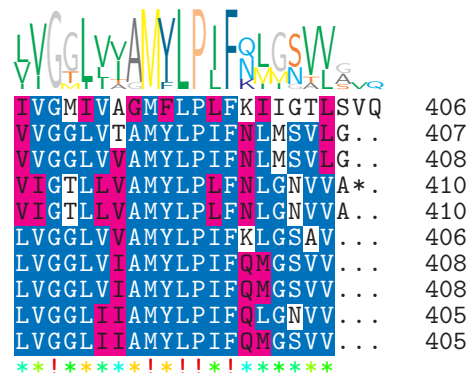
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consensus

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