

logo

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.TSGNVLAMTEIEVRRERLDAQHIKIKKKLKKGSISFITK.....ISHRVKKGK	68
NTDB id 1113 AAA85695.1 219..1451()	MKAT.....QTLPLKNYRWKGINSSGKK.VSQQLAISIEIEVRDKLKDQHIQIKKKLKKGSVSLAR.....LTHRVKSK	70
NTDB id 1112 NGFG RS09215 WP 003689811.1	.MAKNGGFSLF AKKEKR FIFEGRHSASDKLVNGEVSATETEEEARKKLAKRGIRPLQITRVK.....T.....SSKRKITQE	70
NTDB id 1252 GC085 RS07730 WP 011213805.1	.MAKNGGFSLF AKKEKR FIFEGRHSASDKLVNGEVSATETEEEARKKLAKRGIRPLQITRVK.....T.....SSKRKITQE	67
NTDB id 1016 ACIAD RS01680 WP 004920476.1	.MDK.....NSPPLILT FHYQG INKAGQK.MEGDIQARSLAI AKADLR KQGIVTNKVVKRKPFLFDR.....KNKKKITQA	69
NTDB id 1059 ABD1 RS01610 WP 000279215.1	MAAK.....KTQVMPV FAYEGV DRKGIK.LKGELPARNMALAKVTLRKKGITRTIREKRKNIFEG....LLKKKV K PPL	69
NTDB id 399056 GIY05 RS25455 WP 003107299.1	MAVK.....KAQMMPT FAYEGV DRKGVK.IKGELPAKNMALAKVTLRKKGVTVRNIREKRKNILEG....L FKKKV TTL	66
NTDB id 1198 PSJM300 03950 AFN76868.1	.MAD.....KALKTSV FVWEGT DKKGT.VKGELSSQNPTLVKAQLRKKGITPVKVRKKGISLLG.....AGKKIKPM	66
consensus	.MAQ.....KAIKNSVFTWGLDRQGA K .IKGELSGVSPALVKAQLRKKGINPQKVRKKSSISLFS.....AGKKIKPM	66

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NTDB id 1029 TT RS02230 WP 011228203.1	DITIFTRQLATMAGVPLVQAFIVAEGLNPNMRRELLMEIKVEGSSSALRKHPQYFDLYCNLVAGEQSGI	144
NTDB id 1402 DSB67 RS12670 WP 010643257.1	DLAIFSRQLATMLGAGLTLLQALATILERQTE ENRKFREIL KQVRTDVEGGMAFSEALS SKH .KIFSRLYVNLVRAGETSGGL	147
NTDB id 1169 A1552VC RS11075 WP 000648511.1	DITIFTRQIATMLMTGVPIVQALKLVSENHKKAE MSILMS VTRAVEAGTPMSKAMRTASTHFDPLYTD LIAT GEQSGNL	148
NTDB id 1113 AAA85695.1 219..1451()	DITILTRQLATMLTTGVPIVQALKLVGDNHRKAEM KSILIAQ ITKSVEAGTPLSKAMRTASAHFDTLYVD LVET GEMSGNL	150
NTDB id 1112 NGFG RS09215 WP 003689811.1	DITVFTRQLSTMINAGLPLMQAFEIVARGHG NPSMTEML MEIRGQVEQGSSLSRAFSNHPKYFDRFYCNLVAAGETGGVL	150
NTDB id 1252 GC085 RS07730 WP 011213805.1	DITVFTRQLSTM IKAGL PLMQAFEIVARGHG NPSMTEML MEIRGQVEQGSSLSRAFSNHPKYFDRFYCNLVAAGETGGVL	147
NTDB id 1016 ACIAD RS01680 WP 004920476.1	DITVFSRQLATMIESGIPLVQAFDIVAKGQSN KRLKDL IETIKHDIETGLTLAESLIKHPLYFNE LF FCNLVDAGEKSGSL	149
NTDB id 1059 ABD1 RS01610 WP 000279215.1	DIAIFTRQLATMMKAGVPLVQAFEIVAEGLE NPAMRD VVLGLKNEVEGGNTFAGALRKYPQYFDN LF CSLIESGEQSGSL	149
NTDB id 399056 GIY05 RS25455 WP 003107299.1	DITIFTRQLATMMKAGVPLVQAFEIVAEGLE NPAMRE VVLGIKGEVEGGSTFASALRKYPQHFDN LF CSLIVESGEQSGAL	146
NTDB id 1198 PSJM300 03950 AFN76868.1	DIALFTRQ MST MMAGVPL LQSF DIISEGFDNPNMRKLV EEIKQ EVAGGNSLANSLRKKPKQYFDSLYCNLV DAGE QSGAL	146
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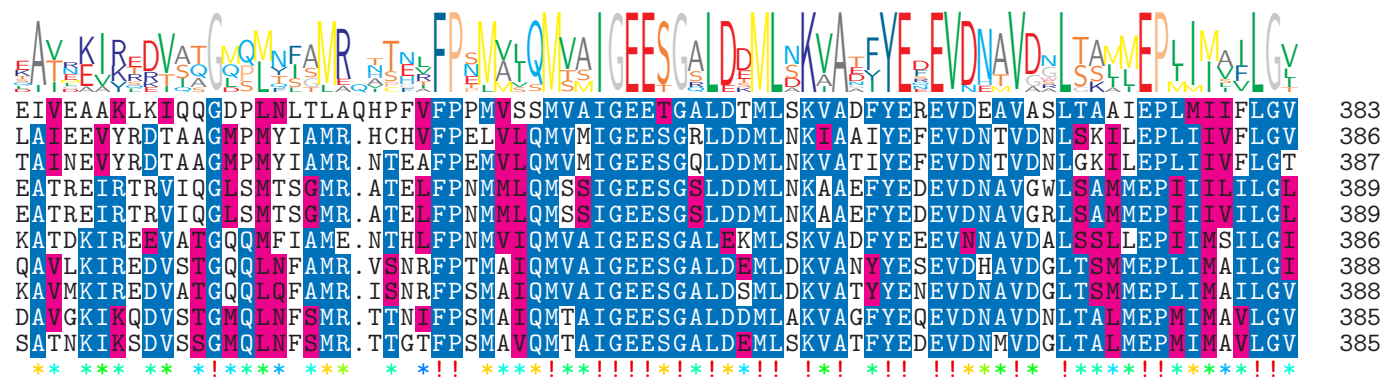
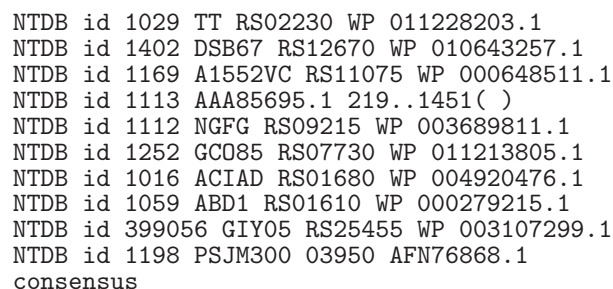
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NTDB id 1029 TT RS02230 WP 011228203.1	EILDRYAYKEKTELRKIKKAMTPYVIVVAVAGVYLLTGWIPQFAQILTDLGSELPLLTRFLIAVSDLLRAATL	224
NTDB id 1402 DSB67 RS12670 WP 010643257.1	SQVFERLATYREKSEQLR AKVIK ALIYPAMVVLVALGVSYLMLTRV IP EF EKMFT GFGADLP FTQ MVLNLSA WTQ NWGP	227
NTDB id 1169 A1552VC RS11075 WP 000648511.1	PEVFERLATYREKSEQLR AKVIK ALIYP SMVVL VALGVSYLMLTMV IP EFESMF KGFGA ELPWFTQ QVL KL SHV QAYSL	228
NTDB id 1113 AAA85695.1 219..1451()	ESLLDKLAIYKEKTQAIRKKVKTALTYPVSVIAVAIGLVFVMMIFVLP AFKEV YANMGAELPALTQTVM DMSD FFVSYGW	230
NTDB id 1112 NGFG RS09215 WP 003689811.1	ESLLDKLAIYKEKTQAIRKKVKTALTYPVSVIAVAIGLVFVMMIFVLP AFKEV YANMGAELPALTQTVM DMSD FFVSYGW	230
NTDB id 1252 GC085 RS07730 WP 011213805.1	DIMLDKVATYKEKIEIKKKIKKALTYP IAVMV VALLVTAGLLIYVVPQFESL FKGFG ADLPAMTRAVITMSEFMQAYWY	227
NTDB id 1016 ACIAD RS01680 WP 004920476.1	EIMLDRVAIVYKEKSEVLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFKDLFTSFGADLP FTQ LVN MSN WMQDYWF	229
NTDB id 1059 ABD1 RS01610 WP 000279215.1	ETMLDRVAIVYKEKSELLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPV FQDL FASFGADLP FTQ LVN MSN WMQDYWF	229
NTDB id 399056 GIY05 RS25455 WP 003107299.1	ETLLDRVATYKEKTEALKAKIKKAMTYPIAVIVVAIVSAILLIKVVPQFQSVFEGFGAELP FTM VINISNVLQEWL	226
NTDB id 1198 PSJM300 03950 AFN76868.1	ETLLDRVATYKEKTEALKAKIKKAMNYPIAVVVLVAVIVTAILLIKVVPQFQDV FAN FGAELP FTL MVIGLSEALQAWWY	226
consensus	ETLLDRVATYKEKTEALKAKIKKAMNYPIAVVVLVAVIVTAILLIKVVPQFQDV FAN FGAELP FTL MVIGLSEALQAWWY	226

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NTDB id 1112 NGCF RS09215 WP 003689811.1
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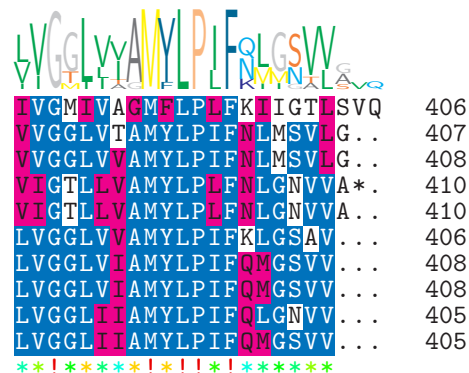
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consensus

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