

logo

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGR.L.VEATIEAEDLRTAARLRDRGLFVAETKEPGKGLQAEVRIPALER	60
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKATQTLPLKKNYRWKGINSGKK.VSGQMLAISEIEVRDKLKDQHIQIKKLKKGVSLSLA....RLTH	63
NTDB id 1402	DSB67 RS12670 WP 010643257.1	MKTIAAPQLKKNYRWKGINSSGKK.TSGNVLAMTEIEVRERLDAQHIKIKKLKKGSSISFIT....KISH	62
NTDB id 1112	NGFG RS09215 WP 003689811.1	MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVSFAFTEEEARKKLAKRGIRPLQITRVK....T....SSKR	65
NTDB id 1113	AAA85695.1 219..1451()	MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVSFAFTEEEARKKLAKRGIRPLQITRVK....T....SSKR	65
NTDB id 1198	PSJM300 03950 AFIN76868.1	MAQKAIAKNSVTFTEGLDRQGAQ.IKGELSGVSPALVKAQLRKQGINPQKVRKKSISLFS.....AGK	61
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKKTQVMPVFAYEGVDRKGIK.LKGELPARNMALAKVTLRKQGITIRTIREKRKNIFEG...LLKK	64
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVKKAQMMPTFAYEGVDRKGIVK.IKGELPAKNMALAKVTLRKQGVTVRNIREKRKNILEG...LFFK	64
NTDB id 322125	EBA04 RS17250 WP 024743356.1	MSVARSAIKKQPVDRNTMSLQTFVWEGADKRGVQ.MKGEQQARNANMLRAELRRQGIIVPSMVKKQKPKPLLG....AAGK	74
NTDB id 1252	GC085 RS07730 WP 011213805.1	MDKNSPPLLTTFHYQGIKNAQKQ.MEGDIQARSLAIKADLRKQGIIVTNKVVKKRKPLFD....RKNK	62
consensus		* * * * *	

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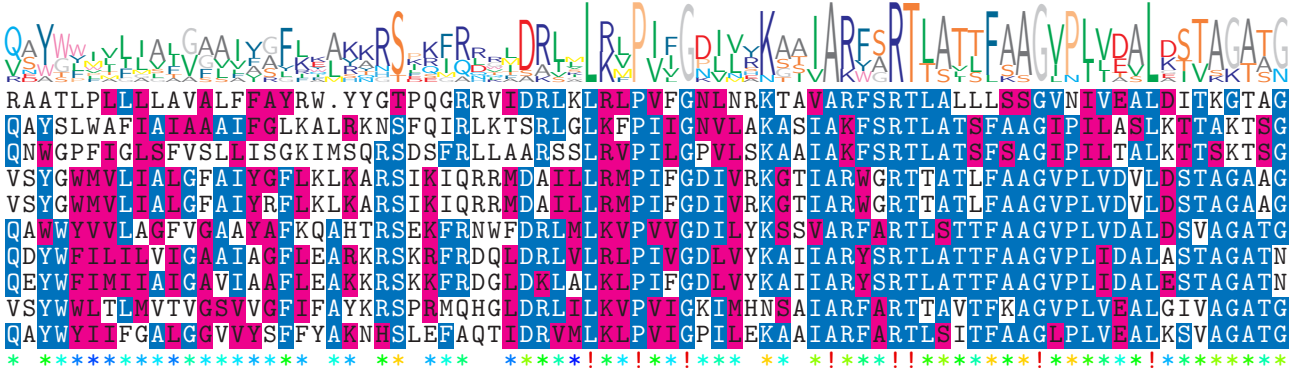
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NTDB id 1029	TT RS02230 WP 011228203.1	TSGGLDLRLDRLEAFLEKELELRGKIRSAMTYPVIVFVFAVGVAFFLLTGIVPQFAQILTDLGSLEPLLLTRFLIAVSDLL	219
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MSGNLPVFERLATYREKSEQLRAKVIKALIYPSMVVLVALGVSYLMLTMVTFEFESMFKGFGAELPWFTQQVLKLSHWV	223
NTDB id 1402	DSB67 RS12670 WP 010643257.1	QSGNLSQVFERLATYREKSEQLRAKVIKALIYPAMVVLVALGVSYLMLTRVIFPEFEKMTFGFGADLPFTQMVLNLSAWT	222
NTDB id 1112	NGFG RS09215 WP 003689811.1	TGCVLESLDLKLATYKEKTQAIRKKVKKTALTYPVSVIAVAIGLVFVMMIFVLPAPFKEVYANMGAELPPLTQTVMDSDF	225
NTDB id 1113	AAA85695.1 219..1451()	TGCVLESLDLKLATYKEKTQAIRKKVKKTALTYPVSVIAVAIGLVFVMMIFVLPAPFKEVYANMGAELPPLTQTVMDSDF	225
NTDB id 1198	PSJM300 03950 AFN76868.1	QSGSLETLDRVATYKEKTEALKAKIKKAMNYPITAVVLVAVIVTAILLIKVVPFQDVANFGAELPAFTLMVIGLSEAL	221
NTDB id 1016	ACIAD RS01680 WP 004920476.1	QSGSLEIMLDRVAVYKEKSEVLKQKIKKAMKYPATVIVVAIVVTIILMVKVPVPFKDLFTSFGADLPFTQLVNNMSNWM	224
NTDB id 1059	ABD1 RS01610 WP 000279215.1	QSGALETMLDRVAVYKEKSELLKQKIKKAMKYPATVIVVAIVVTIILMVKVPVPFQDLFASFGADLPFTQMVVNMSKWM	224
NTDB id 322125	EBA04 RS17250 WP 024743356.1	GAGVLETVLETVATYKENIEALKGKIKKAMFPATVVAIVAIIVSAILLIFVVPQFEQVFKSFGAELPAFTQLLVNASRFM	234
NTDB id 1252	GC085 RS07730 WP 011213805.1	KSGSLEIMLDRVATYKEKTEILKKIKKALTYPIAVMVVALLTAGLLIYVVPQFESLFKGFGADLPAMTRAVITMSEFM	222
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NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 322125 EBA04 RS17250 WP 024743356.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
consensus



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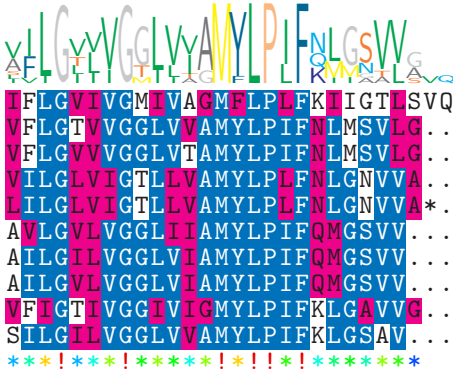


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consensus



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☒ non conserved
☒ similar
☒ ≥ 50% conserved