

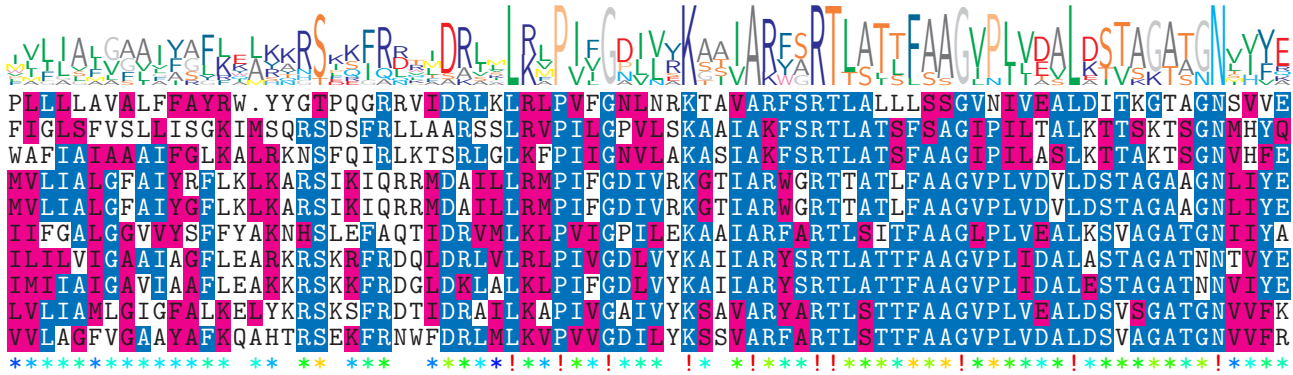
NTDB id	1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGRL.VEATIEAEDLRTAARLLRDRGLFVAETKEPGKGQLAEVRIPALERGPGLK	65
NTDB id	1402	DSB67 RS12670 WP 010643257.1	.MKT.....IAPQLKNYRWKGINSSGKK.TSGNVLAMTEIEVRERLDAQHIKIKKLKKSISFITK....ISHRVKKGK	67
NTDB id	1169	A1552VC RS11075 WP 000648511.1	MKAT.....QTLP LKNYRWKGINSNCKK.VSQMLAIS EIEVRDKLKDQHIIQIKKLKKSVS LLAR.....LTHR VKSK	68
NTDB id	1113	AAA85695.1 219..1451()	.MAKNGGFSLFAKKEKR FIFEGRHASADKL VNGEVSAFT EEEARKKLAKRGIRPLQITRVK.....T...SSK RKITQE	70
NTDB id	1112	NHFG RS09215 WP 003689811.1	.MAKNGGFSLFAKKEKR FIFEGRHASADKL VNGEVSAFT EEEARKKLAKRGIRPLQITRVK.....T...SSK RKITQE	70
NTDB id	1252	GCO85 RS07730 WP 011213805.1	.MDK.....NSPPLLTFHYQGINKAGQK.MEGDIQARSLATAKADLRKGGIVTNKVVKRRKP LFDR.....KNKKITQA	67
NTDB id	1016	ACIAD RS01680 WP 004920476.1	MAAK.....KTQVMPVFAYEGVDRKGIK.LKGELPARNMALAKVT LRKQGITIRTIREKRKN IFEG....LLK KKV KPL	69
NTDB id	1059	ABD1 RS01610 WP 000279215.1	MAVK.....KAQMPTFAYEGVDRKGVK.IKGELPAKNMALAKVT LRKQGVTVRNIREKRKN ILEG....LFK KKV TTL	69
NTDB id	1069248	ACFELF RS24750 WP 400523428.1	.MAD.....KALKTSVFIWEGTDKKGSK.VKGELTGQNPM LVKAQLRKGGINPLKVRKKGIT LFG.....AGKKIKPM	66
NTDB id	1198	PSJM300 O3950 AFN76868.1	.MAQ.....KAIKNSVFTWEGLDRQGA.KIGELSGVSPALVKAQLRKGGINPKVRKKGIS LFS.....AGKKIKPM	66
consensus			*** * * * ** * * * * ! * * * * *	****

[illegible]

NTDB id	1029	TT RS02230 WP 011228203.1	DILIDRLASFLKELELRGKIRSAMTYPVIVFVFAVGVAIFLLTGIVPQFAQILTDLGSELPLLTRFLIAVSDDLRAATL	224
NTDB id	1402	DSB67 RS12670 WP 010643257.1	SQVFERLATYREKSEQLRAKVIKALIYPAMVVVLVALGVSYLMLTRVIPEFEKMFTGFADLP AFTQMVLNLSAWTQNWGP	227
NTDB id	1169	A1552VC RS11075 WP 000648511.1	PEVFERLATYREKSEQLRAKVIKALIYPSMVVLVALGVSYLMITMVIPEFESMFKGFGAELPWFTQQVLKLSHVQAYSL	228
NTDB id	1113	AAA85695.1 219..1451()	ESLLDKLATYKEKTAIRKKVKKTALTYPVSVIAVAIGLVFVMIMIFVLP AFKEVYANMGGAELPA LQTVTMDMSDFFSYGW	230
NTDB id	1112	NGFG RS09215 WP 003689811.1	ESLLDKLATYKEKTAIRKKVKKTALTYPVSVIAVAIGLVFVMIMIFVLP AFKEVYANMGGAELP LQTVTMDMSDFFSYSGW	230
NTDB id	1252	GCO85 RS07730 WP 011213805.1	DIMLDKVATYKEKIEI IKKKIKKALTYP IAVMMVALLVTAGLLIYVVPQFESLFKGFADLPAMTRAVITMSEFMQA YWY	227
NTDB id	1016	ACIAD RS01680 WP 004920476.1	EIMLDRVAIVYKEKSEVLKQKIKKAMKYPATVIVVAIVVTIILMVKVVVPFKDLFTSFGADLP AFTQLVNNMSNWMDYWF	229
NTDB id	1059	ABD1 RS01610 WP 000279215.1	ETMLDRVAIVYKEKSELLKQKIKKAMKYPATVIVVAIVVTIILMVKVVVPFQDLFASF GADLP AFTQM VVNMSKW MQEYWF	229
NTDB id	1069248	ACFELF RS24750 WP 400523428.1	ETLLARVATYKEKTEAL KAKIKKAMTYPIAVVIVALVVSAILLIKVVVPQFQS VFANFGAELPAFTQM VI GLSETLQEWWL	226
NTDB id	1198	PSJM300 03950 AFN76868.1	ETLLDRVATYKEKTEAL KAKIKKAMNYPIAVVVLVAVIVTAILLIKVVVPQFDVFANFGAELPAFTLMVIGLSEALQAWWY	226
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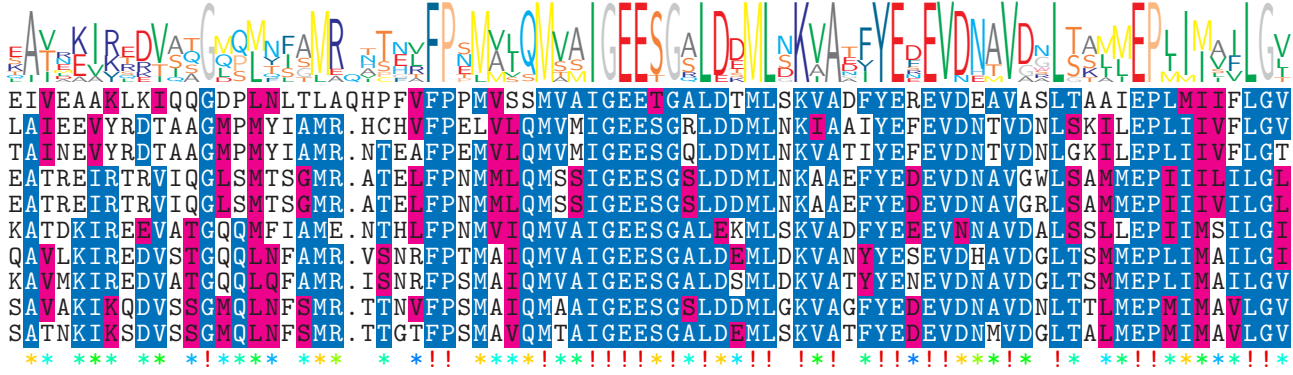
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NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 1069248 ACFELF RS24750 WP 400523428.1
NTDB id 1198 PSJM300 03950 AFN76868.1
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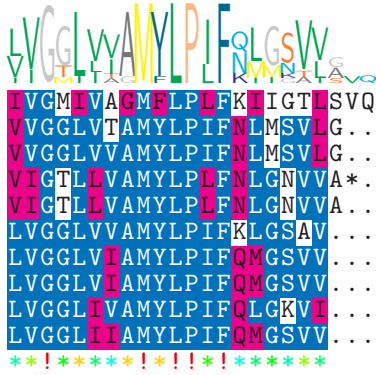


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NTDB id 1198 PSJM300 03950 AFN76868.1
consensus



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☒ non conserved
☒ similar
☒ ≥ 50% conserved