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NTDB id 1402 DSB67 RS12670 WP 010643257.1	MAKNGGFSFLFAKKEKRFIFEGRHSAADKLNVGEVSAFTEEEARKKLAKRGIRPLQITRVK....TSSKRKITQEDITVF	72
NTDB id 1169 A1552VC RS11075 WP 000648511.1	MAKNGGFSFLFAKKEKRFIFEGRHSAADKLNVGEVSAFTEEEARKKLAKRGIRPLQITRVK....TSSKRKITQEDITVF	73
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NTDB id 1112 NGFG RS09215 WP 003689811.1	MAKNGGFSFLFAKKEKRFIFEGRHSAADKLNVGEVSAFTEEEARKKLAKRGIRPLQITRVK....TSSKRKITQEDITVF	75
NTDB id 1252 GC085 RS07730 WP 011213805.1	MAKNGGFSFLFAKKEKRFIFEGRHSAADKLNVGEVSAFTEEEARKKLAKRGIRPLQITRVK....TSSKRKITQEDITVF	72
NTDB id 1016 ACIAD RS01680 WP 004920476.1	MAKNGGFSFLFAKKEKRFIFEGRHSAADKLNVGEVSAFTEEEARKKLAKRGIRPLQITRVK....TSSKRKITQEDITVF	74
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NTDB id 1198 PSJM300 03950 AFN76868.1	MAKNGGFSFLFAKKEKRFIFEGRHSAADKLNVGEVSAFTEEEARKKLAKRGIRPLQITRVK....TSSKRKITQEDITVF	71
consensus	MAKNGGFSFLFAKKEKRFIFEGRHSAADKLNVGEVSAFTEEEARKKLAKRGIRPLQITRVK....TSSKRKITQEDITVF	

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NTDB id 1113 AAA85695.1 219..1451()	TRQLATMIMTGVPIVQALKLVSENHKKAEKMSILMSVTRAVEAGTPMSKAMRTASTHFDPLYTDLIATGEQSGNLSQVFE	155
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NTDB id 1252 GC085 RS07730 WP 011213805.1	TRQLATMIMTGVPIVQALKLVSENHKKAEKMSILMSVTRAVEAGTPMSKAMRTASTHFDPLYTDLIATGEQSGNLSQVFE	152
NTDB id 1016 ACIAD RS01680 WP 004920476.1	TRQLATMIMTGVPIVQALKLVSENHKKAEKMSILMSVTRAVEAGTPMSKAMRTASTHFDPLYTDLIATGEQSGNLSQVFE	154
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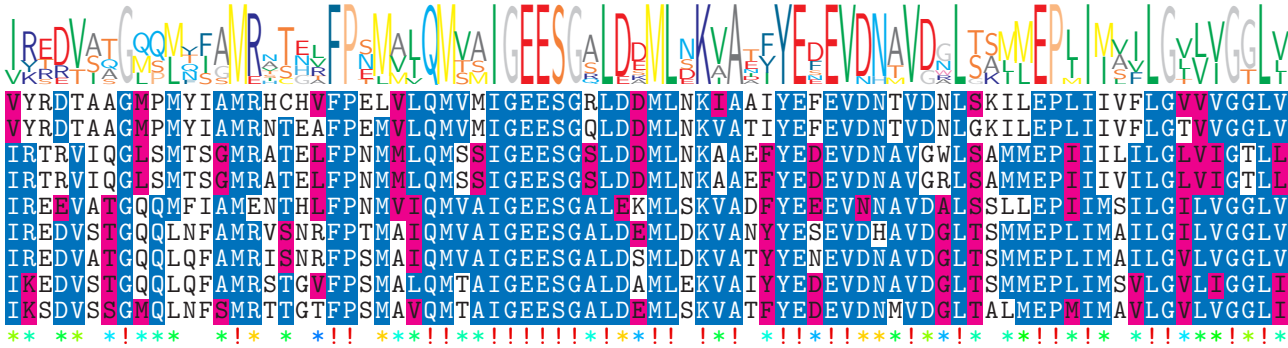
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NTDB id 1402 DSB67 RS12670 WP 010643257.1	FVSLIISGKIMSQRSDSFRLLAARSSLRVPILGPVLSKAAIAKFSRTLATSFSAAGIPILTALKTTSKTSGNMHYQLAIEE	312
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NTDB id 1059 ABD1 RS01610 WP 000279215.1	FVSLIISGKIMSQRSDSFRLLAARSSLRVPILGPVLSKAAIAKFSRTLATSFSAAGIPILTALKTTSKTSGNMHYQLAIEE	314
NTDB id 26146 ABO RS03165 WP 035460653.1	FVSLIISGKIMSQRSDSFRLLAARSSLRVPILGPVLSKAAIAKFSRTLATSFSAAGIPILTALKTTSKTSGNMHYQLAIEE	313
NTDB id 1198 PSJM300 03950 AFN76868.1	FVSLIISGKIMSQRSDSFRLLAARSSLRVPILGPVLSKAAIAKFSRTLATSFSAAGIPILTALKTTSKTSGNMHYQLAIEE	311
consensus	FVSLIISGKIMSQRSDSFRLLAARSSLRVPILGPVLSKAAIAKFSRTLATSFSAAGIPILTALKTTSKTSGNMHYQLAIEE	

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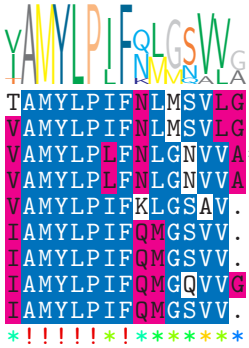


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NTDB id 1198 PSJM300 03950 AFN76868.1
consensus



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☒ non conserved
☒ similar
☒ ≥50% conserved