

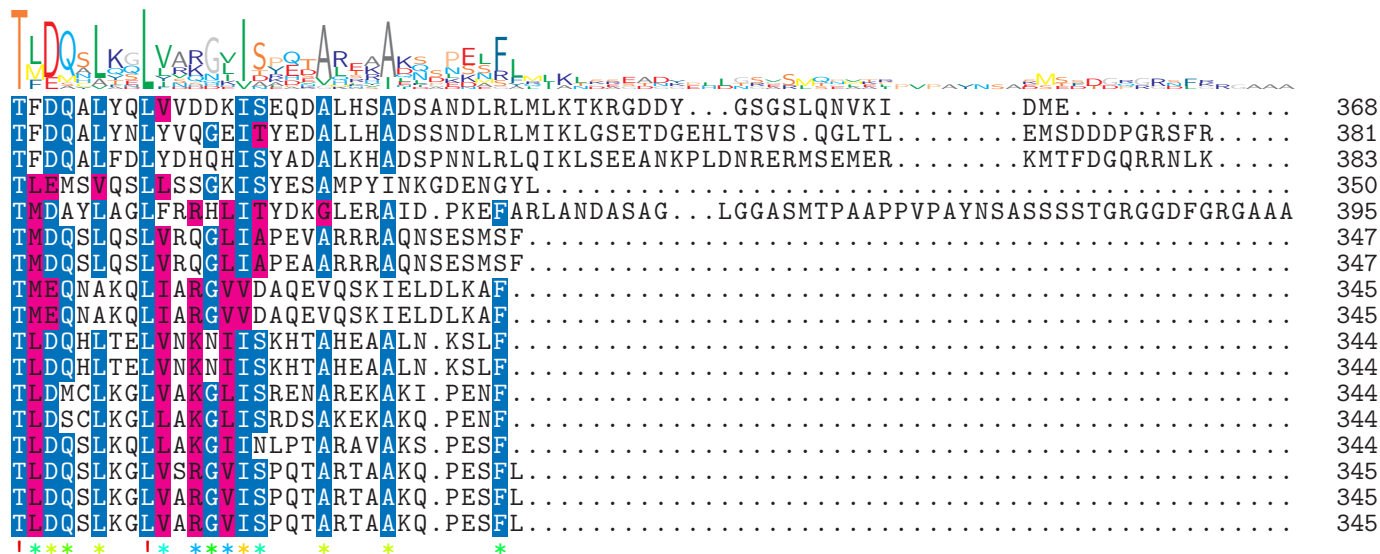
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NTDB id 1177 A1552VC RS01165 WP 000422572.1	LTIVEDPIEFVHEHKKRCIVTQREVGLDTESYEVALKNSLRQAPDMILIGEIRSRRETMEYAMTFAETGHL	233
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NTDB id 1019 ACIAD RS04205 WP 004922051.1	ITIEDPIEFIEHAGCIITQREVGLDTDSFEIALKNTLRQAPDVILIGEIRSRVMDYALAF	236
NTDB id 965509 NSQ84 RS00560 WP 063387121.1	ITLEDPIEYIHSKRSIINQREIGSDTKSFANGLRASLRQDPDIILVGEMRDLETISTA	237
NTDB id 1307 DR RS10055 WP 010888596.1	LTIEDPIEFMHPNKAISIINQREVGSDTLSFDDALRAALRQAPDVILVGEMRDYETIKA	239
NTDB id 1086 NMV RS00245 WP 002218487.1	LTIEDPIEFVHQSKKSLINQRELHQHTLSFANALRSALREDPDVILVGEMRDPETIGL	235
NTDB id 1116 NGFG RS10615 WP 003690200.1	LTIEDPIEFVHQSKKSLINQRELHQHTLSFANALSSALREDPDVILVGEMRDPETIGL	235
NTDB id 1185 GTF74 RS11940 WP 000350195.1	LTIEDPIEFVHSNNKCLINQREVHRDTHSFKNALRSALREDPDVILVGLERDQETISL	235
NTDB id 1176 A1552VC RS01160 WP 000350195.1	LTIEDPIEFVHSNNKCLINQREVHRDTHSFKNALRSALREDPDVILVGLERDQETISL	235
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NTDB id 1259 GC085 RS10305 WP 010947729.1	LTVEDPIEFVHQSKKCLVNQREVHKDTLSFNAALRSALREDPDIILVGLERDLETIRL	235
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NTDB id 1040 HON27 RS13360 WP 000355489.1	LTVEDPIEFVHQSKKCLINQREVHRDTHGFNEALRSALREDPDIILVGEMRDLETIRL	235
consensus	***!!!!*! *****!!!!*! *! *!!!!*!!!!*!!!!*!!!!*! *! *!!!!*!!!!*!!!!*!!!!*! *	

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	TIDRIIDVFPAAEEKMVRMLSESQAIVISQQLKKGGGRVAAHEIMIGTPAIRNLIREKYAQMYSIQTGAAHGMQ	313
NTDB id 1203 PSJM300 01605 AFN76401.1	ALERTIHIVPKEQREQFLLDLSLNLKGVIAQQLLRDKNGKGRHGVFEVLINSPRIADLIIRGELHELKATMARSQEVGMQ	315
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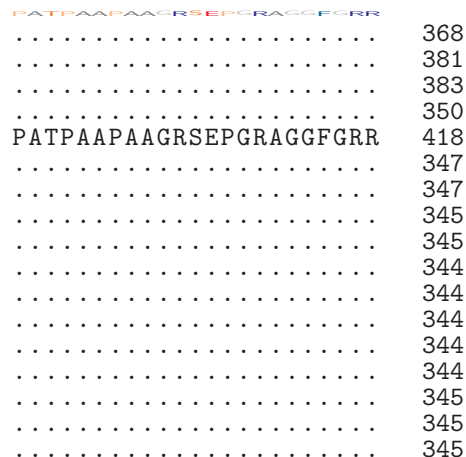
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consensus

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-  non conserved
-  similar
-  > 50% conserved