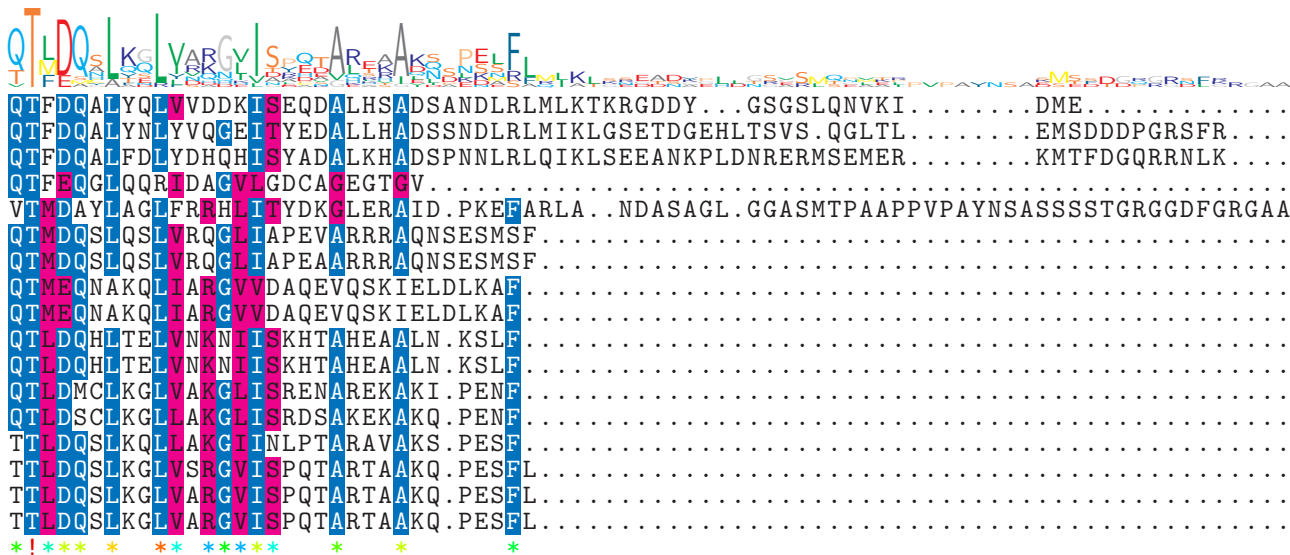


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NTDB id 1177 A1552VC RS01165 WP 000422572.1	QTFDQALYQLVDDKISEQDALHSA	DSANDLRLMLKTKRGDDY...GSGSLQNVKI.....DME.....	368
NTDB id 1203 PSJM300 01605 AFN76401.1	QTFDQALYNLYVQGEITYEDALLHA	DSSNDLRLMIKLGSETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR....	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	QTFDQALFDLYDHQHSYADALKHAD	SPNNLRLQIKLSEEANKPLDNRERMSEMER.....KMTFDGQRRNLK....	383
NTDB id 1153199 LQ139 RS21160 WP 089197423.1	QTFEQGLQQRIDACVIGDCACEGTCV	340
NTDB id 1307 DR RS10055 WP 010888596.1	VTMDAYLAGLFRHLITTYDKGLERA	ID.PKEFARLA..NDASAGL.GGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAA	394
NTDB id 1086 NMV RS00245 WP 002218487.1	QTMDQSLQSLVRQGLIAPVARRRA	QNSSESMSF.....	347
NTDB id 1116 NGFG RS10615 WP 003690200.1	QTMDQSLQSLVRQGLIAPVARRRA	QNSSESMSF.....	347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	QTMEQNAKQLIARCVVDAQEVQSKI	ELDLKAF.....	345
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NTDB id 1259 GCO85 RS10305 WP 010947729.1	QTLDQHLTELVNKNIISKHTAHEA	ALN.KSLF.....	344
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NTDB id 1202 PSJM300 01600 AFN76400.1	QTLDSCCLKGLLAKGLISRDSAKEK	AKQ.PENF.....	344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TTLDDQSLKQLLAKGIINLPTARA	VAKS.PESF.....	344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TTLDDQSLKGLVSRGVISPQTART	AAKQ.PESFL.....	345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TTLDDQSLKGLVARGVISPQTART	AAKQ.PESFL.....	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TTLDDQSLKGLVARGVISPQTART	AAKQ.PESFL.....	345
consensus	*!*** * * * * * * * *		

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	APATPAAPAAAGRSEPGRAGGFGRR	368
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NTDB id 1019 ACIAD RS04205 WP 004922051.1	383
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NTDB id 1020 ACIAD RS04210 WP 011182164.1	344
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NTDB id 1060 ABD1 RS04380 WP 000355489.1	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	345
consensus		



non conserved
 similar
 ≥ 50% conserved