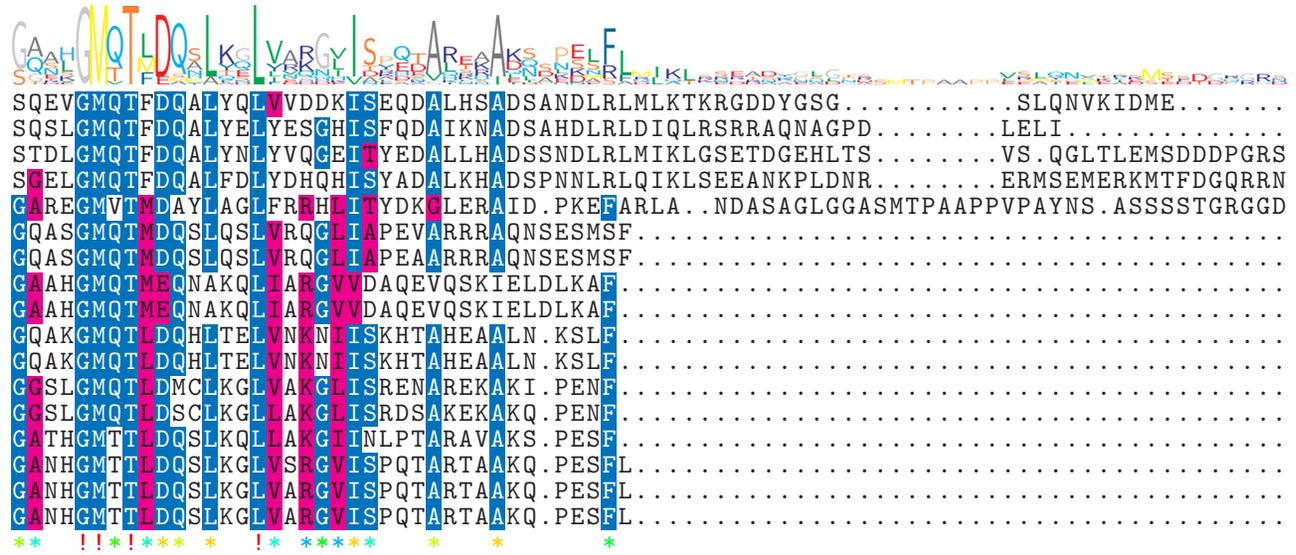


logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	SQEVGMQTFDQALYQLVDDKISEQDALHSAD	SANDLRLMLKTKRGDDYGG	SLQNVKIDME	368
NTDB id 1146947 CKV66 RS08990 WP 085363090.1	SQSLGMQTFDQALYELYESGHISFQDAIKN	ADSAHDLRLDIQLRSRRAQNA	LELI	377
NTDB id 1203 PSJM300 01605 AFN76401.1	STDLMQTFDQALYNLYVQGEITYEDALLH	ADSSNDLRLMIKLGSETDGEHLTS	VS.QGLTLEMSDDDPGRS	379
NTDB id 1019 ACIAD RS04205 WP 004922051.1	SGELCMQTFDQALFDLYDHQHISYADALKH	ADSPNNLRLQIKLSEEANKPLDNR	ERMSEMERKMTFDGQRN	381
NTDB id 1307 DR RS10055 WP 010888596.1	GAAREGMVTMDAYLAGLFRHLITDYDKGL	ERAID.PKEFARLA.NDASAGLGGASMT	PAAPPVPAYNS.ASSSSTGRGGD	388
NTDB id 1086 NMV RS00245 WP 002218487.1	GQASGMQTFDQSLQSLVRQCLIAPEVARRR	AQNSESMSF		347
NTDB id 1116 NGFG RS10615 WP 003690200.1	GQASGMQTFDQSLQSLVRQCLIAPEAARR	AQNSESMSF		347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	GAAHGMQTMENAKQLIARCVVDAQEVQSK	IELDLKAF		345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	GAAHGMQTMENAKQLIARCVVDAQEVQSK	IELDLKAF		345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	GQAKGMQTFDQHLTELVNKNIISKHTAHE	AALN.KSLF		344
NTDB id 1259 GCO85 RS10305 WP 010947729.1	GQAKGMQTFDQHLTELVNKNIISKHTAHE	AALN.KSLF		344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	CGSLCMQTFDQSLKGLVAKGLISRAREKAK	I.PENF		344
NTDB id 1202 PSJM300 01600 AFN76400.1	CGSLCMQTFDQSLKGLVAKGLISRAREKAK	I.PENF		344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	GATHGMTTLDQSLKQLLAKGIINLPTARA	VAKS.PESF		344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	GANHGMTTLDQSLKGLVSRGVISPQTART	AAKQ.PESFL		345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	GANHGMTTLDQSLKGLVARGVISPQTART	AAKQ.PESFL		345
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NTDB id 1040 HON27 RS13360 WP 000355489.1		345

consensus

- X non conserved
- X similar
- X ≥ 50% conserved