

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1
 NTDB id 1203 PSJM300 01605 AFN76401.1
 NTDB id 1019 ACIAD RS04205 WP 004922051.1
 NTDB id 1307 DR RS10055 WP 010888596.1
 NTDB id 1086 NMV RS00245 WP 002218487.1
 NTDB id 1116 NGFG RS10615 WP 003690200.1
 NTDB id 1185 GTF74 RS11940 WP 000350195.1
 NTDB id 1176 A1552VC RS01160 WP 000350195.1
 NTDB id 1283 ACE17W RS02630 WP 010947729.1
 NTDB id 1259 GC085 RS10305 WP 010947729.1
 NTDB id 1020 ACIAD RS04210 WP 011182164.1
 NTDB id 1081 FDQ49 RS05335 WP 006581387.1
 NTDB id 1060 ABD1 RS04380 WP 000355489.1
 NTDB id 1040 HON27 RS13360 WP 000355489.1
 NTDB id 25580 SDE RS19015 WP 011470110.1
 NTDB id 1194 PAKAF RS01995 WP 003084552.1
 NTDB id 1202 PSJM300 01600 AFN76400.1



TFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDY...GSGSLQNVKI.....DME..... 368
 TFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGSETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR..... 381
 TFDQALFDLYDHQHSYADALKHADSPNNLRLQIKLSEEANKPLDNRERMSEMER.....KMTFDGQRRNLK..... 383
 TMDAYLAGLFRHLITYDKGLERAIDPKFARLANDASAGL....GGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA 395
 TMDQSLQSLVRQGLIAPVARRRAQNSESMSF..... 347
 TMDQSLQSLVRQGLIAPVARRRAQNSESMSF..... 347
 TMEQNAKQLIARGVVDVDAQEVQSKIELDLKAF..... 345
 TMEQNAKQLIARGVVDVDAQEVQSKIELDLKAF..... 345
 TLDQHLTELVNKNIISKHTAHEAALNKSLF..... 344
 TLDQHLTELVNKNIISKHTAHEAALNKSLF..... 344
 TLDQSLKQLIAKGIINLPTARAVAKSPESF..... 344
 TLDQSLKGLVSRGVISPQTARTA AKQPESFL..... 345
 TLDQSLKGLVARGVISPQTARTA AKQPESFL..... 345
 TLDQSLKGLVARGVISPQTARTA AKQPESFL..... 345
 TMDQCLADLVERRIISRVDVAKAKMPDQF..... 344
 TLDMCLKGLVAKGLISRREAREKAKIPENF..... 344
 TLDSCCLKGLIAKGLISRDSAKEKAKQPENF..... 344

consensus

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PATPAAPAAGRSEPCRRACCFRR

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 PATPAAPAAGRSEPCRRACCFRR 418
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consensus

non conserved
 similar
 ≥ 50% conserved