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NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 310480 DZA53 RS18155 WP 011408067.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
consensus

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logo

Sequence	Position
ETAINEVYRDTAAGMPMYIAMRNTAEFPEMLVLMVMIGEESGQLDDMLNKVATIYEFYEVNDNTVDNLGKILEPLIIIVFLGT	387
QLAIEEVYRDTAAGMPMYIAMRHCHVPFPELVLMVMIGEESGRLDDMLNKIAIYEFYEVNDNTVDNLSKILEPLIIIVFLGV	386
EATREIRTRVIQGLSMTSGMRATELFPNMMLQMSIGEESGLDDMLNKAAEFYEVNDNAVGRLSAMMEPIIIVILGL	389
EATREIRTRVIQGLSMTSGMRATELFPNMMLQMSIGEESGLDDMLNKAAEFYEVNDNAVGRLSAMMEPIIIVILGL	389
RSATNKIKSDVSSGMQLNFSMRTTGTFFPSMAVQMTAIGEESGALDEMLSKVATFYEVNDNMVDGLTALMEPMIMAVLGV	385
EQAVLKIREDVSTGQQLNFAVRVSNRFPMTAIVMVAIGEESGALDEMLDKVANVYSEVDHVAVDGLTSMMEPLIMAILGI	388
EKAVMKIREDVATGQQLQFAMRISNRFPSMAIVMVAIGEESGALDSMLDKVATYEVNDNAVGLTSMMEPLIMAILGV	388
EAVFRMRDDVSVGYPPVNMKQVNLFPFHMVIQMTAIGEEAGALDAMLFKVAEYFEVNNVAVDALSSLEPLIMVFIGT	398
AKATDKIREEVATGQQMFIAMENHTLFPNMVIMQVAIGEESGALDKMLSKVADFYEEVNNVAVDALSSLEPIIMSILGI	386

[illegible]

- non conserved
- similar
- $\geq 50\%$ conserved