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NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 413618 GT360 RS12145 WP 164649132.1
NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1112 NFGF RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1252 GCO85 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
consensus

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logo

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consensus

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KLRDVATGGMLAMRTHTFPMMQLVAIGEESGLDMLNKVAIFYEVDNAVDSLSSMMEPILIVLGWVGLE
EVYRDTAAGMPMYIAMRHCHVFPELVLMVMIGEESGRLDMLNKLAAIYEFVDNTVDNLGKILEPLIIIVFLGVVVGGL 391
AVYRDTAAGVPYMIAMRNTHAFPEMLVQMVMIGEESGLDMLNKNVASIYEFVDNTVDNLGKILEPLIIIVFLGVVVGGL 392
EVYRDTAAGMPMYIAMRNTEAFPEMLVQMVMIGEESGLDMLNKVATIYEFVDNTVDNLGKILEPLIIIVFLGTVVVGGL 392
EIRTRVIQGLSMTSGMRA TELFPNMMLQMSSIIGEESGLDDMLNKA AEFYEVDNDNAVGRLSAMMEPIIIIVILGLVIGTL 394
EIRTRVIQGLSMTSGMRA TELFPNMMLQMSSIIGEESGLDDMLNKA AEFYEVDNDNAVGWL SAMMEPIIIILGLVLIGTL 394
KIREEVATGQQMFIA MENTHLFPNMVIQMVAIGEESGALDKMLSKVADFYEVEVDNAVDALLSILEPIIMSILGILVGGGL 391
KIKSDVSSGMQLNFSMRITGTFPSMAVQMTAIGEESGALDEMLSKVATFYEDEVDNMVDGLTALMEPMIMAVLGLVLVGGGL 390
KIREDVATGQQQLQFAMRI SNRFPSMAIQMVAIIGEESGALDSMLDKVATYYENEVDNAVDGLTSMMEPLIMAILGLVLVGGGL 393
KIREDVSTGQQLNFA MRVISNRFPMTA IQMVAIGEESGALDEMLDKVANYYESEVDHVAVDGLTSMMEPLIMAILGLVLVGGGL 393

[illegible]

-  non conserved
-  similar
-  > 50% conserved