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logo

NTDB id 414 AAK55818.1 838..2187( )   MLDLL.KQTIFTRDFIFILILLGFILVVTLLLLENRRDNIQLKQINQKVKDLIAGDYSKVLDMQGGSEITNITNNLNDLSEVIRL   84
NTDB id 34382 SDEG RS02835 WP 012766683.1   MTRDIIGNLSAFELAILLLLVFVAF.YFIHLAVRDYRNARIIRMMSHKIRDLINGRYTDMINEKADIELMELADQLNDLSDVFRL   84
NTDB id 375 SMU RS06880 WP 002262929.1   MTNVFESSPLFLRILLAVLIILLFF.YFIFLNYREYKNNNQVKQLNAKVRSLITGHYTDKLKVEDNSDLSELVNNVNDLSEVFRL   84
consensus   !* **  * ** ** * *!****** *** ! ******* ****** !***!! ! !**** * * ****** ***!!!!*!*!!

logo

NTDB id 414 AAK55818.1 838..2187( )   TQENLEQESKRLNSILFYMTDGVLATNRRGQIIMINDTAKKQLGLVKEDVLNRSILELLKIEENYELRDLITQSPELLLDSQDIN   169
NTDB id 34382 SDEG RS02835 WP 012766683.1   THENLAQEKNRLASILAYMSDGVLATDRSGKIIMINETARKQLNLTKEKALEKNITDLLEGDTPYTYRELVSKTPVVTVNSRNDM   169
NTDB id 375 SMU RS06880 WP 002262929.1   THENLAQEKNRLTSILSYMTDGVLATDRSGKITVINDMAQKQLNVTREQALECNILDILDD.DSYTYNDLITKTPEIVLTRRDEY   168
consensus   !*!!!*!!**!! !!! !!*!!!!!!*!*!*!**!!**!*!!!****! *!***!***!* ** !****!****!**** ****

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NTDB id 414 AAK55818.1 838..2187( )   GEYLNLRVRFALIRRESGFISGLVAVLHDTTEQEKEERERRLFVSNVSHELRTPLTSVKSYLEALDEGALCETVAPDFIKVSLDE   254
NTDB id 34382 SDEG RS02835 WP 012766683.1   GEFVTLRLRFALNRRESGFISGLVVVLHDTTEQEKEERERRLFVSNVSHELRTPLTSVKSYLEALDEGAIKEDIAPSFIKVSLDE   254
NTDB id 375 SMU RS06880 WP 002262929.1   DEFITLRIRFALNRRESGFISGLIAVLHDATEQEKEERERRLFVSNVSHELRTPLTSVKSYLEALDDGALTESVAPSFIKVSLDE   253
consensus   *!***!!*!!!!*!!!!!!!!!!!**!!!!*!!!!!!!!!!!
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X non conserved
X similar
X $\geq$50% conserved