```
logo
NTDB id 414 AAK55818.1 838..2187( )   MLDLLKQTIFTRDFIFILLLGFILVVTLLLLENRRDNIQLKQINQKVKDLIAGDYSKVLDMQGGSEITNITNNLNDLSE   80
NTDB id 174505 A2G56 RS09770 WP 062712138.1   MTDQFIF..TFRDLIIAGLMSILVLIYVFLIINDSLNSRRVQQLTEQVQSLISGEYVVNQVTRGSAEVRTLAEHLNDLSE   78
NTDB id 375 SMU RS06880 WP 002262929.1   MTNVFESSPLFLRILLAVLIILLFFYFIFLNYREYKNNNQVKQLNAKVRSLITGHYTDKLKVEDNSDLSELVNNVNDLSE   80
consensus   !****  *  ********************!** * *** ***!** *!**!!*!*!*  * * * **** * ***!!!!!

logo
NTDB id 414 AAK55818.1 838..2187( )   VIRLTQENLEQESKRLNSILFYMTDGVLATNRRGQIIMINDTAKKQLGLVKEDVLNRSILELLKIEENYELRDLITQSPE   160
NTDB id 174505 A2G56 RS09770 WP 062712138.1   VFRLTHENLGQEKNRLSSILSYMSDGVLATDRRGNIIMINDMARQQLNLSEEEDISVNIMTLLE.ETSYTYHDLITKTPE   157
NTDB id 375 SMU RS06880 WP 002262929.1   VFRLTHENLAQEKNRLTSILSYMTDGVLATDRSGKITVINDMAQKQLNVTREQALECNILDILD.DDSYTYNDLITKTPE   159
consensus   !*!!!*!!!*!!**!!*!!!*!!*!!!!!!*!*!*!**!!!*!**!!****!* *  *!***!* ***!** !!!!**!!

logo
NTDB id 414 AAK55818.1 838..2187( )   LLLDSQDINGEYLNLRVRFALIRRESGFISGLVAVLHDTTEQEKEERERRLFVSNVSHELRTPLTSVKSYLEALDEGALC   240
NTDB id 174505 A2G56 RS09770 WP 062712138.1   VTLNRVDEFGEFTVLRINFALIKRESGFISGLVAVLHDTTEQEKEERERRLFVSNVSHELRTPLTSVKSYLEALDEGALK   237
NTDB id 375 SMU RS06880 WP 002262929.1   IVLTRRDEYDEFITLRIRFALNRRESGFISGLIAVLHDATEQEKEERERRLFVSNVSHELRTPLTSVKSYLEALDDGALT   239
consensus   **! * !***!** !!**!!!**!!!!!!!!!*!!!!!*!!!!!!!!!
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X non conserved
X similar
X $\geq$ 50% conserved