```
logo
NTDB id 413 AAK55817.1 141..845( )          ....MKKILIVDDEKPISDIIKFNMTKEGYEVVTA.FNGREALEQFEAEQPDIIILDLMLPEIDGLEVAKTIRKTSSVPI   75
NTDB id 1147137 CKV83 RS06855 WP 018374211.1 MKLTDRKLLILDDNPDILQLVKESLTIAGFTHLTTAHTKKEALDLLRTDQFDLAILDIMLPDGSGFEVLRHIRQHSMMPA   80
NTDB id 372 SMU RS05280 WP 002262198.1      MNYKEKKILILDDNPEILEMVQESLNIAGFNNLTGVQSKKEALEQLGKRSFDLAILDIMLPDGSGFEVLREIRKTSTIPV   80
consensus                                   *   **!*!!*!!**!*********!*  *!*  ***!!!***  ***!**!!!*!!!***!*!!** !!**!**!*

logo
NTDB id 413 AAK55817.1 141..845( )          LMLSAKDSEFDKVIGLELGADDYVTKPFSNRELQARVKALLRRSQPMPVDGQEADSKPQPIQIGDLEIVPDAYVAKKYGE   155
NTDB id 1147137 CKV83 RS06855 WP 018374211.1 LFLSAVSDIEKQYQGFELGADDYIVKPFRPKELELRILSILKRSYPPQEELV.........QLEGCQIDFERAVVLKEEE   151
NTDB id 372 SMU RS05280 WP 002262198.1      LFLSAVSDIEKQYQGFELGADDYIVKPFRPKELELRILSILKRAYPEKDDTL.........ILPTCQVNFSQALITKGEL   151
consensus                                   !*!!!*********!*!!!!!!!**!!!***!!**!****!*!**!   ** *          **  *** ** *** ! **

   <0
logo
NTDB id 413 AAK55817.1 141..845( )          ELDLTHREFELLYHLASHTGQVITREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYMRNNA*   234
NTDB id 1147137 CKV83 RS06855 WP 018374211.1 EISLTAKEFSLLRVLYNHKNKIVTFDTLLTKVWGEQYEGYENTLMAHIRKIRQKIEINPSKPTHLLTVKGLGYQLRVD..   229
NTDB id 372 SMU RS05280 WP 002262198.1      EIQLTAKEYSILKVLYNNQNRIVTFDQLLEKVWGLQYQGYENTMMAHIRKIRQKIEANPSKPECLLTVKGLGYKLKVS..   229
consensus                                   !* !!**!***!**!*** *****!** !!**!!! *! !****!*****!**!*!!! *!!*!* *!!**!*!! ***
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X non conserved

X similar

X $\geq$ 50% conserved