```
logo
NTDB id 37896 MSA RS01135 WP 001008571.1   MNFEKIETAYELILENIQTIENQLKTHIYDALIEQNSYYLGSSCDLDIVVVNNQKLRQLDLSQEEWRRTFQFIFIKSAQTEQLQA   85
NTDB id 392 SMU RS09000 WP 002263442.1     MDFEKIEAAYNLLLDNCQRLETQLHTHLYDALIEQNACYLGAEGADDFIKENNKKLRQLNLSKEEWRRSFQFIFIKATQSQQLQS   85
NTDB id 401 FSA28 RS09370 WP 002291646.1   MDFEKIEAAYNLLLDNCQRLETQLHTHLYDALIEQNACYLGAEGADDFIKENNKKLRQLNLSKEEWRRSFQFIFIKATQSQQLQS   85
consensus                                  !*!!!!!*!!*!*!*!*!**!*!!*!!*!!!!!!!!!**!!!*****!****!*!!!!!*!!*!!!!!*!!!!!!!**!**!!!*

logo
NTDB id 37896 MSA RS01135 WP 001008571.1   NHQFTPDSIGFILLFLLEELTSQETVDVLEIGSGTGNLAQTLLNNSSKELNYMGIEVDDLLIDLSASIAEIIGSSAQFIQEDAVR   170
NTDB id 392 SMU RS09000 WP 002263442.1     NHQFTPDSIGFIILYLLEGLTNKSQLDIIEIGSGTGNLAETIVNNSSKSIDYMGIEVDDLLIDLSASIADVLDSSVHFVQEDAVR   170
NTDB id 401 FSA28 RS09370 WP 002291646.1   NHQFTPDSIGFIILYLLEGLTNKSQLDIIEIGSGTGNLAETIVNNSSKSIDYIGIEVDDLLIDLSASIADVLDSSVHFVQEDAVR   170
consensus                                  !!!!!!!!!!!!!*!*!!!*!!*****!**!!!!!!!!!!!*!**!!!!!***!*!!!!!!!!!!!!!!!!!!****!!**!*!!!!!!

logo
NTDB id 37896 MSA RS01135 WP 001008571.1   PQILKESDVIISDLPVGYYPNDDIAKRYAVSSSKEHTYAHHLLMEQSLKYLKKDGIAIFLAPENLLTSPQSDLLKEWLKGYADVI   255
NTDB id 392 SMU RS09000 WP 002263442.1     PQILKESDIIISDLPVGFYPNDKIAQRYQVAAKLGHTYAHHLLMEQSLKYLKKEGLAIFLAPVDLLTSEQSPLLKKWLQNNVTIL   255
NTDB id 401 FSA28 RS09370 WP 002291646.1   PQILKESDVIISDLPVGFYPNDKIAQRYQVAAKLGHTYAHHLLMEQSLKYLKKEGLAIFLAPVDLLTSEQSPLLKKWLQNNVTIL   255
consensus                                  !!!!!!!!*!!!!!!!!!*!!!!*!!*!!*!****!!!!!!!!!!!!!!!!!!!!!*!*!!!!!!**!!!!*!!*!!!*!!*******

logo
NTDB id 37896 MSA RS01135 WP 001008571.1   AVLTLPETIFGSRQNAKSIFVLKKQAEQKPETFVYPLTDLQNRENMANFIENFQKWSRENSHYSKNMIE   324
NTDB id 392 SMU RS09000 WP 002263442.1     AVISLPEAIFKQKSNIKSIFVLKKQSQQSLESFVYPLTDLQDPEVLRRFMKNFKKWKEDNVF.......   317
NTDB id 401 FSA28 RS09370 WP 002291646.1   AVISLPEAIFKQKSNIKSIFVLKKQSQQSLESFVYPLTDLQDPEVLRRFMKNFKKWKEDNVF.......   317
consensus                                  !!**!!!*!!****!*!!!!!!!!!!**!**!*!!!!!!!!!**!****!**!!*!!***!**
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X non conserved

X similar

X $\geq$ 50% conserved