

logo

NTDB id 161014 APQ13 RS08655 WP 002263442.1
NTDB id 392 SMU RS09000 WP 002263442.1
NTDB id 401 FSA28 RS09370 WP 002291646.1
consensus

MDFEKIEAAYNLLLDNCQRLETQLHTHLYDALIEQNACYLGAEGADDFIKENKKLRQLNLSKEEWRRSFQFIFIKATQS
MDFEKIEAAYNLLLDNCQRLETQLHTHLYDALIEQNACYLGAEGADDFIKENKKLRQLNLSKEEWRRSFQFIFIKATQS
MDFEKIEAAYNLLLDNCQRLETQLHTHLYDALIEQNACYLGAEGADDFIKENKKLRQLNLSKEEWRRSFQFIFIKATQS
!!

80
80
80

logo

NTDB id 161014 APQ13 RS08655 WP 002263442.1
NTDB id 392 SMU RS09000 WP 002263442.1
NTDB id 401 FSA28 RS09370 WP 002291646.1
consensus

QQLQSNHQFTPDSIGFIILYLLEGLTNKSQLDIIIEIGSGTGNAETIVNNSKSIDYMGIEVDDLLIDLASIAADVLDSS
QQLQSNHQFTPDSIGFIILYLLEGLTNKSQLDIIIEIGSGTGNAETIVNNSKSIDYMGIEVDDLLIDLASIAADVLDSS
QQLQSNHQFTPDSIGFIILYLLEGLTNKSQLDIIIEIGSGTGNAETIVNNSKSIDYMGIEVDDLLIDLASIAADVLDSS
!!

160
160
160

logo

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consensus

VHFVQEDAVRPQILKESDIIISDLPVGFYPNDKIAQRYQVAAKLGHTYAHLLMEQSLKYLKKEGLAIFLAPVDLLTSEQ
VHFVQEDAVRPQILKESDIIISDLPVGFYPNDKIAQRYQVAAKLGHTYAHLLMEQSLKYLKKEGLAIFLAPVDLLTSEQ
VHFVQEDAVRPQILKESDIIISDLPVGFYPNDKIAQRYQVAAKLGHTYAHLLMEQSLKYLKKEGLAIFLAPVDLLTSEQ
!!

240
240
240

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consensus

SPLLKKWLQNNVTILAVISLPEAIFKQKSNIKSIFVLKKQSQSLESFVYPLTDLQDPEVLRFRFMKFNFKKWKEDNVF
SPLLKKWLQNNVTILAVISLPEAIFKQKSNIKSIFVLKKQSQSLESFVYPLTDLQDPEVLRFRFMKFNFKKWKEDNVF
SPLLKKWLQNNVTILAVISLPEAIFKQKSNIKSIFVLKKQSQSLESFVYPLTDLQDPEVLRFRFMKFNFKKWKEDNVF
!!

317
317
317

⊠ non conserved
⊠ similar
⊠ ≥ 50% conserved