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logo

NTDB id 58966 QMM10 RS07095 WP 000106346.1  MSTNRKNDMMVYCSFCGKNQEEVQKIIAGNNAFICNECVELAQEIIREELVEEVLADLSEVPKPIELLHILNHYVIGQDRAKRAL   85
NTDB id 379 SMU RS04375 WP 002262816.1      MAGNRTNDVTVHCSFCGKNQDEVKKIIAGNGVFICNECVELSQEIIREELAEEVLADLSEVPKPKELLAILDSYVIGQDRAKRAL   85
consensus                                   !  !! !!* !*!!!!!!!!*!! !!!!!!  !!!!!!!!! !!!!!!!! !!!!!!!!!!!!! !!!*!! *!!!!!!!!!!!!!

logo

NTDB id 58966 QMM10 RS07095 WP 000106346.1  AVAVYNHYKRINFHDTREESEDVDLQKSNILMIGPTGSGKTFLAQTLAKSLNVPFAIADATALTEAGYVGEDVENILLKLLQVAD   170
NTDB id 379 SMU RS04375 WP 002262816.1      AVAVYNHYKRISFTESQ.DDQDVDLQKSNILMIGPTGSGKTFLAQTLAKSLNVPFAIADATALTEAGYVGEDVENILLKLIQAAD   169
consensus                                   !!!!!!!!!!! !*** **  !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*! !!

logo

NTDB id 58966 QMM10 RS07095 WP 000106346.1  FNIERAERGIIYVDEIDKIAKKSENVSITRDVSGEGVQQALLKIIEGTVASVPPQGGRKHPQQEMIQVDTKNILFIVGGAFDGIE   255
NTDB id 379 SMU RS04375 WP 002262816.1      YNVERAERGIIYVDEIDKIAKKGENVSITRDVSGEGVQQALLKIIEGTVASVPPQGGRKHPQQEMIQIDTKNILFIVGGAFDGIE   254
consensus                                   *!*!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 58966 QMM10 RS07095 WP 000106346.1  EIVKQRLGEKVIGFGQNNKAIDENSSYMQEIIAEDIQKFGIIPELIGRLPVFAALEQLTVDDLVRILKEPRNALVKQYQTLLSYD   340
NTDB id 379 SMU RS04375 WP 002262816.1      DIVKQRLGEKIIGFGQNNKKIDDQSSYMQEIISEDIQKFGLIPEFIGRLPVLAALEQLTVDDLVKILTEPKNALVKQYQTLLSYD   339
consensus                                   *!!!!!!!!!*!!!!!!!! !!**!!!!!!!! !!!!!!!*!!! !!!!!! !!!!!!!!!!!!*!! !!*!!!!!!!!!!!!!!!

logo

NTDB id 58966 QMM10 RS07095 WP 000106346.1  DVELEFDDEALQEIANKAIERKTGARGLRSIIEETMLDVMFEVPSQENVKLVRITKETVDGTDKPILETA.   410
NTDB id 379 SMU RS04375 WP 002262816.1      GVELEFDQEALQAIAQKAIERKTGARGLRSIIEETMLDLMFEIPSQEDVTCVRITKKAVEGTDKPILETAS   410
consensus                                    !!!!!! !!!! !!*!!!!!!!!!!!!!!!!!!!
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