```
logo

NTDB id 1123926 EL297 RS06370 WP 082308618.1  MRRKMLNVPKGSYDGMKGFTIIEFLVAGLLSMIVLIAVGSSYFTSRKLNDAANERLAEQQDLRNAATLIVRDARMAGSFG   80
NTDB id 1138 NGFG RS02430 WP 003694978.1      MKRKMLNVPKGGYDGMKGFTIVEFLVAGLLSVIVLIAVVSSYFTSRKLNDAANERLAEQQDLRNAATLIVRDARMAGSFG   80
consensus                                     !*!!!!!!!!! !!!!!!!!!*!!!!!!!!!*!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1123926 EL297 RS06370 WP 082308618.1  CFNMSEHTKDDIVDSSNQTQPASAKPGVKQENPLFSLKRS...DMEQLIPVTESADIKYSGFTQRLNALVFQYGIDDLDA  157
NTDB id 1138 NGFG RS02430 WP 003694978.1      CFNMSEHIGNDVVFN............VAQKNALFSLKRNSTNSTNKLIPITESPNINYQNFFQVSSALIFQYGIDDVDA  148
consensus                                     !!!!!!!   !*!  ************! ! !*!!!!!! ***    !!!*!!!* ! !  ! !   !!*!!!!!!!*!!

logo

NTDB id 1123926 EL297 RS06370 WP 082308618.1  SAETVVVSSCSKIAKPGKKISTLQEAKSALQITNDD.KQNGNITRQRHVVNAYAVGRIADEEGLFRFQLDDKGKWGNPQL  236
NTDB id 1138 NGFG RS02430 WP 003694978.1      SADTTVVSSCAAISKPGKQIPTLENAKKELKIQNSDKEQNGNIARQRHVVNAYAVGKIAGEEGLFRFQLDDKGKWGNPQL  228
consensus                                     !!*! !!!!!  ! !!!! !*!!  !!  ! ! ! !* !!!!! !!!!!!!!!!!!*!! !!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1123926 EL297 RS06370 WP 082308618.1  LVKKVKRMNVRYIYVSGCPEDEDAGKEEQFKYTNEFDKSKNAVTPAGVEVLLDSGLDAKIAASSDNSIYAYRINATIRGG  316
NTDB id 1138 NGFG RS02430 WP 003694978.1      LAKKVKRMRVRYIYVSGCPEDEDAGKEEQFKYTDKFDS...SVTPAGVEVLLDSGSDAKIAASSDNIIYAYRINATIRGG  305
consensus                                     ! !!!!!! !!!!!!!!!!!!!!!!!!!!!!!!  !! *** !!!!!!!!!!!!! !!!!!!!!!! !!!!!!!!!!!!!

logo

NTDB id 1123926 EL297 RS06370 WP 082308618.1  NVCANRTL   324
NTDB id 1138 NGFG RS02430 WP 003694978.1      NVCANRTL   313
consensus                                     !!!!!!!!
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X non conserved
X similar
X ≥ 50% conserved