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logo

NTDB id 328231 EGH14 RS02755 WP 047920986.1   MKRKMLNVPKGGYDGMKGFTIVEFLVAGLLSIIVLIAVVSSYFTSRKLNDAANERLAIQQDLRNAATLIVRDARMAGSFG   80
NTDB id 1138 NGFG RS02430 WP 003694978.1      MKRKMLNVPKGGYDGMKGFTIVEFLVAGLLSVIVLIAVVSSYFTSRKLNDAANERLAEQQDLRNAATLIVRDARMAGSFG   80
consensus                                     !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 328231 EGH14 RS02755 WP 047920986.1   CFNMSEHNKDDIVDSSNQTQFNSAKPDAKQENPLFPLKRSSM..DKQLIPVAESTDIKYPGFTQRLNALVFQYGIDDLDA   158
NTDB id 1138 NGFG RS02430 WP 003694978.1      CFNMSEHIGNDVVFNV............AQKNALFSLKRNSTNSTNKLIPITESPNINYQNFFQVSSALIFQYGIDDVDA   148
consensus                                     !!!!!!!   !*!   ************ ! !*!!*!!! ! **   !!!* !!* ! !* ! !   !!*!!!!!!!*!!

logo

NTDB id 328231 EGH14 RS02755 WP 047920986.1   SAETVVVSSCSKIAKPGKKISTLQEAKSALQITNDD.KQNGNITRQKHVVNAYAVGRIADEEGLFRFQLDDKGKWGNPQL   237
NTDB id 1138 NGFG RS02430 WP 003694978.1      SADTTVVSSCAAISKPGKQIPTLENAKKELKIQNSDKEQNGNIARQRHVVNAYAVGKIAGEEGLFRFQLDDKGKWGNPQL   228
consensus                                     !!*! !!!!!  ! !!!! !*!!  !!  ! ! ! !* !!!!! !!*!!!!!!!!!*!! !!!!!!!!!!!!!!!!!!!!

logo

NTDB id 328231 EGH14 RS02755 WP 047920986.1   LAKKIKRMKVLYIYVSGCPEDEDAGKEEQFKYTDKFDSSVTPAGVEVLLDSGLNAKIAASSDNSIYAYRINATIRGGNVC   317
NTDB id 1138 NGFG RS02430 WP 003694978.1      LAKKVKRMRVRYIYVSGCPEDEDAGKEEQFKYTDKFDSSVTPAGVEVLLDSGSDAKIAASSDNIIYAYRINATIRGGNVC   308
consensus                                     !!!!*!!!*! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!  !!!!!!!!! !!!!!!!!!!!!!!!!

logo

NTDB id 328231 EGH14 RS02755 WP 047920986.1   ANRTL   322
NTDB id 1138 NGFG RS02430 WP 003694978.1      ANRTL   313
consensus                                     !!!!!
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X non conserved
X similar
X ≥ 50% conserved