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logo

NTDB id 26853 NMO RS04195 WP 015815451.1  MRRKMLNVPKGSYDGMKGFTIIEFLVAGMLSMIVLMAVGSSYFTSRKLNDAANERLAIQQDLRNAATLIVRDARMAGSFGCFNMS   85
NTDB id 1138 NGFG RS02430 WP 003694978.1  MKRKMLNVPKGGYDGMKGFTIVEFLVAGLLSVIVLIAVVSSYFTSRKLNDAANERLAEQQDLRNAATLIVRDARMAGSFGCFNMS   85
consensus                                 !*!!!!!!!!! !!!!!!!!!*!!!!!!*!!*!!!*!! !!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 26853 NMO RS04195 WP 015815451.1  EHTEKNVVSDVAQKNRLFSLKGSSANKLIPNGTDNKLIPITESLDIGYQGFTQRLNALVFQYGIDDLDASAETVVVSSCSKIAKP   170
NTDB id 1138 NGFG RS02430 WP 003694978.1  EHIGNDVVFNVAQKNALFSLKRNSTN......STNKLIPITESPNINYQNFFQVSSALIFQYGIDDVDASADTTVVSSCAAISKP   164
consensus                                 !!    !!  !!!!! !!!!!  ! !******* !!!!!!!!!* ! !! ! !   !!*!!!!!!!*!!!!*! !!!!!  ! !!

logo

NTDB id 26853 NMO RS04195 WP 015815451.1  GKKISTLQEAKSALQITNDD.KQNGNITRQRYVVNAYAVGRIAGEEGLFRFQLDDKGKWGNPQLLVKKINKMDIRYIYVSDCPED   254
NTDB id 1138 NGFG RS02430 WP 003694978.1  GKQIPTLENAKKELKIQNSDKEQNGNIARQRHVVNAYAVGKIAGEEGLFRFQLDDKGKWGNPQLLAKKVKRMRVRYIYVSGCPED   249
consensus                                 !! !*!!  !!  ! ! ! !* !!!!! !!!*!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 26853 NMO RS04195 WP 015815451.1  DDAGKEEKFKYTGTFDSSTNAVTPAGVEVLLSSGTDTKIAASSDNHIYAYRIDATIRGGNVCANRTL   321
NTDB id 1138 NGFG RS02430 WP 003694978.1  EDAGKEEQFKYTDKFDSS...VTPAGVEVLLDSGSDAKIAASSDNIIYAYRINATIRGGNVCANRTL   313
consensus                                 *!!!!!! !!!!  !!!!***!!!!!!!!!!! !!*! !!!!!!!!!*!!!!!!! !!!!!!!!!!!!!!!
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X non conserved
X similar
X ≥ 50% conserved