```
logo
NTDB id 32121 NMV RS07480 WP 014574102.1  MRRKMLNVPKGSYDGMKGFTIIEFLVAGMLSMIVLMAVGSSYFTSRKLNDAANERLAIQQDLRNAATLIVRDARMAGSFGCFNMS   85
NTDB id 1138 NGFG RS02430 WP 003694978.1  MKRKMLNVPKGGYDGMKGFTIVEFLVAGLLSVIVLIAVVSSYFTSRKLNDAANERLAEQQDLRNAATLIVRDARMAGSFGCFNMS   85
consensus                                 !*!!!!!!!!! !!!!!!!!!*!!!!!!*!!*!!!*!! !!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 32121 NMV RS07480 WP 014574102.1  EHTEKNVVSDVAQKNRLFSLKGSSANKLIPNGTDNKLIPITESLDIGYQGFTQRLNALVFQYGIDDLDASAETVVVSSCSKIAKQ  170
NTDB id 1138 NGFG RS02430 WP 003694978.1  EHIGNDVVFNVAQKNALFSLKRNSTN......STNKLIPITESPNINYQNFFQVSSALIFQYGIDDVDASADTTVVSSCAAISKP  164
consensus                                 !!    !!  !!!!! !!!!!  ! !******* !!!!!!!!!* ! !! ! !   !!*!!!!!!!*!!!!*! !!!!!  ! !*

logo
NTDB id 32121 NMV RS07480 WP 014574102.1  GKKISTLQEAKSALQITDD.DKQNGNITRQRYVVNAYAVGRIAGEDGLFRFQLDDKGKWGNPQLLVKKINKMDIRYIYVSDCPED  254
NTDB id 1138 NGFG RS02430 WP 003694978.1  GKQIPTLENAKKELKIQNSDKEQNGNIARQRHVVNAYAVGKIAGEEGLFRFQLDDKGKWGNPQLLAKKVKRMRVRYIYVSGCPED  249
consensus                                 !! !*!!  !!  ! !   *  !!!!! !!!*!!!!!!!!*!!!!*!!!!!!!!!!!!!!!!!!!!! !!* *! *!!!!!! !!!!

logo
NTDB id 32121 NMV RS07480 WP 014574102.1  DDAGKEEKFKYTGTFDSSTNAVTPAGVEVLLSSGTDTKIAASSDNHIYAYRIDATIRGGNVCANRTL   321
NTDB id 1138 NGFG RS02430 WP 003694978.1  EDAGKEEQFKYTDKFDSS...VTPAGVEVLLDSGSDAKIAASSDNIIYAYRINATIRGGNVCANRTL   313
consensus                                 *!!!!!! !!!!  !!!!***!!!!!!!!!!! !!*! !!!!!!!!*!!!!!! !!!!!!!!!!!!!!
```

X non conserved
X similar
X $\geq$ 50% conserved