```
logo

NTDB id 21647 NMA RS05550 WP 010981161.1   MRRKMLNVPKGNYDGMKGFTIIEFLVAGMLSMIVLMAVGSSYFTSRKLNDAANERLSAQQDLRNAATLIVRDARMAGGFGCFNMS   85
NTDB id 1138 NGFG RS02430 WP 003694978.1   MKRKMLNVPKGGYDGMKGFTIVEFLVAGLLSVIVLIAVVSSYFTSRKLNDAANERLAEQQDLRNAATLIVRDARMAGSFGCFNMS   85
consensus                                  !*!!!!!!!!! !!!!!!!!!*!!!!!!*!!*!!!*!! !!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!! !!!!!!!

logo

NTDB id 21647 NMA RS05550 WP 010981161.1   EHTKNDIIVDPSKQTQHVPVKPGAKQENPLFSLEWANTNNTNNNTAKLIPIAESTDIKYPGFAQARPALIFQYGIDDLDASAETV   170
NTDB id 1138 NGFG RS02430 WP 003694978.1   EHIGNDVVFNVA.............QKNALFSLKR....NSTNSTNKLIPITESPNINYQNFFQVSSALIFQYGIDDVDASADTT   153
consensus                                  !!  !!**  * *************! !*!!!!  ****!* ! ! !!!!! !!* ! !* ! !  *!!!!!!!!!!*!!!!*!

logo

NTDB id 21647 NMA RS05550 WP 010981161.1   VVSSCSKIAKPGKKISTLQEAKSALQITNDD.KQNGNITRQRHVVNAYAVGRIAGEEGLFRFQLDDKGKWGNPQLLVKKIRHMKV   254
NTDB id 1138 NGFG RS02430 WP 003694978.1   VVSSCAAISKPGKQIPTLENAKKELKIQNSDKEQNGNIARQRHVVNAYAVGKIAGEEGLFRFQLDDKGKWGNPQLLAKKVKRMRV   238
consensus                                  !!!!!  ! !!!! !*!!  !!  ! ! ! !* !!!!! !!!!!!!!!!!!*!!!!!!!!!!!!!!!!!
```