```
logo
NTDB id 146287 AA977 RS06690 WP 064435046.1  MKKSLCLSFFLAFSNPLQALVIELLEEIKTSPHKGTFKAKVLDSKEPRQVLGVYNISPHKKLTLTITHISTAIVYQPLDE   80
NTDB id 1207 C694 RS07860 WP 000749005.1     MKKSLCLSFFLTFSNPLQALVIELLEEIKTSPHKGTFKAKVLDSKKPRQVLGVYNISPHKKLTLTITHISTAIVYQPLDE   80
consensus                                    !!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 146287 AA977 RS06690 WP 064435046.1  KLSLETTLNPDRSIIPRNTQIIFSS.ELKEPRPNQIPSLNAPMQKPQNKPSSSQQLPQNFSYPESKLGPKNSKNSLLQPL  159
NTDB id 1207 C694 RS07860 WP 000749005.1     KLSLETTLNPNRPTIPRNTQIVFSSKELKESHPHQMPSLNAPMQKPQNKPHSSQQPSQNFSYPEPKLGSKNSKNSLLQPL  160
consensus                                    !!!!!!!!!!! !* !!!!!!!*!!!*!!!!**!*!*!!!!!!!!!!!!!!*!!!!**!!!!!!!*!!!*!!!!!!!!!!!!

logo
NTDB id 146287 AA977 RS06690 WP 064435046.1  TTPSKISSANETKTPTNDTKPPLKHSSEDQENNLFITPPTEKTPPNNNASHADTHENNESNEDSKDNAEKQAVRDPNVKE  239
NTDB id 1207 C694 RS07860 WP 000749005.1     AIPSKISPTNETQTPTNDTKPPLKHSSEDQESNLFITPPTEKTLPNNTS.NADISENNESN.ENKDNVEKQAIRDANIKE  238
consensus                                      !!!!!* !!!  !!!!!!!!!!!!!!!!!!!  !!!!!!!!!!!*!!!   **!! *!!!!!!** !!!  !!!!*!!*!*!!

logo
NTDB id 146287 AA977 RS06690 WP 064435046.1  FACGKWVYDDENLQAYRPSILKRVDENKQTTTDITPCDYSMAENKSGKIITPYTKISVQKTEPLEEPQTFEAKNNFAILQ  319
NTDB id 1207 C694 RS07860 WP 000749005.1     FACGKWVYDDENLQAYRPSILKRVDEDKQTATDITPCDYSTAENKSGKIITPYTKISVHKTEPLEEPQTFEAKNNFAILQ  318
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!! !!!  !!!!!!!!! !!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 146287 AA977 RS06690 WP 064435046.1  ARSTTEKCKRARARKDGTIRQCYLIEEPLKQAWESEYEITTQLVKAIYERPKQDDQVEPTFYETSELAYSSTRKSEITHN  399
NTDB id 1207 C694 RS07860 WP 000749005.1     ARSSTEKCKRARARKDGTTRQCYLIEEPLKQAWESEYEITTQLVKAIYERPKQDDQVEPTFYETSELAYSSTRKSEITHN  398
consensus                                    !!!*!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!
```

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