```
logo

NTDB id 350693 CKAES1R RS10490 WP 003091163.1  MWELVQAGGWMMLPIILSSIAATAIVAERLWTLRKSRVAPPELLGQVWKQIKGKQLSAQKLKDLRANSPLGEILAAGLAN   80
NTDB id 1200 PSJM300 12655 AFN78593.1          MWELVKAGGWIMLPIILCSIAALGIIVERLWTLRPSRVTPPHLLGQVWKWIKDKQLSNQKLKELRADSPLGEILAAGLAN   80
consensus                                      !!!!! !!!!*!!!!!!*!!!! *!* !!!!!!!*!!! !!*!!!!!!! !! !!!! !!!!*!!! !!!!!!!!!!!!!

logo

NTDB id 350693 CKAES1R RS10490 WP 003091163.1  SKHGREIMKECIEEAASGVIHELERYLNALGTIAAMAPLLGLLGTVFGMIQIFGAFMDNGMANSPMLAAGISKALITTAA  160
NTDB id 1200 PSJM300 12655 AFN78593.1          SKRGREIMKECIEEAARVVHDLERYLNALGTIAGIAPLLGLLGTVLGMIEIFSAFMGSGMANAPLLAGGISKALITTAA  160
consensus                                      !!*!!!!!!!!!!!!!  !*!*!!!!!!!!!!!!**!!!!!!!!!! !!! !! !!!  !!!! !*!!*!!!!!!!!!!!

logo

NTDB id 350693 CKAES1R RS10490 WP 003091163.1  GLLVAIPAVFFHRYLLRRVDELVIAMEQEAIKLVEVVQGDREVDYVEEGKA   211
NTDB id 1200 PSJM300 12655 AFN78593.1          GLMVGIPALFFHRFLLRRVDELVVGMEQEAIKLVEVVQGDRDVDLGEGKA.   210
consensus                                      !!*!*!!!*!!!!*!!!!!!!!!**!!!!!!!!!!!!!!!!!*!!  !   *
```

X non conserved

X similar

X $\geq$ 50% conserved