```
logo

NTDB id 182679 A214 RS23420 WP 003091163.1  MWELVQAGGWMMLPIILSSIAATAIVAERLWTLRKSRVAPPELLGQVWKQIKGKQLSAQKLKDLRANSPLGEILAAGLANSKHGR   85
NTDB id 1200 PSJM300 12655 AFN78593.1       MWELVKAGGWIMLPIILCSIAALGIIVERLWTLRPSRVTPPHLLGQVWKWIKDKQLSNQKLKELRADSPLGEILAAGLANSKRGR   85
consensus                                   !!!!! !!!!*!!!!!!*!!!! *!* !!!!!!!*!!! !!*!!!!!!! !! !!!! !!!!*!!! !!!!!!!!!!!!!!!*!!

logo

NTDB id 182679 A214 RS23420 WP 003091163.1  EIMKECIEEAASGVIHELERYLNALGTIAAMAPLLGLLGTVFGMIQIFGAFMDNGMANSPMLAAGISKALITTAAGLLVAIPAVF  170
NTDB id 1200 PSJM300 12655 AFN78593.1       EIMKECIEEAARVVHDLERYLNALGTIAGIAPLLGLLGTVLGMIEIFSAFMGSGMANAPLLAGGISKALITTAAGLMVGIPALF  170
consensus                                   !!!!!!!!!!!  !*!*!!!!!!!!!!!!**!!!!!!!!!! !!! !! !!!  !!!! !*!!*!!!!!!!!!!!!!!*!*!!!*!

logo

NTDB id 182679 A214 RS23420 WP 003091163.1  FHRYLLRRVDELVIAMEQEAIKLVEVVQGDREVDYVEEGKA   211
NTDB id 1200 PSJM300 12655 AFN78593.1       FHRFLLRRVDELVVGMEQEAIKLVEVVQGDRDVDLGEGKA.  210
consensus                                   !!!*!!!!!!!!!**!!!!!!!!!!!!!!!!!!*!!  !   *
```

X non conserved

X similar

X $\geq$ 50% conserved