```
logo
NTDB id 302116 Y31 RS14435 WP 003091163.1  MWELVQAGGWMMLPIILSSIAATAIVAERLWTLRKSRVAPPELLGQVWKQIKGKQLSAQKLKDLRANSPLGEILAAGLANSKHGR   85
NTDB id 1200 PSJM300 12655 AFN78593.1      MWELVKAGGWIMLPIILCSIAALGIIVERLWTLRPSRVTPPHLLGQVWKWIKDKQLSNQKLKELRADSPLGEILAAGLANSKRGR   85
consensus                                  !!!!! !!!!*!!!!!!*!!!! *!* !!!!!!!*!!! !!*!!!!!!! !! !!!! !!!!*!!! !!!!!!!!!!!!!!!*!!

logo
NTDB id 302116 Y31 RS14435 WP 003091163.1  EIMKECIEEAASGVIHELERYLNALGTIAAMAPLLGLLGTVFGMIQIFGAFMDNGMANSPMLAAGISKALITTAAGLLVAIPAVF  170
NTDB id 1200 PSJM300 12655 AFN78593.1      EIMKECIEEAARVVHDLERYLNALGTIAGIAPLLGLLGTVLGMIEIFSAFMGSGMANAPLLAGGISKALITTAAGLMVGIPALF  170
consensus                                  !!!!!!!!!!!  !*!*!!!!!!!!!!!!**!!!!!!!!!! !!! !! !!!  !!!!! !*!!*!!!!!!!!!!!!!!*!*!!!*!

logo
NTDB id 302116 Y31 RS14435 WP 003091163.1  FHRYLLRRVDELVIAMEQEAIKLVEVVQGDREVDYVEEGKA   211
NTDB id 1200 PSJM300 12655 AFN78593.1      FHRFLLRRVDELVVGMEQEAIKLVEVVQGDRDVDLGEGKA.   210
consensus                                  !!!*!!!!!!!!!**!!!!!!!!!!!!!!!!!*!!  !   *
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X non conserved

X similar

X ≥50% conserved