```
logo
NTDB id 111495 A16 RS07980 WP 000155594.1  MTIRYPNGKRYNQASQP..HKTPIKKHTYSNRGMSLEEELNETNEYYLTHNIACVHKKPTPLQIVKVDYPARSAAVVKEAYFKQP   83
NTDB id 117 BSU 22310 NP 390112.1          .MIRYPNGKTFQPKHSVSSQNSQKRAPSYSNRGMTLEDDLNETNKYYLTNQIAVIHKKPTPVQIVNVHYPKRSAAVIKEAYFKQS   84
consensus                                  * !!!!!!! *** * **** ** *  *!!!!!!*!!**!!!!! !!!!**!!**!!!!!!*!!! !*!! !!!!!*!!!!!!!*

logo
NTDB id 111495 A16 RS07980 WP 000155594.1  STTDYNGVYKGKYIDFEAKETKNKTSFPLQNFHLHQIEHMKQVIAHNGIAFVIIKFTLFDELYLLDAKHIITFWNRQNTGGRKSI   168
NTDB id 117 BSU 22310 NP 390112.1          STTDYNGIYKGRYIDFEAKETKNKTSFPLQNFHDHQIEHMKQVKAQDGICFVII..SAFDQVYFLEADKLFYFWDRKEKNGRKSI   167
consensus                                  !!!!!!!*!!!*!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!! !* !!*!!!!*** !! *! !*! **  !! !    !!!!!

logo
NTDB id 111495 A16 RS07980 WP 000155594.1  TKEEIVEHGSLLSCGYHPRIDYIRVLDTVYFS.......   200
NTDB id 117 BSU 22310 NP 390112.1          RKDELEETAYPISLGYAPRIDYISIIEQLYFSPSSGAKG   206
consensus                                   !*!* !** **!*!!*!!!!!! *** *!!!*******
```

X non conserved

X similar

X $\geq$50% conserved