```
logo

NTDB id 506 SM12261 RS00975 WP 000265657.1     ........MWKKQKVKAGVLLYAVTMAAIFSLLLQFYLNRQVAHYQDYTLNKEKLVAFAMAKRTKDKVEQESGEQAFNLG   72
NTDB id 109590 ACGHT2 RS00770 WP 238374707.1  MVWRGHLFTLLKQQVKAGILLIYALLMAAIFALLLQFYLGRVVAMERQHQAQLSASQAYLMAELTKDLAKEDTGHLTFDQG   80
NTDB id 400 FSA28 RS09375 WP 002263441.1      ........MLFMKKIKAGVLLYALFMSAIFSLFLQFYLNRVVATERQNQAQLSASKAYLIADLTQDLAKDNFGQLTFNHG   72
NTDB id 391 SMU RS09005 WP 002263441.1        ........MLFMKKIKAGVLLYALFMSAIFSLFLQFYLNRVVATERQNQAQLSASKAYLIADLTQDLAKDNFGQLTFNHG   72
consensus                                               ***  **!!!*!*!!**! !!!*! !!!!!*!*!!**************!***!**! !******!***!**!

logo

NTDB id 506 SM12261 RS00975 WP 000265657.1     QVSYQNKKTSLVTTVRTPK.SQYEFIFPSVKIKEEKTDKKEKVVTDSSNQAEKKKSEEKVEKKENF..   137
NTDB id 109590 ACGHT2 RS00770 WP 238374707.1  VTDYNFQDNRLQVMVKLPNQQEFDFVFQSVQKEKAN.............................   116
NTDB id 400 FSA28 RS09375 WP 002263441.1      RSSYHKQEGELLIKVHLTDGQSYQYRFNCVTENKEKGKF..SVANKTFTKS.........IEKRHFQK   129
NTDB id 391 SMU RS09005 WP 002263441.1        RSSYHKQEGELLIKVHLTDGQSYQYRFNCVTENKEKGKF..SVANKTFTKS.........IEKRHFQK   129
consensus                                      ***!******!***!***********!**!*********  **********         ********
```

X non conserved

X similar

X $\geq$ 50% conserved