```
logo
                                                  ..MWKKQKVKAGVLLYAVTMAAIFSLLLQFYLNRQVAHYQDYTLNKEKLVAFAMAKRTKDKVEQESGEQAFNLGQ    73
NTDB id 506 SM12261 RS00975 WP 000265657.1
NTDB id 1111664 HMPREF9319 RS00415 WP 003062925.1  MFTLLKRQVKAGILIYALLMAAIFALLLQFYLGRVVAMERQHQAQLSASQAYLMAELTKDLAKEDTGHLTFDQGV    75
NTDB id 400 FSA28 RS09375 WP 002263441.1           ..MLFMKKIKAGVLLYALFMSAIFSLFLQFYLNRVVATERQNQAQLSASKAYLIADLTQDLAKDNFGQLTFNHGR    73
NTDB id 391 SMU RS09005 WP 002263441.1             ..MLFMKKIKAGVLLYALFMSAIFSLFLQFYLNRVVATERQNQAQLSASKAYLIADLTQDLAKDNFGQLTFNHGR    73
consensus                                            *** ***!!!*!*!!**! !!!*! !!!!!*!*!!*************!***!**! !******!***!**!*

logo
NTDB id 506 SM12261 RS00975 WP 000265657.1         VSYQNKKTSLVTTVRTPKS.QYEFIFPSVKIKEEKTDKKEKVVTDSSNQAEKKKSEEKVEKKENF..    137
NTDB id 1111664 HMPREF9319 RS00415 WP 003062925.1  TDYNFQDNRLQVMVKLPNQQEFDYVFQSVQKEKAN.............................    110
NTDB id 400 FSA28 RS09375 WP 002263441.1           SSYHKQEGELLIKVHLTDGQSYQYRFNCVTENKEKGKF..SVANKTFTKS.........IEKRHFQK    129
NTDB id 391 SMU RS09005 WP 002263441.1             SSYHKQEGELLIKVHLTDGQSYQYRFNCVTENKEKGKF..SVANKTFTKS.........IEKRHFQK    129
consensus                                          **!******!***!***********!**!*********  **********         ********
```

X non conserved

X similar

X $\geq$ 50% conserved