```
logo
NTDB id 506 SM12261 RS00975 WP 000265657.1   ..MWKKQKVKAGVLLYAVTMAAIFSLLLQFYLNRQVAHYQDYTLNKEKLVAFAMAKRTKDKVEQESGEQAFNLGQVSYQNKKTSL   83
NTDB id 20326 SMA 0098 CCF01389.1            MFTLLKRQVKAGILIYALLMAAIFALLLQFYLGRVVAMERQHQAQLSASQAYLMAKLTKDLAKEDTGHLTFDQGVADYNFQDNRL   85
NTDB id 400 FSA28 RS09375 WP 002263441.1     ..MLFMKKIKAGVLLYALFMSAIFSLFLQFYLNRVVATERQNQAQLSASKAYLIADLTQDLAKDNFGQLTFNHGRSSYHKQEGEL   83
NTDB id 391 SMU RS09005 WP 002263441.1       ..MLFMKKIKAGVLLYALFMSAIFSLFLQFYLNRVVATERQNQAQLSASKAYLIADLTQDLAKDNFGQLTFNHGRSSYHKQEGEL   83
consensus                                      *** ***!!!*!*!!**! !!!*! !!!!!*!*!!*************!***! *! !******!***!**!***!******!

logo
NTDB id 506 SM12261 RS00975 WP 000265657.1   VTTVRTP.KSQYEFIFPSVKIKEEKTDKK..EKVVTDSSNQAEKKKSEEKVEKKENF..   137
NTDB id 20326 SMA 0098 CCF01389.1            QVVVKLSNQQEFDYVF.........................................   101
NTDB id 400 FSA28 RS09375 WP 002263441.1     LIKVHLTDGQSYQYRFNCVTENKEKGKFSVANKTFTKSI.............EKRHFQK   129
NTDB id 391 SMU RS09005 WP 002263441.1       LIKVHLTDGQSYQYRFNCVTENKEKGKFSVANKTFTKSI.............EKRHFQK   129
consensus                                    ***!***********!**********************             *******
```

X non conserved

X similar

X $\geq$ 50% conserved