```
logo
NTDB id 40025 SGGBAA2069 RS00785 WP 009853235.1  MFTLLKRQVKAGILIYALLMAAIFALLLQFYLGRVVAMERQHQAQLSASQAYLMAELTRDLAKEDTGHLTFDQGVADYNF   80
NTDB id 400 FSA28 RS09375 WP 002263441.1         ..MLFMKKIKAGVLLYALFMSAIFSLFLQFYLNRVVATERQNQAQLSASKAYLIADLTQDLAKDNFGQLTFNHGRSSYHK   78
NTDB id 391 SMU RS09005 WP 002263441.1           ..MLFMKKIKAGVLLYALFMSAIFSLFLQFYLNRVVATERQNQAQLSASKAYLIADLTQDLAKDNFGQLTFNHGRSSYHK   78
consensus                                          *!*****!!!*!*!!!*!*!!!*!*!!!!!*!!!!*!!!*!!!!!!!!!*!!!*!*!!*!!!!***!*!!!**!***!**
```

```
logo
NTDB id 40025 SGGBAA2069 RS00785 WP 009853235.1  QDNQLQVVVKLPNQQEFDYVFQRA.........................   104
NTDB id 400 FSA28 RS09375 WP 002263441.1         QEGELLIKVHLTDGQSYQYRFNCVTENKEKGKFSVANKTFTKSIEKRHFQK   129
NTDB id 391 SMU RS09005 WP 002263441.1           QEGELLIKVHLTDGQSYQYRFNCVTENKEKGKFSVANKTFTKSIEKRHFQK   129
consensus                                        !***!***!*!***!***!*!******************************
```

X non conserved

X similar

X $\geq$ 50% conserved