```
logo

NTDB id 48891 SGGB RS00660 WP 009853235.1  MFTLLKRQVKAGILIYALLMAAIFALLLQFYLGRVVAMERQHQAQLSASQAYLMAELTRDLAKEDTGHLTFDQGVADYNFQDNQL   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 48891 SGGB RS00660 WP 009853235.1  QVVVKLPNQQEFDYVFQRA.........................   104
NTDB id 400 FSA28 RS09375 WP 002263441.1   LIKVHLTDGQSYQYRFNCVTENKEKGKFSVANKTFTKSIEKRHFQK   129
NTDB id 391 SMU RS09005 WP 002263441.1     LIKVHLTDGQSYQYRFNCVTENKEKGKFSVANKTFTKSIEKRHFQK   129
consensus                                  ***!*!***!***!*!******************************
```

X non conserved

X similar

X $\geq$ 50% conserved