```
logo                                                  MDDHAYTKELQPTYENLSKAVYTVNRHAKTAPNPKYLYLLKKRALQKLVKEGKGKKIGLHFSKNPRFSQQQSDVLISIGD

NTDB id 370358 BvL003 RS06890 WP 003154622.1          MDDHAYSKELQPTIENLSKAVYTVNRHAKTAPNPKYLYLLKKRALQKLVKEGKGKKIGLHFSNNPRFSQQQSDVLISIGD   80
NTDB id 88 BSU 14020 NP 389285.1                      MDDHAYTKDLQPTVENLSKAVYTVNRHAKTAPNPKYLYLLKKRALQKLVKEGKGKKIGLHFSKNPRFSQQQSDVLISIGD   80
consensus                                             !!!!!!*!*!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!

logo                                                  YYFHMPPTKEDFQHLPHLGTLNQSYRNPKAQMSLTKAKNYLQKYVGMKEKPLYPNRQQPAYHKPVFKKLGESYF
                                                                  E                                 AA
NTDB id 370358 BvL003 RS06890 WP 003154622.1          YYFHMPPTKEDFQHLPHLGTLNQSYRNPKAQMSLTKAKNVLQKYVGLKEKPLAANRQQPAYHKPVFKKLGESYF   154
NTDB id 88 BSU 14020 NP 389285.1                      YYFHMPPTKEDFEHLPHLGTLNQSYRNPKAQMSLTKAKHLLQEYVGMKEKPLVPNRQQPAYHKPVFKKLGESYF   154
consensus                                             !!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!**!! !!!*!!!!! *!!!!!!!!!!!!!!!!!!!!!
```

X non conserved

X similar

X $\geq$ 50% conserved