```
logo
NTDB id 276029 C6N19 RS03215 WP 016806074.1  MFNFLNFQHLIQLFSPCSLCELGMREKYSLCKDCWEQLPWLKQTIQRNNHSVLVACNYAYPVNRIIQQFKYEQKLHYQIL   80
NTDB id 1042 H0N27 RS01930 WP 000472273.1    MFKFLNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQTL   80
NTDB id 1071 ABD1 RS15375 WP 000472269.1     MFKFLNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQTL   80
consensus                                    !!*!!!*!*!*!!*!!!*!!!*!*!!!!!!!*!!!!!!!!!!!!!!!!**!!!!!!*!!!!*!!!!!!!!!!!!!!!!*!
```

```
logo
NTDB id 276029 C6N19 RS03215 WP 016806074.1  LQEILKQLKFPKVQAIVPMPISNQRLIERGFNQSLLLANILSRHLKIPVWQPIQRLNEHSQKGLTRLERFENIEQQFLPH   160
NTDB id 1042 H0N27 RS01930 WP 000472273.1    LAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFVAL   160
NTDB id 1071 ABD1 RS15375 WP 000472269.1     LAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFVAL   160
consensus                                    !*!*!*!!!!!!!!!!!!!!!!!*!!!*!!!!!!!!!!!!!*!***!!!!!!!!*!!!!!!!!!!!!*!!!!!!!!!!!!!***
```

```
logo
NTDB id 276029 C6N19 RS03215 WP 016806074.1  HQEKRRYRRVLIIDDVITTGSSVHALSQALKQLGCTSIHTVCLAATPKS..   209
NTDB id 1042 H0N27 RS01930 WP 000472273.1    TQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY   211
NTDB id 1071 ABD1 RS15375 WP 000472269.1     TQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY   211
consensus                                    *!!*!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!**!!!!!******
```

X non conserved
X similar
X $\geq$ 50% conserved