```
logo
NTDB id 176719 AMD27 RS14150 WP 067661680.1  MLKILKEMCQSSVQHLFPCQLCGIDHAQSHSLCADCWNQLPWLKARIVRHERSIQIALHYDFPIDRIIQKYKYEQDLHYQ   80
NTDB id 1042 HON27 RS01930 WP 000472273.1    MFKFLNP..QYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
NTDB id 1071 ABD1 RS15375 WP 000472269.1     MFKFLNP..QYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
consensus                                    !*!*!**  !***!*!*!!*!!*!******!!!**!!*!!!!!!**!*!***!***!*!!**!!*!!!!**!!!!*!!!!

logo
NTDB id 176719 AMD27 RS14150 WP 067661680.1  TLLAQSLLTLRFSKVQAIVPMPISNERLAERGYNQMLIIAKLVSKQLGIPVWQPVIRAAQHSQKGLSRVERMATIETQFQ  160
NTDB id 1042 HON27 RS01930 WP 000472273.1    TLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFV  158
NTDB id 1071 ABD1 RS15375 WP 000472269.1     TLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFV  158
consensus                                    !!!!**!**!*!*!!!!!!!!!!!!**!!*!!!*!!*!**!*!**!!!*!!!!!!!!*!***!!!!!!!!*!!***!!*!!*

logo
NTDB id 176719 AMD27 RS14150 WP 067661680.1  IIRAERRKFKKVLMIDDVVTTGSSMHALAQALEKLGCQQIEMACIAAAEQ...  210
NTDB id 1042 HON27 RS01930 WP 000472273.1    ALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1     ALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
consensus                                    ****!*!****!!*!!!!*!!!!!*!!!!*!!!**!!!**!***!*!!!*****
```

X non conserved

X similar

X $\geq$ 50% conserved