```
logo

NTDB id 1064 ABD1 RS15725 WP 001046490.1      MNKYYILRFNQGFTLIELMVVIVIVAIFASIAIPSYQSYSRRATASAAKSEILKLAEQLEQHKSRNFTYRGFTTTSITLP    80
NTDB id 403812 GK020 RS13915 WP 156189562.1   .......MKKNGFTLIELMVVVVIVAIFVAIAIPSYQAYIRRAETSKAQQELKRIATLLERHKARNFSYKGFDFSAQTL.    72
NTDB id 1006 ACIAD RS15005 WP 004923843.1     .....MQHHRQGFTLIELMVVVVITAIFAAIAIPSYQEYIRRSNLAMAQQEMQKLAEQLERYRSRNFSYKGFDATYLYQD    75
consensus                                          *    **!!!!!!!!!!*!!*!!!**!!!!!! !*!!*  * !**!* **!**!!****!!!*!*!!* * ***

logo

NTDB id 1064 ABD1 RS15725 WP 001046490.1      R......................GGYTIQITDDTTTGNLLTNTAANGQTWVIKATTTDSRNFNFIAKNSGLRCQSLTAT   137
NTDB id 403812 GK020 RS13915 WP 156189562.1   ......................STPRTYSFELKDGTHTTKLLNATDAAGHAWVLKATTTDAKNYNYLITSTGIRCKNLTAE   131
NTDB id 1006 ACIAD RS15005 WP 004923843.1     KSGQTLAKFDSTKQIVQLPLDTSNTKYTLAVVDTTTNNPLLTSANAIGQGWAIKAMSADNNNYSLLLTNTGFSCKNKTWA   155
consensus                                                          **    !** * ! !**  !!* * ! !**!**!!***! *!*****!***!**!***!*

logo

NTDB id 1064 ABD1 RS15725 WP 001046490.1      AVDNDCGGTVTCNVCETNSENWQ   160
NTDB id 403812 GK020 RS13915 WP 156189562.1   NI........SYTGCGTGGEAW.   145
NTDB id 1006 ACIAD RS15005 WP 004923843.1     NI........SSSSCGTGSESW.   169
consensus                                     **         * * !*!**! !
```

X non conserved

X similar

X $\geq$ 50% conserved