```
logo
NTDB id 1067 ABD1 RS15745 WP 002194578.1        MLMHHYKQKAQAGVGLLEVLVALILLAIGVLGYVALQLRAMDASSEALSKSQAILVMRGLAENIRTNSTQASQYPTFVRS   80
NTDB id 313076 AhaeAN4 RS01575 WP 004641261.1   .....MKYNNQKGVGLVEVLVALFLLAVAVLGFAALQLRAVDASLEASKQVLALTIARDLAERMRANPQAVRDGAYVVGA   75
NTDB id 1003 ACIAD RS15025 WP 004923832.1       ....MSKMRAQRGVGMVEVLVALLLLAIGVLGFTALQLRAVSASMEASNQVLALNLARDLTERMRANPQATQCKLYHTEG   76
consensus                                             ! **!*!!!**!!!!!!*!!!**!!!* !!!!!!**!!*!!* ***!* **!*!*!**!*!***      * * *

logo
NTDB id 1067 ABD1 RS15745 WP 002194578.1        YSNYTSDT.PAPTSCFNSLC.TASQLAQFDAYQAARNANQLGMRITMSNCPGVTNTM.VQQRQCLFVFWGKTAPVITTNG  157
NTDB id 313076 AhaeAN4 RS01575 WP 004641261.1   VSVPTSEVVATV....G....DAASIAKQDLNIAALNAKNNGMNLSISECPTG......AARQCIYVAWEETSLSGNRG.  140
NTDB id 1003 ACIAD RS15025 WP 004923832.1       ALAKDTKAYTKASNCSGSVSTQATAQANSDLLFIYNQANTLGMAVIINTCPNATNAGNNLGRQCVYIAWDKTAQNF.TN.  154
consensus                                        *  ***     * * ***   !* *!* !*  ** *!**!! **** !! ***     *!!!****!**!*    **

logo
NTDB id 1067 ABD1 RS15745 WP 002194578.1        TNTSVDVSSCMSN.NGVYVNNSTCLMMEAY  186
NTDB id 313076 AhaeAN4 RS01575 WP 004641261.1   ......IETCIKDG..VYVKGSHCLFLEAY  162
NTDB id 1003 ACIAD RS15025 WP 004923832.1       ......SDTCYKSGGSSYERNTTCLFLEAY  178
consensus                                             ***!** *  *!*****!!**!!!
```

X non conserved

X similar

X ≥ 50% conserved