```
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1103 NMB RS07590 WP 002212976.1         ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1166 A1552VC RS00795 WP 000168289.1     MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 1089519 ACE2AB RS14965 WP 004738307.1  ..MRGINKVILVGSLGANPITKHYPNGNTYVQFSIATSEKYQDKNTGEWIENTEWHRIIAYGRLGEVATQILKKGSKVYV   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1       ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                           **!!!!**!  !! *!*****!*!*** ****!!!!***!***!***!*!!!!***********!*!***!*!!* !!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        EGRIQSRKYQGKDGIERTAYDIVANEM....KMLGGRNENSGGAPYDEGYGQS.........QEAYQRPAQQSRQPAPDA   143
NTDB id 1103 NMB RS07590 WP 002212976.1         EGRIQSRKYQGKDGIERTAYDIVANEM....KMLGGRNENSGGAPYEEGYGQS.........QEAYQRPAQQSRQPASDA   143
NTDB id 1166 A1552VC RS00795 WP 000168289.1     EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQG.GMPAQGG...MNVPAQ....QGSWGQPQQPAKQ....H   148
NTDB id 1089519 ACE2AB RS14965 WP 004738307.1  EGSLRTRQVTDQRGQQGYITEVRANTF....QSLDSLPQAN.PY.......................................   117
NTDB id 1390 A4U84 RS00055 WP 021115958.1       EGKLKTRKWQDQNGQDRYTTEIQGDVL....QMLDSRSSGG.DFGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNN....F   149
consensus                                       !!****!***** !**** ****** *     **!***  *  * *   * *           * **  * ** **

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1         PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1     Q..PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1089519 ACE2AB RS14965 WP 004738307.1  ................................   117
NTDB id 1390 A4U84 RS00055 WP 021115958.1       NGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                          *   **   *    *  **** ******
```

X non conserved

X similar

X $\geq$ 50% conserved