```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 993560 AAHN93 RS12875 WP 124855709.1  ..MASVNKVILVGNLGRDPELRMFPSGGQVANVSVATTEQWRDKQTGERRELTEWHNVVFNDRLAEIAAQYLRKGSSVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                         **!!!!**!*!!*!!**!**!*!**!******!!*!*!*!* *!***!*!!!!*********!!*!**!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQEAYQRP..............AQQSRQ   138
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQEAYQRP..............AQQSRQ   138
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGMNV..PAQQGSWG................Q   140
NTDB id 993560 AAHN93 RS12875 WP 124855709.1  EGRIRTRKWQDQSGQDRYTTEIRA....DVMQMLGGRGSGSS.DGRDGGY......DGGSYSSGG..........YGGGA   137
NTDB id 1390 A4U84 RS00055 WP 021115958.1     EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGGN   153
consensus                                     !!****!!*!** !**!*******     ***!!**!  ***  * * !**  * * *  *              *  *

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        PASDAPSHPQEAPAAPR..RQPVPAAAPVEDI....DDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       PAPDAPSHPQEAPAAPR..RQPVPAAAPVEDI....DDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1    PQQPAKQHQPMQQSAPQQYSQPQYNEPPMD.F....DDDIPF   177
NTDB id 993560 AAHN93 RS12875 WP 124855709.1  PARQQPERPPQA..AP.AAQRPAAPPPPAQSFDQFEDDDIPF   176
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ATRPQPAQKPA...........AQAEPPMDNF....DDDIPF   180
consensus                                     **  ** *** *  **    **  * *!** *    !!!!!!
```

X non conserved

X similar

X $\geq$ 50% conserved