```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 422983 PmVP161 RS06470 WP 005725039.1 ..MAGVNKVIIVGNLGNDPEIRTMPNGEAVANISVATSESWIDKNTNERREVTEWHRIVFYRRQAEVAGEYLRKGSKVYV   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                          **!!!!**!*!!*!!**!*!!*!*!!*****!!!!*!*!*******!*!!!!*********!!*!!*!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSG..GAPYE......EGYGQSQEAYQRPAQQSRQPASDAP   144
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSG..GAPYD......EGYGQSQEAYQRPAQQSRQPAPDAP   144
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQG...GMPAQGGMNVPAQ....QGSWGQPQQPAKQH.....   148
NTDB id 422983 PmVP161 RS06470 WP 005725039.1 EGRLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRNERQQTGGYAPQ...TAAPQYNAPTGGY................   136
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNG...   151
consensus                                      !!****!!*!** !**!******      ***!!**!***    ! *          **   ****  * ** **

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        SHPQEAPAAPRRQPVPAAAPVE...DIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       SHPQEAPAAPRRQPVPAAAPVE...DIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1    .QPMQQSAPQQYSQPQYNEPPMD...FDDDIPF   177
NTDB id 422983 PmVP161 RS06470 WP 005725039.1 ..GAQPSRPATK.PAPQNEPPMDMGFEEDNIPF   166
NTDB id 1390 A4U84 RS00055 WP 021115958.1      .GNATRPQPAQK.PAAQAEPPMDN..FDDDIPF   180
consensus                                        *   ***  * * * **!***    *!*!!!
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X non conserved

X similar

X $\geq$ 50% conserved