```
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 40441 ALIDE2 RS22940 WP 013723264.1 ..MASVNKVIIVGNLGRDPEMRTFPSGDQVANVRIATTDRWRDKNTGENKEATEWHSVVFNGRLAEIVGQYLRKGSQVYV    78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV    78
consensus                                     **!!!!**!*!!*!!**!**!*!**!**!***!!***!*!***!***!*!!!! *******!!**!*!!*!!* !!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSG.....GAP..........YDEGYGQSQEAYQRPAQQSR   137
NTDB id 1103 NMB RS07590 WP 002212976.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSG.....GAP..........YEEGYGQSQEAYQRPAQQSR   137
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGG..MN.VPA....QQGS.......WGQPQQ...   143
NTDB id 40441 ALIDE2 RS22940 WP 013723264.1 EGSLRTRKWTDQSGQERYTTEIRA....DTMQMLGSRAGSG...GGQGGGGY.........DDGGYGGDGGGYEAPRR..A   140
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGGYSDN...YAQNNNFNG   151
consensus                                   !!*****!!**** !**!*******      **!!**!  ***  *  !*            *****    **  * *

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     QPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1     QPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 .PAKQ.HQPMQQSAPQQYSQPQYNE.PPMDFDDDIPF   177
NTDB id 40441 ALIDE2 RS22940 WP 013723264.1 APAPRPAPPAARPAPAPVAQPPRAASGFDDMDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1   GNATRPQPA...QKPAA.....QAEPPMDNFDDDIPF   180
consensus                                    *!        *   ***  **  ** *****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved