```
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1103 NMB RS07590 WP 002212976.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 73857 PAFU01 RS19935 WP 019105608.1 MSARGVNKVILVGNLGQDPEVRYMPNGTAVTSLSLATSETWRDKQTGENREKTEWHRVVLFGKLAEIAGEYLRKGSQVYI   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                   **!!!!**!*!!*!!**!*!!*!*!! ****!!!!*!*!* *!***!*!!!!********!!*!!*!!*!!* !!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGG........APYDEGYGQSQEAYQRPAQQSRQP.....   139
NTDB id 1103 NMB RS07590 WP 002212976.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGG........APYEEGYGQSQEAYQRPAQQSRQP.....   139
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDF.......GGNQGSGWNQAPAQTNY..NQGGYSDNYAQ   145
NTDB id 73857 PAFU01 RS19935 WP 019105608.1 EGKLRTREWQDSEGTKRYTTEIVV.DVNGTMQMLGGAKSSTN..TSGNNKPSGQQSGWGQPQQPQPS..Q..........   145
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVPAQQGSWGQPQQPAKQ..H..........   148
consensus                                   !!****!**!* *! *!*******      **!!***  ***         * *  ***! *   *   **

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     APDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF................   174
NTDB id 1103 NMB RS07590 WP 002212976.1     ASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF................   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1   NNNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF................   180
NTDB id 73857 PAFU01 RS19935 WP 019105608.1 ......A......QQTAPSNPANEPPMDFDDDIPFLGHGYGVCRKAIYAIS   184
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ......QPMQQSAPQQYSQPQYNEPPMDFDDDIPF................   177
consensus                                          *   **   *   **  !****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved