```
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 1107071 ACLKM1 RS00765 WP 005095120.1 ..MRGVNKVILVGTLGKDPETKTFANGGSLTQFSIATSDSWTDKSTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                        **!!!!**! !!*!!*******!*** ****!!!**!*!* *!***!*!!!!**********!*!**!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYE....................EGYGQSQ.EAYQRP   132
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYD....................EGYGQSQ.EAYQRP   132
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGM.PAQGGMNVP...................AQQGSWGQP   141
NTDB id 1107071 ACLKM1 RS00765 WP 005095120.1 EGSLRTRQWTDQNGQERYTTEIRG....DQMQMLDSNRSQGEQ.GDN.SFNQPRFNNNNQASNNQSGGYGNNPQSGYGSA   152
NTDB id 1390 A4U84 RS00055 WP 021115958.1     EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDF.GGN........................QGSGWNQA   128
consensus                                     !!****!***** !**!*******       **!!***  ***  * *                     **** *  *** *

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        AQQSRQPA.........SDAPSHPQEAPAAPRRQ.PVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       AQQSRQPA.........PDAPSHPQEAPAAPRRQ.PVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1    QQPAKQ................HQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1107071 ACLKM1 RS00765 WP 005095120.1 PQQGGFN.....NNQG..GGYGNNNPSGFAPKSAPA.PTAAPAADLDDDLPF   196
NTDB id 1390 A4U84 RS00055 WP 021115958.1     PAQTNYNQGGYSDNYAQNNNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                      *****             *  * *    *!*     ****!****!!!*!!
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X non conserved

X similar

X $\geq$ 50% conserved