```
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 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 1100643 ACLIX2 RS08255 WP 416630190.1 MANGSVNKVILIGNLGRDPEIRYLPSGGAVANLAVATSEKWRDKQTGENREKTEWHRVVLFGKLADIASGYLCKGSQVYI   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                         **!!!!**!*!!*!!**!**!*!*!!*****!!!!*!*!* *!***!*!!!!********!**!**!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYEEGYGQS.........QEAYQRPAQQSRQPASDA   143
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYDEGYGQS.........QEAYQRPAQQSRQPAPDA   143
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSGGDFG.GNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNG..   151
NTDB id 1100643 ACLIX2 RS08255 WP 416630190.1 EGQLQTREWDDNSV.KRYTTEIVVK.AGGSMQMLGGTIKSAGSH.PAQL.............................   126
NTDB id 1166 A1552VC RS00795 WP 000168289.1   EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMP.AQGG...MNVPAQ....QGSWGQPQQPAKQHQP..   150
consensus                                     !!****!***** * *!******    *  **!!***  *** *    * *           * **  * ** **

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..GNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1100643 ACLIX2 RS08255 WP 416630190.1 .....S..KPPDQ..PHDNQSPMDFD.NIPF   147
NTDB id 1166 A1552VC RS00795 WP 000168289.1   ..MQQSAPQQYSQ..PQYNEPPMDFDDDIPF   177
consensus                                         ****  * *    *  *****!**!!!
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X non conserved

X similar

X $\geq$ 50% conserved