```
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 85822 I0P84 RS14465 WP 214001696.1  ..MRGVNKVILVGTLGRDPETKTFPNGGSLTQFSIATSDSWTDKNTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                       **!!!!**! !!*!!*****!*!*** ****!!!**!*!***!***!*!!!!**********!*!**!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQE..................AYQRPAQQ  135
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQE..................AYQRPAQQ  135
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQG..GMNV......................PAQQ  135
NTDB id 85822 I0P84 RS14465 WP 214001696.1 EGSLRTRQWTDQSGQERYTTEIRG....EQMQMLDSGRPQGE.QGDN...GFNQPRFNNNNQGGGYSNNPQGGYGSAPQQ  150
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSDN..........  142
consensus                                   !!*****!***** !**!*******      **!!***  ***  * * * ! **                   **  ****

logo
NTDB id 1103 NMB RS07590 WP 002212976.1      SRQPASD.........APSH.PQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     SRQPAPD.........APSH.PQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 GSWGQPQQP.......AKQHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 85822 I0P84 RS14465 WP 214001696.1 GGFNNNQGSGYGNNNQNNQGGYAPKS....AP.APVAAPAADLDDDLPF   194
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..........YAQN.NNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                                * ** *   **    *   ****!****!!!*!!
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X non conserved

X similar

X $\geq$ 50% conserved