```
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 1006858 ABLB98 RS13280 WP 005156773.1  ..MRGVNKVILVGTLGKDPETKTFPNGGSLTQFSIATSESWTDKNTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                          **!!!!**!  !!*!!*****!*!*** ****!!!!*!*!***!***!*!!!!*********!*!**!!*!!* !!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGA.....................PYDEGYGQSQEAYQR   131
NTDB id 1103 NMB RS07590 WP 002212976.1         EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGA.....................PYEEGYGQSQEAYQR   131
NTDB id 1166 A1552VC RS00795 WP 000168289.1     EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQ..GGMNVPAQQG..................SWGQ   140
NTDB id 1006858 ABLB98 RS13280 WP 005156773.1  EGSLRTRQWTDQNGQERYSTEIRG....DQMQMLDTRQQSEQGGGD....FNQPRFNNNNQGGGYQNTGYNNNQNGYGQ   150
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQG......GYS...DNYAQ   145
consensus                                      !!****!***** !**!*******       **!!**!* * ****       *     *             ** * ******

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        PAQQSRQPAPDAPSHPQEAPAAPRRQPV.PAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1         PAQQSRQPASDAPSHPQEAPAAPRRQPV.PAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1     PQQPAKQHQ......PM.QQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1006858 ABLB98 RS13280 WP 005156773.1  GGGFAGGNQGNYAGSPQAGNGFNTPKSAPQPATAPADLDDDLPF   194
NTDB id 1390 A4U84 RS00055 WP 021115958.1      NNNFNGGNATR....PQP.....AQKPAAQAEPPMDNFDDDIPF   180
consensus                                      ***  ** *      !* * ***   *  *** *****!!!*!!
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X non conserved

X similar

X $\geq$ 50% conserved