```
logo
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 35142 CAP2UW1 RS00635 WP 012806939.1 ..MASVNKVILVGNLGADPETRYSANGDAICNIRMATTDRFRDKASGELKDSTEWHRVVFFGKLAETAGQYLKKGRQIYV   78
NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
consensus                                       **!!!!**!*!! !!**!****!*!******!!*****!* *!*****!!!!********!!*!!*!!*!!   *!*

logo
NTDB id 1166 A1552VC RS00795 WP 000168289.1   EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGM..NVPAQQGSWG.........QPQQPAKQHQ   149
NTDB id 1390 A4U84 RS00055 WP 021115958.1     EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGGYSDNYAQ....NNNFN   150
NTDB id 35142 CAP2UW1 RS00635 WP 012806939.1 EGRIRTNKWQDKDGNERYTTEIIA....NEMKMLGSREGMGSPA..........SSESEYG..GSMPSSAA.........   133
NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGA..........PYEEGYG..QSQEAYQRPAQQSRQPA   140
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGA..........PYDEGYG..QSQEAYQRPAQQSRQPA   140
consensus                                    !!*****!*!***!**!*******     ****!!**!  *** *          * *  **   * * *  * *  **

logo
NTDB id 1166 A1552VC RS00795 WP 000168289.1   ......PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1     GGNATR....PQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 35142 CAP2UW1 RS00635 WP 012806939.1 ........PASGAARSAPAKKTPSFDDMDDDIPF   159
NTDB id 1103 NMB RS07590 WP 002212976.1      SDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     PDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
consensus                                       * *  **** ** *  *  *****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved