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logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY   75
NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY   75
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY   77
NTDB id 32312 MCHL RS12205 WP 012253693.1   .MAGSVNKVILVGNLGRDPETRRLSSGDPVVNLRIATSESWKDKASGERKEKTEWHSVVIYNDNLAKVAEQYLRKGSKVY   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY   79
consensus                                     **!!!!**!*!!*!!**!****!**! ****!!!!*!*!* *!***!*!!!! ***** **!**!**!!*!!* !!

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEG..YG...........QSQEAYQRPAQQS..   136
NTDB id 1103 NMB RS07590 WP 002212976.1      LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEG..YG...........QSQEAYQRPAQQS..   136
NTDB id 1390 A4U84 RS00055 WP 021115958.1    VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD..F................GGNQGSGWNQAPAQTNYN   135
NTDB id 32312 MCHL RS12205 WP 012253693.1   IEGQLQTRKWTDQSGAEKYTTEVVLQRFRGEMTILDGRGGGGGGDFAGEDEGGGQISRGGDRGGDRGGDRGGR.......   152
NTDB id 1166 A1552VC RS00795 WP 000168289.1 IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVP...........AQQGSWGQP.......   141
consensus                                   *!!****!!**** ! *********      ** *!**!  ****    *    *           *  *  *****

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     .....RQPAP..DAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1      .....RQPAS..DAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    QGGYSDNYAQNNNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 32312 MCHL RS12205 WP 012253693.1   .....DDYGSSRSGGGGGG...GDRRPSSGGGAKPNYDLDDDIPF   189
NTDB id 1166 A1552VC RS00795 WP 000168289.1 .....QQPAK.QHQPMQQS...APQQYSQPQYNEPPMDFDDDIPF   177
consensus                                     ***    **      **  **    ** !****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved