```
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY    77
NTDB id 225359 B5M07 RS10190 WP 120351221.1 .MAGSVNKVILIGNLGRDPEVRSFQNGGKVCNLRIATSETWKDRNTGERREKTEWHSVAIFQEGLVRIAEQYLKKGSKVY    79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY    79
consensus                                       **!!!!**!*!!*!**!****!**!*****!!!!*!*!***!***!*!!!! ***** *****!**!!*!!* !!

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     LEGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGA.....PY....DEGYGQSQE..AYQR.........   131
NTDB id 1103 NMB RS07590 WP 002212976.1      LEGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGA.....PY....EEGYGQSQE..AYQR.........   131
NTDB id 1390 A4U84 RS00055 WP 021115958.1    VEGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSGGDFGG........NQGSGWNQAPAQTNYNQGGYSDNYAQ   145
NTDB id 225359 B5M07 RS10190 WP 120351221.1 IEGQLQTRKWQDQSGADRYSTEVVLQGFGGTLTMLDGPGGAGSGGGGGGGGYGGGGGGGYGGGG................   143
NTDB id 1166 A1552VC RS00795 WP 000168289.1 IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVP.AQQGSWGQPQ................   142
consensus                                   *!!****!!*!** ! *!******    *  * !!***  ******     *       **** *    **

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     ..PAQQSRQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1      ..PAQQSRQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    NNNFNGGNATRP....QPAQ......KPAAQAEPPMDNFDDDIPF   180
NTDB id 225359 B5M07 RS10190 WP 120351221.1 .NDYGGGYDSGPSSS.GGGS......GGGGGGGGGSRDLDDEIPF   180
NTDB id 1166 A1552VC RS00795 WP 000168289.1 .QP...AKQHQPMQQSAPQQ......YSQPQYNEPPMDFDDDIPF   177
consensus                                    ** * *** **   * **        *    ********!!*!!!
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X non conserved

X similar

X ≥ 50% conserved