```
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 1086294 ACLCD0 RS14315 WP 394408950.1  .MAGSVNKVILIGNLGRDPEVRSFQNGGKVCNLRIATSETWKDRNTGERREKTEWHSVAIFQEGLVRVAEQYLKKGSKVY   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY   79
consensus                                          **!!!!**!*!!*!!**!****!**!*****!!!!*!*!***!***!*!!!! ***** *****!**!!*!!* !!

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        LEGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYDEG........YGQSQEAYQRPAQQ.SRQPAPD   142
NTDB id 1103 NMB RS07590 WP 002212976.1         LEGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYEEG........YGQSQEAYQRPAQQ.SRQPASD   142
NTDB id 1390 A4U84 RS00055 WP 021115958.1       VEGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSGGDFG.......GNQGSGWNQAPAQTNYNQGGYSDNYAQN   146
NTDB id 1086294 ACLCD0 RS14315 WP 394408950.1  IEGQLQTRKWQDQSGADRYSTEVVLQGFGGTLTMLDGPGGGSGGGGGGGGGYGGGGGGGYGGGGNDYG.............   146
NTDB id 1166 A1552VC RS00795 WP 000168289.1    IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVPAQQGSWGQPQQPAK...........QH   148
consensus                                      *!!****!!*!** ! *!******    *  * !!***  *****    *        *** *  **      * * *

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        APSH.PQEAPAAPR.RQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1         APSH.PQEAPAAPR.RQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1       NNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1086294 ACLCD0 RS14315 WP 394408950.1  ........YDSGPS.SSGGGGGGGSRDLDDEIPF   171
NTDB id 1166 A1552VC RS00795 WP 000168289.1    QP.....MQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                       *    *    *!  *    ********!!*!!!
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X non conserved

X similar

X ≥ 50% conserved