```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY   75
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY   75
NTDB id 1161613 ACKPCT RS06155 WP 090073711.1  .MAGSVNKVILVGNLGADPDIRRTQDGRPICNLSVATSESWKDRNSGERRERTEWHRVVIFNEGLCRIAEQYLRKGSKVY   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY   79
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY   77
consensus                                          **!!!!**!*!! !!**!****!*******!!!!*!*!**!***!*!!!!*****  *****!**!!*!!* !!

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQEAYQRPAQQS.........RQPASD   142
NTDB id 1131 NGFG RS05740 WP 003695064.1       LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQEAYQRPAQQS.........RQPAPD   142
NTDB id 1161613 ACKPCT RS06155 WP 090073711.1  LEGQLQTRKWQDQSGQDRYSTEVVLQGFNGNLTMLDNRGEGGGGGGFG.GGQSGGGDFGGPGG.....GGYGGGN..PGG   151
NTDB id 1166 A1552VC RS00795 WP 000168289.1    IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQG.GM......NVPAQ....QGSWGQPQQPAKQ   147
NTDB id 1390 A4U84 RS00055 WP 021115958.1      VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQ.GSG....WNQAPAQTNYNQGGYSDNYAQNNN   148
consensus                                      *!!****!!*!** !**!******       * !!**! ********  ! *     ** !**     * *    ***

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        APSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       APSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1161613 ACKPCT RS06155 WP 090073711.1  FGGGG.QGPSGGGQP...SGGFRDEMDDDIPF   179
NTDB id 1166 A1552VC RS00795 WP 000168289.1    HQ..PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1      FNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                          *  ****  *    ********!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved