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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 194845 Ga0102493 RS13050 WP 034903062.1  .MAGSLNKVMLIGNLGADPEVRTFQNGGKVCNLRVATSESWKDRNTGERQERTEWHSVAIFNEGLAGVAERYLKKGSKVY    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....NEMKMLGGRNENSGGAP.YEEGYGQS....................QEAYQR   131
NTDB id 1131 NGFG RS05740 WP 003695064.1         LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAP.YDEGYGQS....................QEAYQR   131
NTDB id 194845 Ga0102493 RS13050 WP 034903062.1  LEGQLQTRKWQDQSGNDRYSTEVVLRGPNSVMTMLDGAPGAGGGGGGRSGGFGGGGAGYGDRGAGGQPAGAGGAGGGWNQ   159
NTDB id 1166 A1552VC RS00795 WP 000168289.1      IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGM...............NVPAQ....QGSWGQ   140
NTDB id 1390 A4U84 RS00055 WP 021115958.1        VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWN............QAPAQTNYNQGGYSD   141
consensus                                        *!!****!!*!** !**!*******     ** !!***  ******    ***             **     *****

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NTDB id 1103 NMB RS07590 WP 002212976.1          PAQQSRQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1         PAQQSRQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 194845 Ga0102493 RS13050 WP 034903062.1  GGGGS....................GGGSGGGNYDDLDDDIPF   182
NTDB id 1166 A1552VC RS00795 WP 000168289.1      PQQPAKQ....HQ..PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1        NYAQNNN....FNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                        ******        *   **   *    ********!!!!!!
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X non conserved

X similar

X ≥ 50% conserved