logo

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NTDB id 1103 NMB RS07590 WP 002212976.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1131 NGFG RS05740 WP 003695064.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 42315 SLG RS18240 WP 014077997.1      .MAGSVNKVILVGNLGADPEVKSFSNGGRICNIRIATSDNWTDRQSGERRERTEWHNVVIRSDGLIGIAEKYLRKGKKVY    79
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY    77
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY    79
consensus                                         **!!!!**!*!! !!*******!********!!!**!*!* *!***!*!!!!****** *****!**!!*!!  !!
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logo

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NTDB id 1103 NMB RS07590 WP 002212976.1       LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQS.........QEAYQRPAQQSRQPASD   142
NTDB id 1131 NGFG RS05740 WP 003695064.1      LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQS.........QEAYQRPAQQSRQPAPD   142
NTDB id 42315 SLG RS18240 WP 014077997.1      IEGSLQTRKWQDQNGQDRYTTEIVLSGPGAVMTMLDGGPGGGSGEG.GGGGGGRR........NEGWGAGGDD...MGGG   147
NTDB id 1390 A4U84 RS00055 WP 021115958.1     VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGG.NQGSGWNQAPAQTNYNQGGYSDNY.A...QNNN   148
NTDB id 1166 A1552VC RS00795 WP 000168289.1   IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPA.QGGM...NVPAQ....QGSWGQPQ.Q...PAKQ   147
consensus                                     *!!****!!*!** !**!*******     ** !!***  ******   ! *           ****  ** *   **
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logo

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NTDB id 1103 NMB RS07590 WP 002212976.1       APSHPQEAPAAPRRQPVP.AAAPVEDID.DDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1      APSHPQEAPAAPRRQPVP.AAAPVEDID.DDIPF   174
NTDB id 42315 SLG RS18240 WP 014077997.1      FGGGAGA.RSSGGARPASGASPFNDGFDDDEIPF   180
NTDB id 1390 A4U84 RS00055 WP 021115958.1     FNGGNAT.RPQPAQKPAAQAEPPMDNFD.DDIPF   180
NTDB id 1166 A1552VC RS00795 WP 000168289.1   HQ..PMQ.QSAPQQYSQPQYNEPPMDFD.DDIPF   177
consensus                                         *     **   * * *  *****! !*!!!
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X non conserved

X similar

X ≥ 50% conserved