logo



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NTDB id 26098 SZO RS05495 WP 012515548.1   ..MIKLLLVEDDLSLSNSIF.DFLDDFADVTQVFDGDEGLYEAESGIYDLILLDLMLPEKDGFQVLKALREKDIKIPVLIMTAKE   82
NTDB id 362 SMU RS05200 WP 002262214.1     ..MIKLLLVEDDLSLSNSIF.DFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE   82
NTDB id 258 KZH43 RS03475 WP 000590640.1   ..MIKILLVEDDLGLSNSVF.DFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE   82
NTDB id 217 SPD RS03775 WP 000590640.1     ..MIKILLVEDDLGLSNSVF.DFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE   82
NTDB id 183 SPR RS03560 WP 000590640.1     ..MIKILLVEDDLGLSNSVF.DFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE   82
NTDB id 147 SP RS03905 WP 000590640.1      ..MIKILLVEDDLGLSNSVF.DFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE   82
NTDB id 606 V4T04 RS01910 WP 003130756.1   MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKE.KDTPIIMMTARD   84
NTDB id 471 HSISS4 RS01430 WP 002883757.1  .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQE.KDTYIMMMTARD   83
consensus                                    ****!**!!** !***** **********!*****!**!!**!*****!!!!!!!!!!!**!!*!***!**********!!!**
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logo



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NTDB id 26098 SZO RS05495 WP 012515548.1   SLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRVGKF.....NDKSLKFGNLTVDTTRKEVQVNGQAIELLGKELDLLIYIL   162
NTDB id 362 SMU RS05200 WP 002262214.1     GLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKF.....NENNLTYGELTVDTATNTTMVNGKEVELLGKEFDLLVYFL   162
NTDB id 258 KZH43 RS03475 WP 000590640.1   SLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKF.....NENTLTYGNIVVNLSTNTVKVEDTPVELLGKEFDLLVYFL   162
NTDB id 217 SPD RS03775 WP 000590640.1     SLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKF.....NENTLTYGNIVVNLSTNTVKVEDTPVELLGKEFDLLVYFL   162
NTDB id 183 SPR RS03560 WP 000590640.1     SLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKF.....NENTLTYGNIVVNLSTNTVKVEDTPVELLGKEFDLLVYFL   162
NTDB id 147 SP RS03905 WP 000590640.1      SLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKF.....NENTLTYGNIVVNLSTNTVKVEDTPVELLGKEFDLLVYFL   162
NTDB id 606 V4T04 RS01910 WP 003130756.1   STMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAV.ERAENTSFRDLVIDKTNRTVHRGKKVIDLTRREYDLLLTLM   168
NTDB id 471 HSISS4 RS01430 WP 002883757.1  SIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAAKNAPAKASTYRDLKLDVQNRTVVRGDEAIPLTKREFDLLNTLL   168
consensus                                  ***!***!***!!!!!**!!!**!!!**!**!***!****     ************ ****** * *  **!***!*!!!****
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logo



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NTDB id 26098 SZO RS05495 WP 012515548.1   QNQNVILPKSQIFDRIWGFDSDTTISVVEVYVSKIRKKLKGTDFVNHLQTLRSVGYILKANE   224
NTDB id 362 SMU RS05200 WP 002262214.1     QNQNVILPKSQIFDRIWGFDSDTTISVVEVYVSKIRKKLKDTTFGKNLQTLRSVGYILKDAN   224
NTDB id 258 KZH43 RS03475 WP 000590640.1   QNQNVILPKTQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGTTFAENLQTLRSVGYLLKDVQ   224
NTDB id 217 SPD RS03775 WP 000590640.1     QNQNVILPKTQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGTTFAENLQTLRSVGYLLKDVQ   224
NTDB id 183 SPR RS03560 WP 000590640.1     QNQNVILPKTQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGTTFAENLQTLRSVGYLLKDVQ   224
NTDB id 147 SP RS03905 WP 000590640.1      QNQNVILPKTQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGTTFAENLQTLRSVGYLLKDVQ   224
NTDB id 606 V4T04 RS01910 WP 003130756.1   QNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKIDVEGQDSYIQTVRGLGYVMRERK   230
NTDB id 471 HSISS4 RS01430 WP 002883757.1  SNMNQVMTREELLLQVWKYDDAIETNVVDVYIRYLRGKIDVPGKESYIQTVRGMGYVIREK.   229
consensus                                  *!**************!**********!!*!!****!*!*******  **!!*!**!!**** *
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X non conserved

X similar

X ≥ 50% conserved