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logo

NTDB id 606 V4T04 RS01910 WP 003130756.1      MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKE.KDTPIIM   79
NTDB id 1139392 DQM69 RS04270 WP 002961524.1  ..MIKILLVEDDLSLSNSVYDFLDDFA.EVTQVFDGEEGLFEAETGVYDLILLDLMLPEKDGFTVLKELRDKGVATPVLI   77
NTDB id 362 SMU RS05200 WP 002262214.1        ..MIKLLLVEDDLSLSNSIFDFLDDFA.DVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLI   77
NTDB id 258 KZH43 RS03475 WP 000590640.1      ..MIKILLVEDDLGLSNSVFDFLDDFA.DVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLI   77
NTDB id 217 SPD RS03775 WP 000590640.1        ..MIKILLVEDDLGLSNSVFDFLDDFA.DVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLI   77
NTDB id 183 SPR RS03560 WP 000590640.1        ..MIKILLVEDDLGLSNSVFDFLDDFA.DVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLI   77
NTDB id 147 SP RS03905 WP 000590640.1         ..MIKILLVEDDLGLSNSVFDFLDDFA.DVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLI   77
consensus                                      **!*!**!!***!*******!**** *******!**!!*!!****!!!!!!!!!!!!**!!*!***!!*****!!***

logo

NTDB id 606 V4T04 RS01910 WP 003130756.1      MTARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAVERAENTSFRDLVIDKTNRTVHRGKKVIDLTRR  159
NTDB id 1139392 DQM69 RS04270 WP 002961524.1  TTAKEALDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKI....NDKLLTYGDLECDTSTNTVTVNGKIVELLGK  153
NTDB id 362 SMU RS05200 WP 002262214.1        MTAKEGLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKF....NENNLTYGELTVDTATNTTMVNGKEVELLGK  153
NTDB id 258 KZH43 RS03475 WP 000590640.1      MTAKESLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKF....NENTLTYGNIVVNLSTNTVKVEDTPVELLGK  153
NTDB id 217 SPD RS03775 WP 000590640.1        MTAKESLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKF....NENTLTYGNIVVNLSTNTVKVEDTPVELLGK  153
NTDB id 183 SPR RS03560 WP 000590640.1        MTAKESLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKF....NENTLTYGNIVVNLSTNTVKVEDTPVELLGK  153
NTDB id 147 SP RS03905 WP 000590640.1         MTAKESLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKF....NENTLTYGNIVVNLSTNTVKVEDTPVELLGK  153
consensus                                     *!!******!***!***!!!!!*!!!!**!!!**!**!***!****    *****************!*********!***

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NTDB id 606 V4T04 RS01910 WP 003130756.1      EYDLLLTLMQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKIDVEGQDSYIQTVRGLGYVMRERK  230
NTDB id 1139392 DQM69 RS04270 WP 002961524.1  EFDLLVYFLQNQNVILPKTQIFDRIWGFDSDTTVSVVEVYVSKIRKKLKGTDFEKCLQTLRSVGYILKDVK  224
NTDB id 362 SMU RS05200 WP 002262214.1        EFDLLVYFLQNQNVILPKSQIFDRIWGFDSDTTISVVEVYVSKIRKKLKDTTFGKNLQTLRSVGYILKDAN  224
NTDB id 258 KZH43 RS03475 WP 000590640.1      EFDLLVYFLQNQNVILPKTQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGTTFAENLQTLRSVGYLLKDVQ  224
NTDB id 217 SPD RS03775 WP 000590640.1        EFDLLVYFLQNQNVILPKTQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGTTFAENLQTLRSVGYLLKDVQ  224
NTDB id 183 SPR RS03560 WP 000590640.1        EFDLLVYFLQNQNVILPKTQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGTTFAENLQTLRSVGYLLKDVQ  224
NTDB id 147 SP RS03905 WP 000590640.1         EFDLLVYFLQNQNVILPKTQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGTTFAENLQTLRSVGYLLKDVQ  224
consensus                                     !*!!!****!!*******************!!****!***!!*!!****!*!*********!!*!**!!******
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X non conserved

X similar

X $\geq$ 50% conserved