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logo

NTDB id 606 V4T04 RS01910 WP 003130756.1        MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKE.KDTPIIM   79
NTDB id 1067667 ACFSN5 RS03975 WP 165213431.1   ..MIKILLVEDDLSLSNSVFDFLDDFA.EVTQVFDGEEGLFEAETGVYDLILLDLMLPEKDGFTVLKELRDKGIPTPVLI   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
NTDB id 362 SMU RS05200 WP 002262214.1          ..MIKLLLVEDDLSLSNSIFDFLDDFA.DVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLI   77
consensus                                         **!*!**!!***!*******!**** ********!**!!*!!****!!!!!!!!!!!!**!!*!***!!*****!!***

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NTDB id 606 V4T04 RS01910 WP 003130756.1        MTARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAVERAENTSFRDLVIDKTNRTVHRGKKVIDLTRR   159
NTDB id 1067667 ACFSN5 RS03975 WP 165213431.1   TTAKEALDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKIN....DKLLTYGNLDCDTSTNTVTVDGKVVELLGK   153
NTDB id 258 KZH43 RS03475 WP 000590640.1        MTAKESLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKFN....ENTLTYGNIVVNLSTNTVKVEDTPVELLGK   153
NTDB id 217 SPD RS03775 WP 000590640.1          MTAKESLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKFN....ENTLTYGNIVVNLSTNTVKVEDTPVELLGK   153
NTDB id 183 SPR RS03560 WP 000590640.1          MTAKESLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKFN....ENTLTYGNIVVNLSTNTVKVEDTPVELLGK   153
NTDB id 147 SP RS03905 WP 000590640.1           MTAKESLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKFN....ENTLTYGNIVVNLSTNTVKVEDTPVELLGK   153
NTDB id 362 SMU RS05200 WP 002262214.1          MTAKEGLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKFN....ENNLTYGELTVDTATNTTMVNGKEVELLGK   153
consensus                                       *!!******!***!***!!!!!*!!!!**!!!**!**!***!*****    ****************!*********!***

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NTDB id 606 V4T04 RS01910 WP 003130756.1        EYDLLLTLMQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKIDVEGQDSYIQTVRGLGYVMRERK   230
NTDB id 1067667 ACFSN5 RS03975 WP 165213431.1   EFDLLVYFLQNQNVILPKTQIFDRIWGFDSDTTVSVVEVYVSKIRKKLKGTDFEQNLQTLRSVGYILKDVQ   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
NTDB id 362 SMU RS05200 WP 002262214.1          EFDLLVYFLQNQNVILPKSQIFDRIWGFDSDTTISVVEVYVSKIRKKLKDTTFGKNLQTLRSVGYILKDAN   224
consensus                                       !*!!!****!!*******************!!****!**!!*!!****!*!*********!!*!**!!******
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X non conserved

X similar

X $\geq$ 50% conserved