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logo

NTDB id 606 V4T04 RS01910 WP 003130756.1      MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKE.KDTPIIM   79
NTDB id 258 KZH43 RS03475 WP 000590640.1      ..MIKILLVEDDLGLSNSVFDFLDDF.ADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLI   77
NTDB id 217 SPD RS03775 WP 000590640.1        ..MIKILLVEDDLGLSNSVFDFLDDF.ADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLI   77
NTDB id 183 SPR RS03560 WP 000590640.1        ..MIKILLVEDDLGLSNSVFDFLDDF.ADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLI   77
NTDB id 147 SP RS03905 WP 000590640.1         ..MIKILLVEDDLGLSNSVFDFLDDF.ADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLI   77
NTDB id 173352 A0021 RS04750 WP 067062089.1  ..MIKILLVEDDLGLSNSVFDFLDDF.ADVMQVFDGEEGIYEAESGVYDLILLDLMLPEKNGFQVLKELREKGVATPVLI   77
NTDB id 362 SMU RS05200 WP 002262214.1        ..MIKLLLVEDDLSLSNSIFDFLDDF.ADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLI   77
consensus                                       **!*!**!!***!*******!***  ********!**!**!!*!**!!!!!!!!!!!!**!!*!***!!****!!***

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NTDB id 606 V4T04 RS01910 WP 003130756.1      MTARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAVERAENTSFRDLVIDKTNRTVHRGKKVIDLTRR   159
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 173352 A0021 RS04750 WP 067062089.1  MTAKEGLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKFN....EKVLSYAELSVDTSTNTTLVKGEEVELLGK   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 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 173352 A0021 RS04750 WP 067062089.1  EFDLLVYFLQNKNVILPKSQIFDRIWGFDSDTTISVVEVYVSKIRKKLKGTTFAKNLQTLRSVGYILKNAE   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