```
logo

NTDB id 362 SMU RS05200 WP 002262214.1        ..MIKLLLVEDDLSLSNSI.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE   82
NTDB id 121061 DK41 RS05155 WP 000590620.1    ..MIKILLIEDDLSLSNSV.FDFLDDFADVMQIFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE   82
NTDB id 258 KZH43 RS03475 WP 000590640.1      ..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE   82
NTDB id 217 SPD RS03775 WP 000590640.1        ..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE   82
NTDB id 183 SPR RS03560 WP 000590640.1        ..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE   82
NTDB id 147 SP RS03905 WP 000590640.1         ..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE   82
NTDB id 606 V4T04 RS01910 WP 003130756.1      MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKE.KDTPIIMMTARD   84
NTDB id 374 SMU RS06885 WP 002262930.1        ..MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRKN.SHVPIIMLSAKD   82
consensus                                       **!*!***!**  *****  ***************!***!*******!!!*!!!!!!!**!**!****!******!*****!**

logo

NTDB id 362 SMU RS05200 WP 002262214.1        GLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKFN.....E.....NNLTYGELTVDTATNTTMVNGKEVELLGKEFDL   157
NTDB id 121061 DK41 RS05155 WP 000590620.1    SIDDKGQGFDLGADDYLTKPFYLEELKMRIQALLKRSGKFN.....D.....NSLIYGDIRVDMSTNSTFVNQTEVELLGKEFDL   157
NTDB id 258 KZH43 RS03475 WP 000590640.1      SLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKFN.....E.....NTLTYGNIVVNLSTNTVKVEDTPVELLGKEFDL   157
NTDB id 217 SPD RS03775 WP 000590640.1        SLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKFN.....E.....NTLTYGNIVVNLSTNTVKVEDTPVELLGKEFDL   157
NTDB id 183 SPR RS03560 WP 000590640.1        SLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKFN.....E.....NTLTYGNIVVNLSTNTVKVEDTPVELLGKEFDL   157
NTDB id 147 SP RS03905 WP 000590640.1         SLDDKGHGFELGADDYLTKPFYLEELKMRIQALLKRSGKFN.....E.....NTLTYGNIVVNLSTNTVKVEDTPVELLGKEFDL   157
NTDB id 606 V4T04 RS01910 WP 003130756.1      STMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAVERAENT.SFRDLVI.....DKTNRTVHRGKKVIDLTREYDL   163
NTDB id 374 SMU RS06885 WP 002262930.1        SEFDKVIGLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGDLQILPDAFVAKKRGTEVELTHREFEL   167
consensus                                     ***!***!***!!!!!*!!!!***!!**!**!***!*****     *      ****** * * ********  * **!***!**!

logo

NTDB id 362 SMU RS05200 WP 002262214.1        LVYFLQNQNVILPKSQIFDRIWGFDSDTTISVVEVYVSKIRKKLKDT.TFGKNLQTLRSVGYILKDAN..   224
NTDB id 121061 DK41 RS05155 WP 000590620.1    LVYFLQNQNVILPKSQIFDRIWGFDSDTTISVVEVYVSKVRKKLKGT.LFSENLQTLRSVGYILKHVETD   226
NTDB id 258 KZH43 RS03475 WP 000590640.1      LVYFLQNQNVILPKTQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGT.TFAENLQTLRSVGYLLKDVQ..   224
NTDB id 217 SPD RS03775 WP 000590640.1        LVYFLQNQNVILPKTQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGT.TFAENLQTLRSVGYLLKDVQ..   224
NTDB id 183 SPR RS03560 WP 000590640.1        LVYFLQNQNVILPKTQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGT.TFAENLQTLRSVGYLLKDVQ..   224
NTDB id 147 SP RS03905 WP 000590640.1         LVYFLQNQNVILPKTQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGT.TFAENLQTLRSVGYLLKDVQ..   224
NTDB id 606 V4T04 RS01910 WP 003130756.1      LLTLMQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKIDV.EGQDSYIQTVRGLGYVMRERK..   230
NTDB id 374 SMU RS06885 WP 002262930.1        LHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYMKSYD..   235
consensus                                     !*******************!!*********!*!*****!*!****  *******!*!**!!******
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X non conserved

X similar

X $\geq$ 50% conserved