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logo
NTDB id 54637 K6K11 RS06690 WP 002913894.1  .......MSEQIVTPDTAALTVPNKDAKINLLDLNRQQMREFFKNMGEKPFRADQVMKWMYHYCCDDFDEMTDINKVLRSKLKEV   78
NTDB id 1457 ASU2 RS00455 WP 014990913.1    MSEQTQTCASEIQAM..NAAVQHPKSEKINLMNLTRQEMRELFAEMGEKPFRADQLMKWIYHFGEDNFDNMSNINKVLREKLKQI   83
NTDB id 1392 A4U84 RS02730 WP 010786053.1   .......MSEQIQAT..DASCNSTKTEKINLLNLNRQQMRELFAEMGEKPFRADQLMKWIYHFGEDNFDNMTNINKVLREKLKRI   76
consensus                                          ****!**     !   *  !*!!!!**!*!!*!!!*!**!!!!!!!!!*!!!*!!***!*!!*!**!!!!!!*!!! *

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NTDB id 54637 K6K11 RS06690 WP 002913894.1  AEIRAPEVVEEQRSTDGTIKWAIAVGDQRVETVYIPEEDRATLCVSSQVGCALECKFCSTAQQGFNRNLRVSEIIGQVWRAAKIV  163
NTDB id 1457 ASU2 RS00455 WP 014990913.1    AEIKAPEVSVEQRSSDGTIKWAMQVGDQQIETVYIPEADRATLCVSSQVGCALACKFCSTAQQGFNRNLTVSEIIGQVWRASKII  168
NTDB id 1392 A4U84 RS02730 WP 010786053.1   AEIKAPEVAVEQRSADGTIKWAMWVGDQQIETVYIPEDDRATLCVSSQVGCALACTFCSTAQQGFNRNLSVSEIIGQVWRASKII  161
consensus                                   !!!*!!!! *!!!!*!!!!!!!* !!!!**!!!!!!!*!!!!!!!!!!!!!!!!*!*!!!!!!!!!!!!!!*!!!!!!!!!!!*!!*

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NTDB id 54637 K6K11 RS06690 WP 002913894.1  GAVKTTGVRPITNVVMMGMGEPLLNLNNVVPAMEIMLDDFGFGLSKRRVTLSTSGVVPALDKLGDMIDVALAISLHAPNDTIRDE  248
NTDB id 1457 ASU2 RS00455 WP 014990913.1    GNFGVTGVRPITNVVMMGMGEPLLNLNNVIPAMEIMLDDFAYGLSKRRVTLSTAGVVPALDIMREKIDVALAISLHAPNDELRDE  253
NTDB id 1392 A4U84 RS02730 WP 010786053.1   GNFGVTGIRPITNVVMMGMGEPLLNMNNVIPAMEIMLDDFAYGLSKRRVTLSTSGVVPALDKMREQIDVALAISLHAPNDELRNE  246
consensus                                   !****!!*!!!!!!!!!!!!!!!!!!!!*!!!*!!!!!!!!!!!**!!!!!!!!!!!!*!!!!!!!**** !!!!!!!!!!!!!!!**!*!

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NTDB id 54637 K6K11 RS06690 WP 002913894.1  IVPINKKYNIETFLNSVRGYISKSNANQGRVTIEYVMLDHVNDGTEHAHELAALLKDTPCKINLIPWNPFPGAPYGRSSNSRIDR  333
NTDB id 1457 ASU2 RS00455 WP 014990913.1    IMPINKKYNIKMLMDSVHKYLEVSNANHGKVTIEYVLLDHVNDGTEHAHQLAEVLKNTPCKINLIPWNPFPEAPYGKSSNSRVDR  338
NTDB id 1392 A4U84 RS02730 WP 010786053.1   LIPLNKKYNIKMLIDSVNKYLEVSNANHGKVTIEYVMLSHVNDDVEHAHQLADVLKNTPSKINLIPWNPFPEAPYAKSSNTRIDR  331
consensus                                   **!*!!!!!!****!! *!***!!!!*!*!!!!!!*!*!!!!**!!!!*!!*!!*!!*!!!!!!!!!!!*!!!**!!!!*!*!!

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NTDB id 54637 K6K11 RS06690 WP 002913894.1  FSKVLMEYGFTTIVRKTRGDDIDAACGQLAGDVIDRTKRTLRKRMQGEAIDVKAV  388
NTDB id 1457 ASU2 RS00455 WP 014990913.1    FQKTLMEYGFTVIVRKTRGDDIDAACGQLAGDVIDRTKRTMEKRKFGKGIAVQNH  393
NTDB id 1392 A4U84 RS02730 WP 010786053.1   FQKTLMEYGFTVTVRKTRGDDIDAACGQLAGDVIDRTKRTFEKRKFGEGIAVQNQ  386
consensus                                   !*!*!!!!!!!**!!!!!!!!!!!!!!!!!!!
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