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logo
NTDB id 38782 SN31241 RS17975 WP 000003206.1  .......MSEQIVTPESSTPVVLNNETKINLLDLNRQQMREFFKNLGEKPFRADQVMKWMYHYCCDNFDEMTDINKVLRG   73
NTDB id 1457 ASU2 RS00455 WP 014990913.1      MSEQTQTCASEIQAM..NAAVQHPKSEKINLMNLTRQEMRELFAEMGEKPFRADQLMKWIYHFGEDNFDNMSNINKVLRE   78
NTDB id 1392 A4U84 RS02730 WP 010786053.1     .......MSEQIQAT..DASCNSTKTEKINLLNLNRQQMRELFAEMGEKPFRADQLMKWIYHFGEDNFDNMTNINKVLRE   71
consensus                                            ****!**    * **  **!!!!**!*!!*!!!*!***!!!!!!!!!*!!!*!!***!!!!*!**!!!!!!*

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NTDB id 38782 SN31241 RS17975 WP 000003206.1  KLKEVAEIRAPEVVEEQRSSDGTIKWAIAVGDQRVETVYIPEDDRATLCVSSQVGCALECKFCSTAQQGFNRNLRVSEII   153
NTDB id 1457 ASU2 RS00455 WP 014990913.1      KLKQIAEIKAPEVSVEQRSSDGTIKWAMQVGDQQIETVYIPEADRATLCVSSQVGCALACKFCSTAQQGFNRNLTVSEII   158
NTDB id 1392 A4U84 RS02730 WP 010786053.1     KLKRIAEIKAPEVAVEQRSADGTIKWAMWVGDQQIETVYIPEDDRATLCVSSQVGCALACTFCSTAQQGFNRNLSVSEII   151
consensus                                     !!! *!!!*!!!! *!!!!*!!!!!!!* !!!!**!!!!!!!*!!!!!!!!!!!!!!!!*!*!!!!!!!!!!!!!*!!!!!

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NTDB id 38782 SN31241 RS17975 WP 000003206.1  GQVWRAAKIVGAAKVTGQRPITNVVMMGMGEPLLNLTNVVPAMEIMLDDFGFGLSKRRVTLSTSGVVPALDKLGLMIDVA   233
NTDB id 1457 ASU2 RS00455 WP 014990913.1      GQVWRASKIIGNFGVTGVRPITNVVMMGMGEPLLNLNNVIPAMEIMLDDFAYGLSKRRVTLSTAGVVPALDIMREKIDVA   238
NTDB id 1392 A4U84 RS02730 WP 010786053.1     GQVWRASKIIGNFGVTGIRPITNVVMMGMGEPLLNMNNVIPAMEIMLDDFAYGLSKRRVTLSTSGVVPALDKMREQIDVA   231
consensus                                     !!!!!!*!!*!***!!!*!!!!!!!!!!!!!!!!!!**!!*!!!!!!!!!!**!!!!!!!!!!!!*!!!!!!!**** !!!!

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NTDB id 38782 SN31241 RS17975 WP 000003206.1  LAISLHAPNDTIRDEIVPINKKYNIETFLGAVRRYLEKSNANQGRVTIEYVMLDHVNDGTEHAHQLAELLKETPCKINLI   313
NTDB id 1457 ASU2 RS00455 WP 014990913.1      LAISLHAPNDELRDEIMPINKKYNIKMLMDSVHKYLEVSNANHGKVTIEYVLLDHVNDGTEHAHQLAEVLKNTPCKINLI   318
NTDB id 1392 A4U84 RS02730 WP 010786053.1     LAISLHAPNDELRNELIPLNKKYNIKMLIDSVNKYLEVSNANHGKVTIEYVMLSHVNDDVEHAHQLADVLKNTPSKINLI   311
consensus                                     !!!!!!!!!!**!*!**!*!!!!!!***!**! *!!!*!!!!*!*!!!!!!*!*!!!!**!!!!!!!**!!*!!*!!!!!

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NTDB id 38782 SN31241 RS17975 WP 000003206.1  PWNPFPGAPYGRSSNSRIDRFSKVLMSYGFTTIVRKTRGDDIDAACGQLAGDVIDRTKRTLRKRMQGEVIDIKAI   388
NTDB id 1457 ASU2 RS00455 WP 014990913.1      PWNPFPEAPYGKSSNSRVDRFQKTLMEYGFTVIVRKTRGDDIDAACGQLAGDVIDRTKRTMEKRKFGKGIAVQNH   393
NTDB id 1392 A4U84 RS02730 WP 010786053.1     PWNPFPEAPYAKSSNTRIDRFQKTLMEYGFTVTVRKTRGDDIDAACGQLAGDVIDRTKRTFEKRKFGEGIAVQNQ   386
consensus                                     !!!!!!*!!!**!!!*!*!!!*!*!!*!!!!**!!!!!!!!!!!!!!!!!!
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