```
logo
NTDB id 106844 A4211 RS09065 WP 010907446.1  .MLEQQTCAEKAANTVNTVELSTDQAKKINLMNLTRQQMREFFKELGEKPFRADQLVKWIYHFGEDNFDNMTNINKKLRD   79
NTDB id 1457 ASU2 RS00455 WP 014990913.1     MSEQTQTCASEIQA..MNAAVQHPKSEKINLMNLTRQEMRELFAEMGEKPFRADQLMKWIYHFGEDNFDNMSNINKVLRE   78
NTDB id 1392 A4U84 RS02730 WP 010786053.1    .......MSEQIQA..TDASCNSTKTEKINLLNLNRQQMRELFAEMGEKPFRADQLMKWIYHFGEDNFDNMTNINKVLRE   71
consensus                                         ***** ***     * *** ***!!!!*!!*!!*!!!*!*!*!!!!!!!!!!*!!!!!!!!!!!!!!*!!!!*!!*

logo
NTDB id 106844 A4211 RS09065 WP 010907446.1  KLKQVAEIKAPEVAVEQRSADGTIKWAMQVGDQQVETVYIPEADRATLCVSSQVGCALACTFCSTAQQGFNRNLTVSEII  159
NTDB id 1457 ASU2 RS00455 WP 014990913.1     KLKQIAEIKAPEVSVEQRSSDGTIKWAMQVGDQQIETVYIPEADRATLCVSSQVGCALACKFCSTAQQGFNRNLTVSEII  158
NTDB id 1392 A4U84 RS02730 WP 010786053.1    KLKRIAEIKAPEVAVEQRSADGTIKWAMWVGDQQIETVYIPEDDRATLCVSSQVGCALACTFCSTAQQGFNRNLSVSEII  151
consensus                                    !!!**!!!!!!!!*!!!!!*!!!!!!!!*!!!!!*!!!!!!!*!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!*!!!!!

logo
NTDB id 106844 A4211 RS09065 WP 010907446.1  GQVWRASKIIGNFGVTGVRPITNVVMMGMGEPLLNVANVVPAMEIMLDDFAYGLSKRRVTLSTSGVVPALDKLSEMIDVA  239
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 106844 A4211 RS09065 WP 010907446.1  LAISLHAPNDELRDEIVPINKKYNIKMLMDSVNRYLSVSNANHGKVTIEYVMLDHVNDGVEHAHQLAQVLKNTPCKINLI  319
NTDB id 1457 ASU2 RS00455 WP 014990913.1     LAISLHAPNDELRDEIMPINKKYNIKMLMDSVHKYLEVSNANHGKVTIEYVLLDHVNDGTEHAHQLAEVLKNTPCKINLI  318
NTDB id 1392 A4U84 RS02730 WP 010786053.1    LAISLHAPNDELRNELIPLNKKYNIKMLIDSVNKYLEVSNANHGKVTIEYVMLSHVNDDVEHAHQLADVLKNTPSKINLI  311
consensus                                    !!!!!!!!!!!!!!*!**!*!!!!!!!!!!*!!!**!!*!!!!!!!!!!!!!!!!*!*!!!!**!!!!!!!*!!!!!!*!!!!!

logo
NTDB id 106844 A4211 RS09065 WP 010907446.1  PWNPFPEAPYAKSSNSRIDRFQKTLMEYGFTVIVRKTRGDDIDAACGQLAGDVIDRTKRTAQKKQFGQEIAVRNH  394
NTDB id 1457 ASU2 RS00455 WP 014990913.1     PWNPFPEAPYGKSSNSRWDRFQKTLMEYGFTVIVRKTRGDDIDAACGQLAGDVIDRTKRTMEKRKFGKGIAVQNH  393
NTDB id 1392 A4U84 RS02730 WP 010786053.1    PWNPFPEAPYAKSSNTRIDRFQKTLMEYGFTVTVRKTRGDDIDAACGQLAGDVIDRTKRTFEKRKFGEGIAVQNQ  386
consensus                                    !!!!!!!!!!*!!!!*!*!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!
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