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logo
NTDB id 61889 SIR RS16155 WP 021003029.1    MIHFYFFLVGSIVASFLGLVVDRFPEQSIIVPSSHCNTCGQKLTPRDLVPILSQLINRLRCRFCGEKIPIRYILLELTGGCLFLV   85
NTDB id 496 SM12261 RS02330 WP 000608383.1  MINIYFFLVGSILASFLGLVIDRFPEQSIIQPASHCDSCHTPLRPLDLIPILSQVFNRFRCRYCKVRYPVWYAFFELGLGLLFLA   85
NTDB id 524 SMSK321 RS08155 WP 000565601.1  MIDIYFFLVGSILASFLGLVIDRFPEQSIISPASHCDSCQTRLRPLDLIPILSQVFNHFRCRYCKAPYPVWYALLELGLGLLFLS   85
NTDB id 156 SP RS08975 WP 000565004.1       MIDFYFFLVGSILASFLGLVIDRFPEQSIISSASHCDSCQTPLRPLDLIPILSQVFNRFRCRYCKVRYPVWYALFELSLGLLFLL   85
NTDB id 266 KZH43 RS07930 WP 000565010.1    MIDFYFFLVGSILASFLGLVIDRFPEQSIISSASHCDSCQTRLRPLDLIPILSQVFNRFCCRYCKVRYPVWYALFELGLGLLFLL   85
NTDB id 225 SPD RS08500 WP 000565010.1      MIDFYFFLVGSILASFLGLVIDRFPEQSIISSASHCDSCQTRLRPLDLIPILSQVFNRFCCRYCKVRYPVWYALFELGLGLLFLL   85
NTDB id 191 SPR RS08095 WP 000565010.1      MIDFYFFLVGSILASFLGLVIDRFPEQSIISSASHCDSCQTRLRPLDLIPILSQVFNRFCCRYCKVRYPVWYALFELGLGLLFLL   85
consensus                                   !!**!!!!!!!!*!!!!!!!*!!!!!!!!!!***!!!**!***!*!*!!*!!!!!**!***!!*!****!**!***!!**!*!!!*

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NTDB id 61889 SIR RS16155 WP 021003029.1    TSLGYLNINQLILLVSGMTLSLYDQRHQEYPLLIWLFSHVLLLFLTGFNLLMAIFLALGILTHFIDLRIGAGDFLFLASCSAIFN   170
NTDB id 496 SM12261 RS02330 WP 000608383.1  WSLGWLSLGQVILITAGLTLGIYDFRHQEYPLLVWMTFHLILMAFCGWSLIMLFFLVLGILAHFIDIRMGAGDFLFLASCSIVFS   170
NTDB id 524 SMSK321 RS08155 WP 000565601.1  YSWGRLSLGQVILITAGLTLGIYDFRHQEYPLLVWMTFHLILMASSGWNLVMVFFLALGILAHFIDIRMGAGDFLFLASCALVFS   170
NTDB id 156 SP RS08975 WP 000565004.1       YSWGWLSLGQVVLITAGLTLGIYDFHHQEYPLLVWMTFQLILIASSGWNLVMVSFLILGILAHFIDIRMGAGDFLFLASCALVFS   170
NTDB id 266 KZH43 RS07930 WP 000565010.1    YSWELLSLSQVILITAGLTLGIYDFRHQEYPLLVWMTFHLILIASSGWNLVMVSFLILGILAHFIDIRMGAGDFLFLASCALVFS   170
NTDB id 225 SPD RS08500 WP 000565010.1      YSWELLSLSQVILITAGLTLGIYDFRHQEYPLLVWMTFHLILIASSGWNLVMVSFLILGILAHFIDIRMGAGDFLFLASCALVFS   170
NTDB id 191 SPR RS08095 WP 000565010.1      YSWELLSLSQVILITAGLTLGIYDFRHQEYPLLVWMTFHLILIASSGWNLVMVSFLILGILAHFIDIRMGAGDFLFLASCALVFS   170
consensus                                   *!***!**  !**!***!*!!**!!**!!!!!!!*!*****!****!**!*!**!!*!!!**!!!!*!*!!!!!!!!!!!!**!*

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NTDB id 61889 SIR RS16155 WP 021003029.1    LTEVLLVLQISSIIGLFLFCLKSRKDRLAFIPCLFGASSILLILIKFVVFHQNILCLI   227
NTDB id 496 SM12261 RS02330 WP 000608383.1  VTELLILIQFASAMGILAFLLQKKKERLPFVPFLLLAACVIIFGKLLLV........   219
NTDB id 524 SMSK321 RS08155 WP 000565601.1  ATELLILIQFASATGILAFLLQKKKERLPFVPFLLLATCVIIFGKLLLV........   219
NTDB id 156 SP RS08975 WP 000565004.1       VTELLILIQFASATGILAFLLQKKKERLPFVPFLLLATCLIIFGKLLLV........   219
NTDB id 266 KZH43 RS07930 WP 000565010.1    VTELLILIQFASATGILAFLLQKKKERLPFVPFLLLAACVIIFGKLLLV........   219
NTDB id 225 SPD RS08500 WP 000565010.1      VTELLILIQFASATGILAFLLQKKKERLPFVPFLLLAACVIIFGKLLLV........   219
NTDB id 191 SPR RS08095 WP 000565010.1      VTELLILIQFASATGILAFLLQKKKERLPFVPFLLLAACVIIFGKLLLV........   219
consensus                                   *!!*!***!**!**!***!*!***!*!!*!*!*!**!****!**!****
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X non conserved

X similar

X ≥50% conserved