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logo
NTDB id 61533 SII RS06515 WP 020998966.1     MIHFYFFLVGSIVASFLGLVWDRFPEQSIIVPSSHCNTCGQKLTPRDLVPILSQLINRLRCRFCGEKIPIRYILLELTGGCLFLV   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 61533 SII RS06515 WP 020998966.1     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 61533 SII RS06515 WP 020998966.1     LTEVLLVLQISSIIGLFIFCLKSRKDRLAFIPCLFGASSILLILIKFVVFHQNILCLI   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