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logo
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NTDB id 252400 CR542 RS08525 WP 004194621.1   MKTIILFFLGASIGSFLGLVIDRFPEQSIIAPSSHCNACKRRLKAWDLIPVLSQLSTKSKCRYCKAKIPYWYLGLEFLAG   80
NTDB id 496 SM12261 RS02330 WP 000608383.1    MINIYFFLVGSILASFLGLVIDRFPEQSIIQPASHCDSCHTPLRPLDLIPILSQVFNRFRCRYCKVRYPVWYAFFELGLG   80
NTDB id 524 SMSK321 RS08155 WP 000565601.1    MIDIYFFLVGSILASFLGLVIDRFPEQSIISPASHCDSCQTRLRPLDLIPILSQVFNHFRCRYCKAPYPVWYALLELGLG   80
NTDB id 266 KZH43 RS07930 WP 000565010.1      MIDFYFFLVGSILASFLGLVIDRFPEQSIISSASHCDSCQTRLRPLDLIPILSQVFNRFCCRYCKVRYPVWYALFELGLG   80
NTDB id 225 SPD RS08500 WP 000565010.1        MIDFYFFLVGSILASFLGLVIDRFPEQSIISSASHCDSCQTRLRPLDLIPILSQVFNRFCCRYCKVRYPVWYALFELGLG   80
NTDB id 191 SPR RS08095 WP 000565010.1        MIDFYFFLVGSILASFLGLVIDRFPEQSIISSASHCDSCQTRLRPLDLIPILSQVFNRFCCRYCKVRYPVWYALFELGLG   80
NTDB id 156 SP RS08975 WP 000565004.1         MIDFYFFLVGSILASFLGLVIDRFPEQSIISSASHCDSCQTPLRPLDLIPILSQVFNRFRCRYCKVRYPVWYALFELSLG   80
consensus                                     !*****!**!****!!!!!!!!!!!!!!!!***!!!**!***!***!!!!*!!!******!!!!!***!*!!***!***!
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logo
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NTDB id 252400 CR542 RS08525 WP 004194621.1   LVVLLCHFQVLNLTETILILAGLVLTIYDIKHQEYPFAVWLIFTFIALILSQLNWLFCGFLLLAYLTEKWQINIGSGDFL   160
NTDB id 496 SM12261 RS02330 WP 000608383.1    LLFLAWSLGWLSLGQVILITAGLTLGIYDFRHQEYPLLVWMTFHIILMAFCGWSLIMLFFLVLGIIAHFIDIRMGAGDFL   160
NTDB id 524 SMSK321 RS08155 WP 000565601.1    LLFLSYSWGFLSLGQVILITAGLTLGIYDFRHQEYPLLVWMTFHLILMASSGWNLVMVFFLALGILAHFIDIRMGAGDFL   160
NTDB id 266 KZH43 RS07930 WP 000565010.1      LLFLLYSWELLSLSQVILITAGLTLGIYDFRHQEYPLLVWMTFHLILIASSGWNLVMVSFLILGILAHFIDIRMGAGDFL   160
NTDB id 225 SPD RS08500 WP 000565010.1        LLFLLYSWELLSLSQVILITAGLTLGIYDFRHQEYPLLVWMTFHLILIASSGWNLVMVSFLILGILAHFIDIRMGAGDFL   160
NTDB id 191 SPR RS08095 WP 000565010.1        LLFLLYSWELLSLSQVILITAGLTLGIYDFRHQEYPLLVWMTFHLILIASSGWNLVMVSFLILGILAHFIDIRMGAGDFL   160
NTDB id 156 SP RS08975 WP 000565004.1         LLFLLYSWGWLSLGQVVLITAGLTLGIYDFHHQEYPLLVWMTFQLILIASSGWNLVMVSFLILGILAHFIDIRMGAGDFL   160
consensus                                     !**!**** *!*!****!!*!!!!*!*!!!*!!!!!**!!**!**!*************!!*!********!**!*!!!!
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logo
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NTDB id 252400 CR542 RS08525 WP 004194621.1   YLASLALICGFTELLWIIQISSLLGLLVFAIFKP.KSIPYVPLLFLSSIPIILCI....   214
NTDB id 496 SM12261 RS02330 WP 000608383.1    FLASCSIVFSVTELLILIQFASAMGILAFLLQKKKERLPFVPFLLLAACVIIFGKLLLV   219
NTDB id 524 SMSK321 RS08155 WP 000565601.1    FLASCALVFSATELLILIQFASATGILAFLLQKKKERLPFVPFLLLATCVIIFGKLLLV   219
NTDB id 266 KZH43 RS07930 WP 000565010.1      FLASCALVFSVTELLILIQFASATGILAFLLQKKKERLPFVPFLLLAACVIIFGKLLLV   219
NTDB id 225 SPD RS08500 WP 000565010.1        FLASCALVFSVTELLILIQFASATGILAFLLQKKKERLPFVPFLLLAACVIIFGKLLLV   219
NTDB id 191 SPR RS08095 WP 000565010.1        FLASCALVFSVTELLILIQFASATGILAFLLQKKKERLPFVPFLLLAACVIIFGKLLLV   219
NTDB id 156 SP RS08975 WP 000565004.1         FLASCALVFSVTELLILIQFASATGILAFLLQKKKERLPFVPFLLLATCLIIFGKLLLV   219
consensus                                     *!!!******!!!!!**!!**!*!*!*!*!***!*****!*!!*!*!***!!*******
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X non conserved

X similar

X $\geq$ 50% conserved