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logo
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NTDB id 1155353 QNH74 RS02115 WP 012027589.1  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 1155353 QNH74 RS02115 WP 012027589.1  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 1155353 QNH74 RS02115 WP 012027589.1  YLASLALICGFTELLWIIQISSLLGLLVFAIFKP.KSIPYVPLLFLSSIPIILCT....  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 ≥ 50% conserved