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logo

NTDB id 496 SM12261 RS02330 WP 000608383.1      MINIYFFLVGSILASFLGLVIDRFPEQSIIQPASHCDSCHTPLRPLDLIPILSQVFNRFRCRYCKVRYPVWYAFFELGLG    80
NTDB id 524 SMSK321 RS08155 WP 000565601.1      MIDIYFFLVGSILASFLGLVIDRFPEQSIISPASHCDSCQTRLRPLDLIPILSQVFNHFRCRYCKAPYPVWYALLELGLG    80
NTDB id 1170193 ACLV7Z RS02315 WP 071478877.1   MIDFYFFLVGSILASFLGLVIDRFPEQSIIQPASHCDSCQTRLRPLDLIPILSQVLNRFRCRYCKAPYPIWYALFELGLG    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                                       !!**!!!!!!!!!!!!!!!!!!!!!!!!!!!!!**!!!!!!!*!*!!!!!!!!!!!!!!*!*!*!!!!!**!!*!!!**!!*!!

logo

NTDB id 496 SM12261 RS02330 WP 000608383.1      LLFLAWSLGWLSLGQVILITAGLTLGIYDFRHQEYPLLVWMTFHIILMAFCGWSLIMLFFLVLGIIAHFIDIRMGAGDFL   160
NTDB id 524 SMSK321 RS08155 WP 000565601.1      LLFLSYSWGFLSLGQVILITAGLTLGIYDFRHQEYPLLVWMTFHLILMASSGWNLVMVFFLALGILAHFIDIRMGAGDFL   160
NTDB id 1170193 ACLV7Z RS02315 WP 071478877.1   LLFLLYSWELLSLSQVILITAGLTLGIYDFRHQEYPLLVWMTFHLILIACSGWNLVMVFFLVLGIVAHFIDIRMGAGDFL   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                                       !!!!**!***!!!*!!*!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!**!!*!**!!*!*!**!!*!!!*!!!!!!!!!!!!!!

logo

NTDB id 496 SM12261 RS02330 WP 000608383.1      FLASCSIVFSVTELLILIQFASAMGILAFLLQKKKERLPFVPFLLLAACVIIFGKLLLV   219
NTDB id 524 SMSK321 RS08155 WP 000565601.1      FLASCALVFSATELLILIQFASATGILAFLLQKKKERLPFVPFLLLATCVIIFGKLLLV   219
NTDB id 1170193 ACLV7Z RS02315 WP 071478877.1   FLASCALVFSVTELLILIQFASTMGILAFLLQKKKERLPFVPFLLLAACVIIFGKLLLV   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