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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 1170275 ACLV7A RS02060 WP 411863453.1   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 1170275 ACLV7A RS02060 WP 411863453.1   LLFLLYSWELLSLSQVILITAGLTLGIYDFRHQEYPLLVWLTFHLILMAFCGWNLVMVFFLILGIMAHFIDIRMGAGDFL  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 1170275 ACLV7A RS02060 WP 411863453.1   FLASCALVFSATELLILIQFASAMGILAFLLQKKKERLPFVPFLLLAACVIIFGKLLLV   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