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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 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 1142072 DQL01 RS09205 WP 111702693.1   MIDIYFFLVGSILASFLGLVIDRFPEQSIISSASHCDSCQTRLRPLDLIPILSQVFNHFRCRYCKVRYPVWYALFELGLG   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 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 1142072 DQL01 RS09205 WP 111702693.1   LLFLLYSWGWLSLGQVVLITAGLTLSIYDFRHQEYPLLVWMTFHLILIASSGWNLVMVSFLILGILAHFMDIRMGAGDFL  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 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 1142072 DQL01 RS09205 WP 111702693.1   FLASCALVFSVTELLILIQFASATGILAFLLQKKKERLPFVPFLLLATCVIIFGKLLLV   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