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logo                                    

NTDB id 21881 STR RS01120 WP 011226822.1 MRYITVEGLSFQYDAEPVLDNIHYHLDSGEFVTLTGENGAAKSTLIKATLGILKPKKGIVTISDINKDGKKLRMAYLPQQIASFN  85
NTDB id 435 SGO RS09490 WP 012130955.1   MRYITVENLSFYYDKEPVLEHIHYFLDSGEFVTLTGENGAAKTTLIKASLGILQPKHGEVKISKINTQGKKLRIAYLPQQIASFN  85
NTDB id 438 SPR RS10080 WP 001269474.1   MRYITVEDLSFYYDKEPVLEHINYCVDSGEFVTLTGENGAAKTTLIKASLGILQPRIGKVAISKTNTQGKKLRIAYLPQQIASFN  85
consensus                               !!!!!!! !!!*!!*!!!!**!*! *!!!!!!!!!!!!!!!!*!!!!*!!!!*!* ! ! !!**!**!!!!!*!!!!!!!!!!!

logo                                    

NTDB id 21881 STR RS01120 WP 011226822.1 AGFPSTVYEFVKSGRYPHKGWFRRLNKHDLEHVQRALESVGMWENRYKQIGHLSGGQKQRAVIARIFALDPDIFVLDEPTTGMDA  170
NTDB id 435 SGO RS09490 WP 012130955.1   AGFPSTVYEFVKSGRYPRKGWFRRLNEHDEEHIKASLESVGMWEHRDKRIGSLSGGQKQRAVIARMFASDPDIFVLDEPTTGMDA  170
NTDB id 438 SPR RS10080 WP 001269474.1   AGFPSTVYEFVKSGRYPRKGWFRRLNAHDEEHIKASLDSVGMWEHRDKRLGSLSGGQKQRAVIARMFASDPDVFILDEPTTGMDA  170
consensus                               !!!!!!!!!!!!!!!!!*!!!!!!!! !!*!!****!*!!!!!!*!*!**!*!!!!!!!!!!!!!!*!!*!!!*!*!!!!!!!!!!

logo                                    

NTDB id 21881 STR RS01120 WP 011226822.1 GTANTFYELMHHSAHNHGKSVLMITHDPEEVKEYTDRNIHLVRNQKLPWRCFNIHEKDGEFHTHD.   235
NTDB id 435 SGO RS09490 WP 012130955.1   GSKDEFYKLMHHSAHKHGKAVLMITHDPEEVKKYADRNIHLVRNQDSPWRCFNVHESDNGQEVSHA   236
NTDB id 438 SPR RS10080 WP 001269474.1   GSKNEFYELMHHSAHHHGKAVLMITHDPEEVKDYADRNIHLVRNQDSPWRCFNVHEN..GQEVGHA   234
consensus                               !****!!*!!!!!!!! !!!*!!!!!!!!!!!!!*!*!!!!!!!!!!**!!!!!!*!! * **** **
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X non conserved
X similar
X ≥ 50% conserved