```
logo
NTDB id 72206 SMUFR RS09165 WP 002263428.1  MRYITVENLSYQYDAEPVLEGINYHLDSGEFVTLTGENGAAKSTLIKATLGILKPKIGSVIFSKKNVSGKPLRLAYLPQQIASFN   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 72206 SMUFR RS09165 WP 002263428.1  AGFPSTVYEFVKSGRYPRNGWFRRLTKHDDEHIKVSLESVGMWENCQKPIGSLSGGQKQRAVIARMFASDPDIFVLDEPTTGMDT   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 72206 SMUFR RS09165 WP 002263428.1  GTTETFYKLMHHSAHEHDKAVLMITHDPEEVKAYADRNIHLVRNQNLPWRCFNVHEKDREEENNVK   236
NTDB id 435 SGO RS09490 WP 012130955.1      GSKDEFYKLMHHSAHKHGKAVLMITHDPEEVKKYADRNIHLVRNQDSPWRCFNVHESDNGQEVSHA   236
NTDB id 438 SPR RS10080 WP 001269474.1      GSKNEFYELMHHSAHHHGKAVLMITHDPEEVKDYADRNIHLVRNQDSPWRCFNVHEN..GQEVGHA   234
consensus                                   !***!!*!!!!!!!! !*!!!!!!!!!!!!!!! !!!!!!!!!!!!!**!!!!!!!!! * **!* **
```

X non conserved
X similar
X $\geq$ 50% conserved