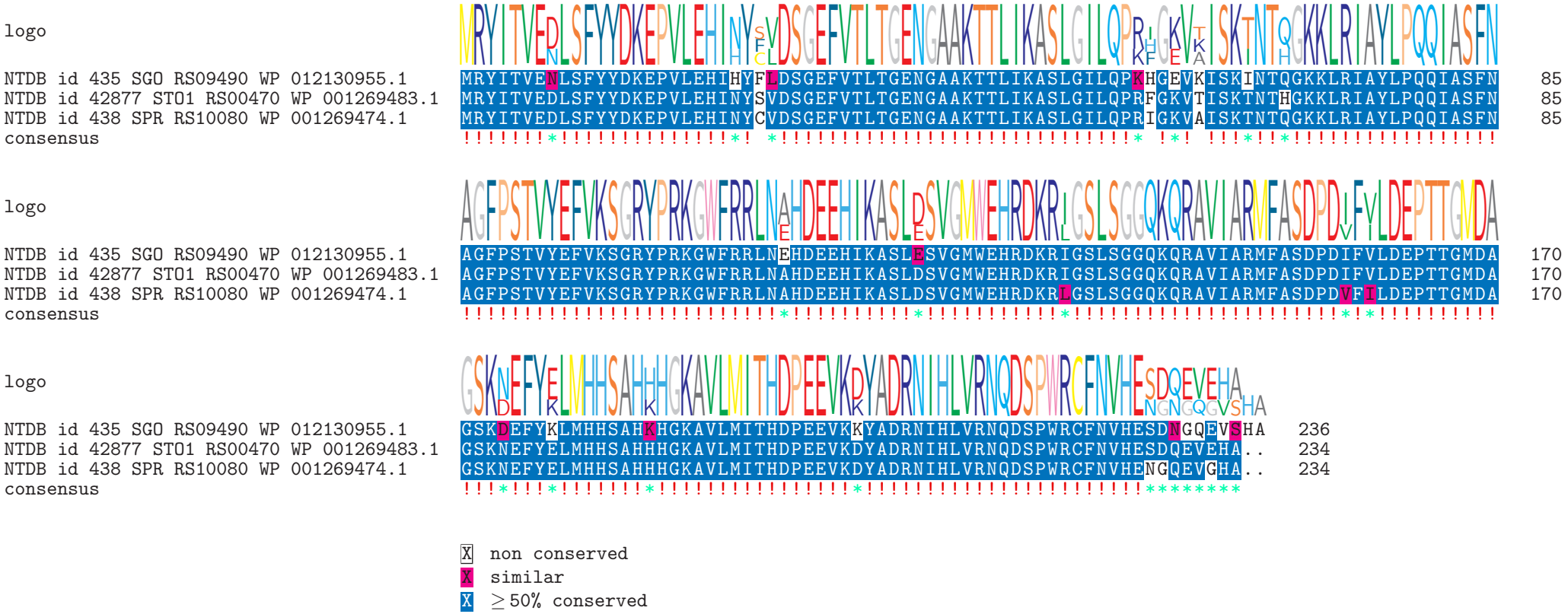
logo
NTDB id 435 SGO RS09490 WP 012130955.1 MRYITVENLSFYYDKEPVLEHIHYFLDSGEFVTLTGENGAAKTTLIKASLGILQPKHGEVKISKINTQGKKLRIAYLPQQIASFN 85
NTDB id 42877 STO1 RS00470 WP 001269483.1 MRYITVEDLSFYYDKEPVLEHINYSVDSGEFVTLTGENGAAKTTLIKASLGILQPRFGKVTISKTNTHGKKLRIAYLPQQIASFN 85
NTDB id 438 SPR RS10080 WP 001269474.1 MRYITVEDLSFYYDKEPVLEHINYCVDSGEFVTLTGENGAAKTTLIKASLGILQPRIGKVAISKTNTQGKKLRIAYLPQQIASFN 85
consensus
logo
NTDB id 435 SGO RS09490 WP 012130955.1 AGFPSTVYEFVKSGRYPRKGWFRRLNEHDEEHIKASLESVGMWEHRDKRIGSLSGGQKQRAVIARMFASDPDIFVLDEPTTGMDA 170
NTDB id 42877 STO1 RS00470 WP 001269483.1 AGFPSTVYEFVKSGRYPRKGWFRRLNAHDEEHIKASLDSVGMWEHRDKRIGSLSGGQKQRAVIARMFASDPDIFVLDEPTTGMDA 170
NTDB id 438 SPR RS10080 WP 001269474.1 AGFPSTVYEFVKSGRYPRKGWFRRLNAHDEEHIKASLDSVGMWEHRDKRLGSLSGGQKQRAVIARMFASDPDVFILDEPTTGMDA 170
consensus
logo
NTDB id 435 SGO RS09490 WP 012130955.1 GSKDEFYKLMHHSAHKHGKAVLMITHDPEEVKKYADRNIHLVRNQDSPWRCFNVHESDNGQEVSHA 236
NTDB id 42877 STO1 RS00470 WP 001269483.1 GSKNEFYELMHHSAHHHGKAVLMITHDPEEVKDYADRNIHLVRNQDSPWRCFNVHESDQEVEHA.. 234
NTDB id 438 SPR RS10080 WP 001269474.1 GSKNEFYELMHHSAHHHGKAVLMITHDPEEVKDYADRNIHLVRNQDSPWRCFNVHENGQEVGHA.. 234
consensus
X non conserved
X similar
X ≥ 50% conserved