```
logo
NTDB id 369690 D069 RS13065 WP 105937480.1  MLLILFFLGLIFGSFFYTAACRIPLRISVVSPRSACSFCRLPLSWGELVPVVSYILQRGRCRNCRAKLSVMYPAAECWTACLFTA   85
NTDB id 103 BSU 28070 NP 390685.2           MLSILFIFGLILGSFYYTAGCRIPLHLSIIAPRSSCPFCRRTLTPAELIPILSFLFQKGKCKSCGHRISFMYPAAELVTACLFAA   85
consensus                                   !! !!!  !!! !!!*!!!*!!!!!**!** !!! !*!!! *!***!!*!**!** !*!*!* ! ***! !!!!!!* !!!!! !

logo
NTDB id 369690 D069 RS13065 WP 105937480.1  AGIHFGFSKELLVALLFLSLLMIVTMTDLQYMLIPDKVLLFFLPLFIAGRMFSPLDSWYAGFAGAVCGFFLLVFIMFVSKGGIGA  170
NTDB id 103 BSU 28070 NP 390685.2           AGIRFGISLELFPAVVFISLLIIVAVTDIHFMLIPNRILIFFLPFLAAARLISPLDSWYAGLLGAAAGFLFLAVIAAITHGGVGG  170
consensus                                   !!!*!! ! !! *!**!*!!!*!! *!!***!!!! **!*!!!!   !*!* !!!!!!!!!  !! *!!  !  !  ***!!*!*

logo
NTDB id 369690 D069 RS13065 WP 105937480.1  GDVKLFGVIGLTLGVKLVLIAFFLSVMIGAVYGMCAAARGRLGKKQPFPFAPAISAGSALSYLYGEELFSFYIKLASGGA  250
NTDB id 103 BSU 28070 NP 390685.2           GDIKLFAVIGFVLGVKMLAAAFFFSVLIGALYGAAAVLTGRLAKRQPLPFAPAIAAGSILAYLYGDSIISFYIKMALG..  248
consensus                                   !!*!!!*!!!  !!!!**  !!! !!*!!!*!! *!   !!!*!*!! !!!!!! !!! ! !!!!* * !!!!!*! !**
```

X non conserved

X similar

X $\geq$ 50% conserved