```
logo
NTDB id 420165 GYN06 RS10705 WP 250333191.1  ..........................MPTDPDSRL.RGNDEAILLAEGQKSAVTEYYLNNGIWPENNGAAGVASA.STI   51
NTDB id 1109 OK783 RS10670 WP 157147416.1    MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGIWPKDNTSAGVASA.STI   79
NTDB id 1119 NGFG RS11465 WP 017147169.1     MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGKWPENNTSAGVASPPTDI   80
consensus                                    ****************************!***!********!!!!!!!!!!!!!!!!!!!*!*!!**!**!!!!!* **!

logo
NTDB id 420165 GYN06 RS10705 WP 250333191.1  KGKYVQKVEVAKGVVTAQMASTGVNNEIKGKKLSLWARRQDGSVKWFCGQPVTRANDAAKDAVTADAKDAIETKHLPSTC  131
NTDB id 1109 OK783 RS10670 WP 157147416.1    KGKYVQKVEVAKGVVTAQMASTGVNKEIQDKKLSLWAKRQDGSVKWFCGQPVTRTGDND..DTVADANNAIDTKHLPSTC  157
NTDB id 1119 NGFG RS11465 WP 017147169.1     KGKYVKEVEVKNGVVTATMLSSGVNNEIKGKKLSLWGRRENGSVKWFCGQPVTRADDDT..VADAKDGKEIDTKHLPSTC  158
consensus                                    !!!!!**!!!**!!!!!*!*!*!!!*!!**!!!!!!**!**!!!!!!!!!!!!!!* !       !**  *!*!!!!!!!!

logo
NTDB id 420165 GYN06 RS10705 WP 250333191.1  RDKHSDT.   138
NTDB id 1109 OK783 RS10670 WP 157147416.1    RDESSAT.   164
NTDB id 1119 NGFG RS11465 WP 017147169.1     RDKASDAK   166
consensus                                    !!* !**
```

X non conserved

X similar

X ≥ 50% conserved