```
logo
NTDB id 420259 GYN04 RS10785 WP 250328381.1  ..........................MPTDPDSRL.RGNDEAILLAEGQKSAVTEYYLNHGIWPKDNTSAGVASA.SDI   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 420259 GYN04 RS10785 WP 250328381.1  KGKYVKEVKVEKGVVTAEMASTGVNKEIQGKKLSLWGRRENGSVKWFCGQPVTRAAGKAGADADAVTADATANGINTKHL  131
NTDB id 1109 OK783 RS10670 WP 157147416.1    KGKYVQKVEVAKGVVTAQMASTGVNKEIQDKKLSLWAKRQDGSVKWFCGQPVTRTGDNDDTV......ADANNAIDTKHL  153
NTDB id 1119 NGFG RS11465 WP 017147169.1     KGKYVKEVEVKNGVVTATMLSSGVNNEIKGKKLSLWGRRENGSVKWFCGQPVTRADDDTVAD......AKDGKEIDTKHL  154
consensus                                    !!!!!**!*! *!!!!! !*!*!!!*!!**!!!!!!**!**!!!!!!!!!!!!!!***   **       *  ***!*!!!!

logo
NTDB id 420259 GYN04 RS10785 WP 250328381.1  PSTCRDKHSDT.   142
NTDB id 1109 OK783 RS10670 WP 157147416.1    PSTCRDESSAT.   164
NTDB id 1119 NGFG RS11465 WP 017147169.1     PSTCRDKASDAK   166
consensus                                    !!!!!!* !**
```

X non conserved

X similar

X $\geq$ 50% conserved