```
logo
NTDB id 163537 AT706 RS12375 WP 236822450.1  MFHLLGAQQNQKLKRRRFFCPVCGGELAVKLGLQKAPHFAHKQNKSCSIDIEPESAYHLEGKRQLYVWLKTQRASPILEP   80
NTDB id 123 BSU 11530 NP 389035.1            MFHLLGAQQNQKLKRRRFFCPVCGGELAVKLGLQKAPHFAHKQNKSCAIDIEPESAYHLEGKRQLYVWLKTQRASPILEP   80
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 163537 AT706 RS12375 WP 236822450.1  YIRTINQRPDVMARIKEHMLAVEYQCATIAPNVFQKLTEGFKQEGIIPQWIMGYSRLKRTASSFYQLSAFHWQFINASPY  160
NTDB id 123 BSU 11530 NP 389035.1            YIRTINQRPDVMARIKEHMLAVEYQCATIAPDVFQKRTEGFKQEGIIPQWIMGYSRLKRTASSFYQLSTFHWQFINASPY  160
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!

logo
NTDB id 163537 AT706 RS12375 WP 236822450.1  RELICYCPERRSFLRLSHIIPFYTNHSYSSVQTIPIHRAGVDDLFFTEPKPSIQYSGWTKAIHRFRHKPHRFNSKEINRL  240
NTDB id 123 BSU 11530 NP 389035.1            RELICYCPERRSFLRLSHIIPFYTNHSYSSVQTIPIHRAGAGDLFFTEPKPSIQYSGWTKAIHRFRHKPHRFNSKETNRL  240
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!  !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!

logo
NTDB id 163537 AT706 RS12375 WP 236822450.1  RLLFYEKRQTPFSFLPTEVFVPVSKGAVFKSPVFVWQGFLYLFMTDLGGKRAPIRFSAVLQQCKLHIHNKNIALRFECSE  320
NTDB id 123 BSU 11530 NP 389035.1            RLLFYEKRQTPFSFLPTEVFVPVRKGAVFKSPVFVWQGFLYLFMTDLGGKRAPIRFSAVLQQCKLHIHNKNIALRSECSE  320
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 163537 AT706 RS12375 WP 236822450.1  ECLSEAVKQYIDFLCKKGFLRETQKEVYVLNQPASGIHSMQDLIERDRSCFIE  373
NTDB id 123 BSU 11530 NP 389035.1            ECLSEAVKQYIDFLCKKGFLRETQKEVYVLNQPAGGIHSMQDLIERDRSCFIE  373
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved