```
logo
NTDB id 108253 AACH71 RS06580 WP 252153584.1   MFHLLGAQQSQKLKRKRFFCPVCGEELAVKLGLQKAPHFAHKQNKSCSIDIEPESAYHLEGKRQLYVWLKTQKASPILEP   80
NTDB id 123 BSU 11530 NP 389035.1              MFHLLGAQQNQKLKRRRFFCPVCGGELAVKLGLQKAPHFAHKQNKSCAIDIEPESAYHLEGKRQLYVWLKTQRASPILEP   80
consensus                                      !!!!!!!!! !!!!!*!!!!!!!! !!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!

logo
NTDB id 108253 AACH71 RS06580 WP 252153584.1   YIKEINQRPDVLARIKERMLAVEYQCATISPDVFQKRTEGFKQAGIIPQWIMGHSRLQRTASSFYQLSAFHWQFINANPY   160
NTDB id 123 BSU 11530 NP 389035.1              YIRTINQRPDVMARIKEHMLAVEYQCATIAPDVFQKRTEGFKQEGIIPQWIMGYSRLKRTASSFYQLSTFHWQFINASPY   160
consensus                                      !!* !!!!!!!*!!!!!*!!!!!!!!!!! !!!!!!!!!!!!! !!!!!!!!!!*!!! !!!!!!!!!!! !!!!!!!! !!

logo
NTDB id 108253 AACH71 RS06580 WP 252153584.1   RELICYCPERRSFLRLSHIIPFYTTHSYSSVQTIPIHRAGIDDLFFTKSKSPIQYSGWAKAIHRFRHKPHRFSSKETNRL   240
NTDB id 123 BSU 11530 NP 389035.1              RELICYCPERRSFLRLSHIIPFYTNHSYSSVQTIPIHRAGAGDLFFTEPKPSIQYSGWTKAIHRFRHKPHRFNSKETNRL   240
consensus                                      !!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!  !!!!!! *!**!!!!!! !!!!!!!!!!!!!! !!!!!!!

logo
NTDB id 108253 AACH71 RS06580 WP 252153584.1   KQLFYEKRQTPFSFLPTEVFVPVVKGAVFKSPVFVWQGFLYLFITDLGEKHAPIRFSAVLQQCKLHIHKKNIALRFECSE   320
NTDB id 123 BSU 11530 NP 389035.1              RLLFYEKRQTPFSFLPTEVFVPVRKGAVFKSPVFVWQGFLYLFMTDLGGKRAPIRFSAVLQQCKLHIHNKNIALRSECSE   320
consensus                                      * !!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!*!!!!! !*!!!!!!!!!!!!!!!!!!! !!!!!! !!!!

logo
NTDB id 108253 AACH71 RS06580 WP 252153584.1   ECLSEAVKQYIDFLCKKGFLRETQKEVYVLNQPADGIRSMQDLIERDRSCFIE   373
NTDB id 123 BSU 11530 NP 389035.1              ECLSEAVKQYIDFLCKKGFLRETQKEVYVLNQPAGGIHSMQDLIERDRSCFIE   373
consensus                                      !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!*!!!!!!!!!!!!!!!!
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X non conserved
X similar
X ≥ 50% conserved