```
logo

NTDB id 1063193 ACG4I6 RS01310 WP 000792032.1  MKNKPLGLWLMSLLVLVGCKANQDDLTSYVAQVEREARKEVTKLKPILEFEATPYQQHKGREPFVLPKEALVQNQPLAKA   80
NTDB id 1174 A1552VC RS12110 WP 000792032.1    MKNKPLGLWLMSLLVLVGCKANQDDLTSYVAQVEREARKEVTKLKPILEFEATPYQQHKGREPFVLPKEALVQNQPLAKA   80
NTDB id 1407 DSB67 RS13985 WP 010643498.1      MRNKSLLM.VCMVSLLSGCQANDESLTDFIRGVENQARRDVAKLQPTEEYVVVSYEPEILRAPFELPKEATIATQPVARQ   79
consensus                                      !*!!*!*********!*!!*!!***!!*****!!**!!**!*!!*!**!*****!*****!*!!*!!!!!****!!*!**

logo

NTDB id 1063193 ACG4I6 RS01310 WP 000792032.1  DCWQPVPRAKNGPLERYPLHQLRLKGVMSSGGSISALVQTPAGSVVKVKAGQYVGINNGKVTRVDDNYLLINETLPDGLG  160
NTDB id 1174 A1552VC RS12110 WP 000792032.1    DCWQPVPRAKNGPLERYPLHQLRLKGVMSSGGSISALVQTPAGSVVKVKAGQYVGINNGKVTRVDDNYLLINETLPDGLG  160
NTDB id 1407 DSB67 RS13985 WP 010643498.1      DCWQPPSRKRTDKLERFPLNQLRLKGVMGMGSSVSGLVQAPNGTVYKVKPGQYLGRNNGKVTHVSHSYLLINETLPDGLG  159
consensus                                      !!!!!**!*****!!!*!!*!!!!!!!!!**!*!*!*!!!*!*!*!*!!!!*!!!*!*!!!!!!*!***!!!!!!!!!!!!!

logo

NTDB id 1063193 ACG4I6 RS01310 WP 000792032.1  CWNQRNVKLALK   172
NTDB id 1174 A1552VC RS12110 WP 000792032.1    CWNQRNVKLALK   172
NTDB id 1407 DSB67 RS13985 WP 010643498.1      CWQKRKVKLALR   171
consensus                                      !!**!*!!!!!*
```

X non conserved

X similar

X ≥ 50% conserved