```
logo
NTDB id 408783 GPY04 RS10990 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 408783 GPY04 RS10990 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 408783 GPY04 RS10990 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