```
logo
NTDB id 1174 A1552VC RS12110 WP 000792032.1   MKNKPLGLWLMSLLVLVGCKANQDDLTSYVAQVEREARKEVTKLKPILEFEATPYQQHKGREPFVLPKEALVQNQPLAKA   80
NTDB id 973628 AABD47 RS01620 WP 005454781.1 MGNKSLLTV.LVSALLVGCQANDESLTDFIRGVESQARRDVEKLKPSDKYIAVAYAPEVMRAPFELPKEATIATQPVARK   79
NTDB id 1407 DSB67 RS13985 WP 010643498.1    MRNKSLLMV.CMVSLLSGCQANDESLTDFIRGVENQARRDVAKLQPTEEYVVVSYEPEILRAPFELPKEATIATQPVARQ   79
consensus                                    !*!!*!*** *** *!*!!*!!***!!*****!! *!!**! !!*!*******  ! ****!*!!*!!!!!****!!*!*

logo
NTDB id 1174 A1552VC RS12110 WP 000792032.1   DCWQPVPRAKNGPLERYPLHQLRLKGVMSSGGSISALVQTPAGSVVKVKAGQYVGINNGKVTRVDDNYLLINETLPDGLG   160
NTDB id 973628 AABD47 RS01620 WP 005454781.1 DCWQPPNRARSGKLEKFPLSQLRLKGVMGMGNTVSGLVQAPNGTVYKVKPGQYLGRNNGKVTHVTHSYLLINETLPDGLG   159
NTDB id 1407 DSB67 RS13985 WP 010643498.1    DCWQPPSRKRTDKLERFPLNQLRLKGVMGMGSSVSGLVQAPNGTVYKVKPGQYLGRNNGKVTHVSHSYLLINETLPDGLG   159
consensus                                    !!!!!* !*****!!**!! !!!!!!!!**! **!*!!!*!*!*!*!!!*!!!*!*!!!!!!*!***!!!!!!!!!!!!!!

logo
NTDB id 1174 A1552VC RS12110 WP 000792032.1   CWNQRNVKLALK   172
NTDB id 973628 AABD47 RS01620 WP 005454781.1 CWQKRKVKLALR   171
NTDB id 1407 DSB67 RS13985 WP 010643498.1    CWQKRKVKLALR   171
consensus                                    !!**!*!!!!!*
```

X non conserved

X similar

X ≥ 50% conserved