```
logo

NTDB id 116674 AW02 RS00020 WP 024084783.1  MYIQNLELTSYRNYERAELQFENKVNVIIGENAQGKTNLMEAIYVLSMAKSHRTSNDKELIRWDEDYAKIEGRVMKRNGDIPMQL   85
NTDB id 353 BSU 00040 NP 387885.1           MYIQNLELTSYRNYDHAELQFENKVNVIIGENAQGKTNLMEAIYVLSMAKSHRTSNDKELIRWDKDYAKIEGRVMKQNGAIPMQL   85
consensus                                   !!!!!!!!!!!!!!**!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 116674 AW02 RS00020 WP 024084783.1  VISKKGKKGKVNHIEQQKLSQYVGALNTIMFAPEDLNLVKGSPQVRRRFLDMEIGQVSAVYLYDLSLYQKILSQRNHFLKQLQSR   170
NTDB id 353 BSU 00040 NP 387885.1           VISKKGKKGKVNHIEQQKLSQYVGALNTIMFAPEDLNLVKGSPQVRRRFLDMEIGQVSPVYLHDLSLYQKILSQRNHFLKQLQTR   170
consensus                                   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 116674 AW02 RS00020 WP 024084783.1  KQTDRTMLDVLTDQLIEAAAKVVARRLQFTAQLEKWAQPIHSGISRGLEELTLKYHTALDVSDPKDLSKIGNSYQESFSKLKEKE   255
NTDB id 353 BSU 00040 NP 387885.1           KQTDRTMLDVLTDQLVEVAAKVVVKRLQFTAQLEKWAQPIHAGISRGLEELTLKYHTALDVSDPLDLSKIGDSYQEAFSKLREKE   255
consensus                                   !!!!!!!!!!!!!!!*! !!!!! *!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!! !!!!!! !!!! !!!!*!!!

logo

NTDB id 116674 AW02 RS00020 WP 024084783.1  IERGVTLFGPHRDDVLFYVNGRDVQTYGSQGQQRTTALSLKLAEIDLIHEEIGEYPILLLDDVLSELDDYRQSHLLHTIQGRVQT   340
NTDB id 353 BSU 00040 NP 387885.1           IERGVTLSGPHRDDVLFYVNGRDVQTYGSQGQQRTTALSLKLAEIDLIHEEIGEYPILLLDDVLSELDDYRQSHLLHTIQGRVQT   340
consensus                                   !!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 116674 AW02 RS00020 WP 024084783.1  FVTTTSVDGIDHDTLHQAGMFRVENGALVK   370
NTDB id 353 BSU 00040 NP 387885.1           FVTTTSVDGIDHETLRQAGMFRVQNGALVK   370
consensus                                   !!!!!!!!!!!!*!!*!!!!!!!! !!!!!!
```

X non conserved

X similar

X ≥ 50% conserved