```
logo

NTDB id 1362 HI 1008 AAC22669.1             MKTLF.TSVVLCGALVVSSSFAEEKATXQTAQSVVTTQAEAQVAPAVVSDKLNINTATASEIQKSLTGIGAKKAEAIVQYREKHG   84
NTDB id 27627 APL RS07435 WP 005599904.1    MKKFARFSISFIIALLSADSIAKPNNPPVSEPTPVVQQMTTTNKTADKQNQVNINTADAAEIKSKLVGIGEKKAQAIVDYRTKNG   85
NTDB id 1382 A4U84 RS07375 WP 062924234.1   MKKLSRYTFSLLFGLFSATTFAEEKVTTSTEHTPVVMQQVAAT.QKVNANVVNINTATASEIQDKLVGIGAKKAQAIVEYREKNG   84
consensus                                   !!** ******* *!******!*** *  ** **!* !  *    **  * *!!!!!*!*!!* *!*!!!*!!!*!!!*!!*!*!

logo

NTDB id 1362 HI 1008 AAC22669.1             NFXNAEQLLEVQGIGKATLEKNRDRIIF.   112
NTDB id 27627 APL RS07435 WP 005599904.1    AFITIEQLTEVSGIGKATLEKNRERITLE   114
NTDB id 1382 A4U84 RS07375 WP 062924234.1   KFLNVEQLTEVSGIGKATLEKNRDCIVIE   113
consensus                                    !***!!!*!!*!!!!!!!!!!!**!***
```

X non conserved

X similar

X ≥50% conserved