```
logo
NTDB id 1362 HI 1008 AAC22669.1              MKTLF.TSVVLCGALVVSSSFAEEKATXQTAQSVVTTQAEAQVAPAVVSDKLNINTATASEIQKSLTGIGAKKAEAIVQY   79
NTDB id 289215 APPSER1 RS08060 WP 005599904.1 MKKFARFSISFIIALLSADSIAKPNNPPVSEPTPVVQQMTTTNKTADKQNQVNINTADAAEIKSKLVGIGEKKAQAIVDY   80
NTDB id 1382 A4U84 RS07375 WP 062924234.1    MKKLSRYTFSLLFGLFSATTFAEEKVTTSTEHTPVVMQQVAAT.QKVNANVVNINTATASEIQDKLVGIGAKKAQAIVEY   79
consensus                                    !!** ******* *!******!*** *  ** **!* !  *    **  * *!!!!!*!*!!* *!*!!!*!!!*!!!*!
```

```
logo
NTDB id 1362 HI 1008 AAC22669.1              REKHGNFXNAEQLLEVQGIGKATLEKNRDRIIF.   112
NTDB id 289215 APPSER1 RS08060 WP 005599904.1 RTKNGAFITIEQLTEVSGIGKATLEKNRERITLE   114
NTDB id 1382 A4U84 RS07375 WP 062924234.1    REKNGKFLNVEQLTEVSGIGKATLEKNRDCIVIE   113
consensus                                    !*!*! !***!!!*!!*!!!!!!!!!!!**!***
```

X non conserved

X similar

X $\geq$ 50% conserved