```
logo
NTDB id 1136728 DQL47 RS01820 WP 111689845.1  MINNVVLVGRMTKDAELRYTPSQVAVATFTLAVNRAFKSQNGEREADFINCVIWRQQAENLANWAKKGALIGITGRIQTR   80
NTDB id 624 LCA RS00040 WP 011373726.1        MINRVVLVGRLTRDVDLRYTSSGAAVGTFSMAVNRQFTNANGDREADFINCVIWRKSAENFANFTKKGSLVGVDGRLQTR   80
NTDB id 113 BSU 40900 NP 391970.1             MLNRVVLVGRLTKDPELRYTPNGAAVATFTLAVNRTFTNQSGEREADFINCVTWRRQAENVANFLKKGSLAGVDGRLQTR   80
NTDB id 112 BSU 36310 NP 391512.2             MFNQVMLVGRLTKDPDLRYTSAGAAVAHVTLAVNRSFKNASGEIEADYVNCTLWRKTAENTALYCQKGSLVGVSGRIQTR   80
consensus                                     !*!*!*!!!!*!*!**!!!!****!!****!!!!*! *  !**!!!**!!**!!**!!!!*!** *!!*!*!**!!*!!!
```

```
logo
NTDB id 1136728 DQL47 RS01820 WP 111689845.1  NYENQQGQRVYVTEVVADNFQMLESRATREASSSGSYGGGFNGSP...........TANSYSAPTQQ....TPNFGRDES  145
NTDB id 624 LCA RS00040 WP 011373726.1        NYENQQGQRVYVTEVVVDNFSLLESRTTTEQRQGCDGASQNFNSNQSNGSQ.......QSGFTSPQQTGNAPAANN.TQAD  152
NTDB id 113 BSU 40900 NP 391970.1             NYENQQGQRVFVTEVQAESVQFLEPKNGGGSGSG.....GYNEGNSGGGQYFGGGQNDNPFGGNQNNQRRNQGNS.FNDD  154
NTDB id 112 BSU 36310 NP 391512.2             SYENEEGVNVYVTEVLADTVRFMDPKPREKAAD.......................................  113
consensus                                     *!!!**!**!*!!!!**** ****** * *****     ***  ** * *        * *******     **   ***
```

```
logo
NTDB id 1136728 DQL47 RS01820 WP 111689845.1  PF.GGSNPMDISDDDLPF  162
NTDB id 624 LCA RS00040 WP 011373726.1        PFANNGQAIDISDDDLPF  170
NTDB id 113 BSU 40900 NP 391970.1             PFANDGKPIDISDDDLPF  172
NTDB id 112 BSU 36310 NP 391512.2             ..................  113
consensus                                     **** *************
```

X non conserved

X similar

X $\geq$ 50% conserved