```
logo
NTDB id 21430 SERP RS10925 WP 001830679.1  MKIFIVEDDLVIAESLANELSKWNYEVHVVDNFEKIMEDFRRVEPQLVLLDINLPTLNGFHWCQEMRKVSHVPIMFISSRTDNMD   85
NTDB id 9 SA RS13865 WP 000697877.1        MKILIVEDDFVIAESLASELKKWNYGVIVVEQFDDILSIFNQHQPQLVLLDINLPTLNGFHWCQEIRKTSNVPIIFISSRIDNMD   85
consensus                                  !!! !!!!! !!!!!!! !! !!!! !*!!**!* !*  !  * !!!!!!!!!!!!!!!!!!!!!*!! !*!!!*!!!!! !!!!

logo
NTDB id 21430 SERP RS10925 WP 001830679.1  QILAIQMGGDDFIEKPFNLSLTVAKIQALLRRTYDLSIARDEIAVKGCKLIVDEAKLSKDNEHVQLSLTELQILKLLFQNENKYV  170
NTDB id 9 SA RS13865 WP 000697877.1        QIMAIQMGGDDFIEKPFNLSLTIAKIQALLRRTYDLSVANDSLTVKGCTLILDEAKVVYQEQNIQLSLTELQILKLLFQNEDKYV  170
consensus                                  !!*!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!*! ! * !!!! !!*!!!!*     **!!!!!!!!!!!!!!!!! !!!

logo
NTDB id 21430 SERP RS10925 WP 001830679.1  SRTALIEKCWESENFIDDNTLAVNMTRLRKKLLSIGVDDLIETKKNVGYRV   221
NTDB id 9 SA RS13865 WP 000697877.1        SRTALIEKCWESENFIDDNTLAVNMTRLRKKLNTIGVNDFIITKKNVGYKV   221
consensus                                  !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! *!!! ! ! !!!!!!!*!
```

X non conserved

X similar

X $\geq$50% conserved