```
logo

NTDB id 1067186 ACH6FT RS02635 WP 002780169.1  .MKLIIFLLAFTLHVFALEAYKPDKDFAPYFNPLNCSVVLDKFFYTNCYDYKLKGTKAVAYKVEASNLKEKQIKKRARFE   79
NTDB id 1225 CJE RS07175 WP 011049930.1        MKKLIILSLLSTLAFADYTQYKPSEDFAKYFTKQNCSQVLDKFYYINCYDYSLKGTKAVAYRLEADNLKGEQIKKRPRFE   80
NTDB id 1224 CJE RS02830 WP 011049704.1        MKKLILLPLLSTLAFADYTQYKPSEEFAKYFTKQSCSQVLDKFYYLNCYDYNLKGTKAVAYKLEVDNLKGEQIKKRPRFE   80
consensus                                      **!!!** !**!!******!!!***!!*!!****!!*!!!!!*!*!!!!! !!!!!!!!!**!**!!!**!!!!!*!!!

logo

NTDB id 1067186 ACH6FT RS02635 WP 002780169.1  DDTNIPKKYRSTWSDYKNSGYTRGHIASNASFRFSKAAQTSVFLMSNITPQNAQVNATVWNEIEQRERSLAFKFQSVEVL  159
NTDB id 1225 CJE RS07175 WP 011049930.1        DDTNIPKKYRTTWSDYKNSGYDRGHTLSNASMRKTTQAQRSTFLMSNITPQNPQINQRVWNKIEKRERQVALKLGSLEVL  160
NTDB id 1224 CJE RS02830 WP 011049704.1        DDTNIPKKYRTTWSDYKNSGYDRGHTLSNASMRKTTQAQRSTFLMSNITPQNPQINQRVWNKIEKRERQVASKLGSLEVL  160
consensus                                      !!!!!!!!!!*!!!!!!!!!!*!!!**!!!!*!****!!*!*!!!!!!!!!!*!*!**!!!*!!*!!!**! !**!*!!!

logo

NTDB id 1067186 ACH6FT RS02635 WP 002780169.1  NLILYGKNPKFIKNGIAIPSSYVKIIKTSRFEECYEVPNNNIQSEEINDYKISCNKF  216
NTDB id 1225 CJE RS07175 WP 011049930.1        NLVNYDNNPQRIKNNIAIPSSYTKILKGDNFKECYQVPNHDVENENLRIYKVKCDNF  217
NTDB id 1224 CJE RS02830 WP 011049704.1        NLVNYDNNPQRIRNQIAIPSSYIKILKGENFKECYQVPNHEVEDENIRKYKIDCDKI  217
consensus                                      !!**!**!!**!*!*!!!!!!!*!!*!**!*!!!*!!!**** !*** !!* !***
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X non conserved

X similar

X $\geq$ 50% conserved