```
logo
NTDB id 1067047 ACH6FG RS02930 WP 002855187.1  MKIF.IFLLTISLNIFALEPYKPSADFSSYFNNINCSQILDKFFYLNCYDYKLKGTKAVAYKVEASNLKDKQIKKRPRFE   79
NTDB id 1224 CJE RS02830 WP 011049704.1        MKKLILLPLLSTLAFADYTQYKPSEEFAKYFTKQSCSQVLDKFYYLNCYDYNLKGTKAVAYKLEVDNLKGEQIKKRPRFE   80
NTDB id 1225 CJE RS07175 WP 011049930.1        MKKLIILSLLSTLAFADYTQYKPSEDFAKYFTKQNCSQVLDKFYYINCYDYSLKGTKAVAYRLEADNLKGEQIKKRPRFE   80
consensus                                      !!***** !***!*******!!!!**!**!!****!!!*!!!!!*!!!!!! !!!!!!!!!**!**!!!**!!!!!!!!!
```

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logo
NTDB id 1067047 ACH6FG RS02930 WP 002855187.1  DDTNIPKKYRTTWSDYKNSGYTRGHTAPNASFSFSKAAQNSVFLMSNITPQIAQINNKIWNEIEQRERNLAFKFQSIEVL  159
NTDB id 1224 CJE RS02830 WP 011049704.1        DDTNIPKKYRTTWSDYKNSGYDRGHTLSNASMRKTTQAQRSTFLMSNITPQNPQINQRVWNKIEKRERQVASKLGSLEVL  160
NTDB id 1225 CJE RS07175 WP 011049930.1        DDTNIPKKYRTTWSDYKNSGYDRGHTLSNASMRKTTQAQRSTFLMSNITPQNPQINQRVWNKIEKRERQVALKLGSLEVL  160
consensus                                      !!!!!!!!!!!!!!!!!!!!!!!*!!!!**!!!******!!*!*!!!!!!!!!!**!!!***!!*!!*!!!**! !**!*!!!
```

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logo
NTDB id 1067047 ACH6FG RS02930 WP 002855187.1  NLVLYDKEPQYIKNRIAIPSFYVKIIKTPKFKECYQVPNHEVNDENIKQYQINCDKF  216
NTDB id 1224 CJE RS02830 WP 011049704.1        NLVNYDNNPQRIRNQIAIPSSYIKILKGENFKECYQVPNHEVEDENIRKYKIDCDKI  217
NTDB id 1225 CJE RS07175 WP 011049930.1        NLVNYDNNPQRIKNNIAIPSSYTKILKGDNFKECYQVPNHDVENENLRIYKVKCDNF  217
consensus                                      !!!*!!**!!*!*!*!!!!!*!*!!*!**!!!!!!!!!!*!**!!** !** !!**
```

X non conserved
X similar
X ≥ 50% conserved