

MDRLSIYELLISFWVPGVIMIELINFSAEYVFGKAPLITAGNLS DGLLFFAIALFIGCLIHITFWLKKRWYKKLACTPMDKMKND EYITEILPDLKEIYRTNKG 105

[illegible]

TDRIFGLYYAELTHNKK	123
.....	123
.....	84
.....	69
TDRIFGLYYAELTHNKK	219

- X non conserved
- X similar
- X >50% conserved