```
logo
NTDB id 293451 DK843 RS00955 WP 114072265.1   MKRYVVAAMLVMVGLAGCATTETY....DETRGWTVEKLYSEAHDELNSGNYTRAVKLYETLEARFPYGRYAQQAQMDLA   76
NTDB id 1102 NMB RS03635 WP 002244053.1       MKKIL.LTVSLGLALSACATQGTVDKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHAQQSQLDTA   79
NTDB id 1106 NGFG RS01495 WP 003687645.1      MKKIL.LTVSLGLALSACATQGTADKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHARQSQLDTA   79
consensus                                     !!*** ********!**!!!**! ******!**!*!!!!!*!!*!!!!!*!!!!!!!!!!*!!*!!!**!*!*!*!*!*!

logo
NTDB id 293451 DK843 RS00955 WP 114072265.1   YTHYKDGEPELAIASADRFIKLHPTHPNLDYIYYLKGLVYYNDDSGLLAKWAGQDMSERDPRAAREAFAAFRELTTRFPN   156
NTDB id 1102 NMB RS03635 WP 002244053.1       YAYYKDDEKDKALAAIDRFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPN   159
NTDB id 1106 NGFG RS01495 WP 003687645.1      YAYYKDDEKDKALAAIERFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPN   159
consensus                                     !**!!!*!***!*!***!!**!!!*!!!*!!**!!*!!!**!*!***!*!*!*!!*!*!!!*!*!!!**!!*!!**!!!!

logo
NTDB id 293451 DK843 RS00955 WP 114072265.1   STYAPDAKAKMIRLVDALGGNEMHVARYYMKRGAYLAAANRAQGVVKDYANTKYPEEALAIMVAAYDKLQLPQLRDDAHR   236
NTDB id 1102 NMB RS03635 WP 002244053.1       SKYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAQKIIGSYQNTRYVEESLAILELAYKKLDKPRLAADTRR   239
NTDB id 1106 NGFG RS01495 WP 003687645.1      SKYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAKKIIGSYQNTRYVEESLAILELAYKKLDKPQLAADTRR   239
consensus                                     !*!!*!!*!*!**!!!!!!!!!!!*!!!!!!!!!!!!*!!!!!******!*!!*!*!!*!!!***!!*!!**!*!**!**!

logo
NTDB id 293451 DK843 RS00955 WP 114072265.1   VLALNYPQSQYLNKPWKVEEMPWWKLWK   264
NTDB id 1102 NMB RS03635 WP 002244053.1       VLETNFPKSPFLKQPWRSDDMPWWRYWH   267
NTDB id 1106 NGFG RS01495 WP 003687645.1      VLETNFPKSPFLTHAWQPDDMPWWRYWH   267
consensus                                     !!**!*!*!**!   *!* **!!!!**!*
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X non conserved

X similar

X ≥ 50% conserved