```
logo
NTDB id 1016886 ABSY19 RS01495 WP 291119609.1  .......MASALVLSACSTTP.....VDETAGWSPNRIYSEATEERAAGNYEKSVTLYEKLEGRAAGTPLAQQAQLDKAY   68
NTDB id 1102 NMB RS03635 WP 002244053.1        MKKILLTVSLGLALSACATQGTVDKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHAQQSQLDTAY   80
NTDB id 1106 NGFG RS01495 WP 003687645.1       MKKILLTVSLGLALSACATQGTADKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHARQSQLDTAY   80
consensus                                      ***********!*!!!!*!*** ******!**!!****!*!!**!****!!***!*!!!*!!*!******!*!*!!!*!!

logo
NTDB id 1016886 ABSY19 RS01495 WP 291119609.1  AQYKAGEQAQALATLDRFQKLHPTSPAMDYALYLKGLVNFNDNLGLFGSLARQDLSERDQKAAKESFEAFRELVSRFPDS  148
NTDB id 1102 NMB RS03635 WP 002244053.1        AYYKDDEKDKALAAIDRFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNS  160
NTDB id 1106 NGFG RS01495 WP 003687645.1       AYYKDDEKDKALAAIERFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNS  160
consensus                                      !*!!**!***!!!***!!**!!!**!*!!!!!!!*!!!*!!********!!*!!*!*!!*!!**!***!!*!!!*!!!*!

logo
NTDB id 1016886 ABSY19 RS01495 WP 291119609.1  RYSPDARDRMTYIVNSLAQYEVHVARYYHSRGAYVAAINRAQTALTDYVNAPALEEALFIMLRSYEALGLTQLRDDTRRV  228
NTDB id 1102 NMB RS03635 WP 002244053.1        KYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAQKIIGSYQNTRYVEESLAILELAYKKLDKPRLAADTRRV  240
NTDB id 1106 NGFG RS01495 WP 003687645.1       KYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAKKIIGSYQNTRYVEESLAILELAYKKLDKPQLAADTRRV  240
consensus                                      *!**!!**!!***!**!***!**!!!!!**!!!!*!!*!!!******!*!****!!*!*!****!**!****!**!!!!!

logo
NTDB id 1016886 ABSY19 RS01495 WP 291119609.1  MQASYPNSEYLSQGFKSKDDPWWKIW.  254
NTDB id 1102 NMB RS03635 WP 002244053.1        LETNFPKSPFLKQPWRSDDMPWWRYWH  267
NTDB id 1106 NGFG RS01495 WP 003687645.1       LETNFPKSPFLTHAWQPDDMPWWRYWH  267
consensus                                      *****!*!**!*******!*!!!**!*
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X non conserved

X similar

X $\geq$ 50% conserved