```
logo
NTDB id 1129057 FX095 RS10855 WP 027823241.1   MKKIAVMMLMAAWLAGCSTT...SEPADETRGWIVEKLYSEARDELNSGNYTRAIKLYETLEARYPYGRYAQQAQMDLAY   77
NTDB id 1102 NMB RS03635 WP 002244053.1        MKKILLTVSLGLALSACATQGTVDKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHAQQSQLDTAY   80
NTDB id 1106 NGFG RS01495 WP 003687645.1       MKKILLTVSLGLALSACATQGTADKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHARQSQLDTAY   80
consensus                                      !!!!*********!**!*!*** ***!**!**!*!!!!!*!*!!!!!*!!!!!*!!!!*!!*!*!**!*!*!*!*!*!!

logo
NTDB id 1129057 FX095 RS10855 WP 027823241.1   AHFKDQEPALSLAAADRFIKLHPAHPNVDYVYYLKGLVNYNEDGGILSKYTGQDRAERDPKAAREAFTSFRDLVVRFPDS   157
NTDB id 1102 NMB RS03635 WP 002244053.1        AYYKDDEKDKALAAIDRFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNS   160
NTDB id 1106 NGFG RS01495 WP 003687645.1       AYYKDDEKDKALAAIERFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNS   160
consensus                                      !**!!*!****!!!**!!**!!!*!!!*!!**!!*!!!**!!!***!*!***!!***!!!!!*!!!***!**!!*!!!*!

logo
NTDB id 1129057 FX095 RS10855 WP 027823241.1   KYAPDARVKMQNLVDGLAEHELFVARYYMRRSAYLAAANRAQGMIKEFPDSPFVEESFAIMVTAYDKLGKTTLRDDTRRV   237
NTDB id 1102 NMB RS03635 WP 002244053.1        KYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAQKIIGSYQNTRYVEESLAILELAYKKLDKPRLAADTRRV   240
NTDB id 1106 NGFG RS01495 WP 003687645.1       KYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAKKIIGSYQNTRYVEESLAILELAYKKLDKPQLAADTRRV   240
consensus                                      !!!*!!***!**!!!*!***!*******!*!*!!*!!!!!!***!********!!!!*!!***!!*!!*!* !**!!!!!

logo
NTDB id 1129057 FX095 RS10855 WP 027823241.1   LETNFPNSVYLKKPWQPRDVPWYQFW.   263
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 ≥ 50% conserved