```
logo
NTDB id 312961 D1345 RS12250 WP 039753413.1   MKRYVVAAMLVMGLAGCATTETYD....ETRGWTVEKLYSEAHDELNSGNYTRAVKLYETLEARFPYGRYAQQAQMDLAY   76
NTDB id 1102 NMB RS03635 WP 002244053.1       MKKILLTVSLGLALSACATQGTVDKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHAQQSQLDTAY   80
NTDB id 1106 NGFG RS01495 WP 003687645.1      MKKILLTVSLGLALSACATQGTADKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHARQSQLDTAY   80
consensus                                     !!*******!***!**!!!**! !*****!**!*!!!!*!!*!!!!!*!!!!!!!!!!*!!*!!!**!*!*!*!*!*!!

logo
NTDB id 312961 D1345 RS12250 WP 039753413.1   THYKDNEPEQAIAAADRFIKLHPTHPNLDYLYYLKGLVFYNDDSGLLAKWAGQDMSERDPRAAREAFTAFRELVTRFPNS   156
NTDB id 1102 NMB RS03635 WP 002244053.1       AYYKDDEKDKALAAIDRFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNS   160
NTDB id 1106 NGFG RS01495 WP 003687645.1      AYYKDDEKDKALAAIERFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNS   160
consensus                                     **!!!*!***!*!!**!!**!!!*!!!*!!**!!*!!!**!*!***!*!*!*!!*!*!!!*!*!!!**!!*!!!*!!!!!

logo
NTDB id 312961 D1345 RS12250 WP 039753413.1   IYKEDASKKMERLLDALGGNEMHVARYYMKRGAYLAAANRAQGVVKSYANTKYPEEALGIMVAAYDKLSMPQLRDDAKRV   236
NTDB id 1102 NMB RS03635 WP 002244053.1       KYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAQKIIGSYQNTRYVEESLAILELAYKKLDKPRLAADTRRV   240
NTDB id 1106 NGFG RS01495 WP 003687645.1      KYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAKKIIGSYQNTRYVEESLAILELAYKKLDKPQLAADTRRV   240
consensus                                     *!**!!***!**!*!!!!!!!!!*!!!!!!!!!!!!*!!!!!!*****!!*!!*!*!!!*!*!***!!*!!**!*!**!**!!

logo
NTDB id 312961 D1345 RS12250 WP 039753413.1   LALNYPNSEYLKEGWSIEDMPWWKLWK   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