```
logo
NTDB id 240310 CGZ77 RS04900 WP 094030985.1  MKKILLVALLGAALSGCASKGTSDKDAQITQDWSVEKLYAEAQDELNSNNYTRAVKLYELLESRFPQGRYAQQAQLDTAY   80
NTDB id 1106 NGFG RS01495 WP 003687645.1     MKKILLTVSLGLALSACATQGTADKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHARQSQLDTAY   80
NTDB id 1102 NMB RS03635 WP 002244053.1      MKKILLTVSLGLALSACATQGTVDKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHAQQSQLDTAY   80
consensus                                    !!!!!!***!!*!!!*!!**!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!*!!!!!!**!*!*!*!!!!!!

logo
NTDB id 240310 CGZ77 RS04900 WP 094030985.1  AYYKDEEREKALAAVERFQKLHPQHPNMDYALYLKGLILFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVRRYAQS  160
NTDB id 1106 NGFG RS01495 WP 003687645.1     AYYKDDEKDKALAAIERFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNS  160
NTDB id 1102 NMB RS03635 WP 002244053.1      AYYKDDEKDKALAAIDRFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNS  160
consensus                                    !!!!!*!**!!!!!**!!**!!!!!!!!!!!!!!!*!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!***!

logo
NTDB id 240310 CGZ77 RS04900 WP 094030985.1  KYVEDATKRMTKLVDALAGNEMAVARYYAKRGAYLAAANRAQNIVTGFQNTRYVEEALAVMELSYRKMERPQLADDTRRI  240
NTDB id 1106 NGFG RS01495 WP 003687645.1     KYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAKKIIGSYQNTRYVEESLAILELAYKKLDKPQLAADTRRV  240
NTDB id 1102 NMB RS03635 WP 002244053.1      KYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAQKIIGSYQNTRYVEESLAILELAYKKLDKPRLAADTRRV  240
consensus                                    !!**!!!*!!*!!!!!!*!!!!*!!!!!*!!!!!*!!!!!!**!****!!!!!!!!*!!**!!*!*!***!*!!*!!!!!*

logo
NTDB id 240310 CGZ77 RS04900 WP 094030985.1  LEKNFPQSPYLTRSWKDNDAPWWRYWK  267
NTDB id 1106 NGFG RS01495 WP 003687645.1     LETNFPKSPFLTHAWQPDDMPWWRYWH  267
NTDB id 1102 NMB RS03635 WP 002244053.1      LETNFPKSPFLKQPWRSDDMPWWRYWH  267
consensus                                    !!*!!!*!!*!*  !* *!*!!!!!!*
```

X non conserved
X similar
X ≥ 50% conserved