```
logo
NTDB id 24665 TBD RS04105 WP 337998338.1  ......MLAAALALGGCGLL...PEVKDETTGWSAQKLYAEAKDNLNEGNYERAVKLFETLESRYPFGRYAQQAQLEVAYAYYKD   76
NTDB id 1102 NMB RS03635 WP 002244053.1   MKKILLTVSLGLALSACATQGTVDKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHAQQSQLDTAYAYYKD   85
NTDB id 1106 NGFG RS01495 WP 003687645.1  MKKILLTVSLGLALSACATQGTADKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHARQSQLDTAYAYYKD   85
consensus                                 ***********!!!**!***** ******!**!!**!!!!!!*!*!!**!!*!!!!!*!*!!!!*!**!*!*!*!!**!!!!!!!

logo
NTDB id 24665 TBD RS04105 WP 337998338.1  NEPISAVAACDRFIKLHPNHPNVDYAYYLKGLANFNDDLGLLGNLVDQDMSERDPRAARDAFLAFKELATRFPQSIYAADATARM  161
NTDB id 1102 NMB RS03635 WP 002244053.1   DEKDKALAAIDRFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNSKYAADATARM  170
NTDB id 1106 NGFG RS01495 WP 003687645.1  DEKDKALAAIERFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNSKYAADATARM  170
consensus                                 *!***!*!!**!!**!!!*!!!*!!!*!!*!!**!!*!***!**!**!!*!*!!!*!*!*!**!!*!!**!!!*!*!!!!!!!!!

logo
NTDB id 24665 TBD RS04105 WP 337998338.1  KYLVNALANNEVHVAKYYLKRKAYVAAANRAKEVLKTYPEAPALEEALAIMALSYDRLKLPELRDDARRVLTLNFPNSKYLQGV.  245
NTDB id 1102 NMB RS03635 WP 002244053.1   VKLVDALGGNEMSVARYYMKRGAYIAAANRAQKIIGSYQNTRYVEESLAILELAYKKLDKPRLAADTRRVLETNFPKSPFLKQPW  255
NTDB id 1106 NGFG RS01495 WP 003687645.1  VKLVDALGGNEMSVARYYMKRGAYIAAANRAKKIIGSYQNTRYVEESLAILELAYKKLDKPQLAADTRRVLETNFPKSPFLTHAW  255
consensus                                 **!!*!!**!!**!!*!!*!!*!!*!!!!!!!******!******!!*!!!**!*!**!**!  !**!*!!!!**!!!*!**!    *

logo
NTDB id 24665 TBD RS04105 WP 337998338.1  SIKEKAWYRFW.   256
NTDB id 1102 NMB RS03635 WP 002244053.1   RSDDMPWWRYWH   267
NTDB id 1106 NGFG RS01495 WP 003687645.1  QPDDMPWWRYWH   267
consensus                                   ****!*!*!*
```

X non conserved

X similar

X $\geq$ 50% conserved