```
logo

NTDB id 153899 F822 RS12470 WP 025042045.1  MSR.SVALFLVLLLSAC...GLLSKEVKDSKNWSASKYYSEAKSELNEGNYGGAIKLFEALEARYPYGRFAQQAQLEVAYAYYKD   81
NTDB id 1102 NMB RS03635 WP 002244053.1     MKKILLTVSLGLALSACATQGTVDKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHAQQSQLDTAYAYYKD   85
NTDB id 1106 NGFG RS01495 WP 003687645.1    MKKILLTVSLGLALSACATQGTADKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHARQSQLDTAYAYYKD   85
consensus                                   !********!*!*!!!!***!***!*******!!**!*!*!!**!!!**!!**!*!!*!*!!*!*!**!*!*!*!!**!!!!!!!

logo

NTDB id 153899 F822 RS12470 WP 025042045.1  GEPATAIAAADRFIKLHPNHTNVDYAYYLKGLSNFNDDLGLMGIVSEKILNQDMSERDPKASRESFENFKELVSRYPKSKYAADA  166
NTDB id 1102 NMB RS03635 WP 002244053.1     DEKDKALAAIDRFRRLHPQHPNMDYALYLRGLVLFNEDQSFLN....KLASQDWSDRDPKANREAYQAFAELVQRFPNSKYAADA  166
NTDB id 1106 NGFG RS01495 WP 003687645.1    DEKDKALAAIERFRRLHPQHPNMDYALYLRGLVLFNEDQSFLN....KLASQDWSDRDPKANREAYQAFAELVQRFPNSKYAADA  166
consensus                                   *!***!*!!**!!**!!!*!*!*!!!*!!*!!**!!*!*****    !***!!*!*!!!!!*!!****!*!!!*!*!*!!!!!!!

logo

NTDB id 153899 F822 RS12470 WP 025042045.1  VQRMKHLVNVVALNEVQVARYYIKRGGYVAAANRAQYALKEYPQTPATEEALFIMMKSYDALGMTDLRDDAHRVMKKNFPDSRFF  251
NTDB id 1102 NMB RS03635 WP 002244053.1     TARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAQKIIGSYQNTRYVEESLAILELAYKKLDKPRLAADTRRVLETNFPKSPFL  251
NTDB id 1106 NGFG RS01495 WP 003687645.1    TARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAKKIIGSYQNTRYVEESLAILELAYKKLDKPQLAADTRRVLETNFPKSPFL  251
consensus                                   **!!**!!*****!!**!!!!!*!!!*!*!!!!!!!******!**!***!!*!*!****!**!***  !**!**!!***!!!*!*!*

logo

NTDB id 153899 F822 RS12470 WP 025042045.1  EDSTGKSAEPWWRFW.   266
NTDB id 1102 NMB RS03635 WP 002244053.1     KQPWRSDDMPWWRYWH   267
NTDB id 1106 NGFG RS01495 WP 003687645.1    THAWQPDDMPWWRYWH   267
consensus                                      *  ***!!!!*!*
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X non conserved

X similar

X ≥ 50% conserved