```
logo
NTDB id 310559 D0T92 RS07655 WP 151051705.1   MKKILLVISLGLALAGCAGTTGTIDKDAQITQEWTVERLYAEAHDELNSSNYTRAIKLYEILESRFPNGRYAQQAQLDTA   80
NTDB id 1102 NMB RS03635 WP 002244053.1       MKKILLTVSLGLALSACA.TQGTVDKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHAQQSQLDTA   79
NTDB id 1106 NGFG RS01495 WP 003687645.1      MKKILLTVSLGLALSACA.TQGTADKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHARQSQLDTA   79
consensus                                     !!!!!!**!!!!!!**!! !*!!*!!!!!!!*!*!!*!!!!!*!!!!!!!!!!!*!!!!!!!!!!!**!*!*!*!!!!!

logo
NTDB id 310559 D0T92 RS07655 WP 151051705.1   YAYYKDDEPEKALAAVERFQRIHPRHPNMDYALYLKGLILFNEDQSFLNRLASQDWSDRDPKANRDAYQAFSQLIELYPN   160
NTDB id 1102 NMB RS03635 WP 002244053.1       YAYYKDDEKDKALAAIDRFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPN   159
NTDB id 1106 NGFG RS01495 WP 003687645.1      YAYYKDDEKDKALAAIERFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPN   159
consensus                                     !!!!!!!!**!!!!!**!!*!*!!*!!!!!!!!!!*!!*!!!!!!!!!!*!!!!!!!!!!!!!!!!!*!!!!!**!****!!

logo
NTDB id 310559 D0T92 RS07655 WP 151051705.1   SRYAPDAAERMTKLVDALGGNEMAVARYYMKRGAYLAAANRAQKIVERYQNTRFVEESLAIMEASYRKLNKPQLADDTRR   240
NTDB id 1102 NMB RS03635 WP 002244053.1       SKYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAQKIIGSYQNTRYVEESLAILELAYKKLDKPRLAADTRR   239
NTDB id 1106 NGFG RS01495 WP 003687645.1      SKYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAKKIIGSYQNTRYVEESLAILELAYKKLDKPQLAADTRR   239
consensus                                     !*!!*!!**!!*!!!!!!!!!!!!*!!!!!!!!!!!!*!!!!!!*!!***!!!!!*!!!!!!!*!**!*!!*!!*!!*!!!!

logo
NTDB id 310559 D0T92 RS07655 WP 151051705.1   ILAQNFPNSPYLQKSWQENSSIPWWRYWK   269
NTDB id 1102 NMB RS03635 WP 002244053.1       VLETNFPKSPFLKQPWRSD.DMPWWRYWH   267
NTDB id 1106 NGFG RS01495 WP 003687645.1      VLETNFPKSPFLTHAWQPD.DMPWWRYWH   267
consensus                                     *!**!!!*!!*!   !* * **!!!!!!*
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X non conserved

X similar

X ≥ 50% conserved