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logo

NTDB id 39075 NLA RS07955 WP 013449289.1   MKKILLMVSLSLALCACASSSQGTVDKDTQTTQGWGVEKLYAEAQDELGGGNYTRAVKLYEILGSRFPTGRYAQQALLDTAYAYYK   85
NTDB id 1106 NGFG RS01495 WP 003687645.1   MKKILLTVSLGLALSACA.TQGTADKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHARQSQLDTAYAYYK   84
NTDB id 1102 NMB RS03635 WP 002244053.1    MKKILLTVSLGLALSACA.TQGTVDKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHAQQSQLDTAYAYYK   84
consensus                                  !!!!!!*!!!*!!!*!!!  *!!!*!!!*!*!!*!*!!!!!!!!!!!!***!!!!!!!!!!!!*!!!!!*!*!*!**!!!!!!!!!!

logo

NTDB id 39075 NLA RS07955 WP 013449289.1   DDEKDKALAAIDRFRNLYPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVRRFPDSKYAADASAR   170
NTDB id 1106 NGFG RS01495 WP 003687645.1   DDEKDKALAAIERFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNSKYAADATAR   169
NTDB id 1102 NMB RS03635 WP 002244053.1    DDEKDKALAAIDRFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNSKYAADATAR   169
consensus                                  !!!!!!!!!!!*!!!*!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!*!!!!!!!*!!

logo

NTDB id 39075 NLA RS07955 WP 013449289.1   MIKLVDALGGNEMAVARYYMKRGAYIAAANRAQKIIGSYQNTRYVEESLAILELAYQKLGKPQLAADTRRVLETNFPKSPFLTHA   255
NTDB id 1106 NGFG RS01495 WP 003687645.1   MVKLVDALGGNEMSVARYYMKRGAYIAAANRAKKIIGSYQNTRYVEESLAILELAYKKLDKPQLAADTRRVLETNFPKSPFLTHA   254
NTDB id 1102 NMB RS03635 WP 002244053.1    MVKLVDALGGNEMSVARYYMKRGAYIAAANRAQKIIGSYQNTRYVEESLAILELAYKKLDKPRLAADTRRVLETNFPKSPFLKQP   254
consensus                                  !*!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!
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