logo

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NTDB id 251299 CRN81 RS09970 WP 011135741.1  MKRYVVAAMLVMVGLAGCATTETY....DETRGWTVEKLYSEAHDELNSGNYTRAVKLYETLEARFPYGRYAQQAQMDLA   76
NTDB id 1102 NMB RS03635 WP 002244053.1      MKKILL.TVSLGLALSACATQGTVDKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHAQQSQLDTA   79
NTDB id 1106 NGFG RS01495 WP 003687645.1     MKKILL.TVSLGLALSACATQGTADKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHARQSQLDTA   79
consensus                                    !!**** *******!**!!!**! ******!**!*!!!!!*!!*!!!!!*!!!!!!!!!!*!!*!!!**!*!*!*!*!*!
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logo

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NTDB id 251299 CRN81 RS09970 WP 011135741.1  YTHYKDGEPELAIASADRFIKLHPTHPNLDYIYYLKGLVYYNDDSGLLAKWAGQDMSERDPRAAREAFAAFRELTTRFPS   156
NTDB id 1102 NMB RS03635 WP 002244053.1      YAYYKDDEKDKALAAIDRFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPN   159
NTDB id 1106 NGFG RS01495 WP 003687645.1     YAYYKDDEKDKALAAIERFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPN   159
consensus                                    !**!!!*!***!*!***!!**!!!*!!!*!!**!!*!!!**!*!***!*!*!*!!*!*!!!*!*!!!**!!*!!**!!!*
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logo

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NTDB id 251299 CRN81 RS09970 WP 011135741.1  SSYAPDAKAKMIRLVDALGGNEMHVARYYMKRGAYLAAANRAQGVVKDYANTKYPEEALAIMVAAYDKLQLPQLRDDARR   236
NTDB id 1102 NMB RS03635 WP 002244053.1      SKYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAQKIIGSYQNTRYVEESLAILELAYKKLDKPRLAADTRR   239
NTDB id 1106 NGFG RS01495 WP 003687645.1     SKYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAKKIIGSYQNTRYVEESLAILELAYKKLDKPQLAADTRR   239
consensus                                    !*!!*!!*!*!**!!!!!!!!!!!*!!!!!!!!!!!!*!!!!!!******!*!!*!*!!*!!!***!!*!!**!*!**!*!!
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logo

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NTDB id 251299 CRN81 RS09970 WP 011135741.1  VLALNYPQSQYLGKSWTVEEMPWWKLWK   264
NTDB id 1102 NMB RS03635 WP 002244053.1      VLETNFPKSPFLKQPWRSDDMPWWRYWH   267
NTDB id 1106 NGFG RS01495 WP 003687645.1     VLETNFPKSPFLTHAWQPDDMPWWRYWH   267
consensus                                    !!**!*!*!**!    !  **!!!!**!*
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X non conserved

X similar

X ≥ 50% conserved