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logo
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NTDB id 61846 SCR2 RS08560 WP 020998065.1  MNFEKIEQAYSLLLENVQELQNTLLTNFYDALVEQNGVYLDGNTELDIVKKNNETLKSLKLDKEEWRRAYQFLLMKAAQTEPLQA  85
NTDB id 401 FSA28 RS09370 WP 002291646.1   MDFEKIEAAYNLLLDNCQRLETQLHTHLYDALIEQNACYLGAEGADDFIKENNKKLRQLNLSKEEWRRSFQFIFIKATQSQQLQS  85
NTDB id 392 SMU RS09000 WP 002263442.1     MDFEKIEAAYNLLLDNCQRLETQLHTHLYDALIEQNACYLGAEGADDFIKENNKKLRQLNLSKEEWRRSFQFIFIKATQSQQLQS  85
consensus                                  !*!!!!!*!!*!!!*!*!*!***!*!**!!!!*!!!**!!******!**!*!*!**!**!*!*!!!!!!**!!***!!*!***!!*
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logo
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NTDB id 61846 SCR2 RS08560 WP 020998065.1  NHQFTPDSIGFIVTFLIDQLSNKEHIDVLEIGSGMGNLAETILNNTNKNVDYLGLELDDLLIDISASIADVMDAKVSFVQGDAVR  170
NTDB id 401 FSA28 RS09370 WP 002291646.1   NHQFTPDSIGFIILYLLEGLTNKSQLDIIEIGSGTGNLAETIVNNSSKSIDYIGIEVDDLLIDLSASIADVLDSSVHFVQEDAVR  170
NTDB id 392 SMU RS09000 WP 002263442.1     NHQFTPDSIGFIILYLLEGLTNKSQLDIIEIGSGTGNLAETIVNNSSKSIDYMGIEVDDLLIDLSASIADVLDSSVHFVQEDAVR  170
consensus                                  !!!!!!!!!!!!!***!***!*!!***!**!!!!!*!!!!!!!*!!**!**!!*!*!*!!!!!!*!!!!!!!*!**!*!!!*!!!!
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logo
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NTDB id 61846 SCR2 RS08560 WP 020998065.1  PQVLKESDVIISDLPVGFYPDDNIATRYEVASRQEHTYAHHLLMEQSLKYLKKSDGYAIFLAPNDLLTSTQSDLLKNWLQNHAQIV  255
NTDB id 401 FSA28 RS09370 WP 002291646.1   PQILKESDVIISDLPVGFYPNDKIAQRYQVAAKLGHTYAHHLLMEQSLKYLKKEGLAIFLAPVDLLTSEQSPLLKKWLQNNVTIL  255
NTDB id 392 SMU RS09000 WP 002263442.1     PQILKESDIIISDLPVGFYPNDKIAQRYQVAAKLGHTYAHHLLMEQSLKYLKKEGLAIFLAPVDLLTSEQSPLLKKWLQNNVTIL  255
consensus                                  !!*!!!!!*!!!!!!!!!!!!*!*!!*!!*!!****!!!!!!!!!!!!!!!!!!**!*!!!!!*!!!!!*!!*!!!*!!!***!*
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logo
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NTDB id 61846 SCR2 RS08560 WP 020998065.1  AMIALPESLFGNAAYAKTIFVLKKQDEQAVQPFVYALSDLQNQADLLTFSEKFQNWSQESEI  317
NTDB id 401 FSA28 RS09370 WP 002291646.1   AVISLPEAIFKQKSNIKSIFVLKKQSQQSLESFVYPLTDLQDPEVLRRFMKNFKKWKEDNVF  317
NTDB id 392 SMU RS09000 WP 002263442.1     AVISLPEAIFKQKSNIKSIFVLKKQSQQSLESFVYPLTDLQDPEVLRRFMKNFKKWKEDNVF  317
consensus                                  !*!*!!!**!******!*!!!!!!!**!****!!!*!*!!!****!**!***!**!******
```

X non conserved
X similar
X ≥ 50% conserved