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logo
NTDB id 1021 ACIAD RS00975 WP 004930462.1   MITILSSHHYHTIKVWYVLQHSLSGFKKLIHYFGDCETALQPQRLSDWQKLGLHANHLKRLEEVHTLQGQQQFEKMLAQLQTHTD   85
NTDB id 37706 AOLE RS18580 WP 013199175.1   MLTQLSSHHYHTLKVWYLVQHSLVSFKKIVNYFGSCEKAVQPAGLSQWRNLGLHANHLKRVDEFQTAQGQAQFEKLIEQIQQYTD   85
NTDB id 1073 ABD1 RS00865 WP 015451369.1    MLTQLSSHHYHTLKVWYLVQHSLVSFKKIIDYFGNCEKATQPDGLAKWSSLGLHANHLKRVNEFQTTQGQAQFEQLVQQVRQHTD   85
NTDB id 1043 HON27 RS16615 WP 168727007.1   MLTQLSSHQYHTLKVWYLVQHSLVSFKKIIDYFGNCEKATQPDGLAKWSSLGLHANHLKRVNEFQTPQGQAQFEQLVQQVRQHTD   85
consensus                                   !*!*!!!!*!!!*!!!!**!!!!**!!!***!!!*!!*!*!!**! *!**!!!!!!!!!!**!**! !!!*!!! ***!* **!!

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NTDB id 1021 ACIAD RS00975 WP 004930462.1   FIVTPDDPAYPSQLIPYQDRPPILFGQGSIQSLIQPQIAVVGSRKPSPHGRQVAYDFSYFLSEKGFYITSGLAHGIDEAAHQGSL   170
NTDB id 37706 AOLE RS18580 WP 013199175.1   FILTSEDSGYPTQLLPYTDHPPIIFGKGQAQALLQPQIAIVGSRKPSPHGRQVAYDFAYYLSEKGFYVSSGLAYGIDEAAHQGAS   170
NTDB id 1073 ABD1 RS00865 WP 015451369.1    FILTPDDSGYPTQLLPYTDHPPIIFGKGQAQALLQPQIAIVGSRKPSPHGRQVAYDFAYYLSEKGFFISSGLAYGIDEAAHQGAS   170
NTDB id 1043 HON27 RS16615 WP 168727007.1   FILTPDDSGYPTQLLPYTDHPPIIFGKGQAQALLQPQIAIVGSRKPSPHGRQVAYDFAYYLSEKGFFISSGLAYGIDEAAHQGAS   170
consensus                                   !!*!**!**!!*!!*!!*!*!!!*!!*!**!*!*!!!!!*!!!!!!!!!!!!!!!!!!*!*!!!!!!***!!!!*!!!!!!!!!**

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NTDB id 1021 ACIAD RS00975 WP 004930462.1   VHHRAIAVVGTGLDLCYPSGNKALKQHIRDQGGAIVSEFLPGTPPLQAHFPRRNRIVSGLSLGVLVVEATLKSGSLITANKAAEQ   255
NTDB id 37706 AOLE RS18580 WP 013199175.1   AHQRTIAVTGTGLDSTYPAQNKKLAEQILAQNGAIITEFLPGTPPLQQHFPRRNRIVSGLSLGVLVVEATLKSGSLITANKAAEQ   255
NTDB id 1073 ABD1 RS00865 WP 015451369.1    AHQRTIAVTGTGLDTTYPAQNKKLAEYILAQNGAIITEFLPGTPPLQQHFPRRNRIVSGLSLGVLVVEATLKSGSLITANKAAEQ   255
NTDB id 1043 HON27 RS16615 WP 168727007.1   THQRTIAVTGTGLDTTYPAQNKKLAEHILAKNGAIITEFLPGTPPLQQHFPRRNRIVSGLSLGVLVVEATLKSGSLITANKAAEQ   255
consensus                                   *!*!*!!!*!!!!!**!!**!!*!***!****!!**!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

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NTDB id 1021 ACIAD RS00975 WP 004930462.1   GKVVFAIPGHIYSEFHQGCHQLIREGAILVDHPEQIIEDLALPTHWHSQSTASGEANADDSGKTDDADLPDDLLPLYQLLDWQGQ   340
NTDB id 37706 AOLE RS18580 WP 013199175.1   GKTVFAIPGHIYSEFHQGCHQLIREGAILVDHPEQIIEDLALPTQWQSQQQNSTE....EVVVSCAPEIPEHLINLYQSLDWVGQ   336
NTDB id 1073 ABD1 RS00865 WP 015451369.1    GKTVFAIPGHIYSEFHQGCHQLIREGAILIDHPEQIIEDLALPTQWQSQQQNQTE....EAN.TNTPEIREHLIDLYQSLDWVGQ   335
NTDB id 1043 HON27 RS16615 WP 168727007.1   GKTVFAIPGHIYSEFHQGCHQLIREGAILIDHPEQIIEDLALPTQWQSQQQNQTE....ETN.TNTPEIPEHLIDLYQSLDWVGQ   335
consensus                                   !!*!!!!!!!!!!!!!!!!!!!
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