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logo
NTDB id 313885 CDG61 RS16590 WP 087551149.1   MLNPLSQAQIETITLWYLVQHSLSAFYKLKQHFGSVHLAIAPKGLAQWSQLNLHKNHIQRAQEFQTDAGQQKFAELLQSI   80
NTDB id 1043 HON27 RS16615 WP 168727007.1     MLTQLSSHQYHTLKVWYLVQHSLVSFKKIIDYFGNCEKATQPDGLAKWSSLGLHANHLKRVNEFQTPQGQAQFEQLVQQV   80
NTDB id 1073 ABD1 RS00865 WP 015451369.1     MLTQLSSHHYHTLKVWYLVQHSLVSFKKIIDYFGNCEKATQPDGLAKWSSLGLHANHLKRVNEFQTTQGQAQFEQLVQQV   80
NTDB id 1021 ACIAD RS00975 WP 004930462.1    MTTILSSHHYHTIKVWYVLQHSLSGFKKLIHYFGDCETALQPQRLSDWQKLGLHANHLKRLEEVHTLQGQQQFEKMLAQL   80
consensus                                    !***!!*****!***!!**!!!! *!*!****!!****!**!**!**!**!*!!*!!**!**!**! *!! *!*******

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NTDB id 313885 CDG61 RS16590 WP 087551149.1   QSHTDFVLIPESTDYPAQLLPYTSPPPILFGQGNAQALLQPQIAIVGSRKPSPHGRQIAYDFAFYLSEKGFYINSGLAQG   160
NTDB id 1043 HON27 RS16615 WP 168727007.1     RQHTDFILTPDDSGYPTQLLPYTDHPPIIFGKGQAQALLQPQIAIVGSRKPSPHGRQVAYDFAYYLSEKGFFISSGLAYG   160
NTDB id 1073 ABD1 RS00865 WP 015451369.1     RQHTDFILTPDDSGYPTQLLPYTDHPPIIFGKGQAQALLQPQIAIVGSRKPSPHGRQVAYDFAYYLSEKGFFISSGLAYG   160
NTDB id 1021 ACIAD RS00975 WP 004930462.1    QTHTDFIVTPDDPAYPSQLIPYQDRPPILFGQGSIQSLIQPQIAVVGSRKPSPHGRQVAYDFSYFLSEKGFYITSGLAHG   160
consensus                                     *!!!!***!****!!*!!*!!***!!!*!! !**!*!*!!!!!*!!!!!!!!!!!!*!!!!***!!!!!!*!*!!!!*!

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NTDB id 313885 CDG61 RS16590 WP 087551149.1   IDEAAHKGGLKHQRTIAVMGTGLDQTYPTAHTQLRQQITEQRGAVITEFLPGTLPLQQHFPRRNRIVSGLSLGVIVAEAS   240
NTDB id 1043 HON27 RS16615 WP 168727007.1     IDEAAHQGASTHQRTIAVTGTGLDTTYPAQNKKLAEHILAKNGAIITEFLPGTPPLQQHFPRRNRIVSGLSLGVLVVEAT   240
NTDB id 1073 ABD1 RS00865 WP 015451369.1     IDEAAHQGASAHQRTIAVTGTGLDTTYPAQNKKLAEYILAQNGAIITEFLPGTPPLQQHFPRRNRIVSGLSLGVLVVEAT   240
NTDB id 1021 ACIAD RS00975 WP 004930462.1    IDEAAHQGSLVHHRAIAVVGTGLDLCYPSGNKALKQHIRDQGGAIVSEFLPGTPPLQAHFPRRNRIVSGLSLGVLVVEAT   240
consensus                                    !!!!!!*!*  !*!*!!!*!!!!!**!!*****!* *!****!!***!!!!!!*!!!*!!!!!!!!!!!!!!!!*!*!!*

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NTDB id 313885 CDG61 RS16590 WP 087551149.1   LNSGSLITAGLAAEQGKTVFAIPGHIYSEFHQGCHQLIREGAILIDHPEQVIEDLALPTQWMVQQKKET......STVPE   314
NTDB id 1043 HON27 RS16615 WP 168727007.1     LKSGSLITANKAAEQGKTVFAIPGHIYSEFHQGCHQLIREGAILIDHPEQIIEDLALPTQWQSQQQNQT.....EETNTN   315
NTDB id 1073 ABD1 RS00865 WP 015451369.1     LKSGSLITANKAAEQGKTVFAIPGHIYSEFHQGCHQLIREGAILIDHPEQIIEDLALPTQWQSQQQNQT.....EEANTN   315
NTDB id 1021 ACIAD RS00975 WP 004930462.1    LKSGSLITANKAAEQGKVVFAIPGHIYSEFHQGCHQLIREGAILVDHPEQIIEDLALPTHWHSQSTASGEANADDSGKTD   320
consensus                                    !*!!!!!!!**!!!!!!*!!!!!!!!!!!!!!!!
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