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logo
NTDB id 1090397 ACJQW4 RS04395 WP 241626590.1  MLKPLSQVQMDTITLWYLVQHSLSSFYKLEQHFGSIHNAIQPESLLLWPGLGLHKNHLQRAQQFYTPAEQSKFQRIIENI   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 1090397 ACJQW4 RS04395 WP 241626590.1  QQHADFILCYQDPEYPRQLLPYSDKPPILFGQGQLSLLMQPQIAIVGSRKPSPHGRQIAYDFSYYLSEKGFVINSGLAQG  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 1090397 ACJQW4 RS04395 WP 241626590.1  IDEAAHQGALPHQRTLAVMGTGLDQTYPSQHQQLRQQILQQSGTIITEFLPHTPPLQHHFPRRNRIVSALSLGVIVAEAS  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 1090397 ACJQW4 RS04395 WP 241626590.1  LNSGSIITAKLAAEQGKLVFAIPGHIYSEFHQGCHQLIREGAILVDHPEQVLEDLALPTQWQAQQQAHDESL..TTSPIV  318
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      LKSGSLITANKAAEQGKVVFAIPGHIYSEFHQGCHQLIREGAILVDHPEQIIEDLALPTHWHSQSTASGEANADDSGKID  320
consensus                                      !*!!!*!!!**!!!!!!*!!!!!!!!!!!!
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