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logo
NTDB id 422145 G0028 RS01265 WP 180046515.1   MLNFISDVQLDTIKLWYLVQHSNLAFRSLETYFGSVAKAVSSENLLRWSEVNIHKNHLQRAEEFLTDKGQTSFYLCLEKL   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 422145 G0028 RS01265 WP 180046515.1   KQHCDFILLDHDSHYPQQLLPYADHPPILFGQGAWQNLTQPQVAIVGSRKPSPHGKQVAYDFAYYLSEKGFFITSGLAEG  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 422145 G0028 RS01265 WP 180046515.1   IDAAAHAGALKNLRTIAVIGTGLEQVYPTQNRQLHAQILEHAGTIISLYLPNTPPLKSNFPRRNRIVSGLSLGVLVAEAA  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 422145 G0028 RS01265 WP 180046515.1   LESGSLITAQAAAEQGKIIFAIPGHIYSEYHQGCHQLIREGAILVDHPEQIIEDLALPTQWHCENTQKI......EEKEK  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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