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logo

NTDB id 310704 BEN71 RS17900 WP 068975660.1  MLNQIPDDQFNTIKLWYLVQHSLSAFRKLETYFGSIQQAIQAENLARWSELKIHKSHIERANEYLSSNGEAKFQICIDKI   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    MITILSSHHYHTIKVWYVLQHSLSGFKKLIHYFGDCETALQPQRLSDWQKLGLHANHLKRLEEVHTLQGQQQFEKMLAQL   80
consensus                                    !**********!*!*!!**!!!!  *!*!***!!!****!*!***!**!**!**!**!**!**!****!***!*******

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NTDB id 310704 BEN71 RS17900 WP 068975660.1  KQHTDFILLDTDPHYPQQLLAFDNHPPILFGKGSFQNLSQPQIAIVGSRKPSPHGKQVAYDFAYYLSEKGFFITSGLAQG  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 310704 BEN71 RS17900 WP 068975660.1  IDEAAHMGGLKNHRTIAVIGTGLDLIYPSQNKHVQEQILQNSGTVITEYLPETQPIQRNFPYRNRIVSGLSLGVLVAEAQ  240
NTDB id 1043 HON27 RS16615 WP 168727007.1    IDEAAHQGASTHQRTIAVTGTGLDTTYPAQNKKLAEHILAKNGAITEFLPGTPPLQQHFPRRNRIVSGLSLGVLVVEAT  240
NTDB id 1073 ABD1 RS00865 WP 015451369.1     IDEAAHQGASAHQRTIAVTGTGLDTTYPAQNKKLAEYILAQNGAITEFLPGTPPLQQHFPRRNRIVSGLSLGVLVVEAT  240
NTDB id 1021 ACIAD RS00975 WP 004930462.1    IDEAAHQGSLVHHRAIAVVGTGLDLCYPSGNKALKQHIRDQGGAIVSEFLPGTPPLQAHFPRRNRIVSGLSLGVLVVEAT  240
consensus                                    !!!!!!*!*  **!*!!!*!!!!! *!! *!!*****!****!****!*!!*!*!*!**!!*!!!!!!!!!!!!!!!*!!*

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NTDB id 310704 BEN71 RS17900 WP 068975660.1  LGSGSLITAQTAAEQGKIIFAIPGHIYGEHHQGCHQLIREGAILVDHPEQIVEDLALPTQWHCEQHVENTEK..RTEVRL  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    LKSGSLITANKAAEQGKVVFAIPGHIYSEFHQGCHQLIREGAILVDHPEQIIEDLALPTHWHSQSTASGEANADDSGKTD  320
consensus                                    !*!!!!!!!**!!!!!!*!!!!!!!!!!*!*!!!!!!!!!!!!!!!!*!!!!!!*!!!!!!!*!********     ******

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NTDB id 310704 BEN71 RS17900 WP 068975660.1  SPEIPAHLFSLYQLLDWIGQDLDSLVEKSQLNTAELTSYLMELELLGLCIQQFGRYLRCRASH  381
NTDB id 1043 HON27 RS16615 WP 168727007.1    TPEIPEHLIDLYQSLDWVGQNIDQLVVHHNIPVSELTSSLMELELLGLCMQQSGLYLRCRS..  376
NTDB id 1073 ABD1 RS00865 WP 015451369.1     TPEIREHLIDLYQSLDWVGQNIDQLVVHHNIPVSELTSSLMELELLGLCMQQSGLYLRCRS..  376
NTDB id 1021 ACIAD RS00975 WP 004930462.1    DADLPDDLLPLYQLLDWQGQDLDQLAQHYQGSTAELTAQLMELELLGVCIQQSGRYLRCRSPY  383
consensus                                    *******!**!!!  !!!*!!  *!*!*******  !!!**!!!!!!!!*!*!!*!  !!!!!*
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X non conserved

X similar

X ≥ 50% conserved