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logo

NTDB id 375 SMU RS06880 WP 002262929.1    MTNVFESSPLFLRILLAVLIILLFF.YFIFLNYREYKNNNQVKQLNAKVRSLITGHYTDKLKVEDNSDLSELVNNVNDLS    79
NTDB id 1141439 DQN42 RS03875 WP 003072979.1  MIDNIRQ.FVVSKDFVFVLLILGFILVITLLSLENRRDNVRIRQLNQKVKDLIAGDYSEVLDMQGGPEITDITNSINDLS    79
NTDB id 414 AAK55818.1 838..2187( )          MLDLLKQ.TIFTRDFIFILILLGFIVVTLLLLENRRDNIQLKQINQKVKDLIAGDYSKVLDMQGGSEITNITNNUNDLS    79
consensus

logo

NTDB id 375 SMU RS06880 WP 002262929.1    EVFRLTHENLAQEKNRLTSILSYMTDGVLATDRSGKITVINDMAQKQLNVTREQALECNILDILD.DDSYTYNDLITKTP    158
NTDB id 1141439 DQN42 RS03875 WP 003072979.1  EVIRLTHENLEQETKRLSSILSYMTDGVLATNRRGQIIMINDMATKQLGVKESDARKMNILKLLGIEDEYDLRDLITKVP    159
NTDB id 414 AAK55818.1 838..2187( )          EVIRLTQENLEQESKRLNSILFYMTDGVLATNRRGQIIMINDTAKKQLGVKEDVLNRSILELLKIEPNYELRDLITQSP    159
consensus

logo

NTDB id 375 SMU RS06880 WP 002262929.1    EIVLTRRDEYDEFITLRIRFALNRRESGFISGLIAVLHDATEQEKEERERRLFVSNVSHELRTPLTSVKSYLEALDDGAL    238
NTDB id 1141439 DQN42 RS03875 WP 003072979.1  ELTIDSQDENGEFLSLRVRFALIRRESGFISGLVAVLHDTTEQEKEERERRLFVSNVSHELRTPLTSVKSYLEALDEGAL    239
NTDB id 414 AAK55818.1 838..2187( )          ELLLDSQDINGEYLNLRVRFALIRRESGFISGLVAVLHDTTEQEKEERERRLFVSNVSHELRTPLTSVKSYLEALDEGAL    239
consensus

logo

NTDB id 375 SMU RS06880 WP 002262929.1    TESVAPSFIKVSLDETNRMMRMITDLLSLSRIDNQTSHLDVELTNFTAFMNYILDRFDQIQSQQSTNKVYEIIRDYPDKS    318
NTDB id 1141439 DQN42 RS03875 WP 003072979.1  SEPVAPDFIKVSLDETNRMMRMVSDLLSLSRIDNATSHLDIELTNFTAFITFILNRFDKIRSQN.DDKKYEIIRDYPINS    318
NTDB id 414 AAK55818.1 838..2187( )          CETVAPDFIKVSLDETNRMMRMVTDLLHLSCIDNATSHLDVELINFTAFITFILNRFDKMKGQE.KEKKYELVRDYPINS    318
consensus

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NTDB id 375 SMU RS06880 WP 002262929.1    VWIEIDTDKMTQVIDNILNNAIKYSPDGGKVTITMQTTDTQLILSISDQGLGIPKKDLPLIFDRFYRVDKARSRAQGGTG    398
NTDB id 1141439 DQN42 RS03875 WP 003072979.1  TWVEIDTDKMTQVIDNIINNAIKYSPDGGKITVSMKTTETQMILSISDEGLGIPKKDLPKIFDRFYRVDKARSRAQGGSG    398
NTDB id 414 AAK55818.1 838..2187( )          IWMEIDTDKMTQVVDNILNNAIKYSPDGGKITVRMKTTEDQMILSISDHGLGIPKQDLPRIFDRFYRVDRARSRAQGGTG    398
consensus

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NTDB id 375 SMU RS06880 WP 002262929.1    LGLAIAKEIVKQHKGFIWANSEEGEGSTFTIVLPYENDNDAIDEWEEDEDES    450
NTDB id 1141439 DQN42 RS03875 WP 003072979.1  LGLAIAKEIIKQHNGFIWAKSEYGKGSTFTIVLPYDKEAVKDDDWEEDGLEG    450
NTDB id 414 AAK55818.1 838..2187( )          LGLSIAKEIIKQHKGFIWAKSEYGKGSTFTIVLPYDKDAVKEEVWEDFVED*    449
consensus
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