logo
NTDB id 395288 GE559 RS09455 WP 096750174.1 MFSFFKRLIGSKADGAPTDDPDAPTDR.....QDDAEAP....AAEAARPVERVAAPA.....PAPAVPVEPA 59
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG 80
consensus
NTDB id 395288 GE559 RS09455 WP 096750174.1 APAQVAPALPAATADEDEAQAVTAVEIVPPPAPEPAAKKSWIARLRSGLSKTS....SSLTSVFVSTKIDEDLYEELETA 135
NTDB id 1118 NGFG RS11455 WP 003696286.1 E...AAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETV 157
consensus
NTDB id 395288 GE559 RS09455 WP 096750174.1 LLMSDAGVDATEYLLGALREKVKAERLTDPQQVKAALRGLLIALLAPLEKSLMLG.RAQPLVMMIAGVNGAGKTTSIGKL 214
NTDB id 1118 NGFG RS11455 WP 003696286.1 LITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKL 237
consensus
NTDB id 395288 GE559 RS09455 WP 096750174.1 AKHLQSFDQSVLLAAGDTFRAAAREQLAIWGERNNVTVVQQESGDPAAVIFDAVGAARARKINVVMADTAGRLPTQLHLM 294
NTDB id 1118 NGFG RS11455 WP 003696286.1 AKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLM 317
consensus
NTDB id 395288 GE559 RS09455 WP 096750174.1 EELRKVKRVIGKAQADAPHEVLLVIDANTGQNALAQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPVPVYFIGVG 374
NTDB id 1118 NGFG RS11455 WP 003696286.1 EEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVG 397
consensus
logo
NTDB id 395288 GE559 RS09455 WP 096750174.1 EKVEDLQPFNAEEFTDALLG 394
NTDB id 1118 NGFG RS11455 WP 003696286.1 EGIDDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved