logo
NTDB id 40309 BGLA RS17425 WP 013699425.1 MFSFFKRLIGSKADGAPTDDPDAPTD.....RQDDAEAP....AAEA.....AWPVERVAAPAPAVPVEPAAPAQV 62
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALFEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAGEAAE. 84
consensus
logo
NTDB id 40309 BGLA RS17425 WP 013699425.1 APALPAATANDDEAQAVTAVEIVPPPAPEPAAKKSWIARLRSGLSKTS....SSLTSVFVSTKIDEDLYEELETALLMSDAGVDA 143
NTDB id 1118 NGFG RS11455 WP 003696286.1 ..RVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLITGDMGMEA 167
consensus
logo
NTDB id 40309 BGLA RS17425 WP 013699425.1 TEYLLGALREKVKAERLTDPQQVKAALRGLLIALLAPLEKSLMLG.RAQPLVMMIAGVNGAGKTTSIGKLAKHLQSFDQSVLLAA 227
NTDB id 1118 NGFG RS11455 WP 003696286.1 TEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKYFQAQGKSVLLAA 252
consensus
logo
NTDB id 40309 BGLA RS17425 WP 013699425.1 GDTFRAAAREQLAIWGERNNVTVVQQESGDPAAVIFDAVGAARARKINVVMADTAGRLPTQLHLMEELRKVKRVIGKAQADAPHE 312
NTDB id 1118 NGFG RS11455 WP 003696286.1 GDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEIKKVKRVLQKAIPGAPHE 337
consensus
logo
NTDB id 40309 BGLA RS17425 WP 013699425.1 VLLVIDANTGQNALAQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPVPVYFIGVGEKVEDLQPFNAEEFTDALLG 392
NTDB id 1118 NGFG RS11455 WP 003696286.1 IIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGIDDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved