logo
NTDB id 68886 MCB1EB RS10820 WP 045364349.1 MFSFFKKLHKSSTSTSEL...T 19
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG 80
consensus
logo
NTDB id 68886 MCB1EB RS10820 WP 045364349.1 EA..........ENKSKNDTQDKNPPAPELTPDRPTSWVKRLKSGLAKTRNNF....TNLFTNVKIDEDFFEELEAALLM 85
NTDB id 1118 NGFG RS11455 WP 003696286.1 EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT 160
consensus
logo
NTDB id 68886 MCB1EB RS10820 WP 045364349.1 ADTGLEASQFLLNALRQKILAERLNDGEAVKAALRSLLADLLKPLERTLVLD.RAQPLVMMITGVNGAGKTTSIGKLAQH 164
NTDB id 1118 NGFG RS11455 WP 003696286.1 GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY 240
consensus
logo
NTDB id 68886 MCB1EB RS10820 WP 045364349.1 LQRHGQSVLLAAGDTFRAAAHEQLALWGERNQINVISQQSGDPAAVIFDALNAARARKLDILMADTAGRLPTQLHLMDEL 244
NTDB id 1118 NGFG RS11455 WP 003696286.1 FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI 320
consensus
logo
NTDB id 68886 MCB1EB RS10820 WP 045364349.1 KKIKRVIGKAMHDAPHEILLVIDANTGQNALAQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPIPVYFIGVGEQI 324
NTDB id 1118 NGFG RS11455 WP 003696286.1 KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI 400
consensus
logo
NTDB id 68886 MCB1EB RS10820 WP 045364349.1 DDLQPFYAEEFANALLN 341
NTDB id 1118 NGFG RS11455 WP 003696286.1 DDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved