logo
NTDB id 158366 UW163 RS02525 WP 014618044.1 MFSFWKKRKAEPQPAAEPAPPAAAPEAVQAPVPAPVASPT.....PAPAPAPVAVPDTPAAPA 58
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG 80
consensus
logo
NTDB id 158366 UW163 RS02525 WP 014618044.1 ..........ALDMQADDIE.TVPTPPVMEQARKGWMSRLRSGLSKTS....KNLTTLFVGVKVDEALFEELETALLM 121
NTDB id 1118 NGFG RS11455 WP 003696286.1 EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT 160
consensus
logo
NTDB id 158366 UW163 RS02525 WP 014618044.1 ADAGVDATEYLLDELRRRVKAQRIETAESVKTALRDLLVELLHPLEKTMVLG.RDQPMVIMIAGVNGAGKTTSIGKLCKH 200
NTDB id 1118 NGFG RS11455 WP 003696286.1 GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY 240
consensus
logo
NTDB id 158366 UW163 RS02525 WP 014618044.1 FQTYGQSVLLAAGDTFRAAAREQLQVIWGQRNNVTVVAQESGDPAAVIFDAVNAARARGIDIVMADTAGRLPTQMHLMEEL 280
NTDB id 1118 NGFG RS11455 WP 003696286.1 FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI 320
consensus
logo
NTDB id 158366 UW163 RS02525 WP 014618044.1 KKVRRVIGKAMVTAPHETLLVIDANTGQNALAQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPVPVYFIGVGEQV 360
NTDB id 1118 NGFG RS11455 WP 003696286.1 KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI 400
consensus
logo
NTDB id 158366 UW163 RS02525 WP 014618044.1 EDLQPFSAREFADALLG 377
NTDB id 1118 NGFG RS11455 WP 003696286.1 DDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved