logo
NTDB id 228102 B9Z07 RS26210 WP 034174397.1 MFSFFKRFKKTQEPDPAESQSADAQQAAEAAETAPVLDA.....PPAADVPQASAQPAAP.......AVV..MTVTPTND 66
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESE.VAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEA 79
consensus
logo
NTDB id 228102 B9Z07 RS26210 WP 034174397.1 .........GRDEVVETVEI....V.PPPLQDASAKKSWLARLKTGLAKTG....SSITGVFVNTKIDEDLYEELETALL 128
NTDB id 1118 NGFG RS11455 WP 003696286.1 GEAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLI 159
consensus
logo
NTDB id 228102 B9Z07 RS26210 WP 034174397.1 MSDAGVDATEYLLDALREKVRAGRLTDPQQVKDALHDLLVELLTPLEKSLMLGR.AQPLVMMITGVNGAGKTTSIGKLAK 207
NTDB id 1118 NGFG RS11455 WP 003696286.1 TGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAK 239
consensus
logo
NTDB id 228102 B9Z07 RS26210 WP 034174397.1 HLQSFDQSVLLAAGDTFRAAAREQLAVWGERNNVTVVQQESGDPAAVIFDAVSAARARKIDVMMADTAGRLPTQLHLMEE 287
NTDB id 1118 NGFG RS11455 WP 003696286.1 YFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEE 319
consensus
logo
NTDB id 228102 B9Z07 RS26210 WP 034174397.1 LKKVKRVIAKAHDGAPHEVLLVIDANTGQNALTQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPVPVYFIGVGEK 367
NTDB id 1118 NGFG RS11455 WP 003696286.1 IKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEG 399
consensus
logo
NTDB id 228102 B9Z07 RS26210 WP 034174397.1 VEDLQPFSAVEFADALLG 385
NTDB id 1118 NGFG RS11455 WP 003696286.1 IDDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved