logo
NTDB id 351083 E2R23 RS18960 WP 028358576.1 MFSFFKRFKKAPDAAPAEPQQAPDAPRDASRAPASPDAARVDAPAAPQPRDERSENHDRPEHDETPTAEHSRSGAQPAAR 80
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKG......R.....AESAVETVSGAVEQVK 69
consensus
logo
NTDB id 351083 E2R23 RS18960 WP 028358576.1 ATPPASAAASEAPVGQPRAAAARASEPAPSVVMTVTPSSNGRHGVVETVEIVPPPAPEPAAKKSWIARLKSGLAKTSS.. 158
NTDB id 1118 NGFG RS11455 WP 003696286.1 E......TVAEMPSEAGE.AAERVESAKEAVAETV....GEAVG.....QVQEAVATTEEHKLGWAARLKQGLAKSRDKM 133
consensus
logo
NTDB id 351083 E2R23 RS18960 WP 028358576.1 ..SITNVFVNTKIDDALYEELETALLMSDAGIDATEHLLGALREKVRAGRLTDPQQVKDALRELLVELLAPLEKSLVLGR 236
NTDB id 1118 NGFG RS11455 WP 003696286.1 AKSLAGVFGGGQIGEDLYEELETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPE 213
consensus
logo
NTDB id 351083 E2R23 RS18960 WP 028358576.1 .AQPLVMMIAGVNGAGKTTSIGKLAKHLQSFDQSVLLAAGDTFRAAAREQLAIWGERNNVTVVQQESGDPAAVIFDAVSA 315
NTDB id 1118 NGFG RS11455 WP 003696286.1 TKEPFVIMLAGINGAGKTTSIGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQA 293
consensus
logo
NTDB id 351083 E2R23 RS18960 WP 028358576.1 ARARNIDVMMADTAGRLPTQLHLMEELRKVKRVIGKAHDGAPHEVLLVIDANTGQNALAQVKAFDDALGLTGLIVTKLDG 395
NTDB id 1118 NGFG RS11455 WP 003696286.1 AKARGIDIVLADTAGRLPTQLHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDG 373
consensus
logo
NTDB id 351083 E2R23 RS18960 WP 028358576.1 TAKGGILAAIARQRPIPVYFIGIGEKVEDLQPFSAEEFADALLG 439
NTDB id 1118 NGFG RS11455 WP 003696286.1 TAKGGILAALASDRPVPVRYIGVGEGIDDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved