
logo
NTDB id 130135 BG24 RS16045 WP 028358576.1 MFSFFKRFKKAPDAAPAEPQQAPDAPRDASRAPASPDAARVDAPAAPQPRDERSENHDRPEHDETPTAEHSRSGAQPAARATPPA 85
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKG......R.....AESAVETVSGAVEQVKE.... 70
consensus
logo
NTDB id 130135 BG24 RS16045 WP 028358576.1 SAAASEAPVGQPRAAAARASEPAPSVVMTVTPSSNGRHGVVETVEIVPPPAPEPAAKKSWIARLKSGLAKTSS....SITNVFVN 166
NTDB id 1118 NGFG RS11455 WP 003696286.1 ..TVAEMPSEAGE.AAERVESAKEAVAETV....GEAVG.....QVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGG 143
consensus
logo
NTDB id 130135 BG24 RS16045 WP 028358576.1 TKIDDALYEELETALLMSDAGIDATEHLLGALREKVRAGRLTDPQQVKDALRELLVELLAPLEKSLVLGR.AQPLVMMIAGVNGA 250
NTDB id 1118 NGFG RS11455 WP 003696286.1 GQIGEDLYEELETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGA 228
consensus
logo
NTDB id 130135 BG24 RS16045 WP 028358576.1 GKTTSIGKLAKHLQSFDQSVLLAAGDTFRAAAREQLAIWGERNNVTVVQQESGDPAAVIFDAVSAARARNIDVMMADTAGRLPTQ 335
NTDB id 1118 NGFG RS11455 WP 003696286.1 GKTTSIGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQ 313
consensus
logo
NTDB id 130135 BG24 RS16045 WP 028358576.1 LHLMEELRKVKRVIGKAHDGAPHEVLLVIDANTGQNALAQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPIPVYFIGIGE 420
NTDB id 1118 NGFG RS11455 WP 003696286.1 LHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGE 398
consensus
logo
NTDB id 130135 BG24 RS16045 WP 028358576.1 KVEDLQPFSAEEFADALLG 439
NTDB id 1118 NGFG RS11455 WP 003696286.1 GIDDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved