logo
NTDB id 400109 GJG85 RS14860 WP 050014853.1 MFSFFKRFKKTQEPDPAESQSADAQQADEASDAD.....APAPEAP..PAADVPQASAQAAAPAVVVTVTPTNDG 68
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG 80
consensus
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NTDB id 400109 GJG85 RS14860 WP 050014853.1 .....RDEVVETVEI....V.PPPLQDASAKKSWLARLKTGLAKTG....SSITGVFVNTKIDEDLYEELETALLM 130
NTDB id 1118 NGFG RS11455 WP 003696286.1 EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT 160
consensus
logo
NTDB id 400109 GJG85 RS14860 WP 050014853.1 SDAGVDATEYLLDALREKVRAGRLTDPQQVKDALHDLLVELLTPLEKSLMLG.RAQPLVMMIAGVNGAGKTTSIGKLAKH 209
NTDB id 1118 NGFG RS11455 WP 003696286.1 GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY 240
consensus
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NTDB id 400109 GJG85 RS14860 WP 050014853.1 LQSFDQSVLLAAGDTFRAAAREQLAVWGERNNVTVVQQESGDPAAVIFDAVSAARARKIDVMMADTAGRLPTQLHLMEEL 289
NTDB id 1118 NGFG RS11455 WP 003696286.1 FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI 320
consensus
logo
NTDB id 400109 GJG85 RS14860 WP 050014853.1 KKVKRVIAKAHDGAPHEVLLVIDANTGQNALTQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPVPVYFIGVGEKV 369
NTDB id 1118 NGFG RS11455 WP 003696286.1 KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI 400
consensus
logo
NTDB id 400109 GJG85 RS14860 WP 050014853.1 EDLQPFSAVEFADALLG 386
NTDB id 1118 NGFG RS11455 WP 003696286.1 DDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved