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NTDB id 256484 CUJ88 RS01460 WP 007579684.1 MFSFFKRFKGSKAPDDTPETPQADSGADVDVDM.....QDVAEPEA....PAVAR.SQSVPDSADAPEFDPNYV 64
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQE.TAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEA 79
consensus
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NTDB id 256484 CUJ88 RS01460 WP 007579684.1 PDAA.DFEDSDDE.....AAEIVPPPVPEAAEKRSWLTRLRSGLSKT....SSSITGIFVGAKIDEDLYEELETALL 131
NTDB id 1118 NGFG RS11455 WP 003696286.1 GEAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLI 159
consensus
logo
NTDB id 256484 CUJ88 RS01460 WP 007579684.1 MSDAGVDATEFLLEALREKVRAERLTDPQQVKGALRTLLVDLLKPLEKSLTLGH.YQPTVMMIAGVNGAGKTTSIGKLAK 210
NTDB id 1118 NGFG RS11455 WP 003696286.1 TGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAK 239
consensus
logo
NTDB id 256484 CUJ88 RS01460 WP 007579684.1 HMQHFNQSVLLAAGDTFRAAAREQLAIWGQRNNVTVVQQESGDPAAVIFDAVGAARARKINVVMADTAGRLPTQLHLMEE 290
NTDB id 1118 NGFG RS11455 WP 003696286.1 YFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEE 319
consensus
logo
NTDB id 256484 CUJ88 RS01460 WP 007579684.1 LRKVKRVVGKAMDGAPHEVLLVIDANTGQNALAQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPVPVYFIGVGEK 370
NTDB id 1118 NGFG RS11455 WP 003696286.1 IKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEG 399
consensus
logo
NTDB id 256484 CUJ88 RS01460 WP 007579684.1 VEDLQPFSAEEFSDALLG 388
NTDB id 1118 NGFG RS11455 WP 003696286.1 IDDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved