logo
NTDB id 1078201 ACJU26 RS13355 WP 414040117.1 MVFNWFKRNKSPAP...EERGEASETAPVPQELEALSRNSIPEAVEPTEKSAT..AEDVAASPSG.....LLPEP 65
NTDB id 1118 NGFG RS11455 WP 003696286.1 .MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVG.NIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSE 78
consensus
logo
NTDB id 1078201 ACJU26 RS13355 WP 414040117.1 A.........AEAIPEAAAEDALSQPQPA..AASAPTKSMFARLREGLSRSREQFTEGVSRLVLGKKVIDEDLLEDLEA 133
NTDB id 1118 NGFG RS11455 WP 003696286.1 AGEAAERVESAKEAVAETV.GEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAG.VFGGGQIGEDLYEELET 156
consensus
logo
NTDB id 1078201 ACJU26 RS13355 WP 414040117.1 LLLQADLGSAATQEIMSAVTESVRRKELTDPAALKKALRNALLDILRPRAETW.VPEKGRTQVLMMVGINGAGKTTTTGK 212
NTDB id 1118 NGFG RS11455 WP 003696286.1 VLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGK 236
consensus
logo
NTDB id 1078201 ACJU26 RS13355 WP 414040117.1 LAARWKAEGFSLMLAAGDTFRAAAVEQLQGWGQRVQVPVVAQGTGADSASVIFDAFTAARARGADLLIADTAGRLHTQNH 292
NTDB id 1118 NGFG RS11455 WP 003696286.1 LAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTG.DSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLH 315
consensus
logo
NTDB id 1078201 ACJU26 RS13355 WP 414040117.1 LMEELKKVKRVLGKQDPDAPHQVWLVLDAGTGQNALNQAQQFHQAIGLTGICITKLDGTAKGGVVAAIAKTLPIPIRYIG 372
NTDB id 1118 NGFG RS11455 WP 003696286.1 LMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIG 395
consensus
logo
NTDB id 1078201 ACJU26 RS13355 WP 414040117.1 MGEQVADLRPFDPESFVDALFAETKS 398
NTDB id 1118 NGFG RS11455 WP 003696286.1 VGEGIDDLRPFDARAFVDALLD.... 417
consensus
X non conserved
X similar
X ≥ 50% conserved