logo
NTDB id 415025 GT347 RS16370 WP 160553227.1 MFSFFKKKPPAATPEPASTTPPQEDA....APTGALGWLKSQLGLAPAPAETPETTTPPAAPAVAP..TPVVAVPVAAP 74
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVK.......GRAESAVETVSGAVEQVKETVA 73
consensus
logo
NTDB id 415025 GT347 RS16370 WP 160553227.1 PLPPPPAAAPLVPPPPLAPPAPAAA.PVVAAPAPAAQAEEGGTAERSGWMQRLRGGLRKTG....GSIAAVFSDARIDDA 149
NTDB id 1118 NGFG RS11455 WP 003696286.1 EMPSEAGEAA....ERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGED 149
consensus
logo
NTDB id 415025 GT347 RS16370 WP 160553227.1 LYDELEDALLMADTGIKATQHLLDDLRRRVKERRSTEPTAVKALLADAIADLLKPLEKPLVIGQ.HTPTVIMVAGVNGAG 228
NTDB id 1118 NGFG RS11455 WP 003696286.1 LYEELETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPE.TKEPFVIMLAGINGAG 229
consensus
logo
NTDB id 415025 GT347 RS16370 WP 160553227.1 KTTSIGKLTKHLADAKATVLLAAADTFRAAAREQLGVWADRNLVEIIGQQGGDPAAVSFDAVTAGKARGKDVVLVDTAGR 308
NTDB id 1118 NGFG RS11455 WP 003696286.1 KTTSIGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGR 309
consensus
logo
NTDB id 415025 GT347 RS16370 WP 160553227.1 LPTQLHLMQELQKIKRVVQKADNSAPHEVLLVIDGNTGQNALAQVRAFDDALGLTGLIVTKLDGTAKGGVLAAIAQERPI 388
NTDB id 1118 NGFG RS11455 WP 003696286.1 LPTQLHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPV 389
consensus
logo
NTDB id 415025 GT347 RS16370 WP 160553227.1 PVYFIGVGEKLEDLETFNAREFAQALLG 416
NTDB id 1118 NGFG RS11455 WP 003696286.1 PVRYIGVGEGIDDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved