logo
NTDB id 211644 BVD88 RS00895 WP 011799029.1 MLYPEKMSGMAGRHPYGVRSGLRRNGLKLWDIHFRMTRFIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINA 80
NTDB id 1127 NGFG RS10560 WP 003703428.1 .....MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINA 57
NTDB id 1126 NGFG RS09375 WP 003703428.1 .....MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINA 57
NTDB id 1125 NGFG RS07215 WP 003703428.1 .....MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINA 57
NTDB id 1108 NGFG RS06525 WP 003703428.1 .....MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINA 57
consensus
logo
NTDB id 211644 BVD88 RS00895 WP 011799029.1 ASQQELEALPGIGPAKAKAIAEYRAQNGAFKSVDDLTKVKGIGPAVLAKLKDQASVGAPAPKGPAKPATPAAKK 154
NTDB id 1127 NGFG RS10560 WP 003703428.1 ASQQELEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1126 NGFG RS09375 WP 003703428.1 ASQQELEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1125 NGFG RS07215 WP 003703428.1 ASQQELEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1108 NGFG RS06525 WP 003703428.1 ASQQELEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
consensus
X non conserved
X similar
X ≥ 50% conserved