logo
NTDB id 75014 NMA510612 RS01090 WP 010981060.1 MLCPEKMSGMAGQYPYGVRSGLRRNGLKLWDIHFRMTRFIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINA 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 75014 NMA510612 RS01090 WP 010981060.1 ASQQELEALPGIGPAKAKAIAEYRAQNGAFKSVDDLTKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPADKK 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