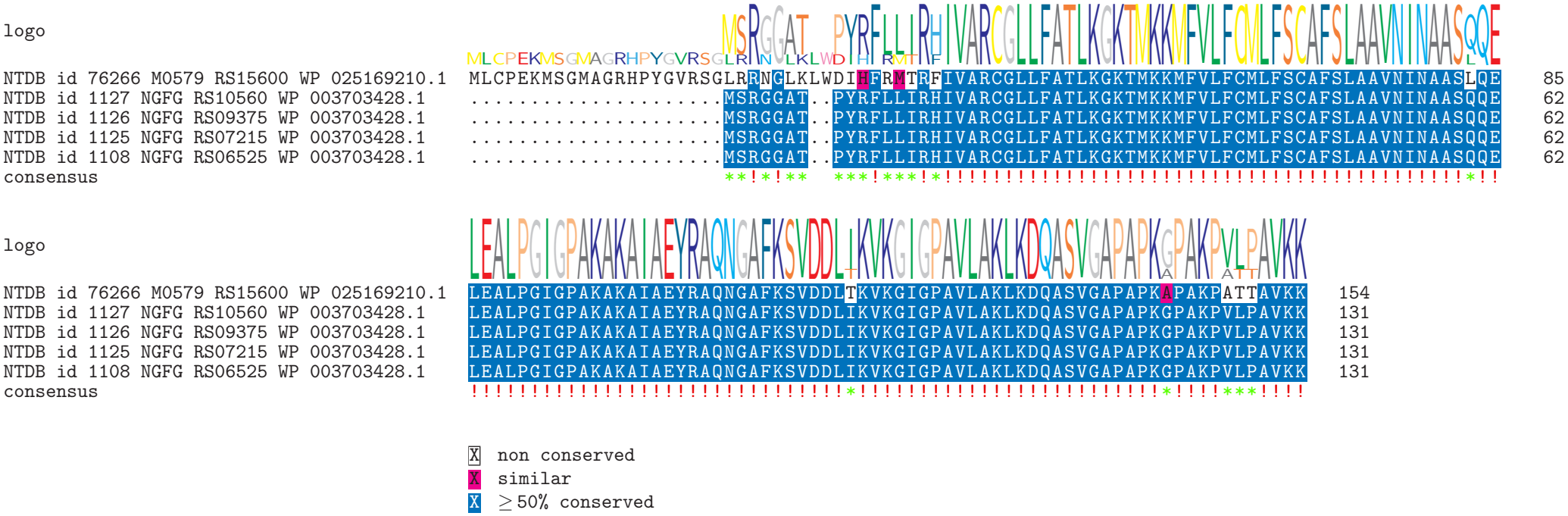
logo
MLCPEKMSGMAGRHPYGVRSGLRRNGLKLWDIHFRMTRFIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASLQE
NTDB id 76266 M0579 RS15600 WP 025169210.1 MLCPEKMSGMAGRHPYGVRSGLRRNGLKLWDIHFRMTRFIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASLQE 85
NTDB id 1127 NGFG RS10560 WP 003703428.1 .....MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQE 62
NTDB id 1126 NGFG RS09375 WP 003703428.1 .....MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQE 62
NTDB id 1125 NGFG RS07215 WP 003703428.1 .....MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQE 62
NTDB id 1108 NGFG RS06525 WP 003703428.1 .....MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQE 62
consensus **!*!** ***!***!*!!*!!
logo
LEALPGIGPAKAKAIAEYRAQNGAFKSVDDLTKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK
NTDB id 76266 M0579 RS15600 WP 025169210.1 LEALPGIGPAKAKAIAEYRAQNGAFKSVDDLTKVKGIGPAVLAKLKDQASVGAPAPKAPAKPATTAVKK 154
NTDB id 1127 NGFG RS10560 WP 003703428.1 LEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1126 NGFG RS09375 WP 003703428.1 LEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1125 NGFG RS07215 WP 003703428.1 LEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1108 NGFG RS06525 WP 003703428.1 LEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
consensus !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!*!!!!***!!!!
X non conserved
X similar
X ≥ 50% conserved