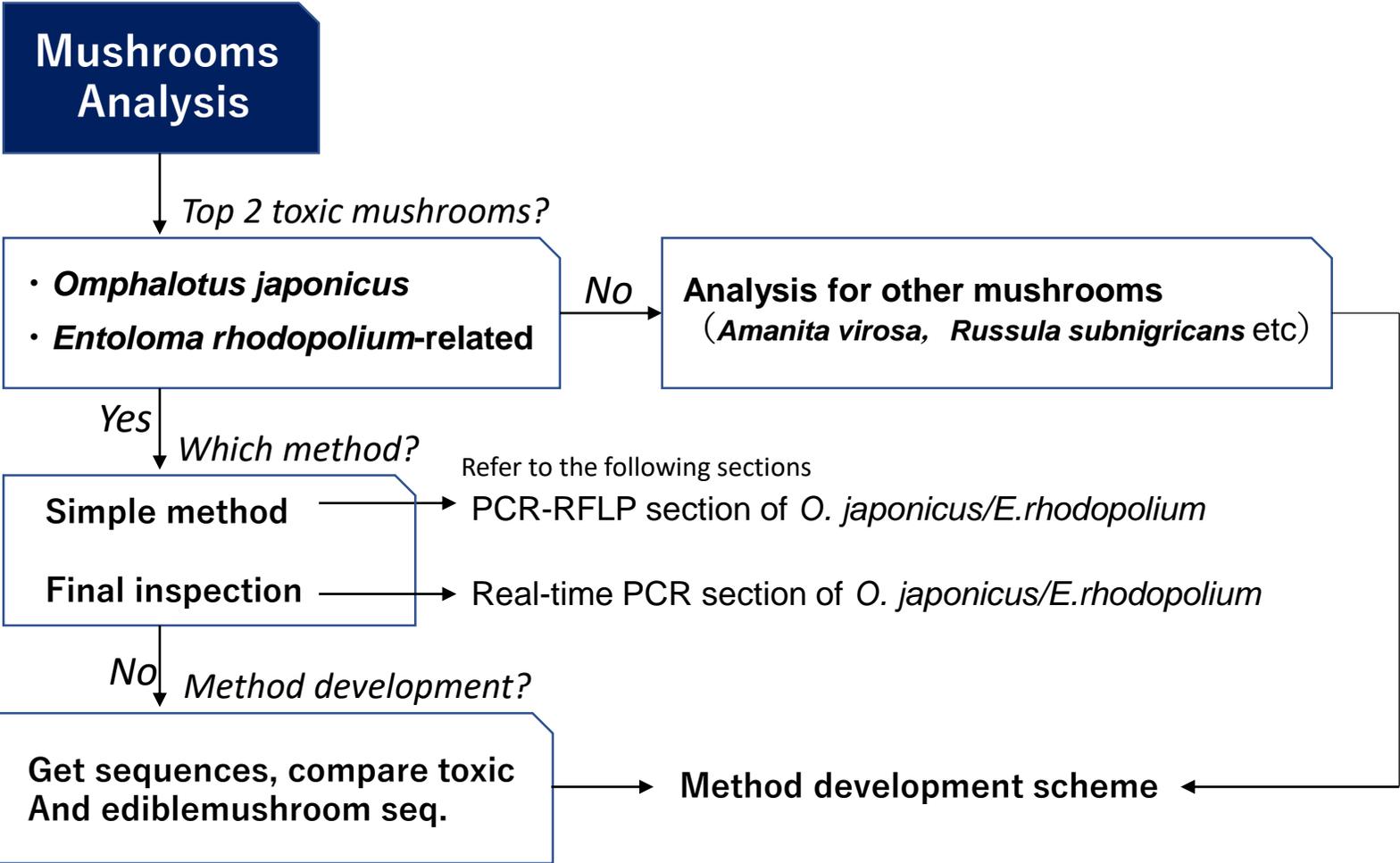
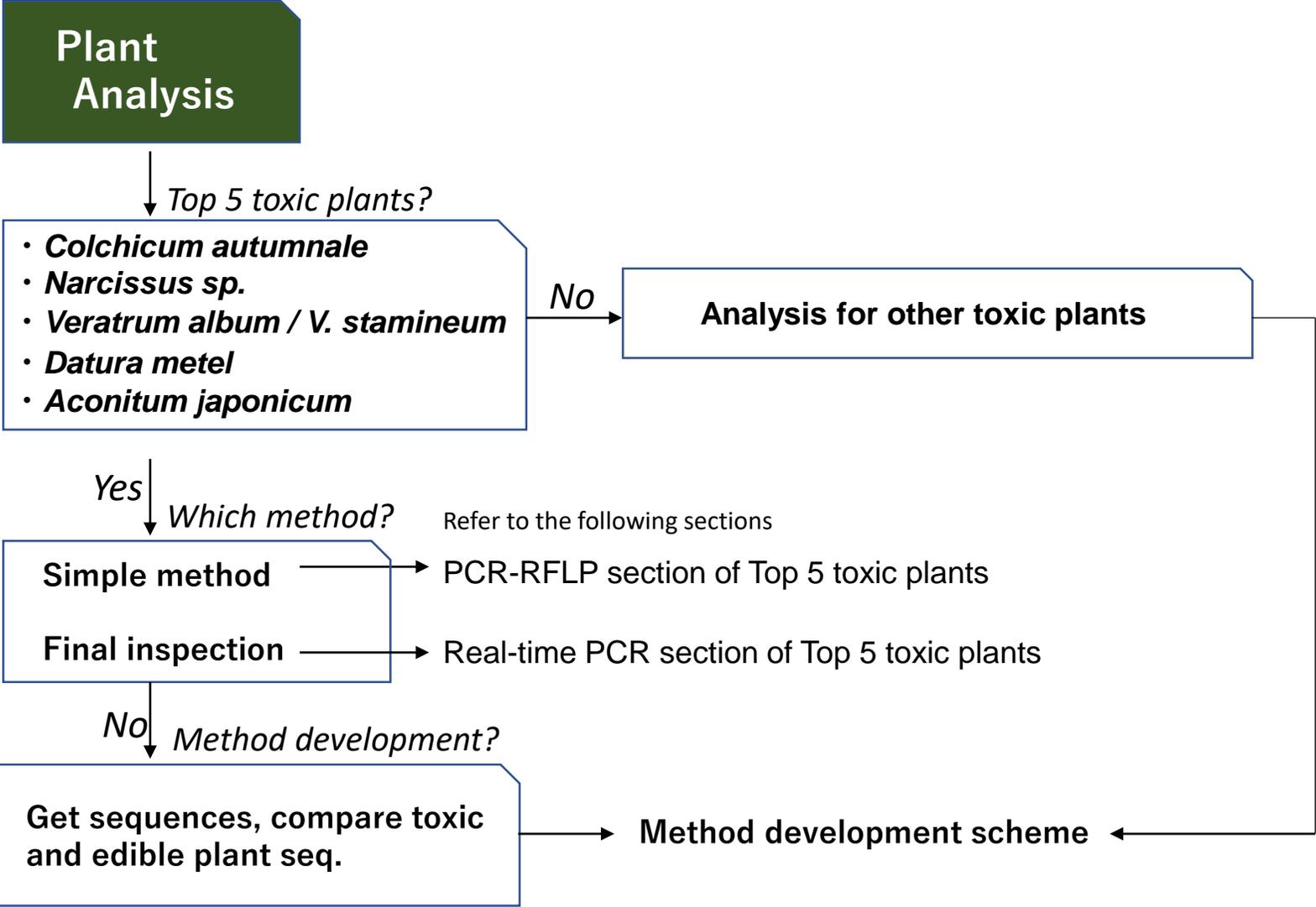


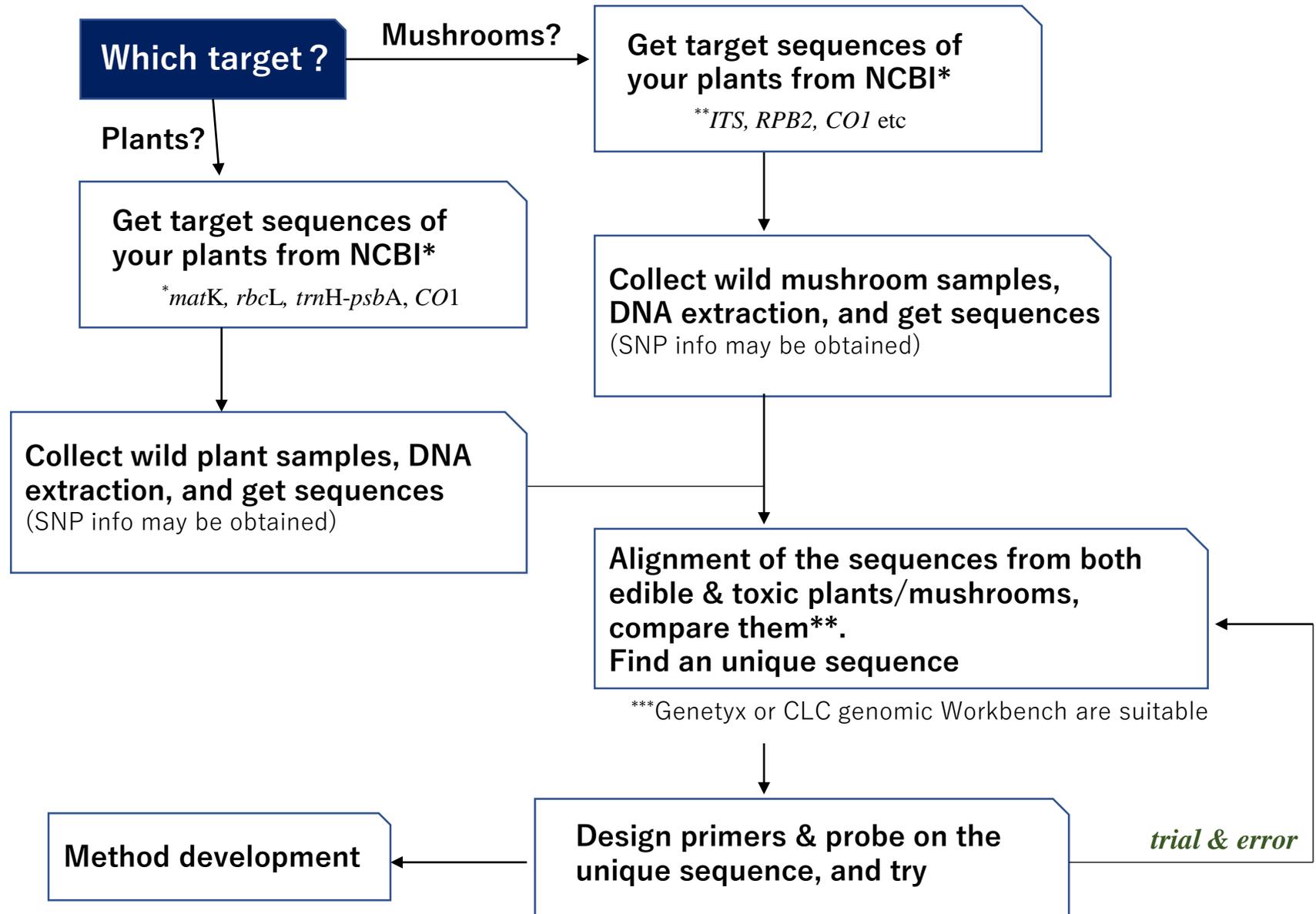
Toxic mushroom analysis scheme



Toxic plants analysis scheme



Method development scheme



Sequence Analysis

Use Muscle as an alignment program

For molecular phylogenetic analysis, software that can run the Maximum Likelihood method should be used

“MEGA” is easy & reliable free software

It is not always necessary to use specialized programs such as RAxML and PAUP, because they can run only on Linux

As commercial software,

Genetyx and CLC Genomic Workbench that can perform comprehensive analysis are recommended

*, In addition to clastalW, you can select Muscle for general gene analysis software such as Genetyx..