

<http://phylo.uoregon.edu/software/phym1-m31>

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oo

.....  
Menu : Tree Searching  
.....

[+] ..... Next sub-menu  
[-] ..... Previous sub-menu  
[Y] ..... Launch the analysis

[O] ..... Optimise tree topology    yes  
[U] ..... Starting tree (BioNJ/parsimony/user tree)    BioNJ  
[S] ..... Tree topology search operations    NNI moves (fast, approximate)

. Are these settings correct ? (type '+', '-', 'Y' or other letter for one to change) ☐