```
[+] ..... Next sub-menu
           [-] ..... Previous sub-menu
           [Y] ..... Launch the analysis
           [0] ..... Optimise tree topology
           [U] ...... Starting tree (BioNJ/parsimony/user tree) BioNJ
           [S] ..... Tree topology search operations Simulated Thermal Annealing
           [R] ..... Add random starting trees no
. Are these settings correct ? (type '+', '-', 'Y' or other letter for one to change) s
```

Menu : Tree Searching