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<http://phylo.uoregon.edu/software/phyml-m3l>

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```
.....
Menu : Mixed Branch Length Model
.....
```

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

[N] ..... Number of Branch lengths per edge: 1
```

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