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                      . . . . . . . . . . . . . . . . . . . . . . . . . . . .
                          Menu : Substitution Model

                      . . . . . . . . . . . . . . . . . . . . . . . . . . . .



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


          [M] . . . . . . . . . . . . . . . . . Model of nucleotide substitution  JC69
          [V] . Proportion of invariable sites (fixed/estimated)  fixed (p-invar = 0.00)
          [R] . . . . . . . One category of substitution rate (yes/no)  no
          [C] . . . . . . . . . . . Number of substitution rate categories  4
          [A] . . . Gamma distribution parameter (fixed/estimated)  estimated
          [G] . . . . . . . . . 'Middle' of each rate class (mean/median)  mean



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