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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)  yes


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