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Menu : Substitution Model  
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[+] ..... Next sub-menu  
[-] ..... Previous sub-menu  
[Y] ..... Launch the analysis
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[M] ..... Model of nucleotide substitution   HKY85  
[F] ..... Optimise equilibrium frequencies    no  
[T] ..... Ts/tv ratio (fixed/estimated)      estimated  
[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
```