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.....
Menu : Input Data
.....
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```
[+] ..... Next sub-menu
[-] ..... Previous sub-menu
[Y] ..... Launch the analysis
```

```
[D] ..... Data type (DNA/AA)   DNA
[I] ..... Input sequences interleaved (or sequential) interleaved
[M] ..... Analyze multiple data sets no
[R] ..... Run ID none
```

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