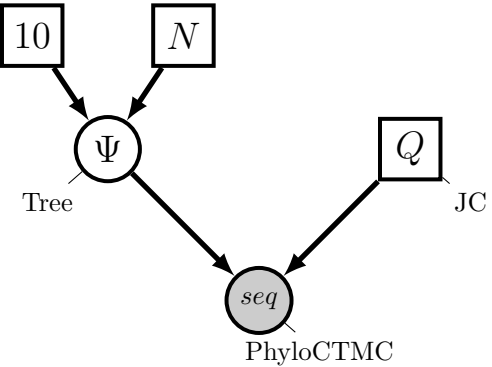


```
psi ~ dnUniformTopologyBranchLength(names, dnExponential(10))


Q_morpho <- fnJC(2)


phyMorpho ~ dnPhyloCTMC( tree=phylogeny,
siteRates=rates_morpho, Q=Q_morpho,
type="Standard", coding="variable" )

phyMorpho.clamp( data )
```