

$$P(\text{[Tree prior]} \mid \text{[Prior information]}) = \frac{P(\text{[genetic sequences]} \mid \text{[genealogy]} \text{ [demographic model]} \text{ [substitution model]} \text{ [molecular clock model]})}{P(\text{[genetic sequences]})}$$


 genetic
sequences


 genealogy


 demographic
model


 substitution
model


 molecular clock
model