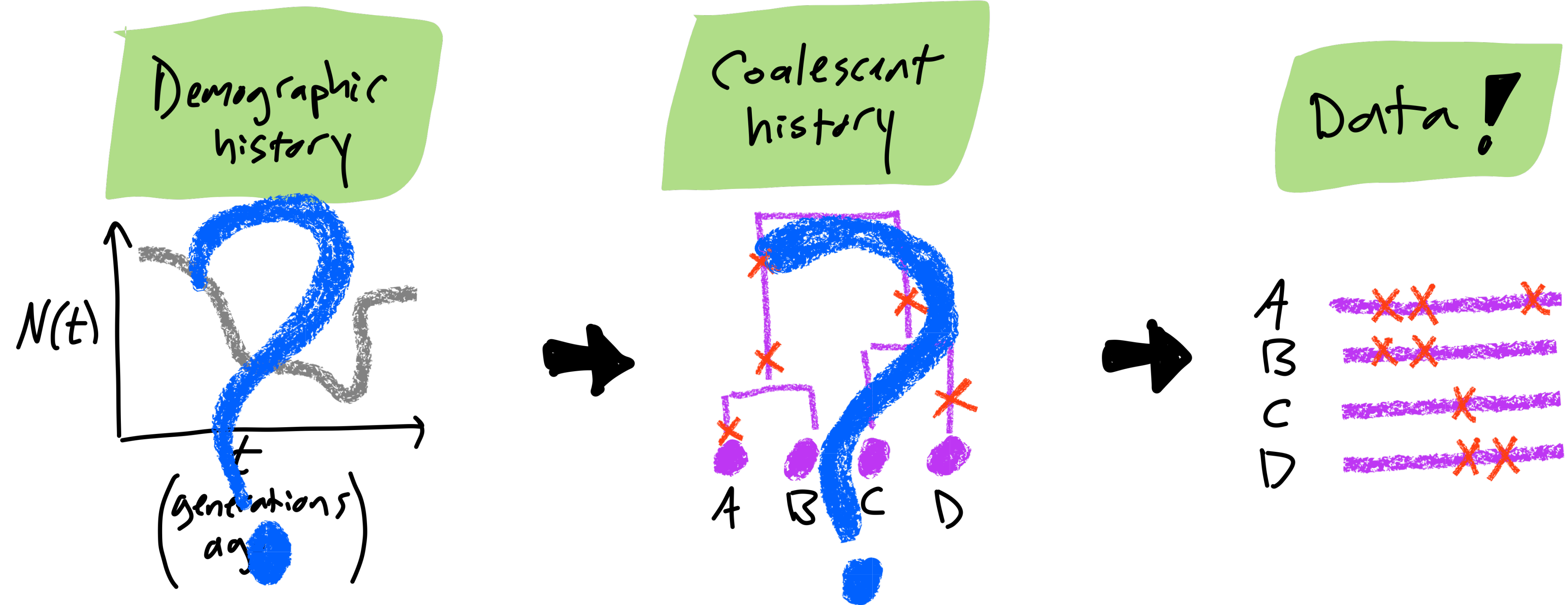


# Coalescent theory

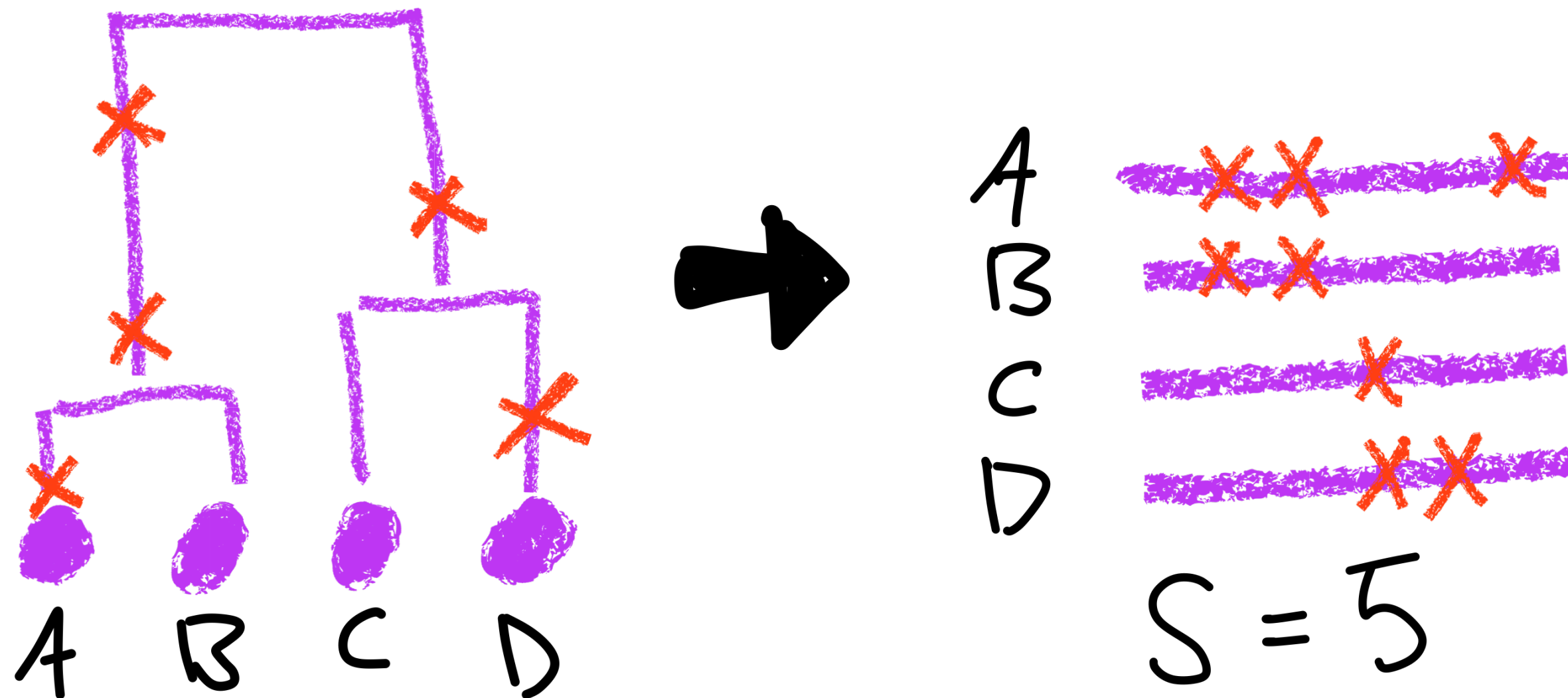
## Genetic diversity



# Coalescent theory

## Genetic diversity

# segregating sites,  $S$ , equals # mutations in the sample's history (infinite sites approximation)



Constant  $N$  case:

$$\begin{aligned} E[S] &= \mu E[T_{\text{total}}] \\ &= \mu \sum_{i=2}^n i E[T_i] \\ &= \mu \sum_{i=2}^n i \frac{2N}{\binom{n}{2}} \\ &= 4\mu N \sum_{i=1}^{n-1} \frac{1}{i} \end{aligned}$$

total branch length