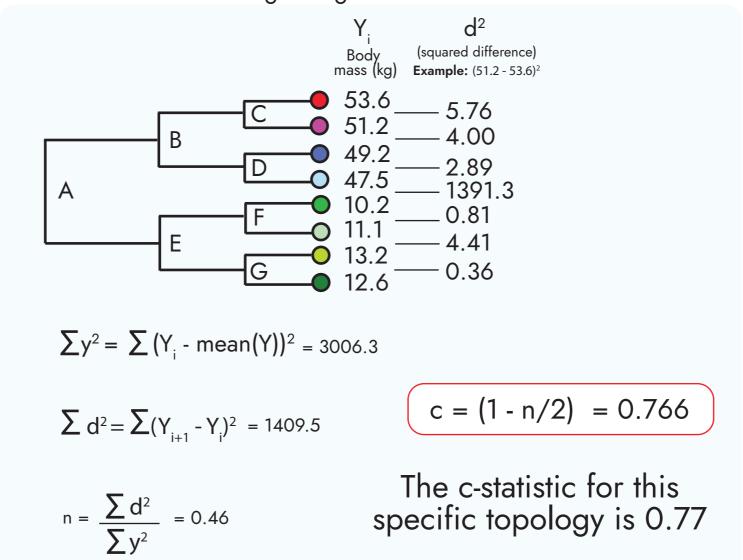
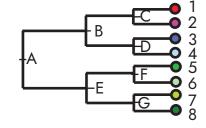
Question:

Is there a phylogenetic signal in body mass?

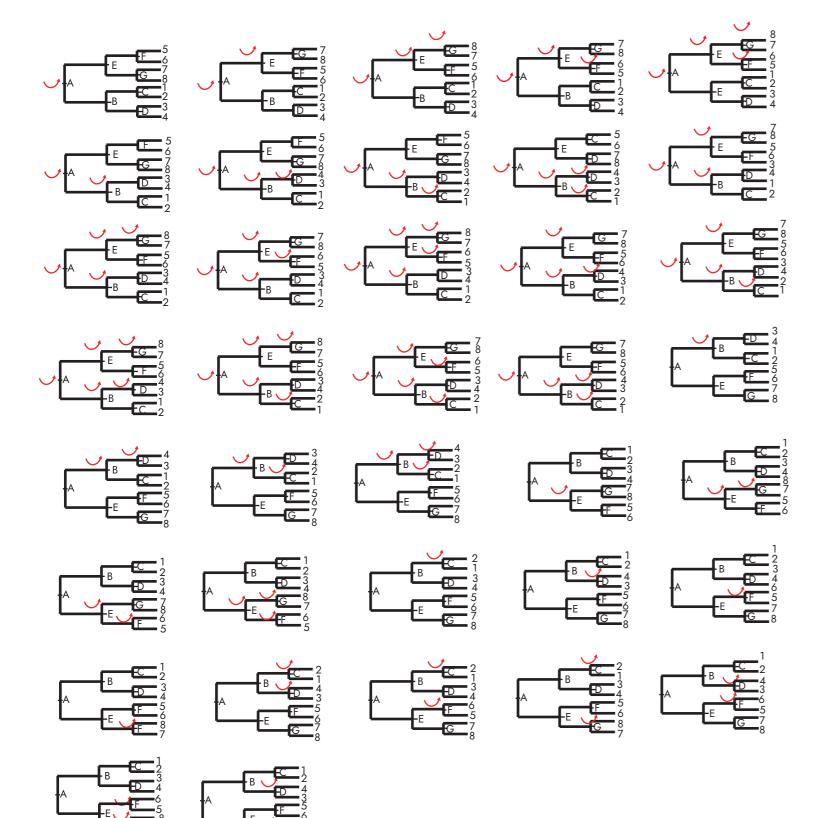


Our original way of showing the topology:



You have to account for the fact that topologies can be shown in multiple ways!

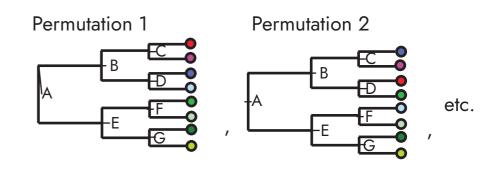
You must account for all (or subsample) the multiple ways that nodes can be rotated



- 1) For each new topology, calculate the d² as above:
- 2) Then average all of the d² across all topologies:

$$1 - \frac{\text{average}(\Sigma d^2)}{2\Sigma^{y^2}} = C_{\text{mean}}$$

3) Then to create a null distribution, permutate trait value across the original topology and calculate a distribution of C:



4) Calculate p-value

