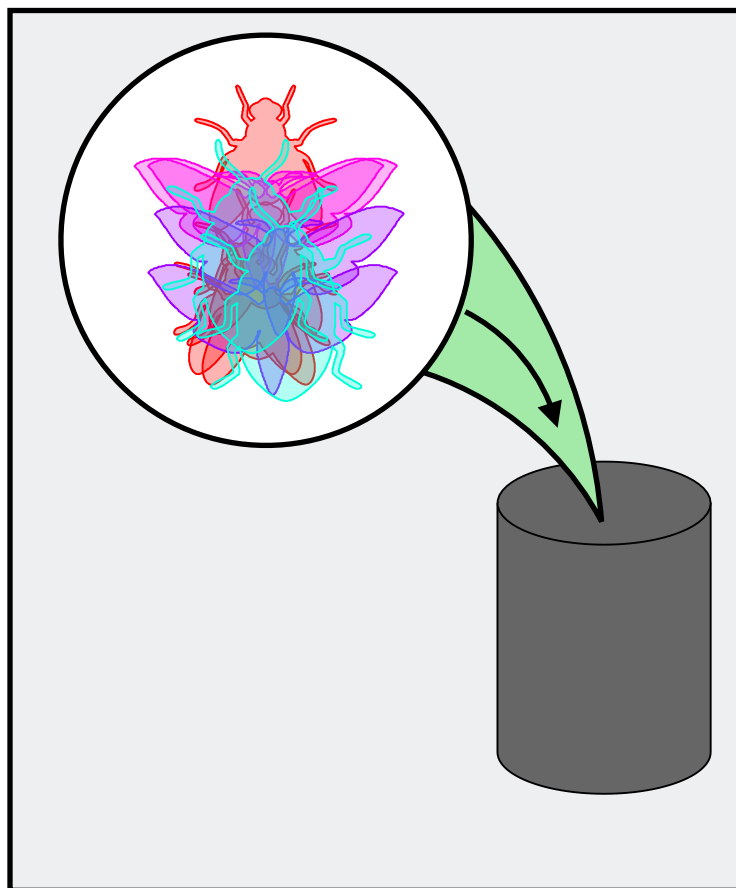
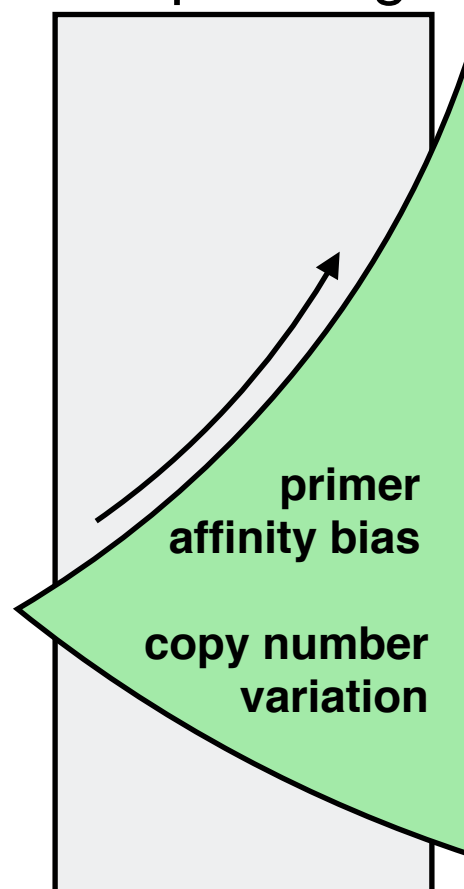


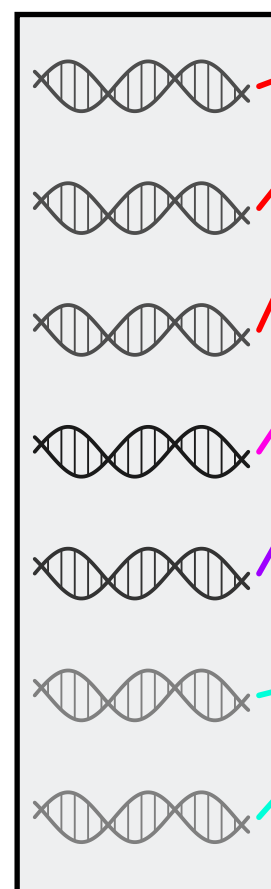
Bulk sample input



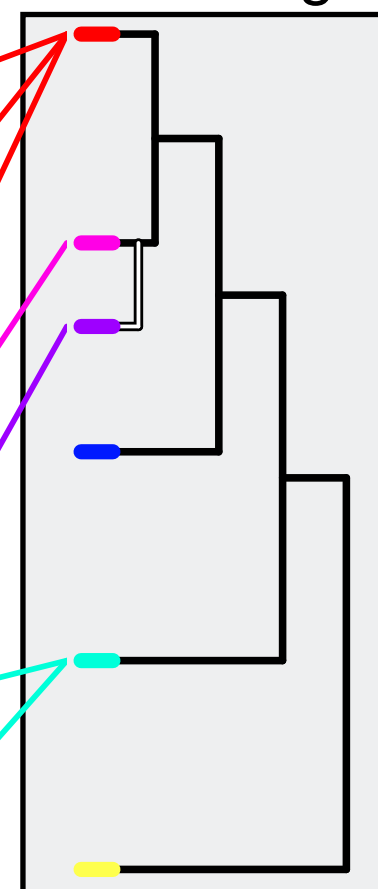
Biased sequencing



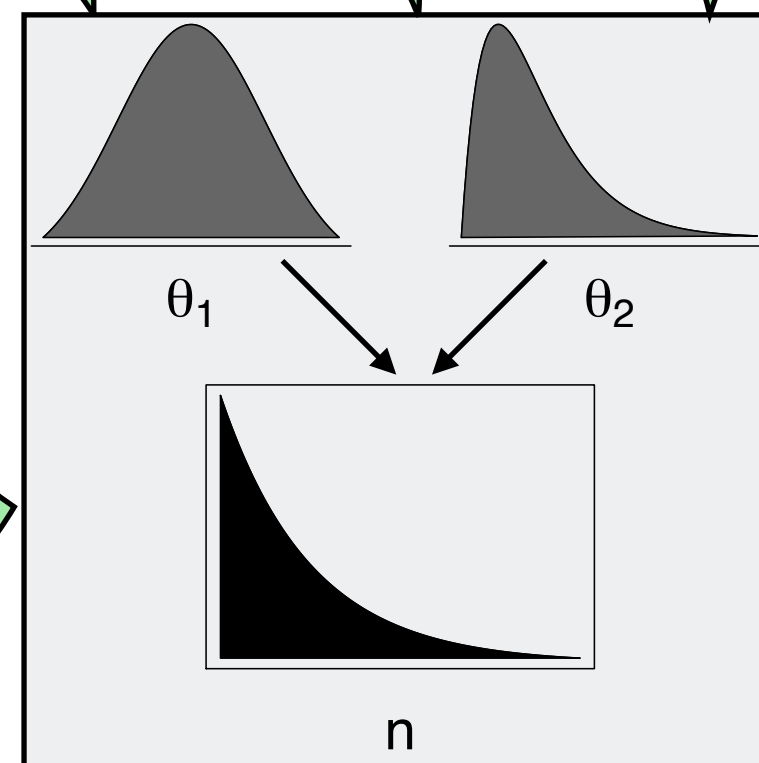
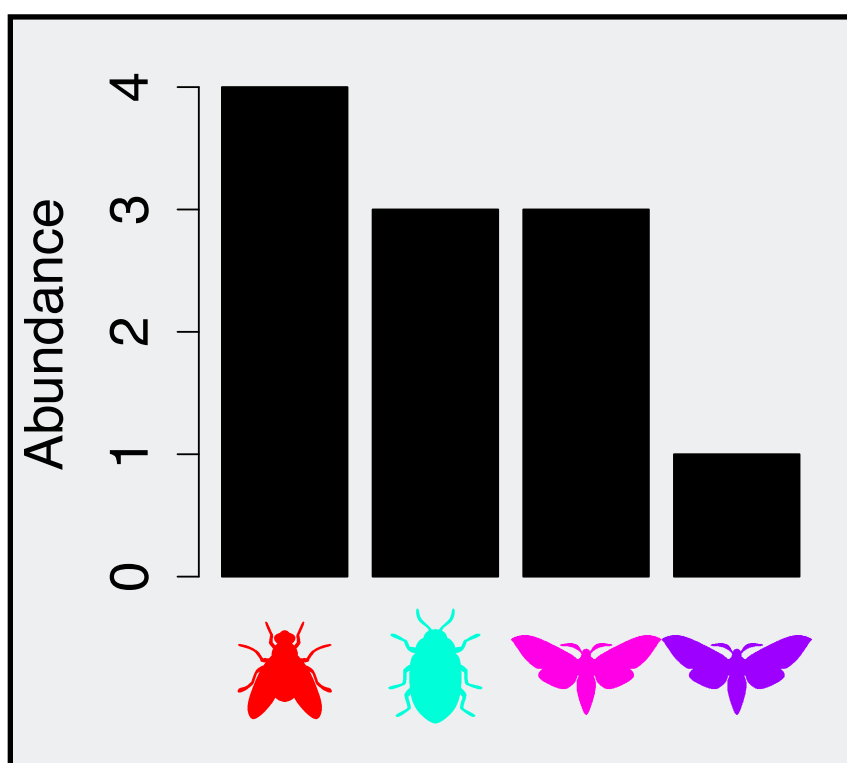
Raw reads



Phylogenetic matching



Unbiased abundances



Hierarchical model