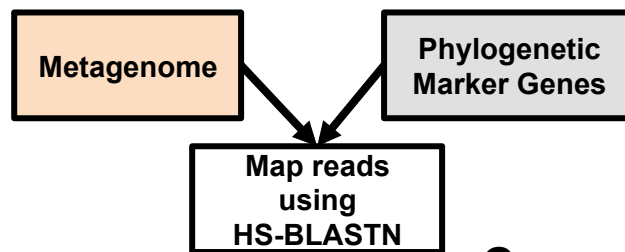
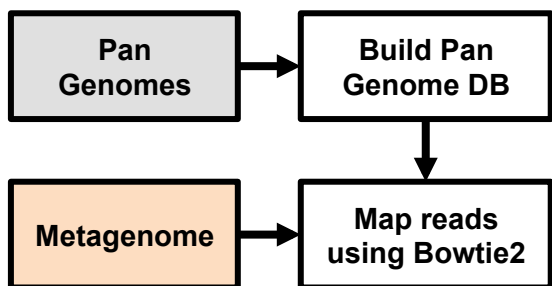


Per Sample

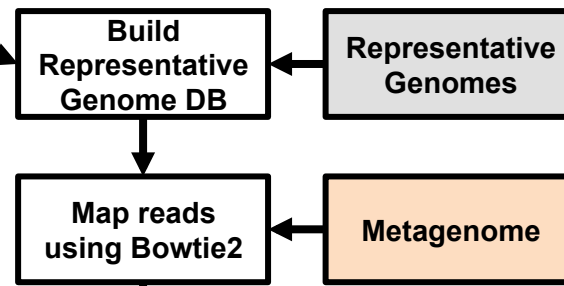
A Estimate Species Abundance



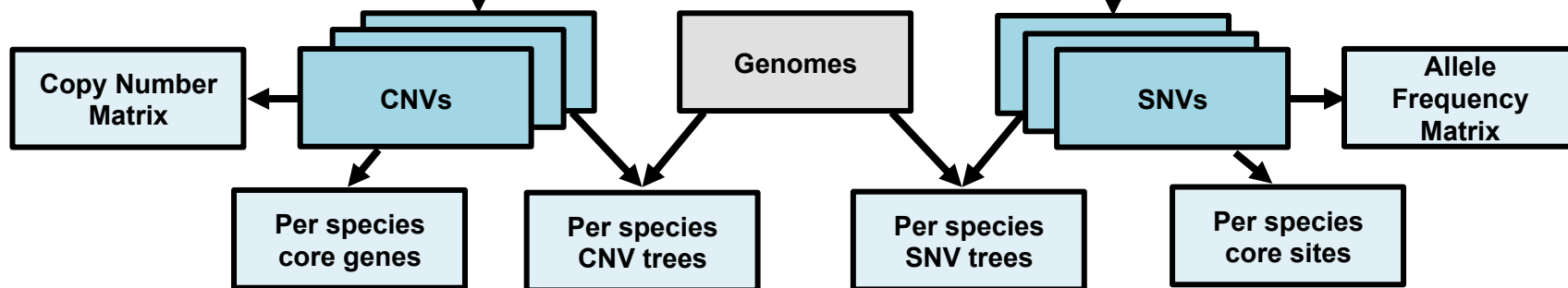
B Predict Gene Presence/Absence In Abundant Species



C Identify Single-Nucleotide Polymorphisms in Abundant Species



D



Multiple Samples