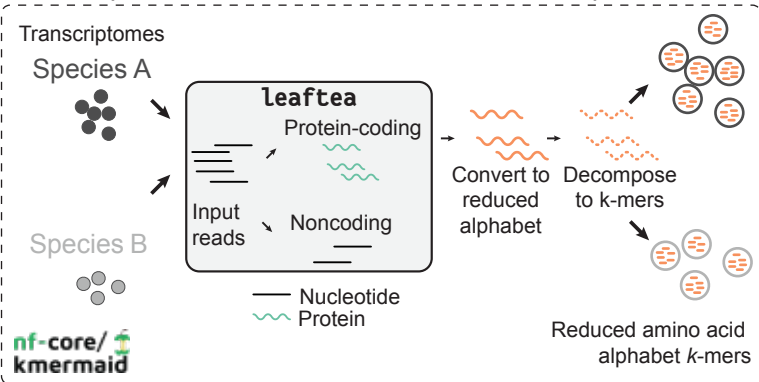
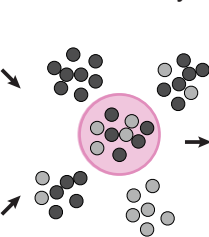


Subsample amino acid k-mers from each transcriptome



Cluster on k-mer similarity



Perform differential k-mer expression and query for functional annotation

