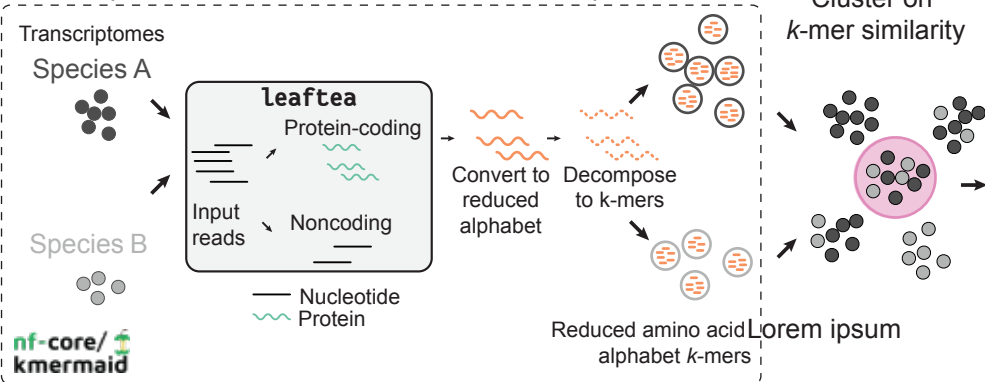


Subsample amino acid k-mers from each transcriptome



Perform differential k-mer expression and query for functional annotation

