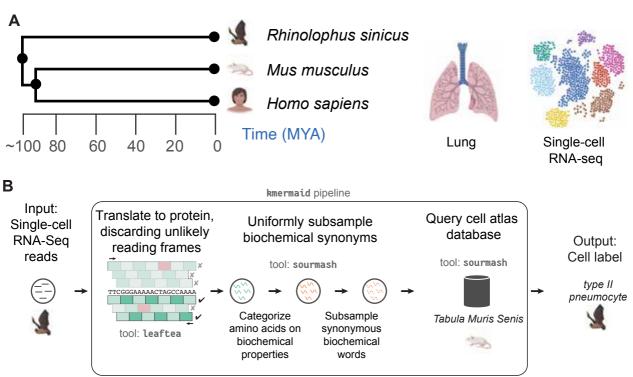
Figure 1: K-mer homology is an alignment-free method for cross-species cell type identification



Minhashed based random subsampling Subsample synonymous encoded sequences