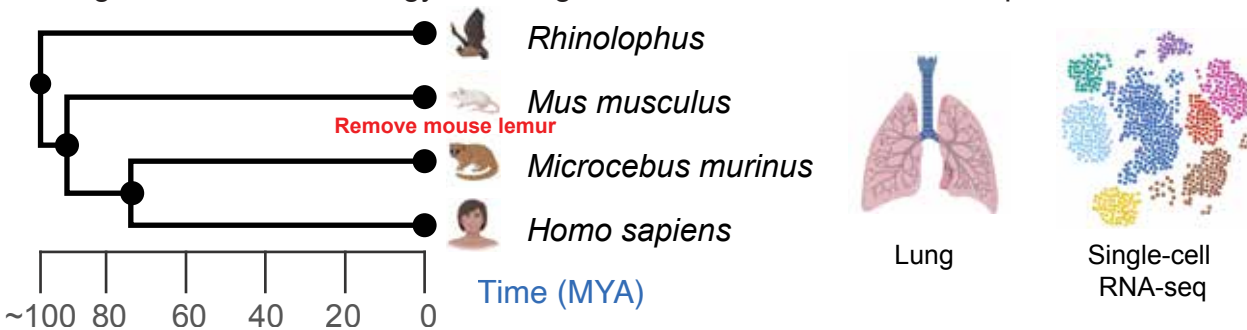


A Figure 1: K-mer homology is an alignment-free and reference-independent method for cross-species analyses



B

