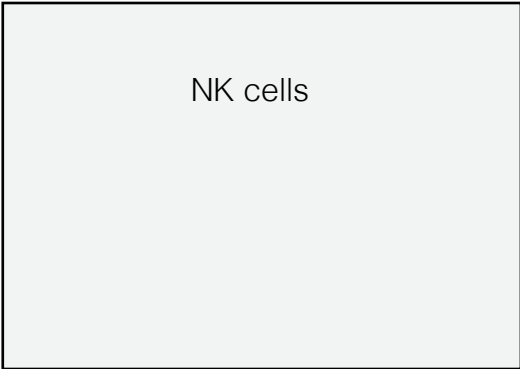
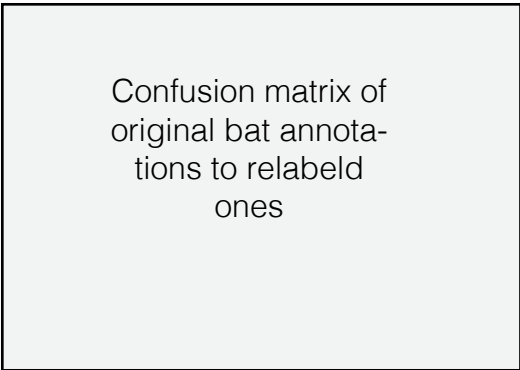


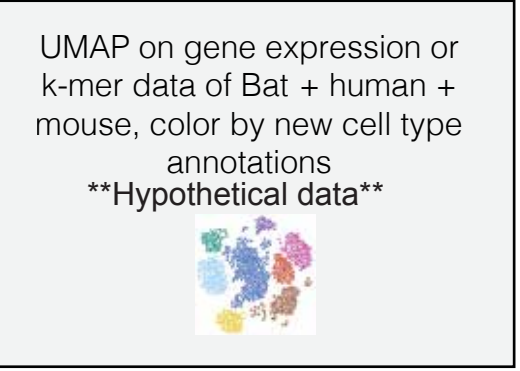
Figure 3: Reduced amino acid k-mers from transcriptomes enabled identification of unannotated immune genes in Chinese horseshoe bat suggesting immune adaptations

A

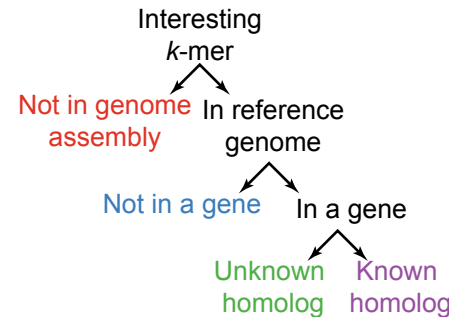
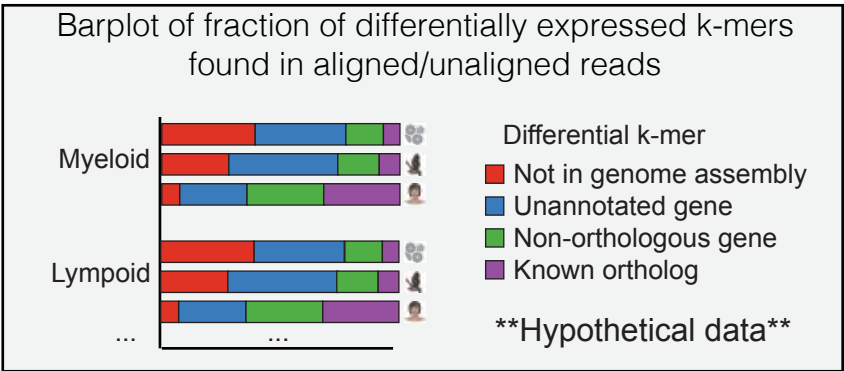


what is the negative control? → random group of cells  
what about bat-specific or human-specific sequences?

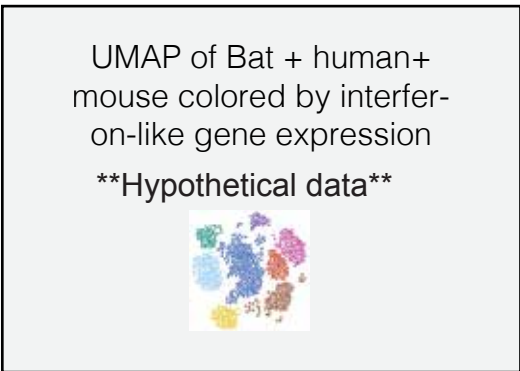
B



C



D



E

