Figure 3 - Find Bat-specific immune adaptations

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

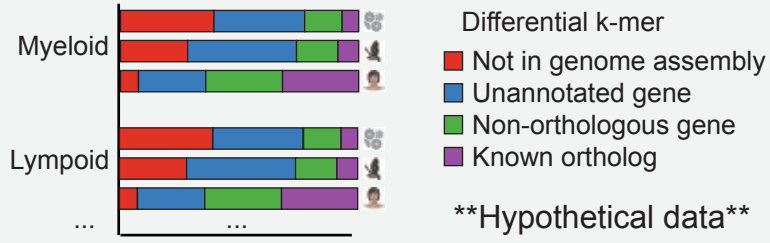
A

UMAP of Bat + human + mouse + lemur

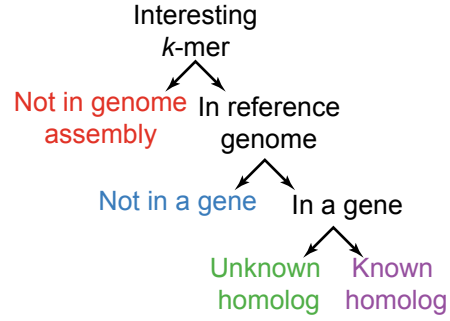
Hypothetical data

B

Barplot of fraction of differentially expressed k-mers found in aligned/unaligned reads



Hypothetical data



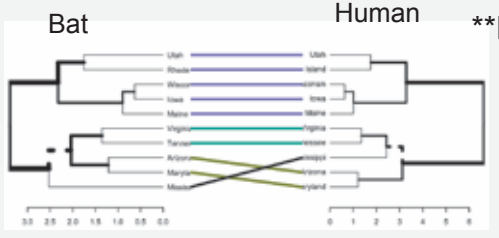
C

UMAP of Bat + human+ mouse + lemur colored by interferon-like gene expression

Hypothetical data

D

Alignment of assembled Bat interferon proteins sequences and known human ones



Hypothetical data