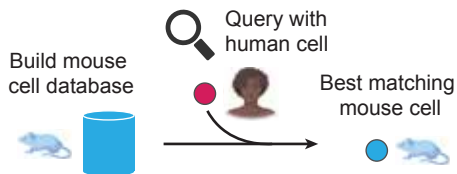


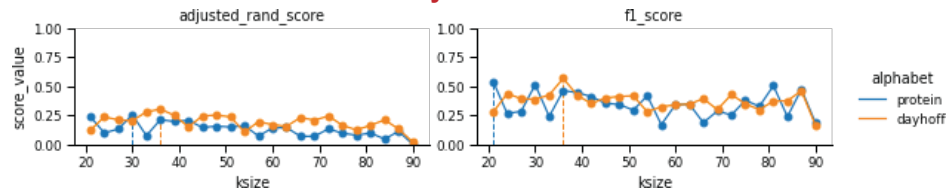
Figure 3: Cell databases from degenerate protein alphabet k-mers can reliably identify cell types across individuals of different species

A



B

Re-do with many random subsets



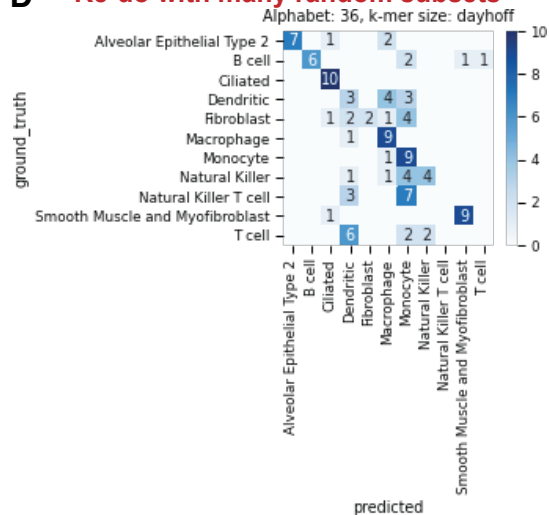
C

Re-do with many random subsets



D

Re-do with many random subsets



E

