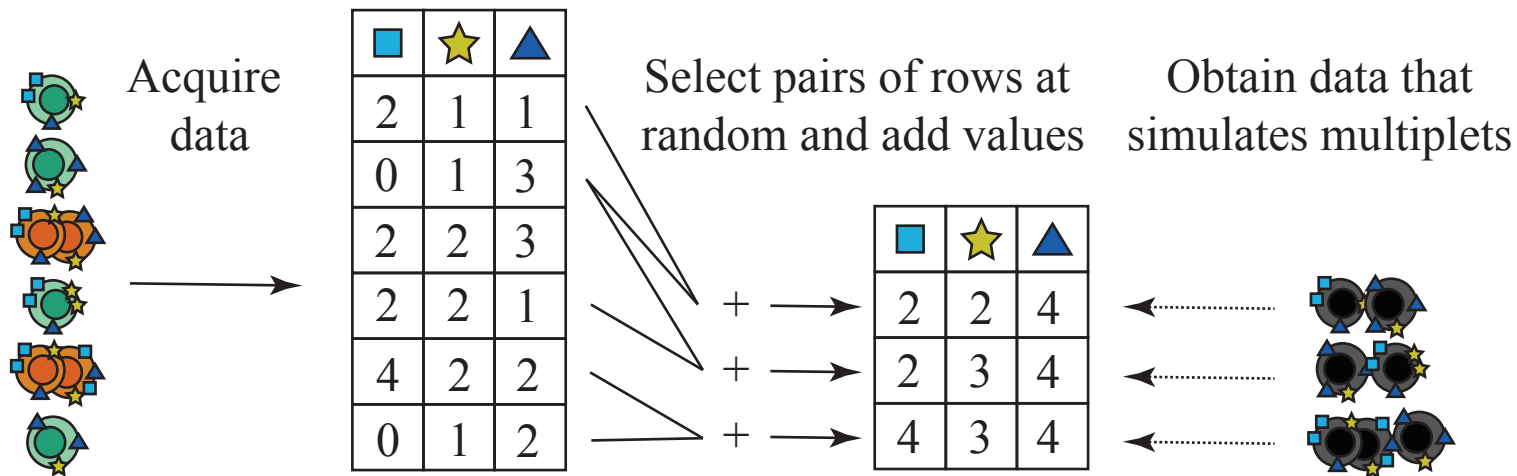
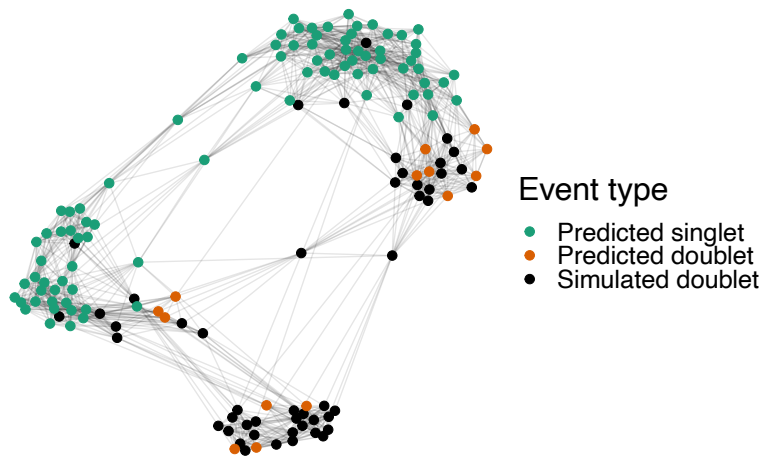


A

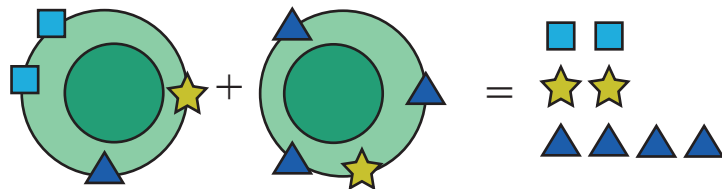


B Call doublets based on % of simulated doublets among neighbors



C

Cytometry



scRNA-seq

