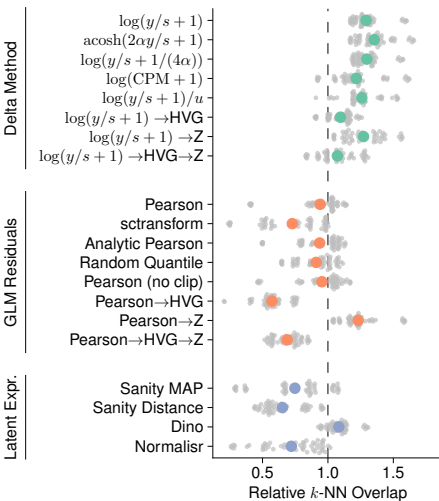


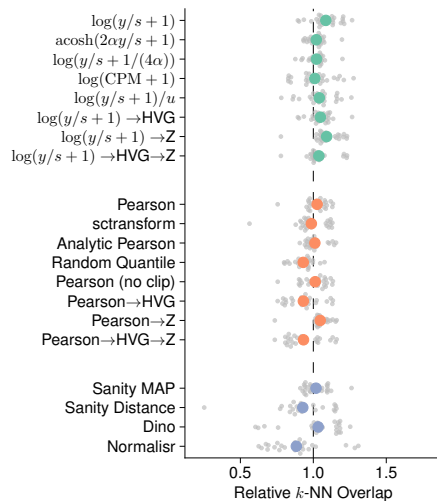
## (A) Consistency

10X gene subset A vs. gene subset B



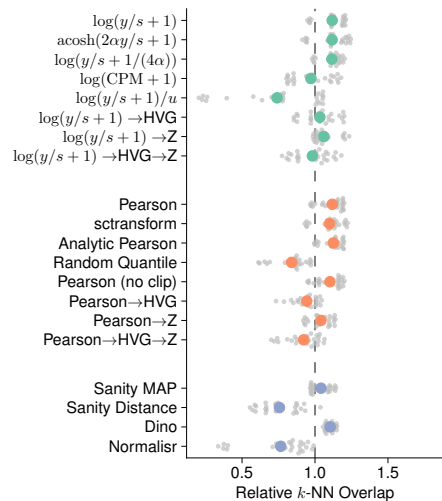
## (B) Simulation

Ground truth vs. simulated counts



## (C) Downsampling

Original vs. downsampled Smart-seq3 data



## (D) $k$ -NN overlap depends on #PCA-dimensions

