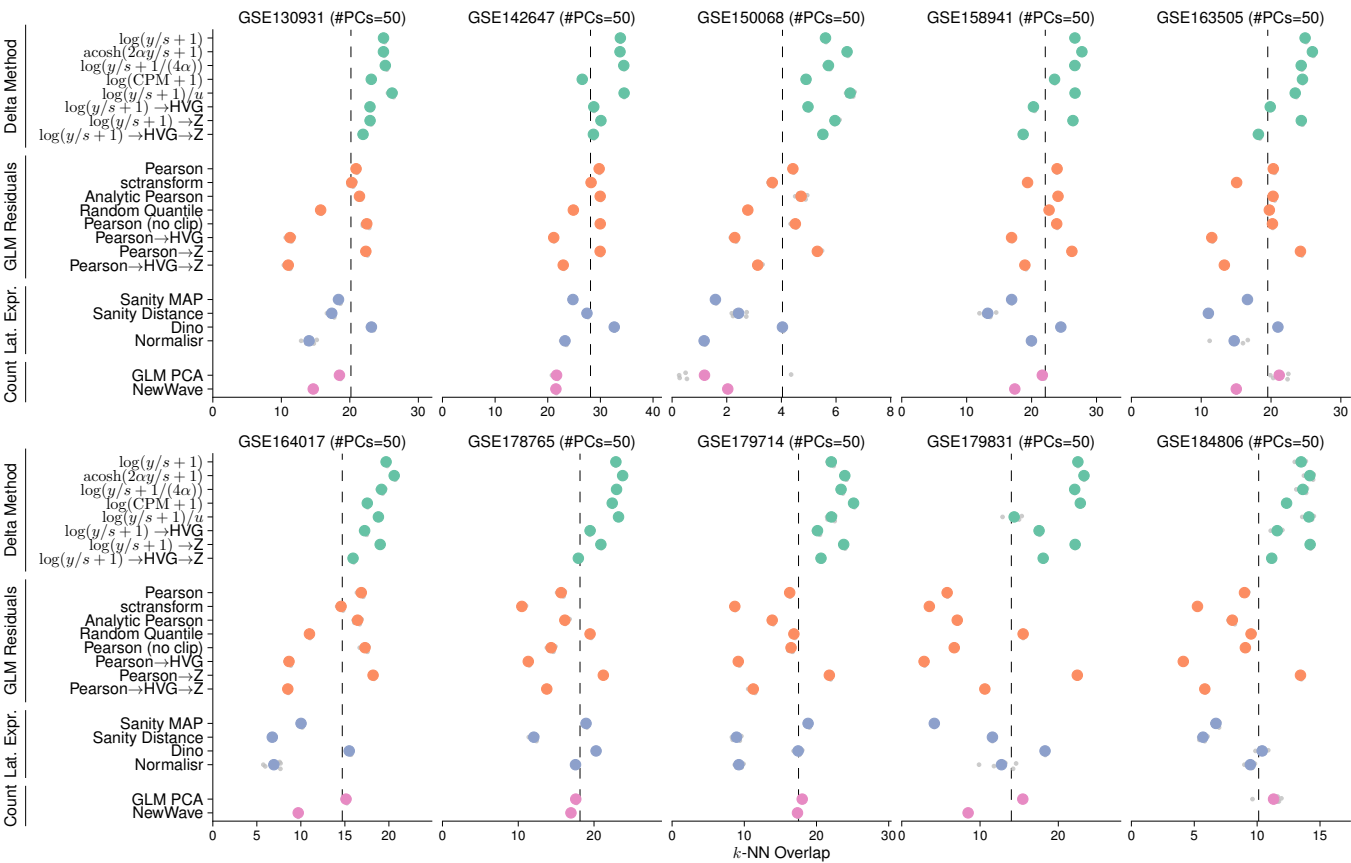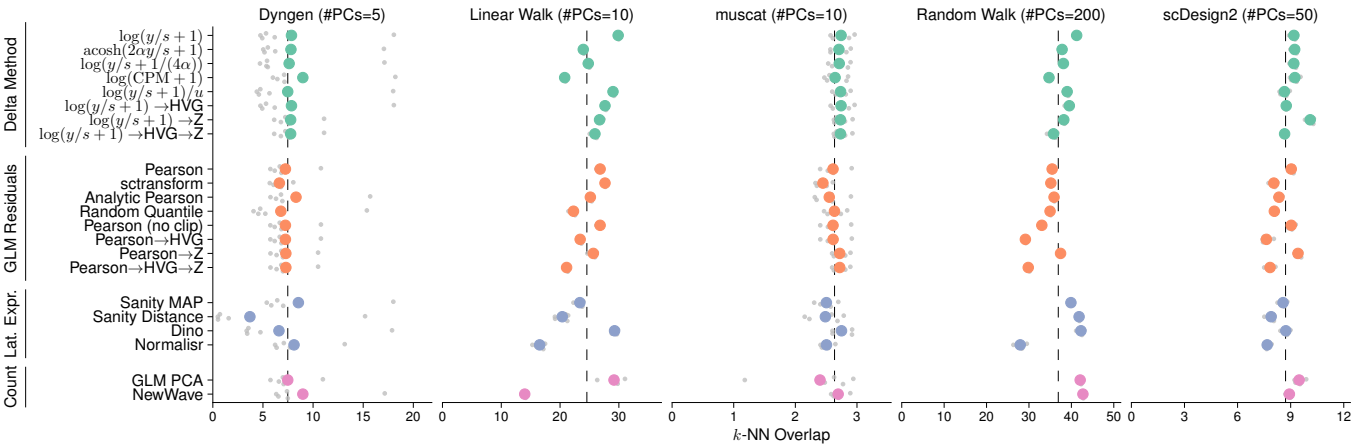**(A) Consistency: 10X gene subset A vs. gene subset B**

(B) Simulation: Ground truth vs. simulated counts

(C) Downsampling: Original vs. downsampled Smart-seq3 data