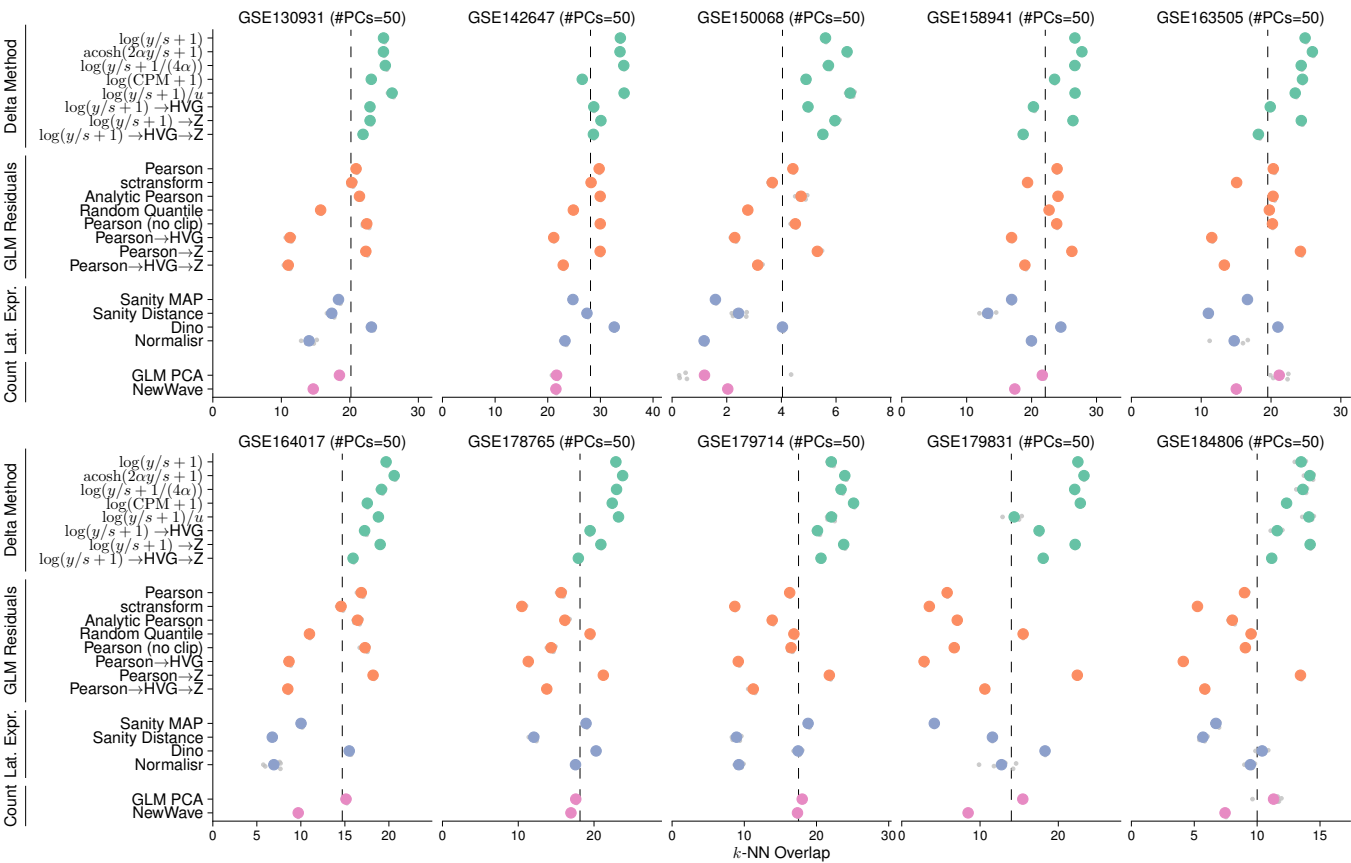
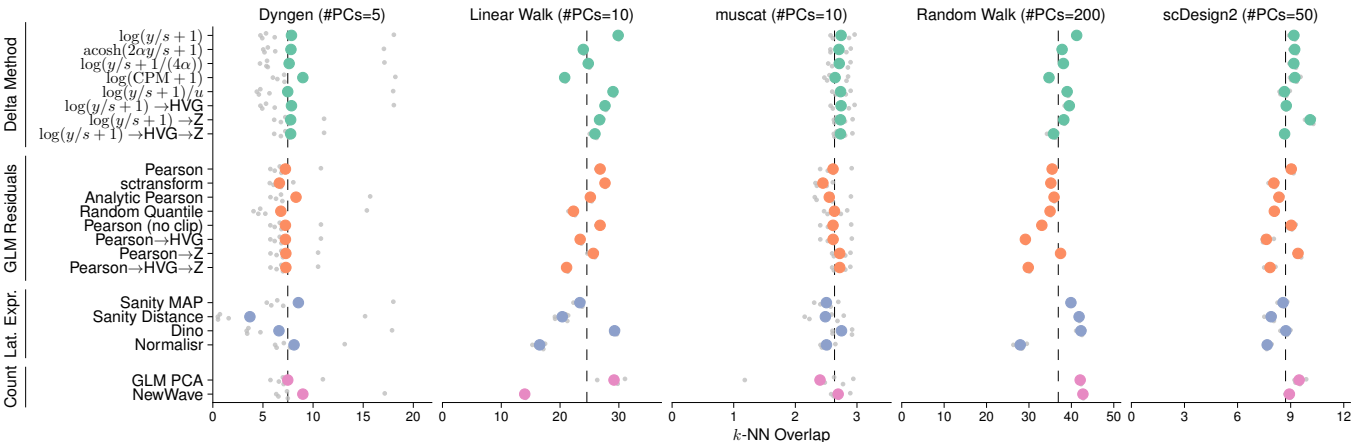


(A) Consistency: 10X gene subset 1 vs. gene subset 2**(B) Simulation: Ground truth vs. simulated counts****(C) Downsampling: Original vs. downsampled Smart-seq3 data**