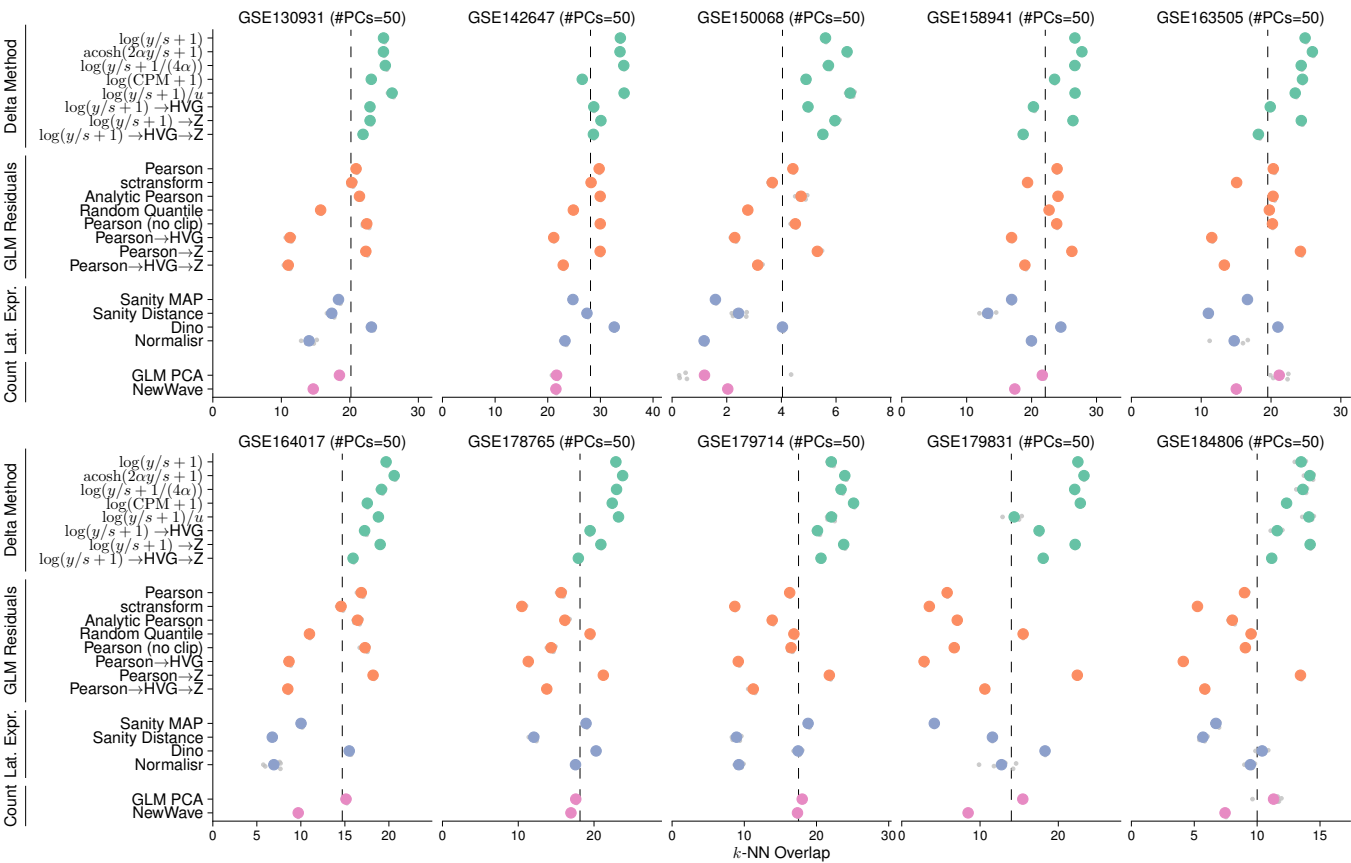
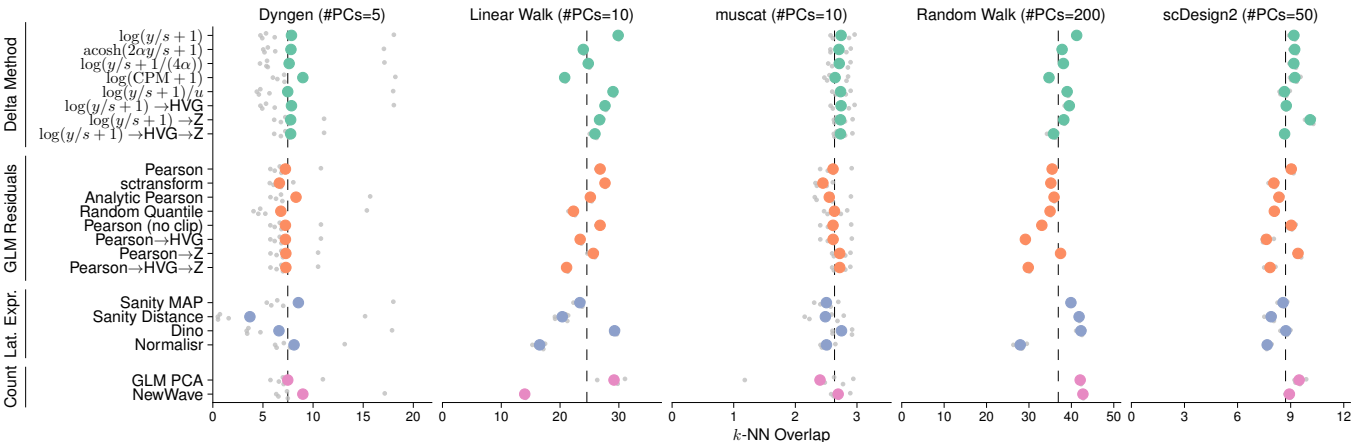


**(A) Consistency: 10X gene subset A vs. gene subset B****(B) Simulation: Ground truth vs. simulated counts****(C) Downsampling: Original vs. downsampled Smart-seq3 data**