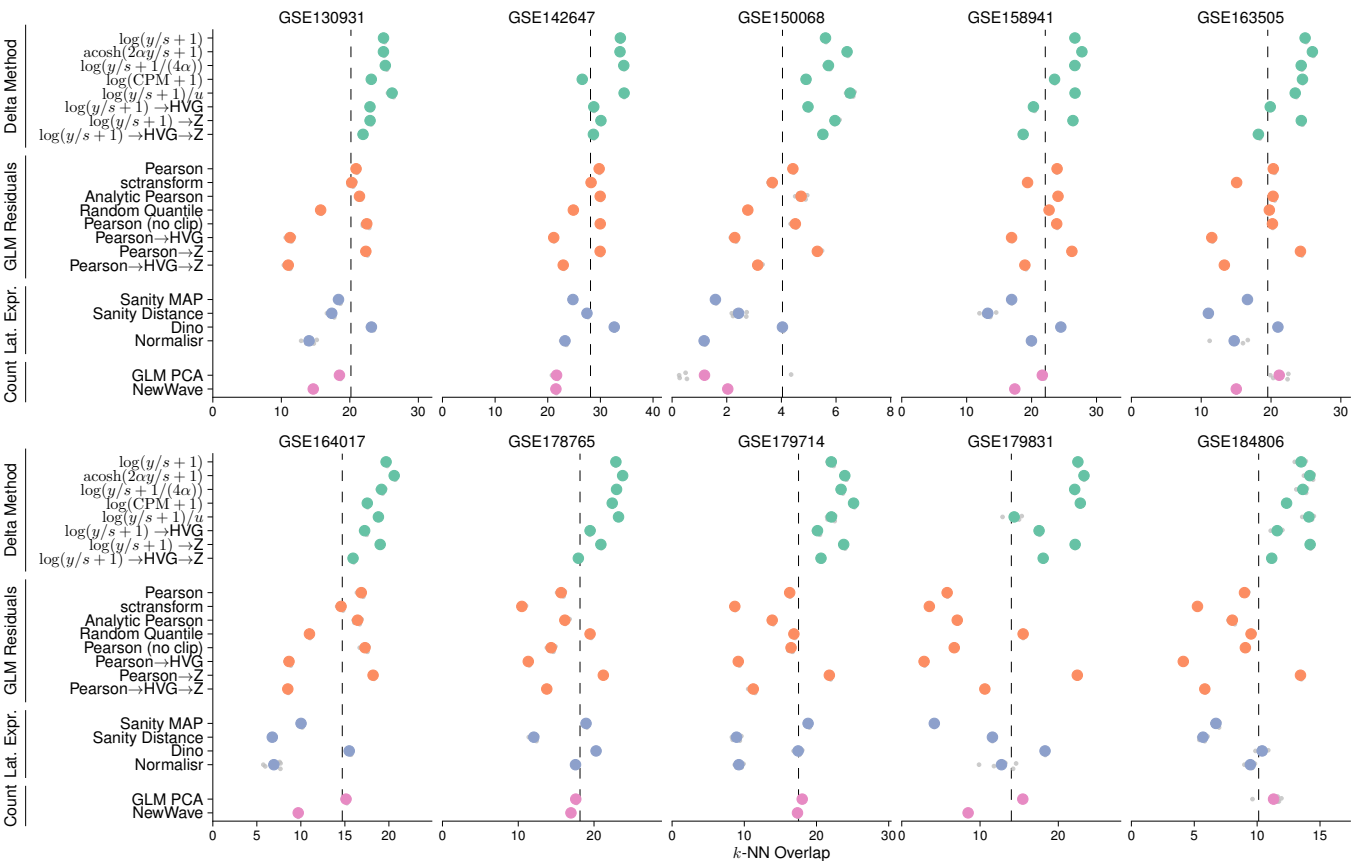
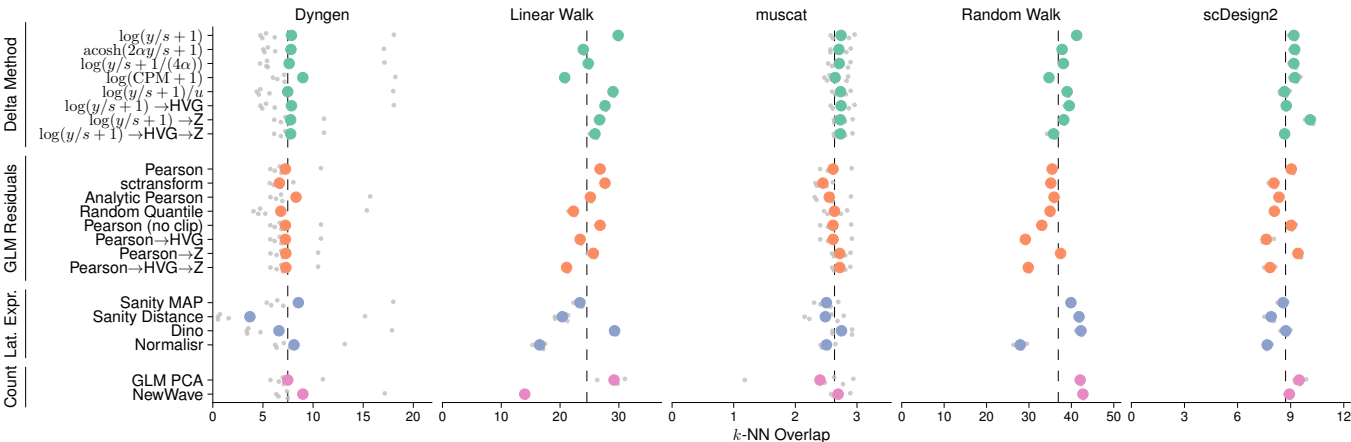


# (A) Consistency: 10X gene subset A vs. gene subset B



# (B) Simulation: Ground truth vs. simulated counts



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

