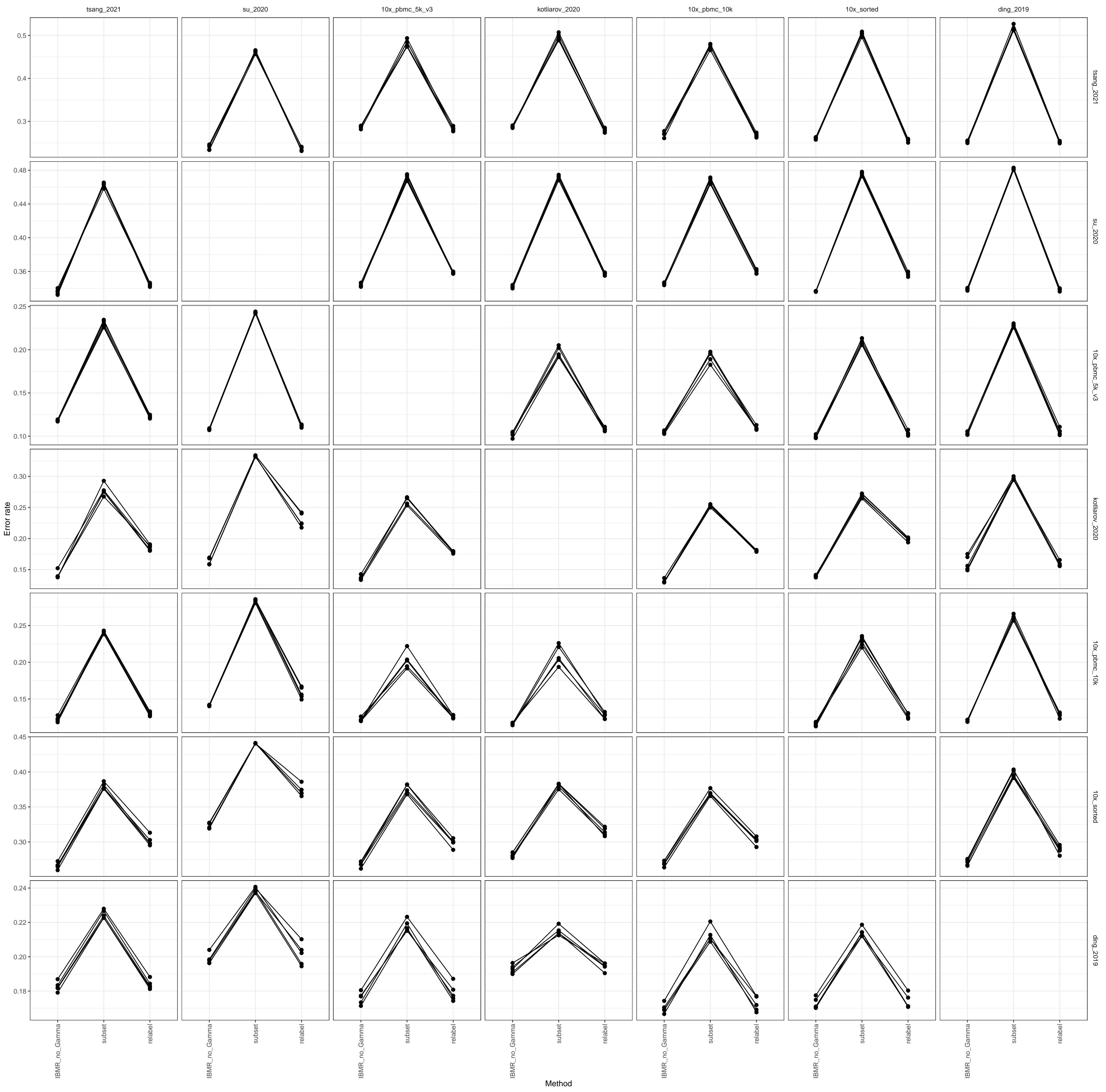
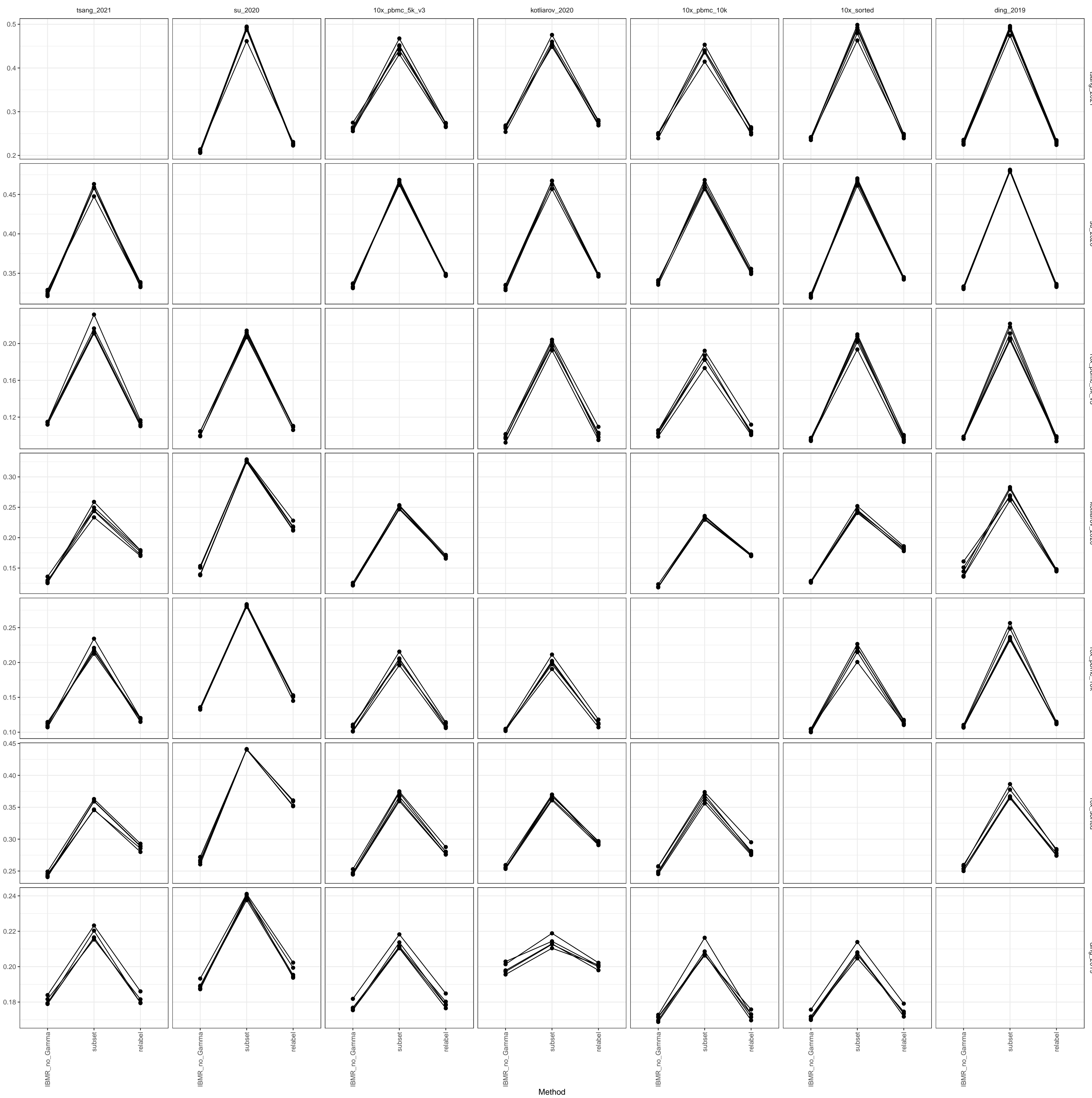


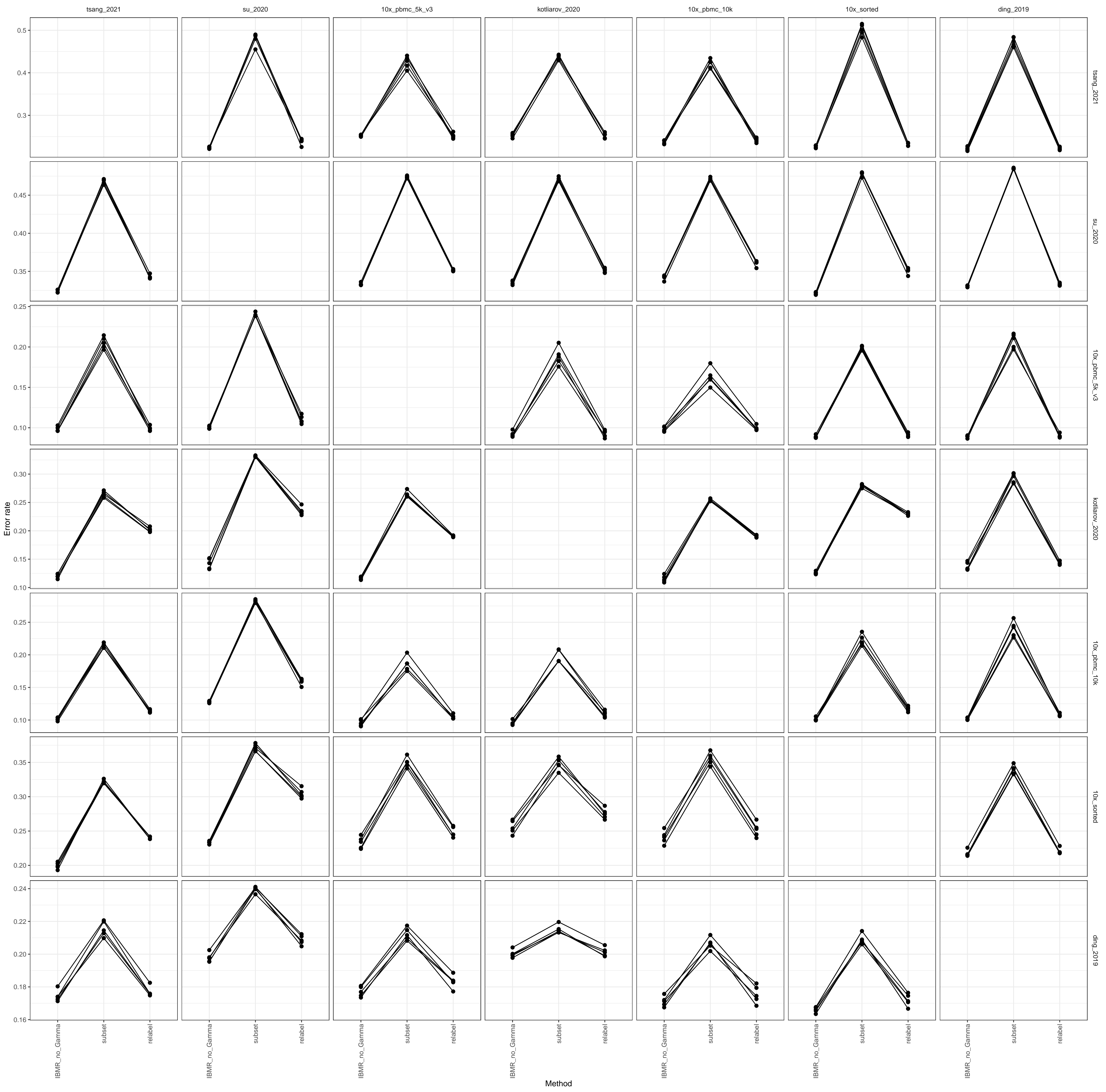
n_sample = 5000; n_genes = 250



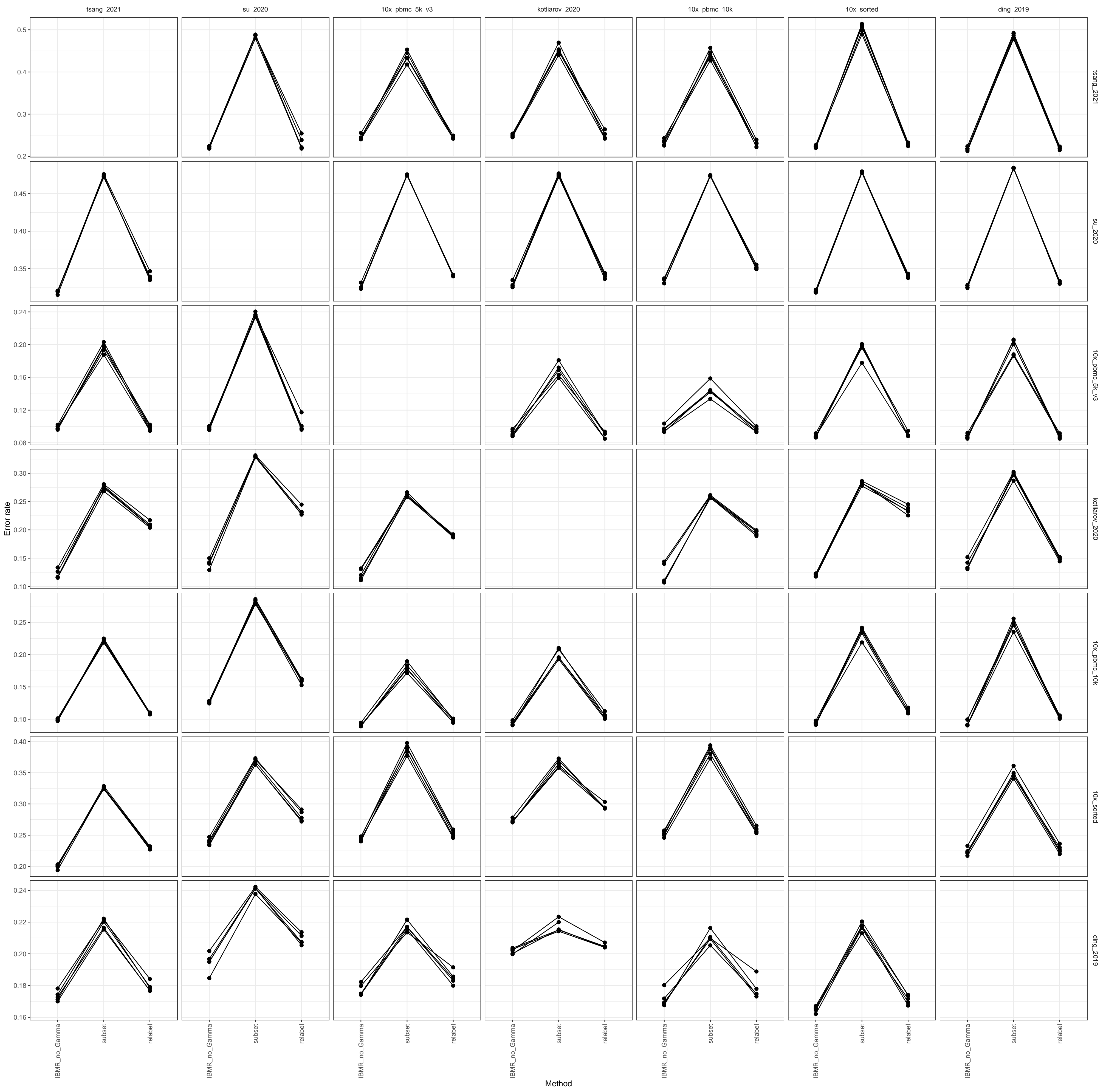
n_sample = 5000; n_genes = 500



n_sample = 5000; n_genes = 1000



n_sample = 5000; n_genes = 2000



n_sample = 10000; n_genes = 1000

