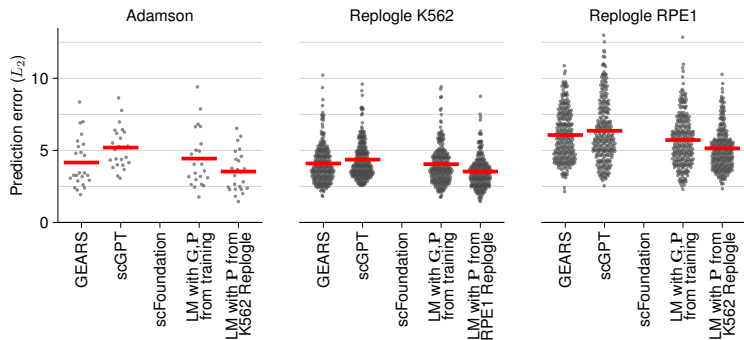
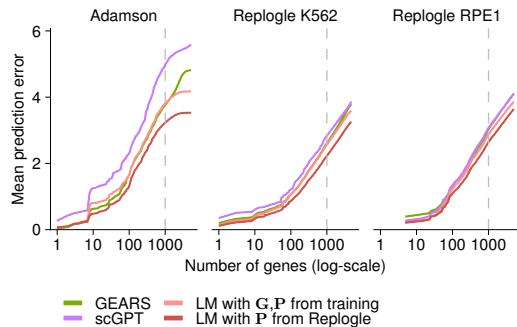


**(A) Single perturbation prediction error****(B) Prediction error stratified by gene set size****(C) Plugging different embeddings into the linear model**