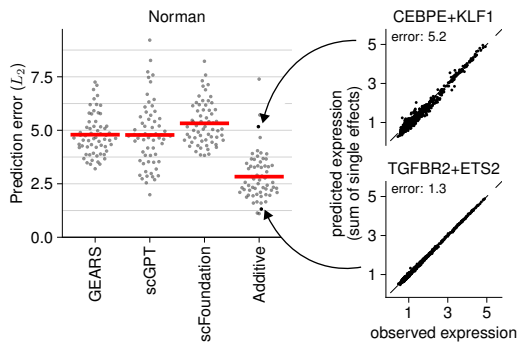
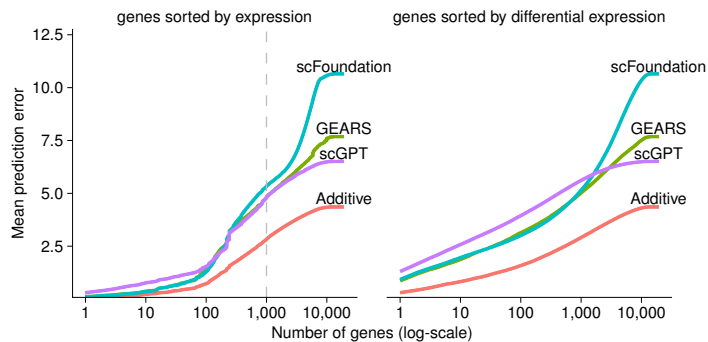


(A) Double perturbation prediction error**(B) Prediction error stratified by gene set size****(C) Prediction of non-additive perturbation effects**