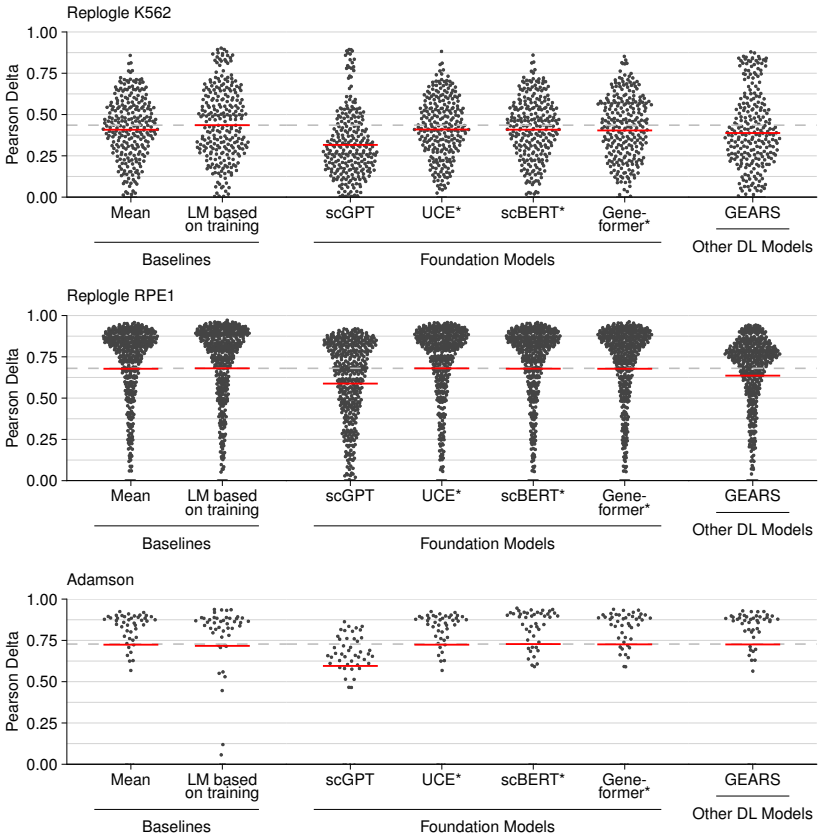
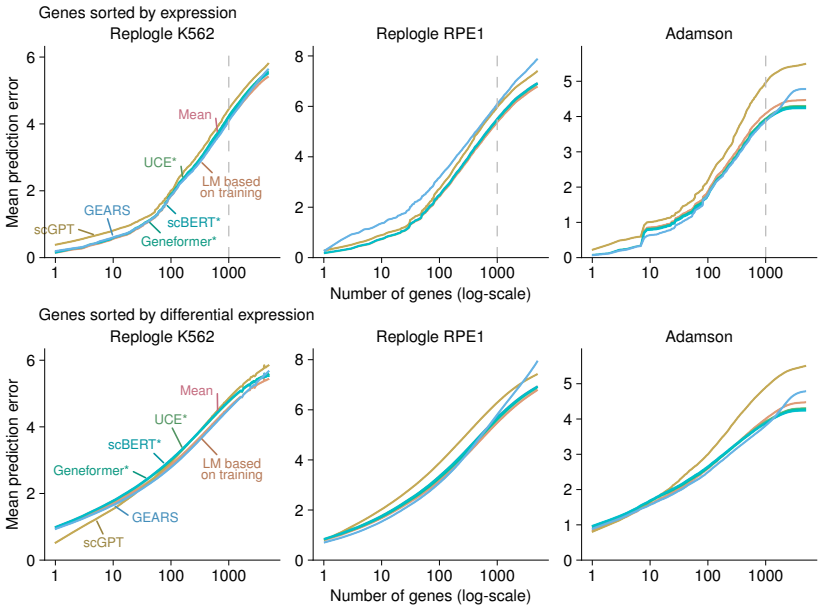
(A) Single unseen perturbation prediction correlation



(B) Prediction error stratified by the considered gene sets