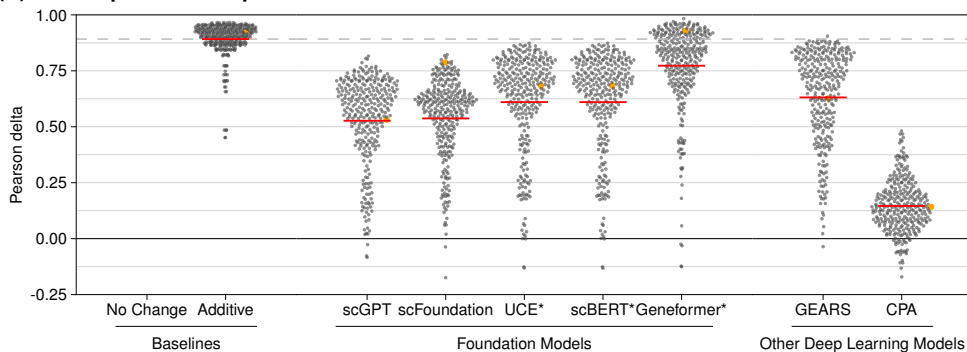**(A) Double perturbation prediction correlation**

(B) Prediction error stratified by the considered gene sets