(A) Double perturbation prediction correlation
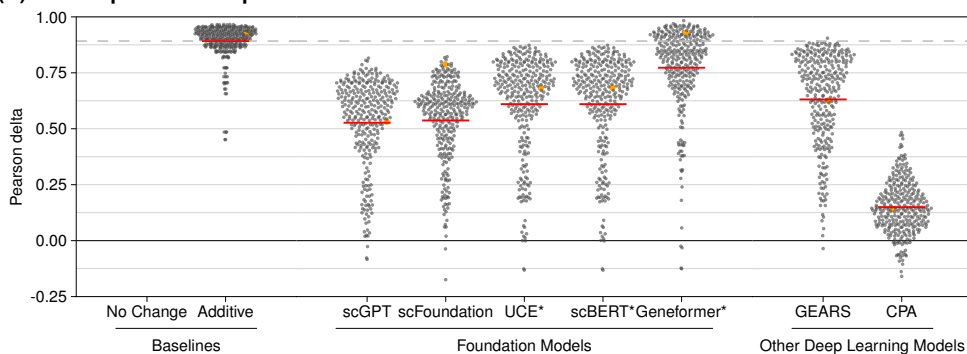

(B) Prediction error stratified by the number of considered genes