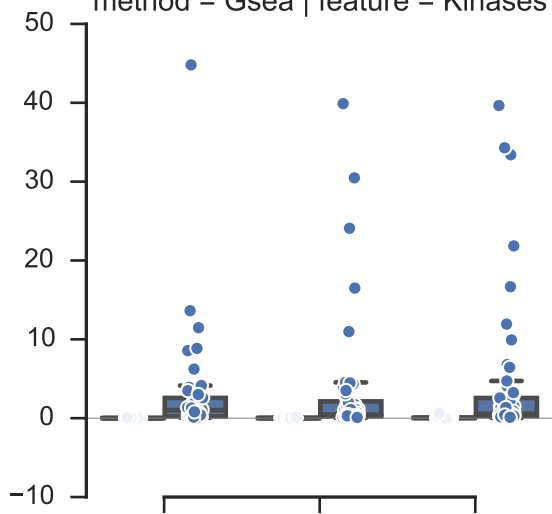
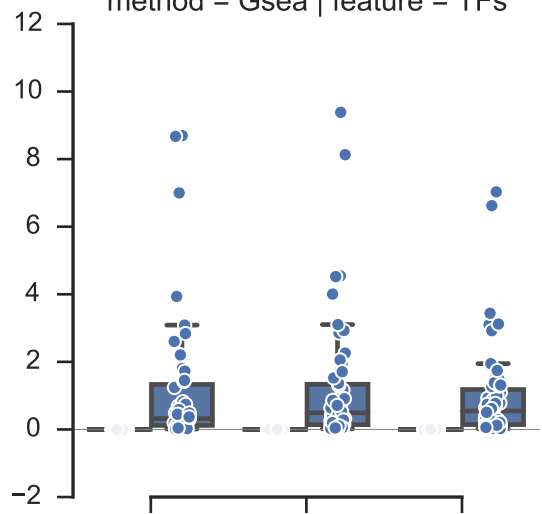


method = Gsea | feature = Kinases

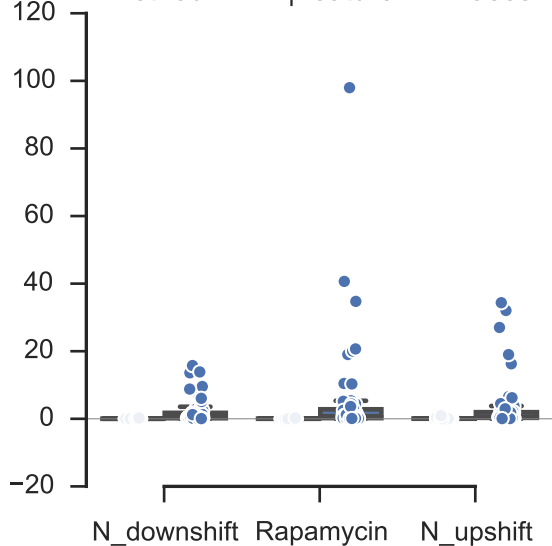


method = Gsea | feature = TFs



● test
● train

method = Lm | feature = Kinases



method = Lm | feature = TFs

