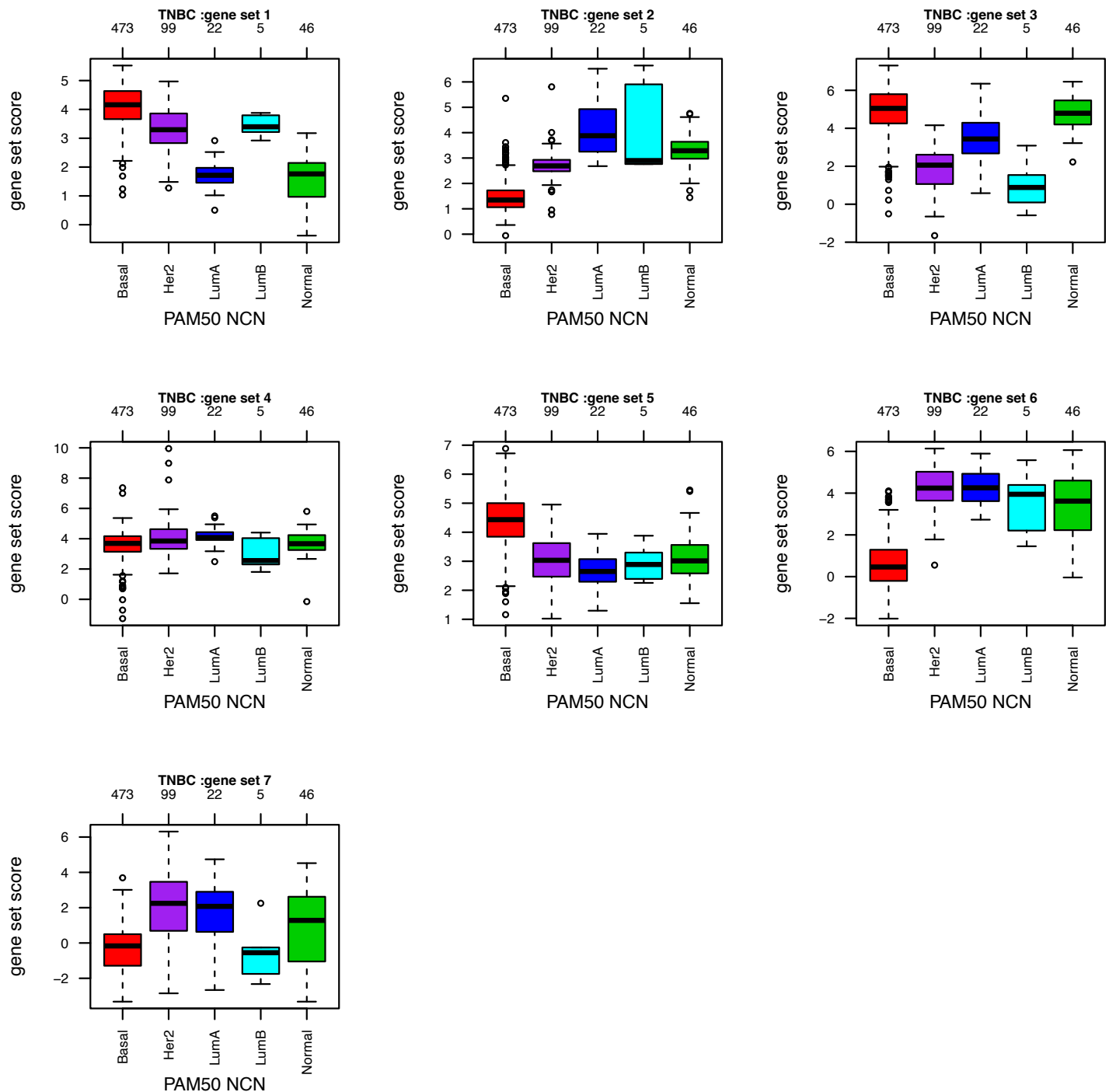
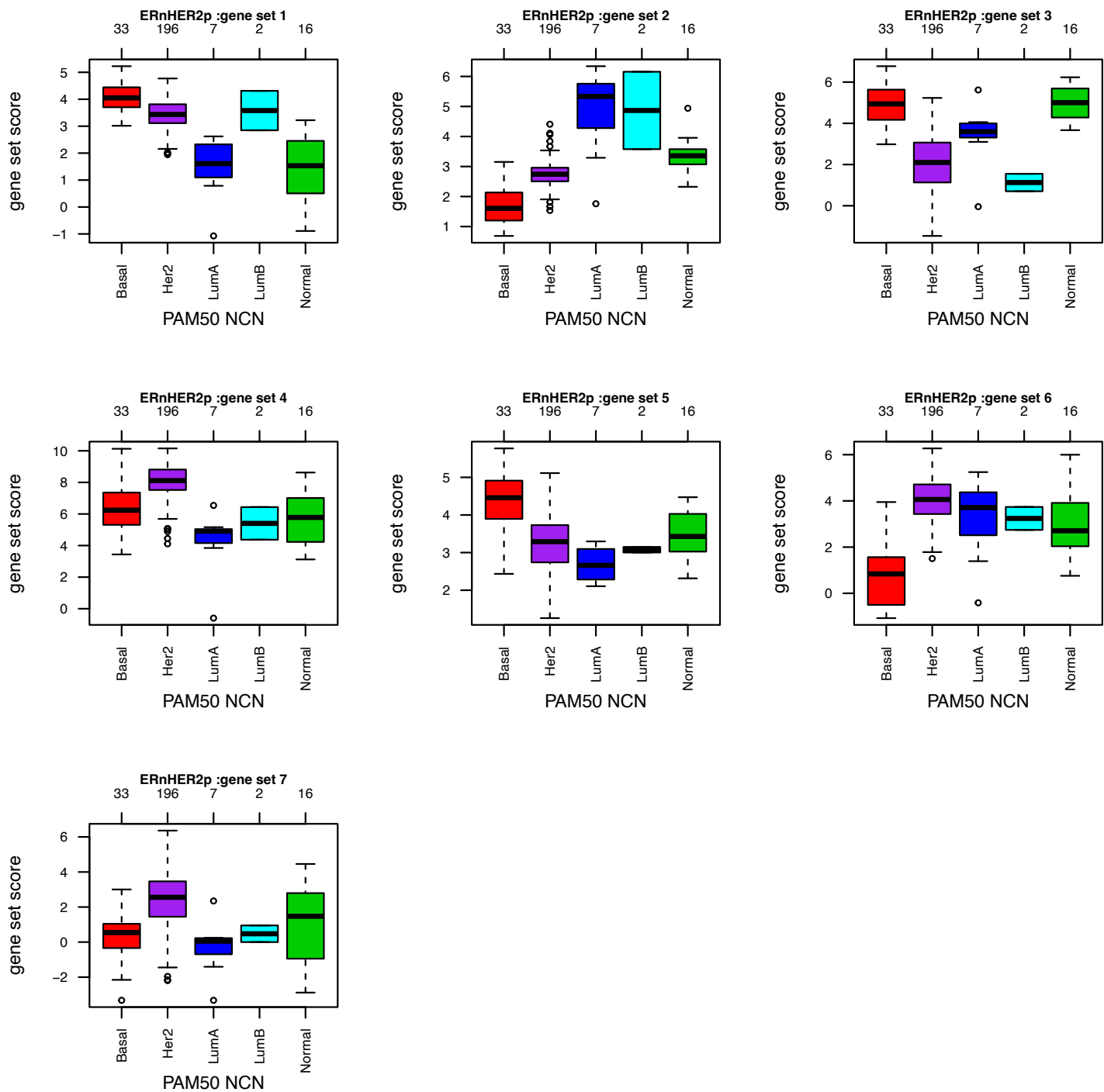


A) TNBC

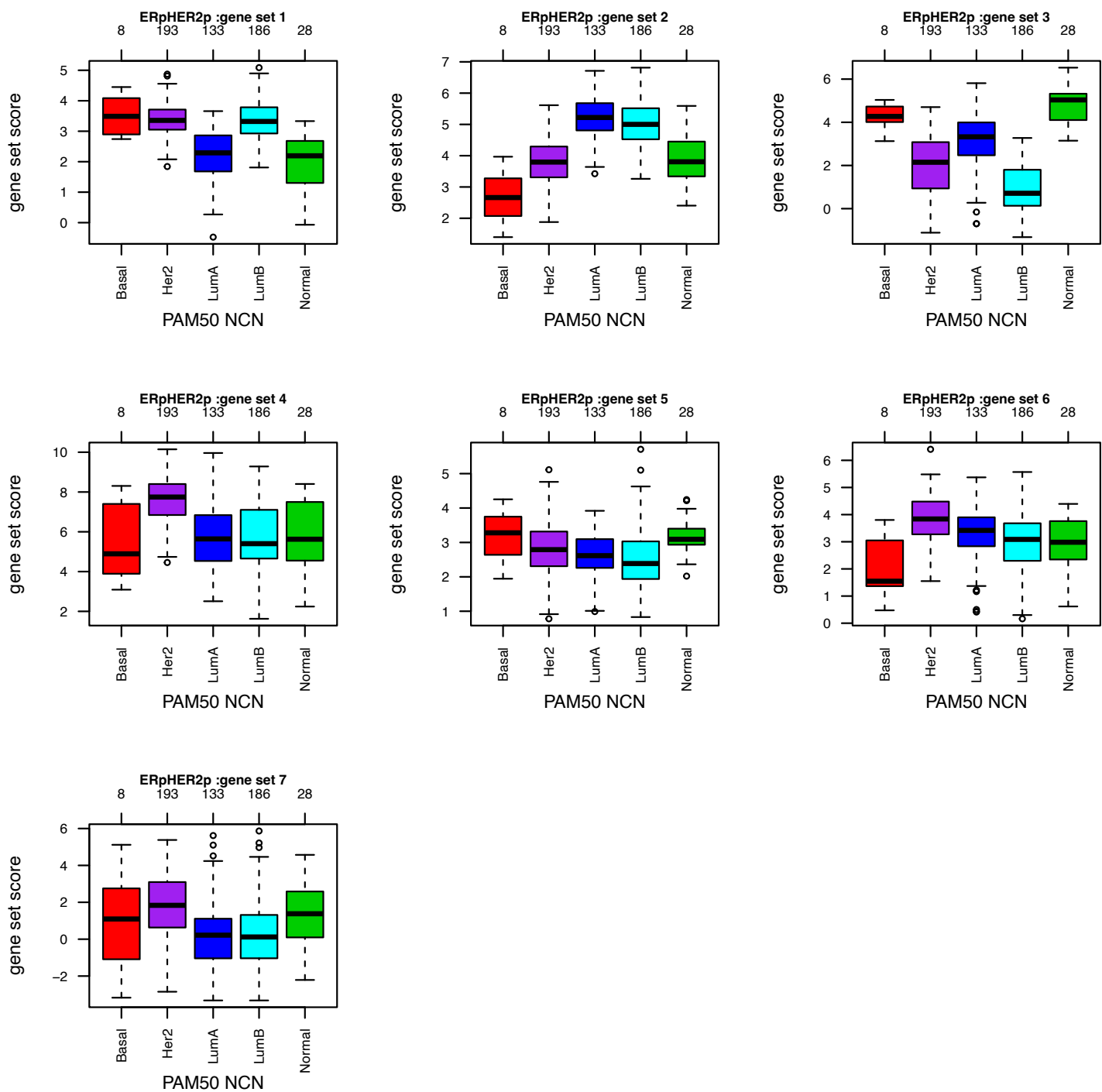


Supplementary Figure S1. Gene set score expression in breast cancer subgroups defined by ER, PR, and HER2 status and stratified by PAM50 subtypes, and versus biological expression metagenes. The gene set score is the average log2 FPKM of genes included in the gene set. (A) TNBC tumors. (B) ERnHER2p tumors (C) ERpHER2p tumors. (D) ERpHER2n tumors. (E) Mean gene set expression (FPKM) of gene set 1, 2, and 3 versus rank scores for specific biological metagenes representative of proliferation, steroid response, and basal/keratin expression as defined by Fredlund et al. BCR 2012. Correlations calculated using Spearman correlation using all 6233 tumors.

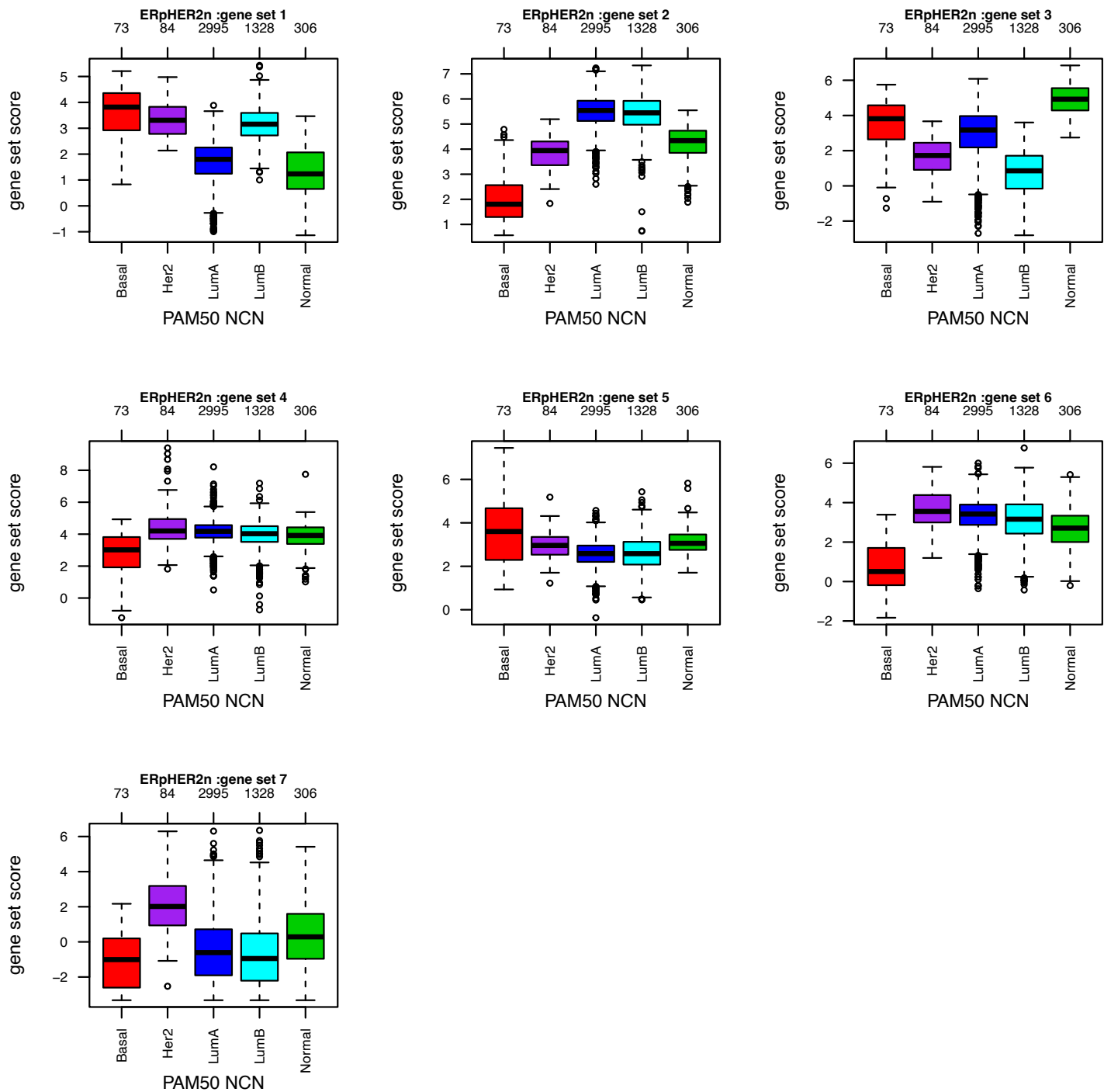
B) ERnHER2p



C) ERpHER2p



D) ERpHER2n



E) Gene set scores vs Fredlund metagenes

