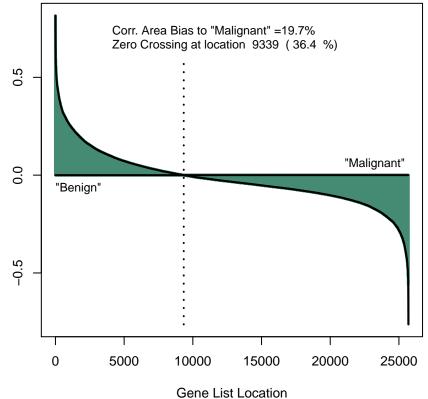
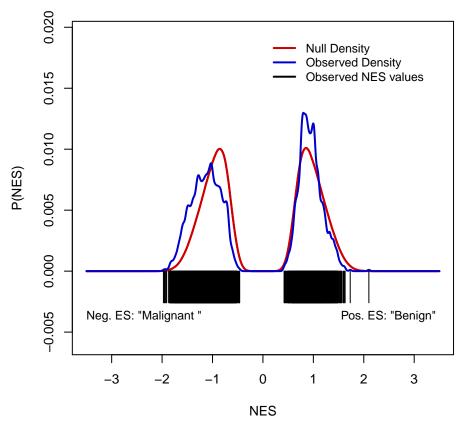
Gene List Correlation (S2N) Profile

Corr. Area Bias to "Malignant" =19.7%

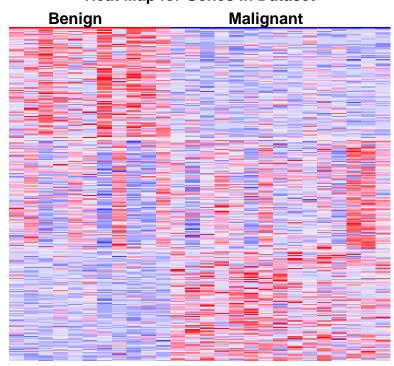


Signal to Noise Ratio (S2N)

Global Observed and Null Densities (Area Normalized)



Heat Map for Genes in Dataset



p-values vs. NES

