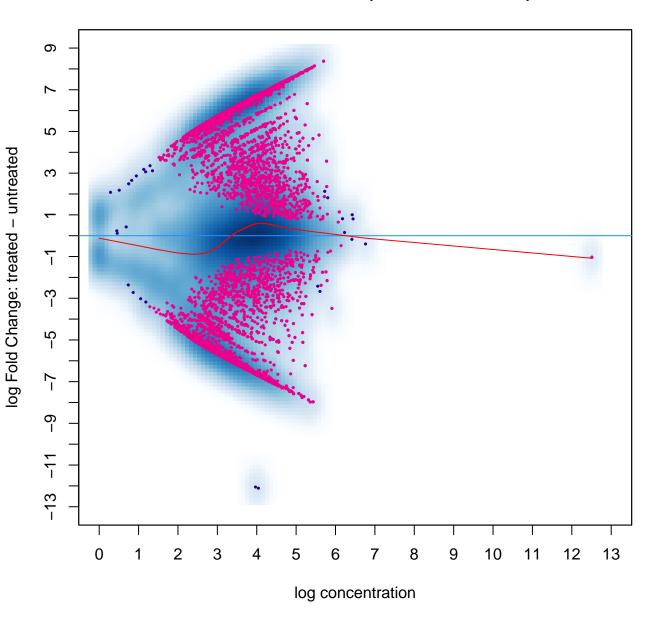
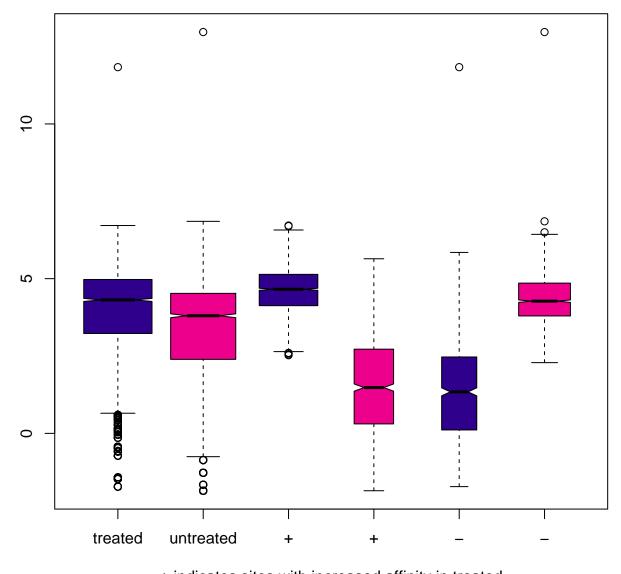


## treated vs. untreated (4150 FDR < 0.050)



Contrast: treated vs. untreated [4150 FDR<=0.050] 7.5 **-**-log10(FDR) Legend FDR >0.05 FDR<=0.05 2.5 -0.0 -5 -10 log2 Fold Change

## treated vs. untreated



log2 normalized reads in binding sites

+ indicates sites with increased affinity in treated - indicates sites with increased affinity in untreated

