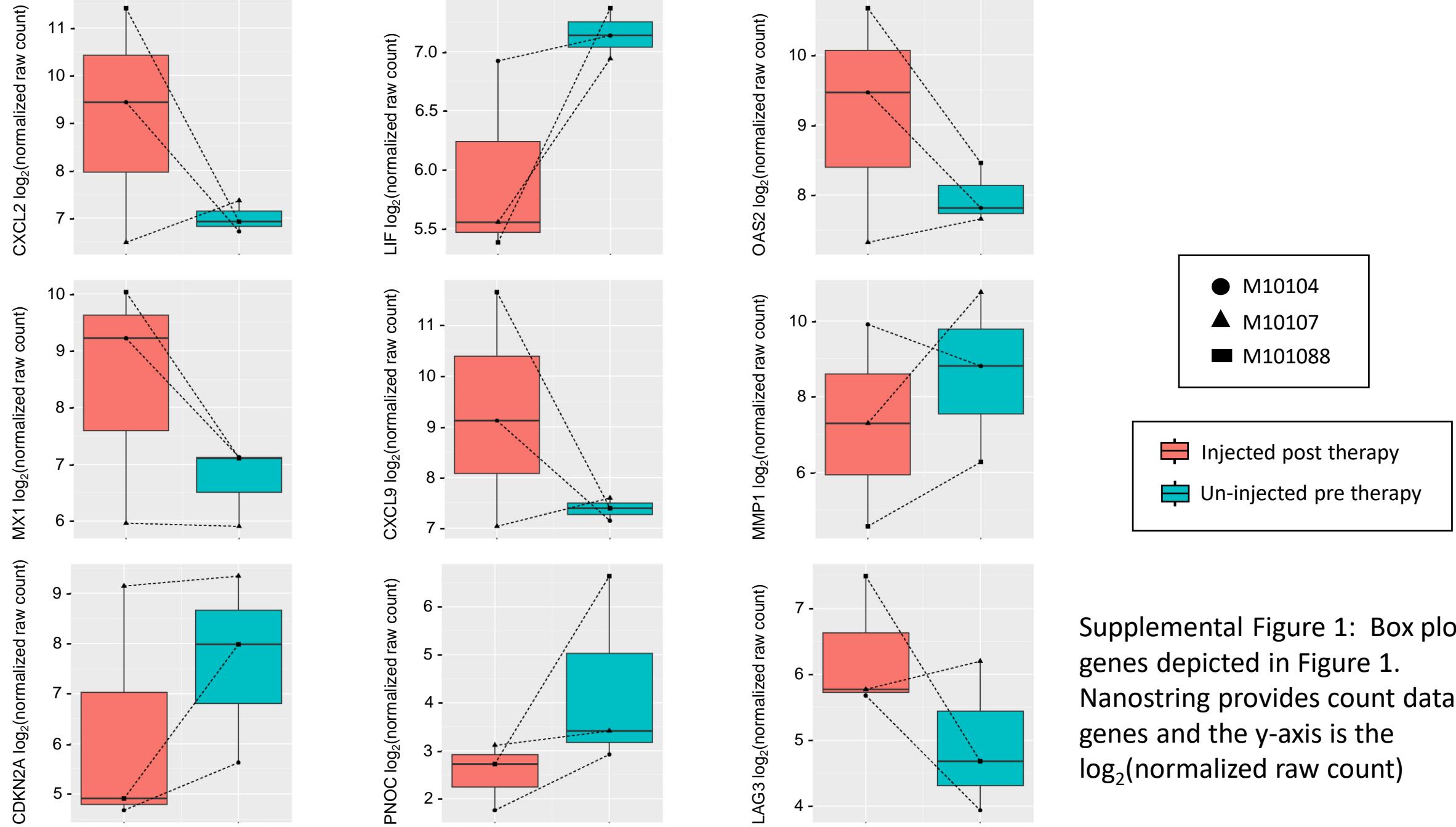
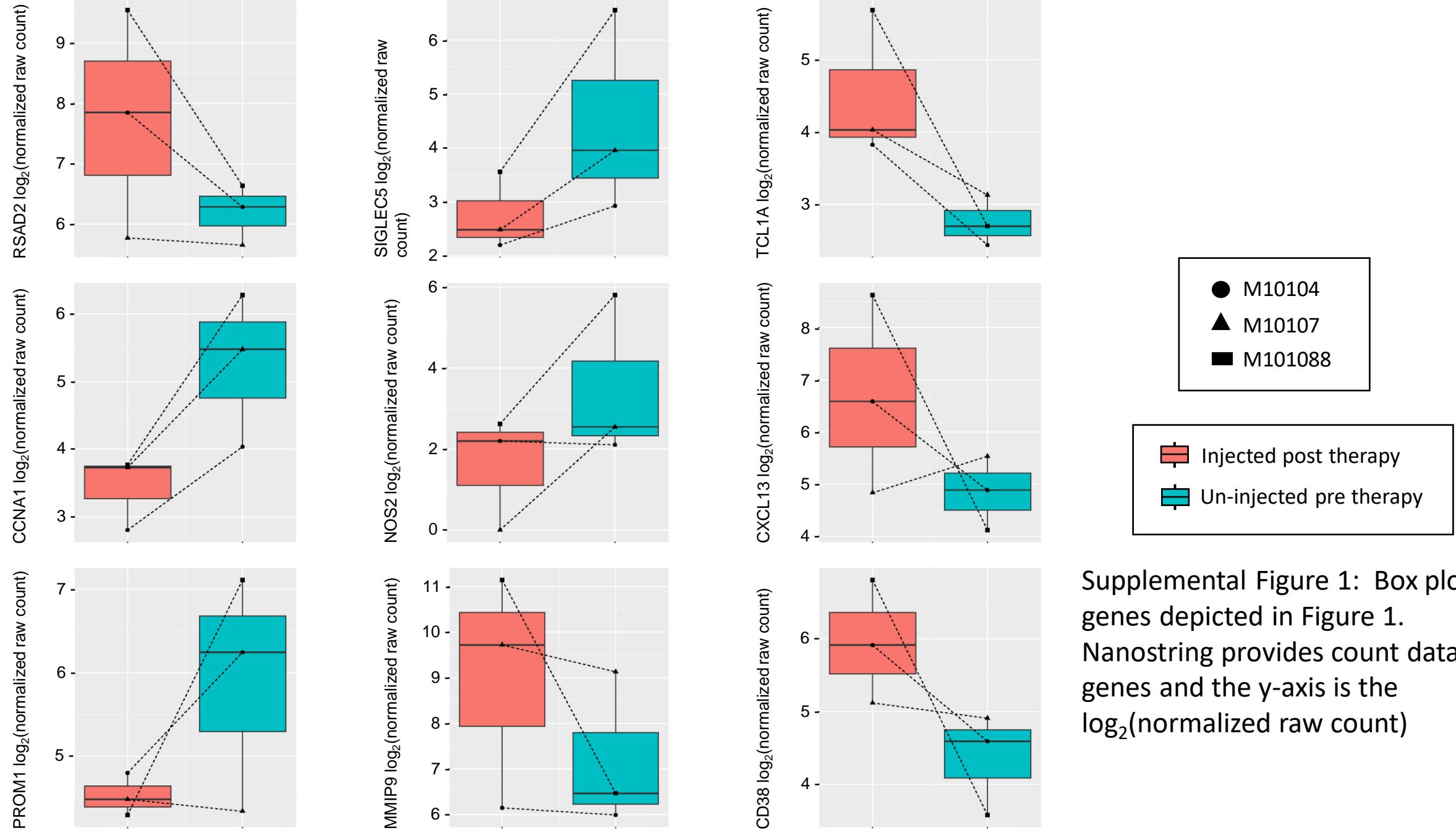


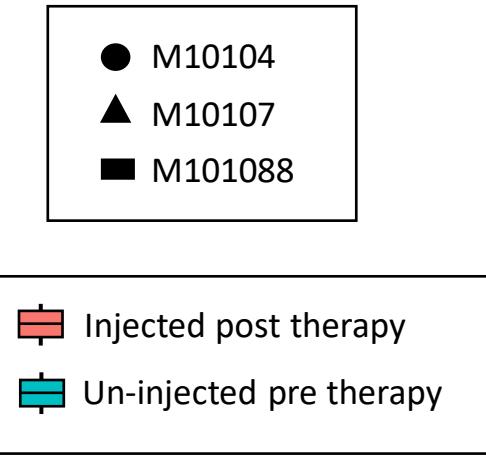
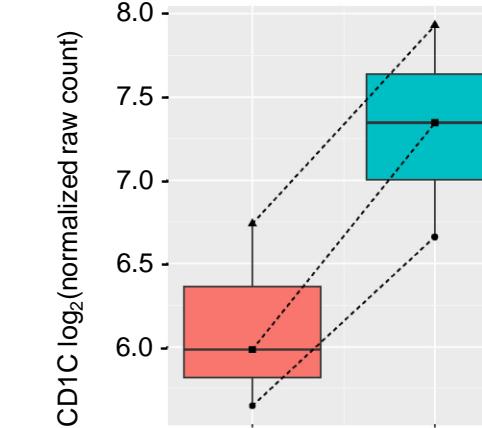
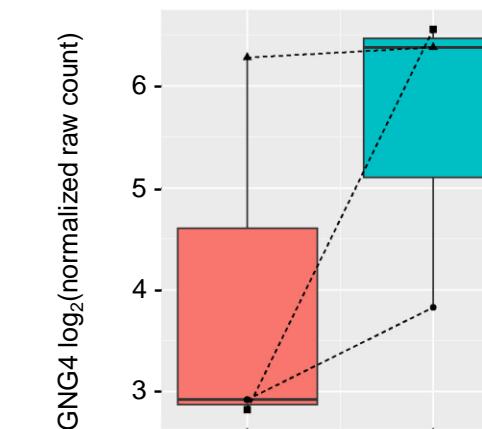
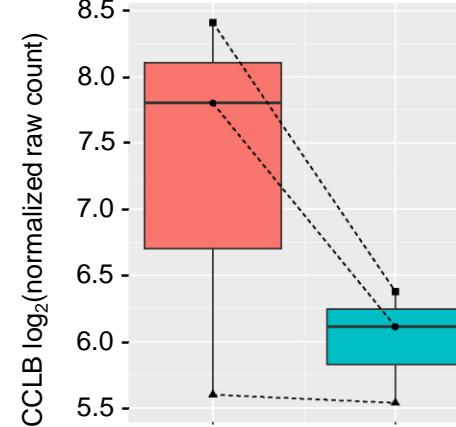
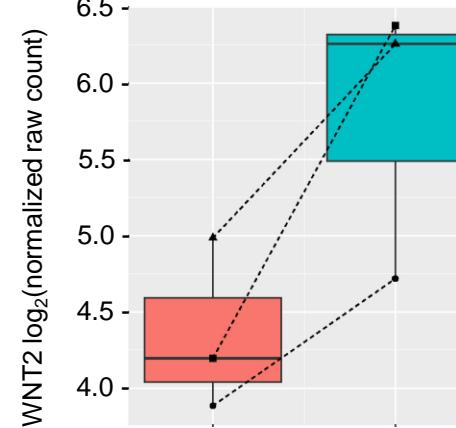
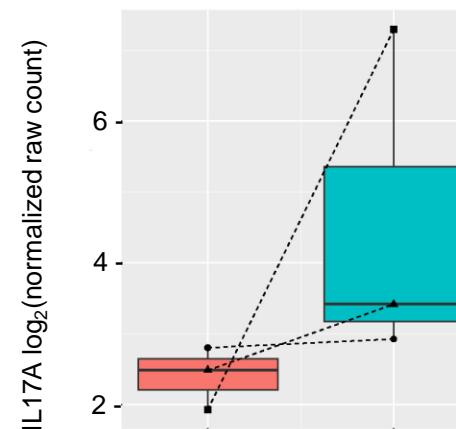
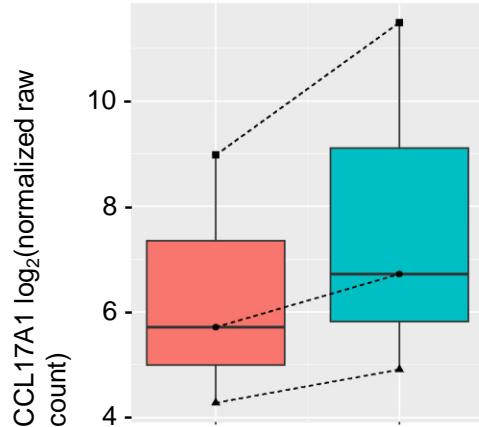
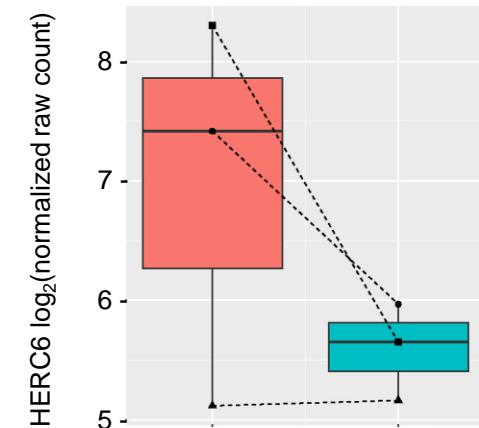
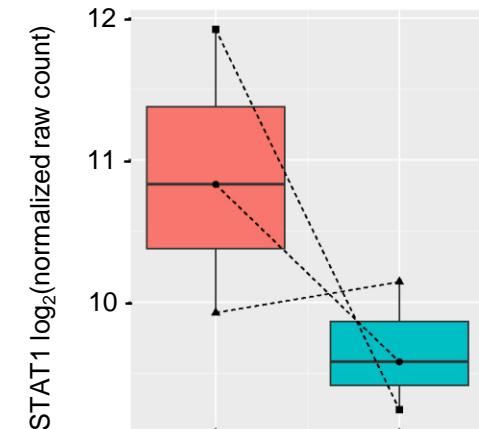
Supplemental Figure 1: Box plot of genes depicted in Figure 1. Nanostring provides count data for genes and the y-axis is the $\log_2(\text{normalized raw count})$



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