

Figure S1. Boxplot of the normalization and base-2 logarithm conversion for the group containing three datasets, GSE76250, GSE86945 and GSE86946.

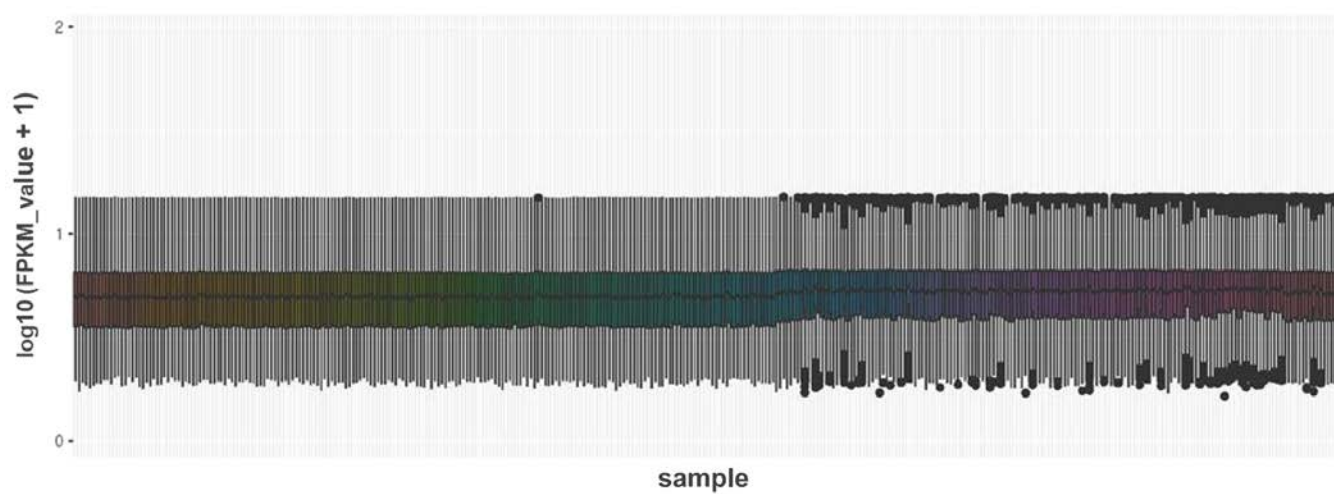


Figure S2. Heatmaps of the top 200 differentially expressed long non-coding RNAs identified from the The Cancer Genome Atlas database.

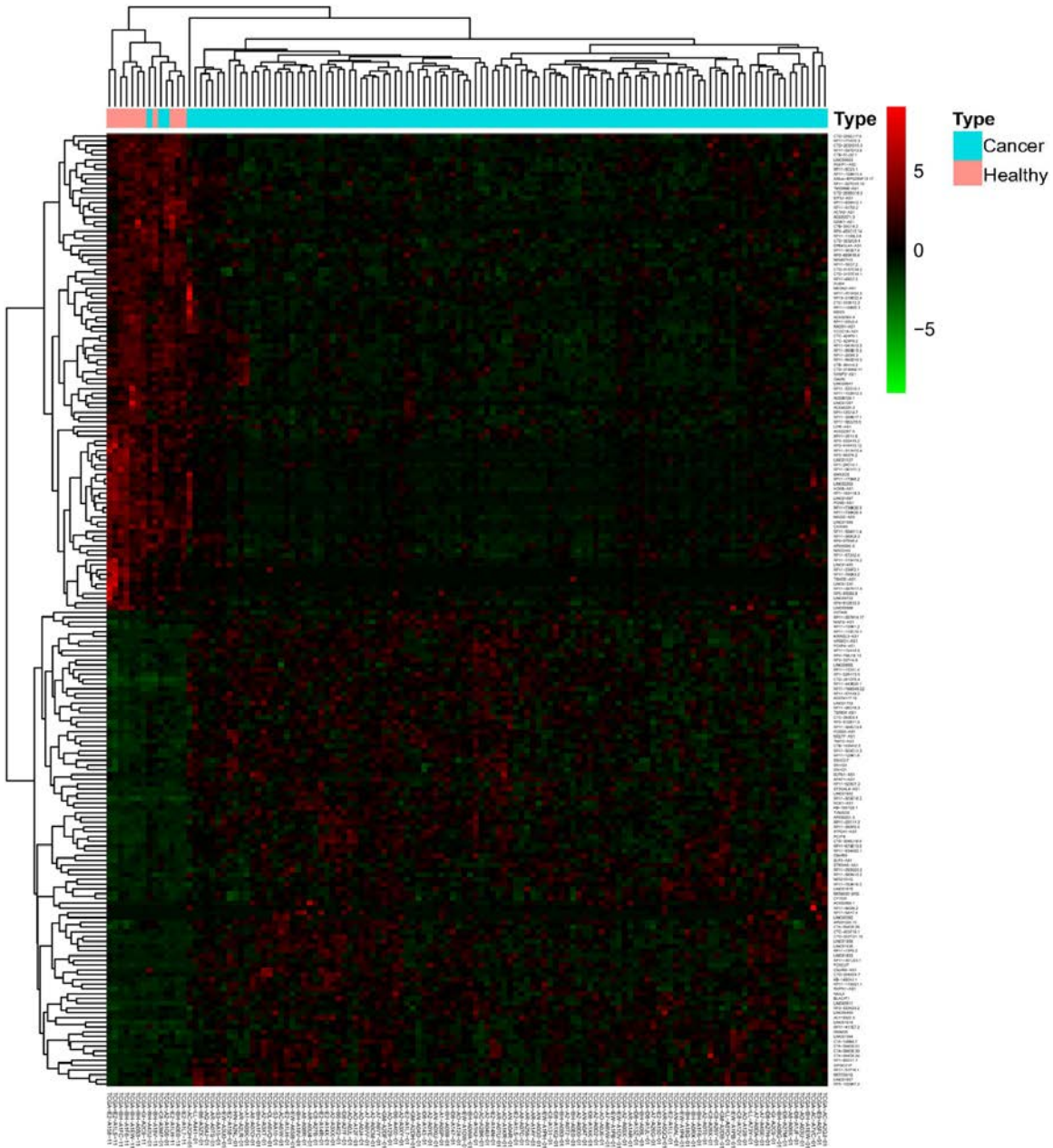


Figure S3. Heatmaps of the top 200 differentially expressed mRNAs identified from the The Cancer Genome Atlas database.

