Figure S6

10⁰

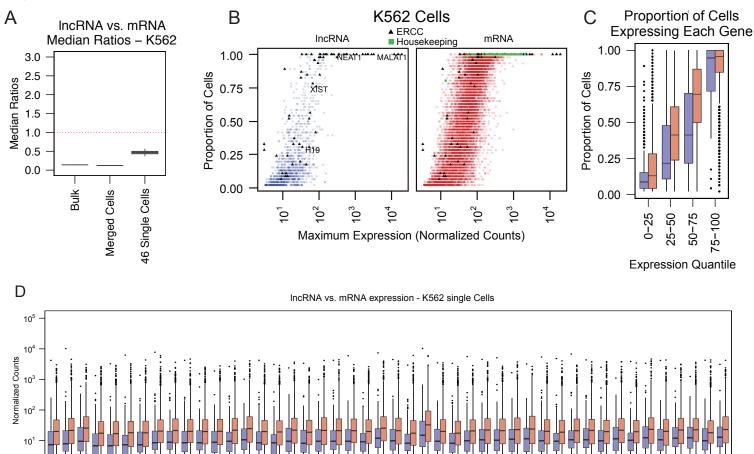


Figure S6. Single cell transcriptomics of IncRNA expression in K562 cell cultures

A) Distributions of median IncRNA expression to median mRNA expression ratios (IncRNA:mRNA) in populations, in silico merged single cells, and single cells from K562 cultures. B) Proportion of K562 cells that expressed each IncRNA (blue) and mRNA (red), separated by maximum expression in single cells. C) Same as in B) but grouped by maximum expression quantile. D) Distributions of non-zero IncRNA (blue) and mRNA (red) expression in 46 single K562 cells. Green squares, housekeeping genes. Black triangles, ERCC Spike-In Controls.