Figure S5. Expression of IncRNAs and mRNAs in single cells and whole tissues

3-4 4-5 5-6 6-7 7-8 8-9

log₂(Normalized Counts+1)

0-0 0-1 1-2 2-3

A) Distributions of non-zero IncRNA (blue) and mRNA (red) expression in 276 single cells from neocortex. The median IncRNA expression and the median mRNA expression for each cell was compared as a ratio and summarized in Figure 3B. B) Distributions of non-zero IncRNA (blue) and mRNA (red) expression in 8 bulk tissue RNA-seq samples using the same set of 1400 IncRNAs and 10929 mRNAs as in A). C) Distributions of non-zero IncRNA (blue) and mRNA (red) expression in in silico merged neocortex single cells. D) Proportion of single neocortex cells that expressed each IncRNA (blue) and mRNA (red), binned by their expression levels in bulk tissues. E) Distributions of housekeeping gene and IncRNA (F) expression levels in neocortex single cells, binned by log2(Normalized Counts + 1). Colors represent number of cells in each bin. Abundant IncRNAs were ranked by their median expression levels across 276 neocortex single cells.