lncRNA group = IDRlessthan0\_01 ::: data cols = Inpa (11 / 109) = (func/total IncRNA) ::: biotype = lincRNA measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.3 0.3 0.2 0.2 -0.2 -0.1 -0.1 0.1 0.0 0.0 0.0 0.0 –2.5 -2 2 -5.0 0 4 2.5 0 minExpr maxExpr nTimesRest 1.5 -0.3 0.4 withinSubset density 1.0 false true 0.2 -0.5 -0.1 -0.0 0.0 -0.0 і –2 -4 2 0 -6 2 0 4 -8 tissSpec 0.6 0.4 0.2 0.0

value of facet label

-2.0 -1.5 -1.0 -0.5