(4/71) = (func/total lncRNA) ::: biotype = antisense measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.4 0.4 0.3 -0.3 -0.2 0.2 -0.2 -0.1 0.1 -0.1 -0.0 0.0 0.0 3 0.0 2 -2.5 -2 -1 -5.0 <u>-3</u> 0 2.5 4 maxExpr minExpr nTimesRest 0.20 0.6 -0.4 withinSubset density 0.10 0.4 false true 0.2 -0.2 -0.05 -0.00 0.0 0.0 0 2 1 2 **-7** -5 -3 3 -9 0 1 tissSpec 0.6 0.4 -0.2 0.0 -

lncRNA group = IDRlessthan0\_01 ::: data cols = Inpa

value of facet label

-2.0 -1.5 -1.0 -0.5