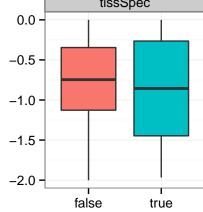
lncRNA group = IDRlessthan0_2 ::: data cols = lpa (31 / 2051) = (func/total IncRNA) ::: biotype = lincRNA measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0 0 **-5** --10 --8 --15 **-**-20 -12 false true false true false true minExpr maxExpr nTimesRest 3 8 -0 -6 withinSubset false -3· 4 · true -6· 2 --9 0 false true false false true true tissSpec



5

0 .

-5 -

0.0

-2.5 -

-5.0 **-**

-7.5 **-**

-10.0 -

log(value)

value of facet label