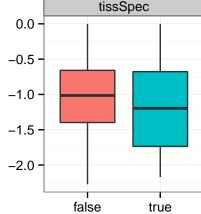
lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (51 / 2754) = (func/total lncRNA) ::: biotype = remove_antisense measures across all samples(cell-type/pulldown) varianceExpr sumExpr averageExpr 5 -0 -0 -5 --10 **--5** --10 --20 · true false false true false true minExpr maxExpr nTimesRest 6 5 · -2.5 withinSubset 4 0 false true –7.5 **-**2 --5 · -10.0 --10 0 false true false false true true tissSpec



0

-10 ·

-5.0 ·

log(value)

value of facet label