IncRNA subsection= foundInDisease measures accross all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 5 -0 0 -0 -5 --10 **-**-5 **-10** -20 **-**-10 false true false true false true minExpr maxExpr nTimesRest 6 -5 -**-3** log(value) withinSubset 4 0 false -6· true -5 2 -9 -10 0 false false false true true true tissSpec 0.0 -0.5 · -1.0 --1.5 --2.0 false true

value of facet label