IncRNA genes as max trans in sample cshltranscript expression data facet by sample with maximum expression color by threeTimesRest cell-type expr only transcripts w/ > 0.8 tissue specifity(max JSD) (Top Expressed Transcript For Each Gene) A549.longNonPolyA A549.longPolyA -5· -10 · -15 AG04450.longNonPolyA AG04450.longPolyA -5· -10 · -15 BJ.longPolyA BJ.longNonPolyA -5 · -10 · -15 GM12878.longNonPolyA GM12878.longPolyA -5 · -10 · -15 H1.HESC.longNonPolyA H1.HESC.longPolyA -5 · -10 -15 HELA.S3.longNonPolyA HELA.S3.longPolyA three Times RestA549.longNonPolyA A549.longPolyA -5 · -10 · AG04450.longNonPolyA AG04450.longPolyA -15 · HEPG2.longNonPolyA HEPG2.longPolyA BJ.longNonPolyA BJ.longPolyA GM12878.longNonPolyA -5 · GM12878.longPolyA -10 · H1.HESC.longNonPolyA -15 H1.HESC.longPolyA HMEC.longNonPolyA HMEC.longPolyA HELA.S3.longNonPolyA HELA.S3.longPolyA HEPG2.longNonPolyA HEPG2.longPolyA log2(value) HMEC.longNonPolyA HMEC.longPolyA HSMM.longNonPolyA HSMM.longPolyA HSMM.longNonPolyAHSMM.longPolyA HUVEC.longNonPolyA -5 · HUVEC.longPolyA -10 · K562.longNonPolyA -15 HUVEC.longPolyA HUVEC.longNonPolyA K562.longPolyA MCF.7.longNonPolyA MCF.7.longPolyA NHEK.longNonPolyA -5 · -10 NHEK.longNonPolyA.1 -15 · NHEK.longPolyA K562.longNonPolyA K562.longPolyA NHEK.longPolyA.1 NHLF.longNonPolyA NHLF.longPolyA -5· SK.N.SH_RA.longNonPolyA -10 SK.N.SH_RA.longPolyA -15 MCF.7.longNonPolyA MCF.7.longPolyA -5 -10 -15 NHEK.longNonPolyA NHEK.longNonPolyA.1 -5 · -10 -15 NHEK.longPolyA NHEK.longPolyA.1 -5 -10 -15 · NHLF.longNonPolyA NHLF.longPolyA -5· -10 -15 SK.N.SH_RA.longNonPolyA SK.N.SH_RA.longPolyA 0 --5 A549A5019NGO-BETCHENDLEBONFERING LEGEN

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