IncRNA genes as max trans in sample Cell Type Expression:(top transcripts per gene) Cell-type max expression of transcript is filled A549.longNonPolyA A549.longPolyA 0.4 0.2 0.0 AG04450.longNonPolyA AG04450.longPolyA 0.4 0.2 0.0 BJ.longNonPolyA BJ.longPolyA 0.4 0.2 0.0 GM12878.longNonPolyA GM12878.longPolyA 0.4 0.2 0.0 H1.HESC.longNonPolyA H1.HESC.longPolyA 0.4 0.2 0.0 HELA.S3.longNonPolyA HELA.S3.longPolyA 0.4 0.2 0.0 HEPG2.longPolyA HEPG2.longNonPolyA 0.4 0.2 0.0 HMEC.longNonPolyA HMEC.longPolyA 0.4 0.2 maxExprCat density 0.0 cell-type-other HSMM.longNonPolyA HSMM.longPolyA cell-type-this 0.4 0.2 0.0 -HUVEC.longNonPolyA HUVEC. Iong Poly A0.4 0.2 -0.0 -K562.longNonPolyA K562.longPolyA 0.4 0.2 0.0 -MCF.7. Iong Non Poly AMCF.7.longPolyA 0.4 0.2 0.0 -NHEK.longNonPolyA NHEK.longNonPolyA.1 0.4 0.2 0.0 -NHEK.longPolyA NHEK.longPolyA.1 0.4 0.2 0.0 -NHLF. long Non Poly ANHLF.longPolyA 0.4 0.2 0.0 -SK.N.SH_RA.longNonPolyA ${\sf SK.N.SH_RA.longPolyA}$ 0.4 0.2 -0.0 -5 -5 -10 -10 5 log(value)