IncRNA genes as max trans in sample Cell Type Expression:(top transcripts per gene) Cell-type specificity of transcript is colored A549.longPolyA A549.longNonPolyA 300 300 200 200 100 100 0 AG04450.longNonPolyA AG04450.longPolyA 300 300 200 200 100 100 BJ. Iong Non Poly ABJ.longPolyA 300 300 200 200 100 100 0 0 GM12878.longNonPolyA GM12878.longPolyA 200 200 -100 100 H1.HESC.longNonPolyA H1.HESC.longPolyA 300 300 200 200 100 100 0 0 HELA.S3.longNonPolyA HELA.S3.longPolyA 300 200 200 100 100 0 HEPG2.longNonPolyA HEPG2.longPolyA 300 200 200 100 100 0 HMEC.longPolyA HMEC. Iong Non Poly A200 -60 150 -40 -100 20 50 ctsCat 0 cell-type-other HSMM.longNonPolyA HSMM.longPolyA cell-type-this 150 -- none 200 100 100 50 · 0 HUVEC.longNonPolyA HUVEC.longPolyA 250 200 200 150 -100 100 -50 K562.longNonPolyA K562.longPolyA 250 200 -200 150 -100 -100 50 0 MCF.7.longPolyA MCF.7. Iong Non Poly A300 200 -200 150 100 100 50 · 0 NHEK. Iong Non Poly ANHEK.longNonPolyA.1 120 250 200 -90 150 60 100 -30 50 0 NHEK.longPolyA NHEK.longPolyA.1 300 200 -200 -150 100 100 50 NHLF.longNonPolyA NHLF.longPolyA100 200 -50 100 0 0 SK.N.SH\_RA.longNonPolyA SK.N.SH\_RA.longPolyA 300 -300 -200 -200 100 100 -5 -10 -10 -5 log(value)