cshl mRNA transcript expression data facetby tissue with maximum expression color by three times rest cell-type expr only transcripts w/ > 0.8 tissue specifity (Top Expressed Transcript For Each Genen) A549.longNonPolyA A549.longPolyA 0 -10 -20 AG04450.longNonPolyA AG04450.longPolyA -10 -20 BJ.longNonPolyA BJ.longPolyA -10 · -20 GM12878.longNonPolyA GM12878.longPolyA -10 -20 H1.HESC.longNonPolyA H1.HESC.longPolyA -10 -20 HELA.S3.longNonPolyA HELA.S3.longPolyA 0 threeTimesRest A549.longNonPolyA -10 A549.longPolyA -20 AG04450.longNonPolyA HEPG2.longNonPolyA HEPG2.longPolyA AG04450.longPolyA BJ.longNonPolyA 0 BJ.longPolyA -10 GM12878.longNonPolyA -20 GM12878.longPolyA HMEC.longNonPolyA HMEC.longPolyA H1.HESC.longNonPolyA H1.HESC.longPolyA 0 HELA.S3.longNonPolyA -10 · HELA.S3.longPolyA HEPG2.longNonPolyA -20 HSMM.longNonPolyA HSMM.longPolyA HEPG2.longPolyA log2(value) HMEC.longNonPolyA HMEC.longPolyA HSMM.longNonPolyA HSMM.longPolyA –20 · HUVEC.longNonPolyA HUVEC.longPolyA HUVEC.longNonPolyA HUVEC.longPolyA K562.longNonPolyA K562.longPolyA -10 · MCF.7.longNonPolyA MCF.7.longPolyA -20 K562.longNonPolyA K562.longPolyA NHEK.longNonPolyA NHEK.longNonPolyA.1 NHEK.longPolyA NHEK.longPolyA.1 -10NHLF.longNonPolyA -20 NHLF.longPolyA MCF.7.longNonPolyA MCF.7.longPolyA none SK.N.SH_RA.longNonPolyA 0 SK.N.SH_RA.longPolyA -10 -20 NHEK.longNonPolyA NHEK.longNonPolyA.1 0 -10 -20 NHEK.longPolyA NHEK.longPolyA.1 0 -10 -20 NHLF.longNonPolyA NHLF.longPolyA 0 -10 -20 ${\sf SK.N.SH_RA.longNonPolyA}$ none 0 -10 · -20 SK.N.SH_RA.longPolyA 0 -A549MGONGON REGISTERS LEED **Ny Nonga**blyA variable