For transcripts found in {cell-type/polyA pulldown} pair SK.N.SH\_RA.longPolyA -SK.N.SH\_RA.longNonPolyA -NHLF.longPolyA -NHLF.longNonPolyA -NHEK.longPolyA.1 -NHEK.longPolyA -NHEK.longNonPolyA.1 NHEK.longNonPolyA -MCF.7.longPolyA -MCF.7.longNonPolyA -K562.longPolyA -K562.longNonPolyA -HUVEC.longPolyA -HUVEC.longNonPolyA -RnaSeqExpr HSMM.longPolyA ctsCat HSMM.longNonPolyA cell-type-this HMEC.longPolyA cell-type-other HMEC.longNonPolyA -HEPG2.longPolyA -HEPG2.longNonPolyA -HELA.S3.longPolyA -HELA.S3.longNonPolyA -H1.HESC.longPolyA · H1.HESC.longNonPolyA -GM12878.longPolyA -GM12878.longNonPolyA -BJ.longPolyA -BJ.longNonPolyA -AG04450.longPolyA -AG04450.longNonPolyA -A549.longPolyA -A549.longNonPolyA -1000 2000 3000 4000 0

transcripts

Cell-type Max Expr