Number of transcripts in a cell-type with 3x RPKM values (Top Expressed Transcript For Each Gene) N= 2905 / 16408 (total) SK.N.SH_RA.longPolyA -SK.N.SH_RA.longNonPolyA -NHLF.longPolyA -NHLF.longNonPolyA -NHEK.longPolyA.1 -Frequency of Cell-type with 3x expression than rest NHEK.longPolyA -NHEK.longNonPolyA.1 -NHEK.longNonPolyA -MCF.7.longPolyA -MCF.7.longNonPolyA -K562.longPolyA -K562.longNonPolyA -HUVEC.longPolyA -HUVEC.longNonPolyA -HSMM.longPolyA -HSMM.longNonPolyA -HMEC.longPolyA -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 -100 200 300 0

cell-type and extraction method