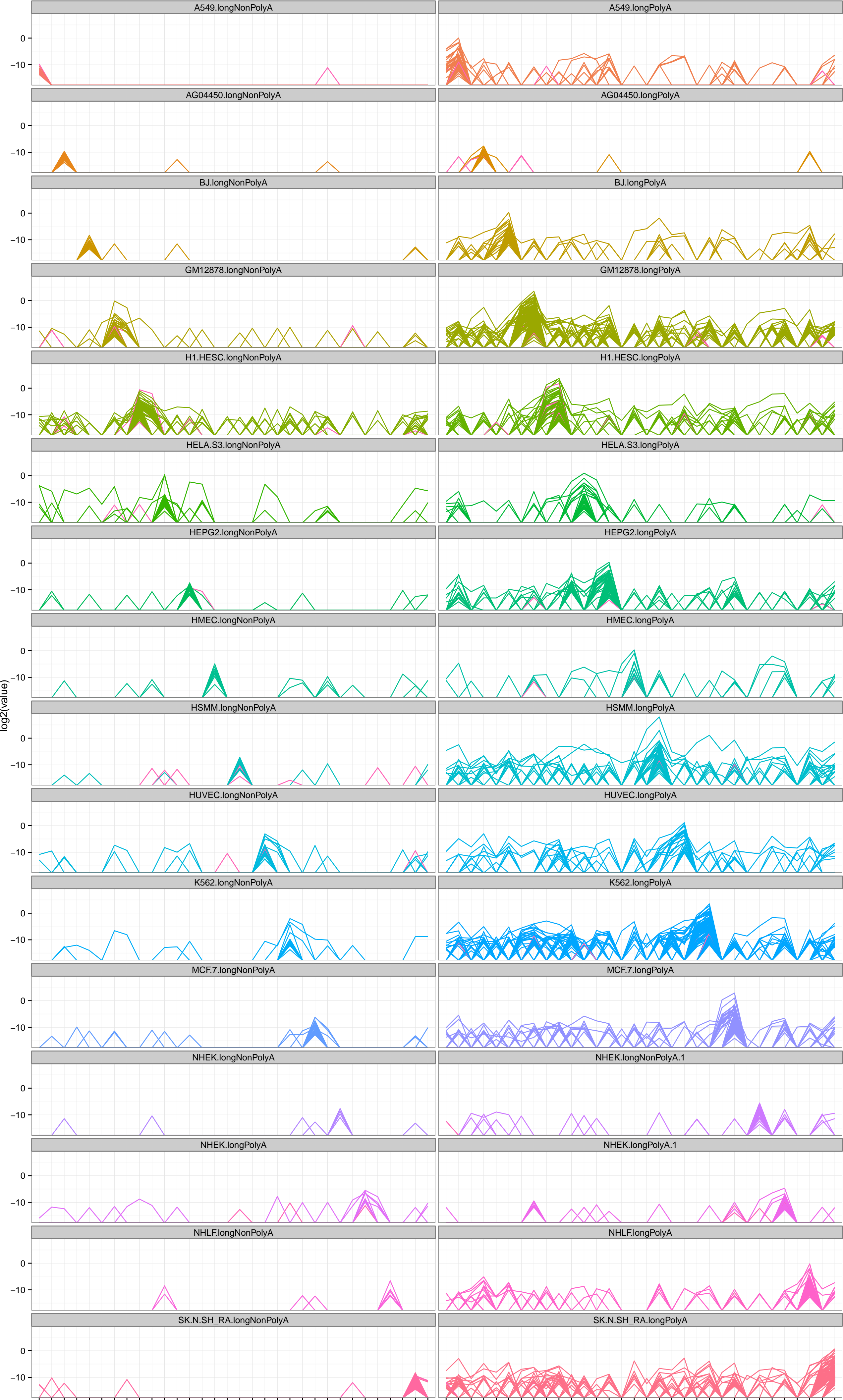


lncRNA genes as max trans in sample
cshl lncRNA transcript expression data
facet by sample with maximum expression
color by threeTimesRest cell-type expr
only transcripts w/ > 0.6 tissue specificity(max JSD)
(Top Expressed Transcript For Each Gene)



threeTimesRest

- A549.longNonPolyA
- A549.longPolyA
- AG04450.longNonPolyA
- AG04450.longPolyA
- BJ.longNonPolyA
- BJ.longPolyA
- GM12878.longNonPolyA
- GM12878.longPolyA
- H1.HESC.longNonPolyA
- H1.HESC.longPolyA
- HELA.S3.longNonPolyA
- HELA.S3.longPolyA
- HEPG2.longNonPolyA
- HEPG2.longPolyA
- HMEC.longNonPolyA
- HMEC.longPolyA
- HSMM.longNonPolyA
- HSMM.longPolyA
- HUVEC.longNonPolyA
- HUVEC.longPolyA
- K562.longNonPolyA
- K562.longPolyA
- MCF.7.longNonPolyA
- MCF.7.longPolyA
- NHEK.longNonPolyA
- NHEK.longPolyA
- NHEK.longNonPolyA.1
- NHEK.longPolyA.1
- NHLF.longNonPolyA
- NHLF.longPolyA
- none
- SK.N.SH_RA.longNonPolyA
- SK.N.SH_RA.longPolyA