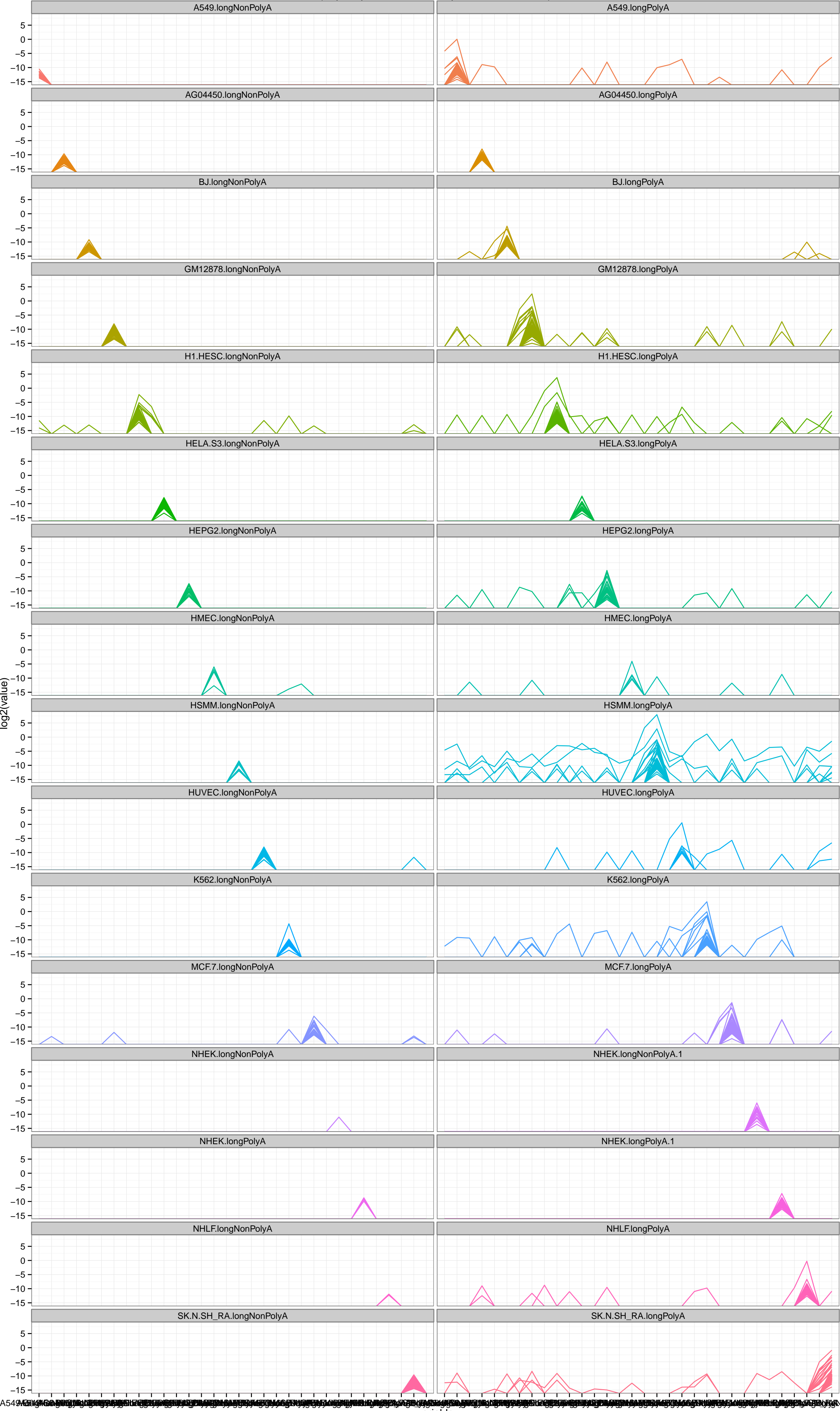


lncRNA genes as max trans in sample  
cshltranscript expression data  
facet by sample with maximum expression  
color by threeTimesRest cell-type expr  
only transcripts w/ > 0.8 tissue specificity(max JSD)  
(Top Expressed Transcript For Each Gene)



- 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.longNonPolyA.1
  - NHEK.longPolyA
  - NHEK.longPolyA.1
  - NHLF.longNonPolyA
  - NHLF.longPolyA
  - SK.N.SH\_RA.longNonPolyA
  - SK.N.SH\_RA.longPolyA