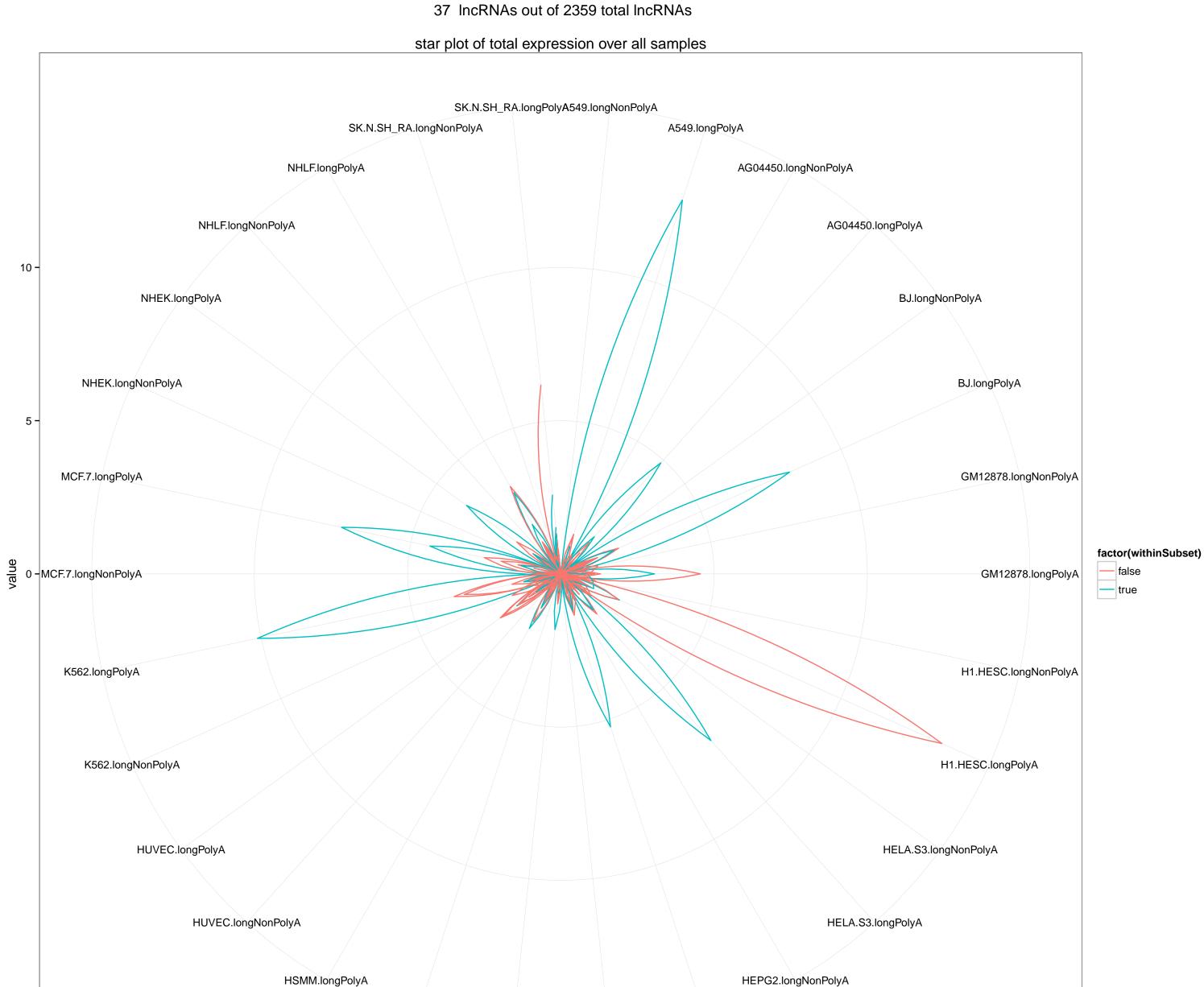
## IncRNA group = allLncRNA ::: data cols = Ipa ( 37 / 2359 ) = (func/total IncRNA) ::: biotype = antisense subset = 37 IncRNAs out of 2359 total IncRNAs



HMEC.longPolyA HMEC.longNonPolyA

HEPG2.longPolyA

HSMM.longNonPolyA