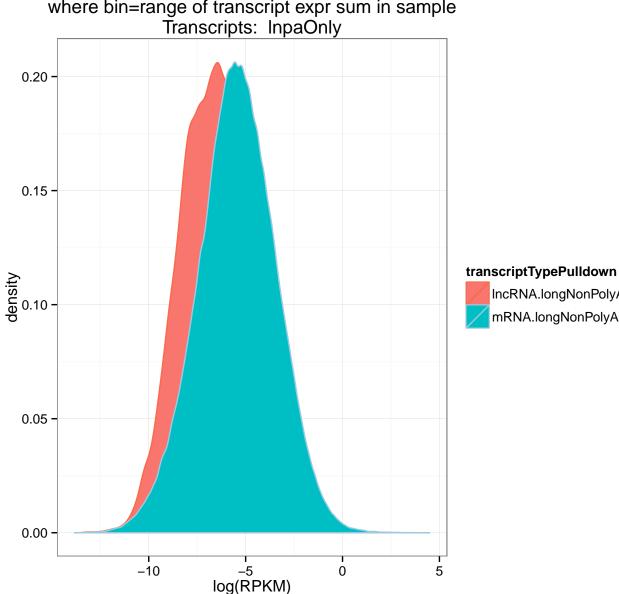
transcript pulldown RPKM comparisons facet over bin where bin=range of transcript expr sum in sample



IncRNA.longNonPolyA mRNA.longNonPolyA