transcript pulldown RPKM comparisons facet over bin where bin=range of transcript expr sum in sample (1e-05,0.000364] (0.000364,0.00075] (0.00075, 0.00136] (0.00136, 0.00235](0.00235, 0.0042] transcriptTypePulldown IncRNA.longNonPolyA count IncRNA.longPolyA mRNA.longNonPolyA (0.0042, 0.0075]mRNA.longPolyA (0.0075, 0.0141](0.0141, 0.0298](0.0298, 0.0847](0.0847,258]log(RPKM) -10