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

(0.000364,0.00075] (0.00075,0.00136] (1e-05,0.000364] (0.00136,0.00235] 150 transcriptTypePulldown IncRNA.longPolyA mRNA.longPolyA -12.5 -12.5 -10.0 -7.5 -12.5 -12.5 log(RPKM)