transcript pulldown RPKM comparisons facet over celltype and bin where bin=range of transcript expr sum in sample (0.0075,0.0141] (0.0141,0.0298] (0.0847,258] (0.00235,0.0042] (0.0042,0.0075] (0.0298,0.0847] 400 -200 -600 -400 -200 -600 -400 -200 -600 -400 -200 -600 -400 -200 -0 <del>-</del> 400 -200 -600 -400 -200 -600 transcriptTypePulldown IncRNA.longPolyA