transcript pulldown RPKM comparisons facet over bin where bin=range of transcript expr sum in sample
Transcripts: both 0.6 (2e-06,0.000517] 0.4 0.2 0.0 (0.000517,0.00137] 0.6 0.4 0.2 0.0 (0.00137,0.00278] 0.75 0.50 0.25 0.00 (0.00278,0.00506] 0.75 0.50 0.25 0.00 1.00 (0.00506, 0.0089]0.75 0.50 0.25 0.00 transcript Type Pull down(0.0089, 0.0155]0.9 IncRNA.longNonPolyA density 0.6 IncRNA.longPolyA mRNA.longNonPolyA 0.3 mRNA.longPolyA 0.0 0.9 (0.0155, 0.028] 0.6 0.3 0.0 1.00 (0.028, 0.0562]0.75 0.50 0.25 -0.00 0.75 (0.0562, 0.15]0.50 0.25 0.00 0.4 (0.15,258]0.2 0.0 0.75 0.50 Ζ 0.25 0.00 -5 log(RPKM) 0 -10 5