transcript pulldown RPKM comparisons facet over bin where bin=range of transcript expr sum in sample
Transcripts: IpaOnly 8.0 (2e-06,0.000517] 0.6 0.4 0.2 0.0 (0.000517,0.00137] 1.0 0.5 0.0 2.0 (0.00137,0.00278] 1.5 1.0 0.5 0.0 2.5 (0.00278,0.00506] 2.0 1.5 1.0 0.5 0.0 (0.00506, 0.0089]2 1 0 (0.0089, 0.0155]2 transcriptTypePulldown density IncRNA.longPolyA mRNA.longPolyA 1 0 2.5 2.0 (0.0155, 0.028]1.5 1.0 0.5 0.0 -1.5 (0.028,0.05 1.0 -562] 0.5 -0.0 1.25 1.00 (0.0562, 0.15]0.75 0.50 0.25 0.00 0.6 (0.15,258]0.4 0.2 -0.0 0.075 -0.050 Z 0.025 -0.000 --5 log(RPKM) -10 0 5