transcript pulldown RPKM comparisons facet over bin where bin=range of transcript expr sum in sample Transcripts: InpaOnly 0.6 (2e-06,0.000517] 0.4 0.2 0.0 1.25 (0.000517,0.00137] 1.00 0.75 0.50 0.25 0.00 1.5 (0.00137,0.00278] 1.0 0.5 0.0 2.0 (0.00278,0.00506] 1.5 1.0 0.5 0.0 2.0 (0.00506, 0.0089]1.5 1.0 0.5 0.0 2.0 (0.0089, 0.0155]transcriptTypePulldown density 1.5 IncRNA.longNonPolyA 1.0 mRNA.longNonPolyA 0.5 0.0 2.0 (0.0155, 0.028]1.5 1.0 0.5 0.0 1.5 (0.028, 0.0562]1.0 0.5 0.0 (0.0562, 0.15]1.0 0.5 0.0 1.00 0.75 (0.15,258]0.50 0.25 0.00 3 2 Z 1 log(RPKM) 0 -10 5