RSEM Read Mapping Distribution longPolyA nucleus Facet by Replicate,Celltype multiplicity=genome aligns per read

8 6 · A549 4 · 2 -0 8 6 · GM12878 4 2 0 8 log10(count) H1-hESC 0 8 6 -HeLa-S3 4 2 · 0 8 6 -HepG2 4 2 2.0 0.0 0.5 log10(multiplicity) 1.0 1.5 1.0 2.0 0.5 1.5 0.0