RSEM Read Mapping Distribution longNonPolyA cytosol Facet by Replicate,Celltype multiplicity=genome aligns per reac

multiplicity=genome aligns per read 8 6 GM12878 2 0 8 6 H1-hESC 4 2 0 6 4 2 0 8 8 . 6 · HepG2 4 · 2 · 0 8 6 K562 2 0 8 6 NHEK 4 2 0 5 2.0 0.0 0.5 log10(multiplicity) 1.0 2.0 0.0 0.5 1.0 1.5 1.5