RSEM Read Mapping Distribution longNonPolyA nucleus Facet by Replicate,Celltype

multiplicity=genome aligns per read 1.0e+08 7.5e+07 5.0e+07 2.5e+07 · 0.0e+00 1.0e+08 7.5e+07 -5.0e+07 -2.5e+07 0.0e+00 1.0e+08 7.5e+07 5.0e+07 2.5e+07 0.0e+00 1.0e+08 7.5e+07 5.0e+07 2.5e+07 0.0e + 001.0e+08 7.5e+07 HepG2 5.0e+07 2.5e+07 0.0e+00 1.0e+08 -7.5e+07 -K562 5.0e+07 -2.5e+07 -0.0e+00 1.0e+08 -7.5e+07 NHEK 5.0e+07 -2.5e+07 -0.0e+00 1.5 2.0 0.5 0.5 1.0 0.0 1.0 1.5 0.0 2.0

log10(multiplicity)