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+001.0e+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 -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+0050 100 150 50 100 150

multiplicity