RSEM Read Mapping Distribution longPolyA cytosol Facet by Replicate,Celltype

multiplicity=genome aligns per read 6e+07 A549 4e+07 -2e+07 -0e+00 6e+07 -4e+07 -2e+07 0e+00 6e+07 -4e+07 -2e+07 -0e+00 6e+07 -HUVEC 4e+07 -2e+07 0e+00 6e+07 4e+07 2e+07 0e+00 6e+07 -4e+07 2e+07 0e+00 6e+07 -IMR90 4e+07 -2e+07 -0e+00 -6e+07 -4e+07 -2e+07 0e+00 6e+07 -4e+07 -2e+07 · 0e+00 -

1.0

0.5

0.0

1.5

2.0

0.5

1.0

1.5

2.0

0.0

log10(multiplicity)