RSeQC: Bam Stat 16 samples Total records 20M 40M OM 10M 30M 50M QC failed OM 10M 20M 30M 40M 50M Duplicates OM 10M 20M 30M 40M 50M Non primary hit OM 10M 20M 50M 30M 40M Unmapped 20M 30M OM 10M 40M 50M Unique 10M 20M 30M 0M 40M 50M Read-1 OM 20M 10M 40M 50M 30M Read-2 OM 20M 10M 40M 50M 30M +ve strand OM 10M 20M 30M 40M 50M OM 10M 20M 30M 40M 50M Non-splice reads 10M OM 20M 30M 50M 40M Splice reads •••• OM 20M 10M 30M 40M 50M Proper pairs OM 10M 20M 30M 40M 50M Different chrom OM 10M 20M 30M 40M 50M