RSeQC: Bam Stat 6 samples Total records 30M OM 5M 10M 15M 20M 25M 35M 40M QC failed 5M 10M 25M 30M OM 15M 20M 35M 40M Duplicates OM 5M 10M 15M 20M 25M 30M 35M 40M Non primary hit 5M OM 10M 15M 20M 25M 30M 35M 40M Unmapped 5M 25M 30M OM 10M 15M 20M 35M 40M Unique = 15M 25M 5M 20M 30M OM 10M 35M 40M Read-1 = 10M 15M OM 5M 20M 25M 30M 35M 40M Read-2 OM 5M 10M 15M 20M 25M 30M 35M 40M +ve strand 5M 10M OM 15M 20M 25M 30M 35M 40M -ve strand 5M 10M OM 15M 30M 35M 20M 25M 40M Non-splice reads 15M 5M 25M 10M 20M 30M OM 35M 40M Splice reads OM 5M 10M 15M 20M 25M 30M 35M 40M Proper pairs 5M 15M 20M 25M 30M 35M OM 10M 40M Different chrom • OM 5M 10M 15M 20M 25M 30M 35M 40M