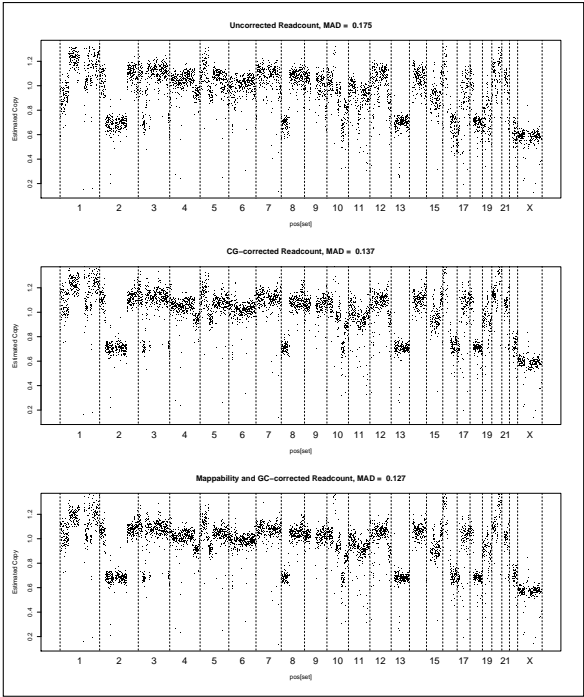
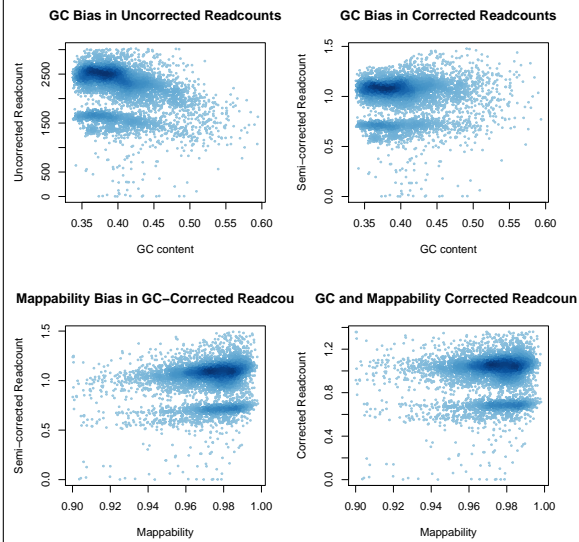


Bioinformatics Workflow for Ultra-Low-Pass Whole-Genome-Sequencing Data

Reference genome

bwa
Indexing reference genome
`ULPwgs::index_ref()`



Raw ULP-WGS data

