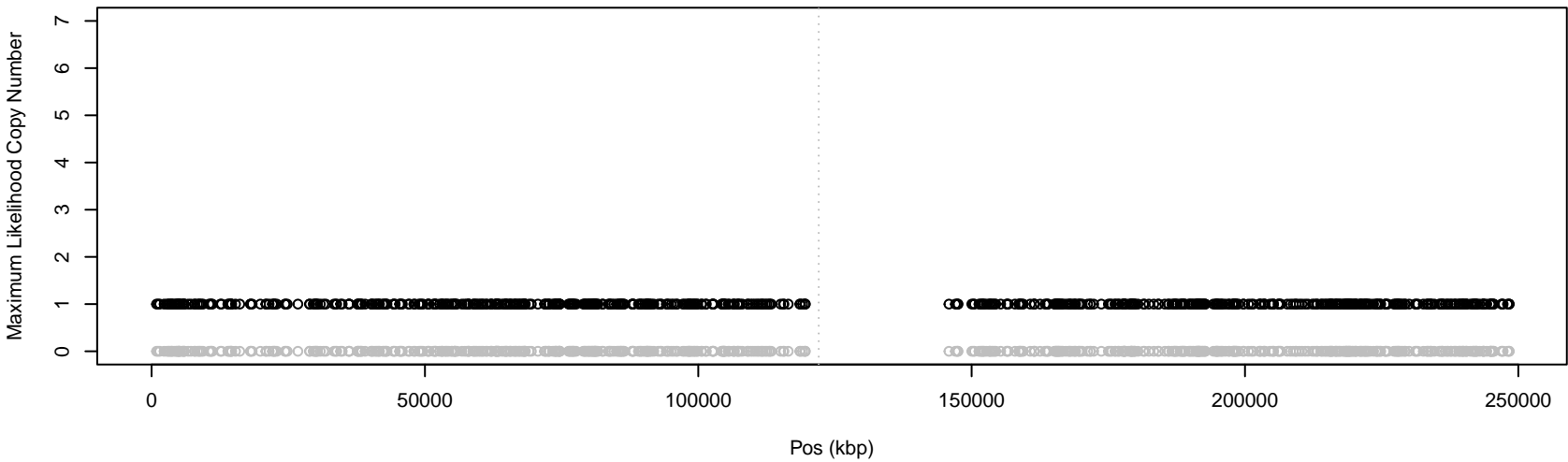
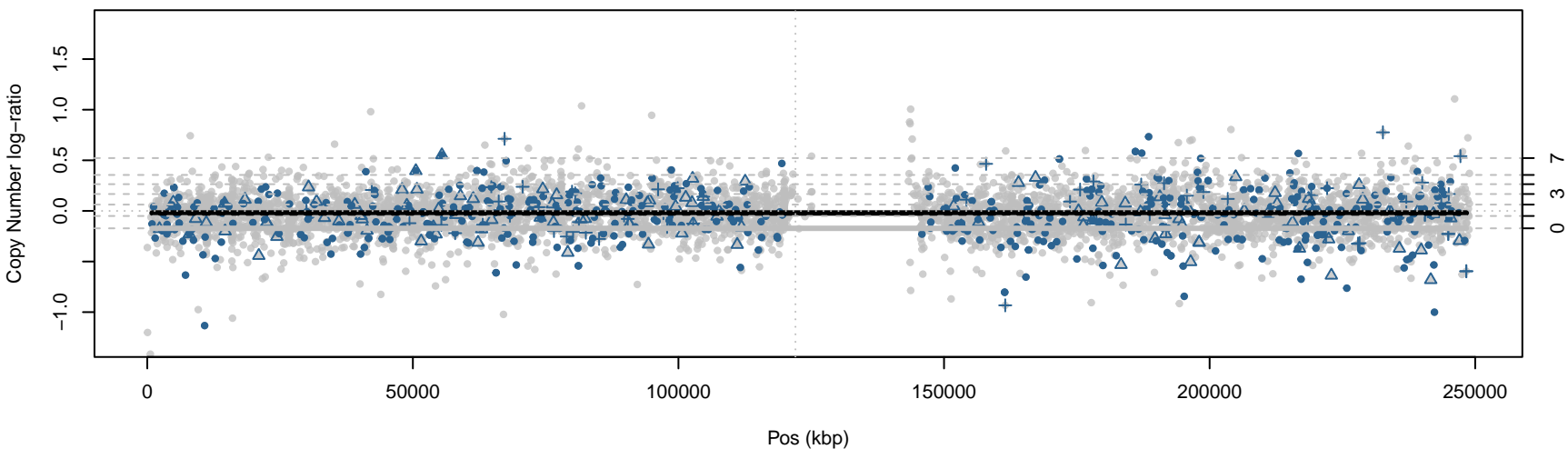
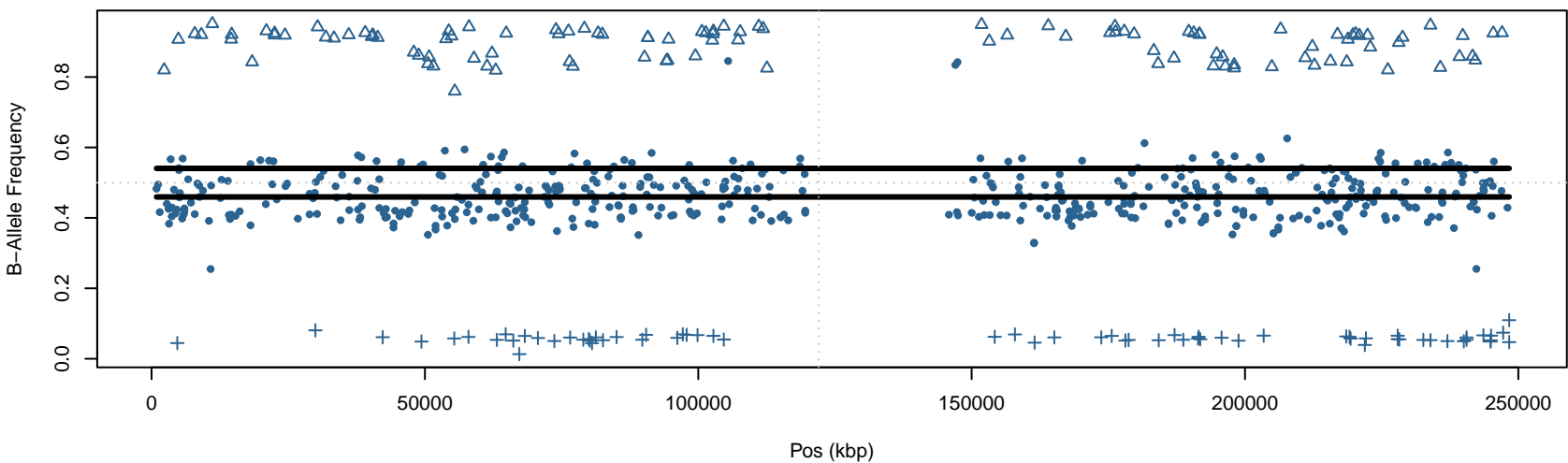
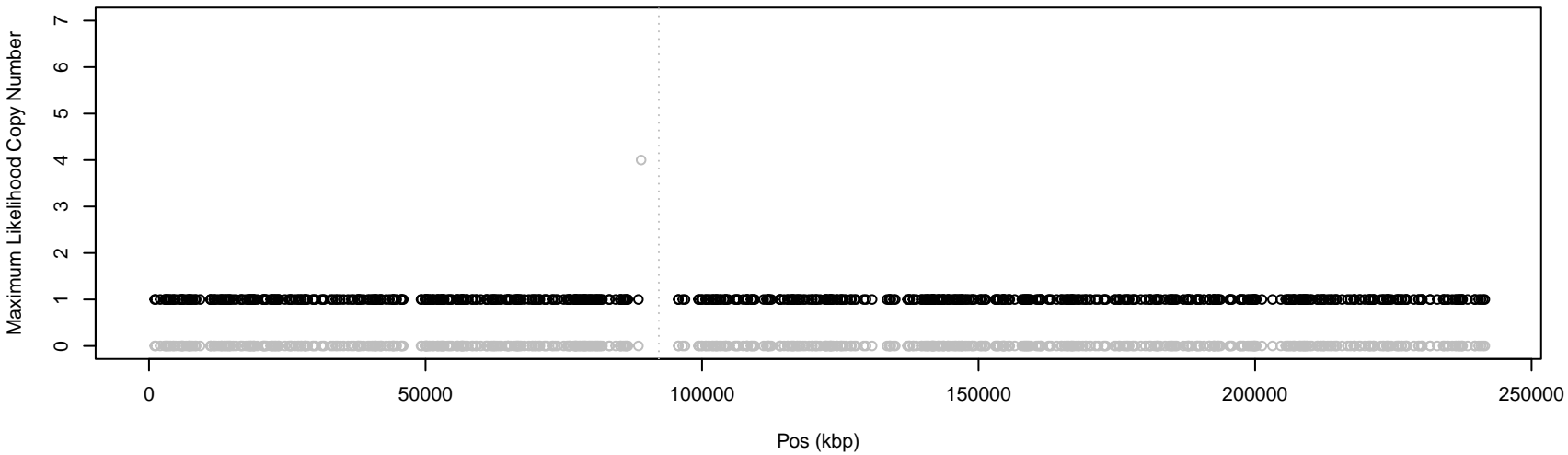
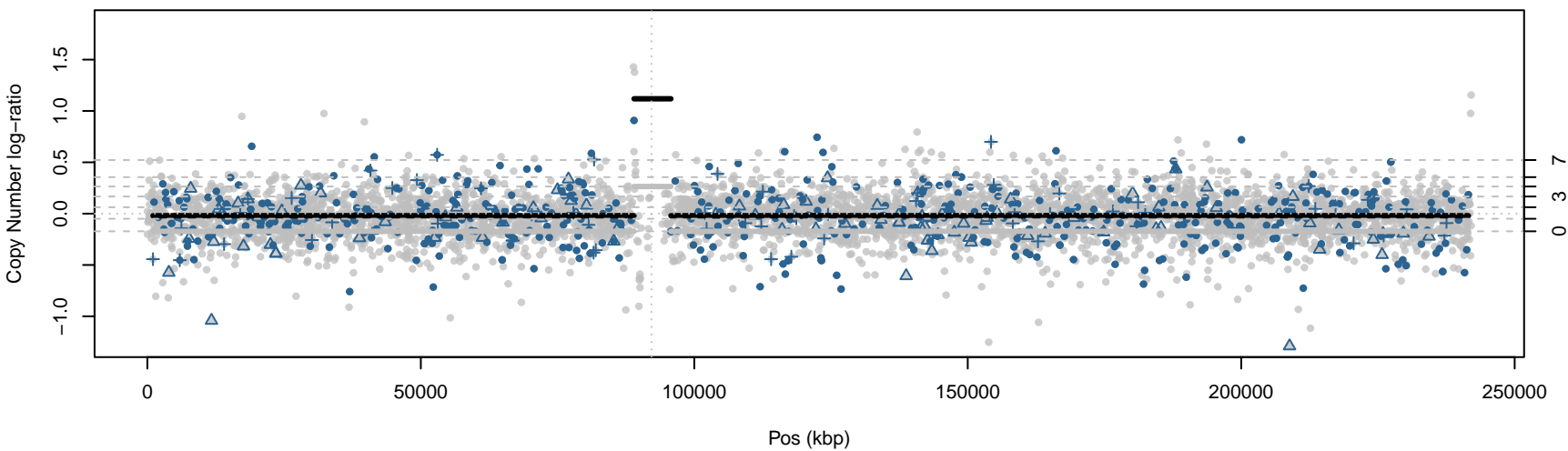
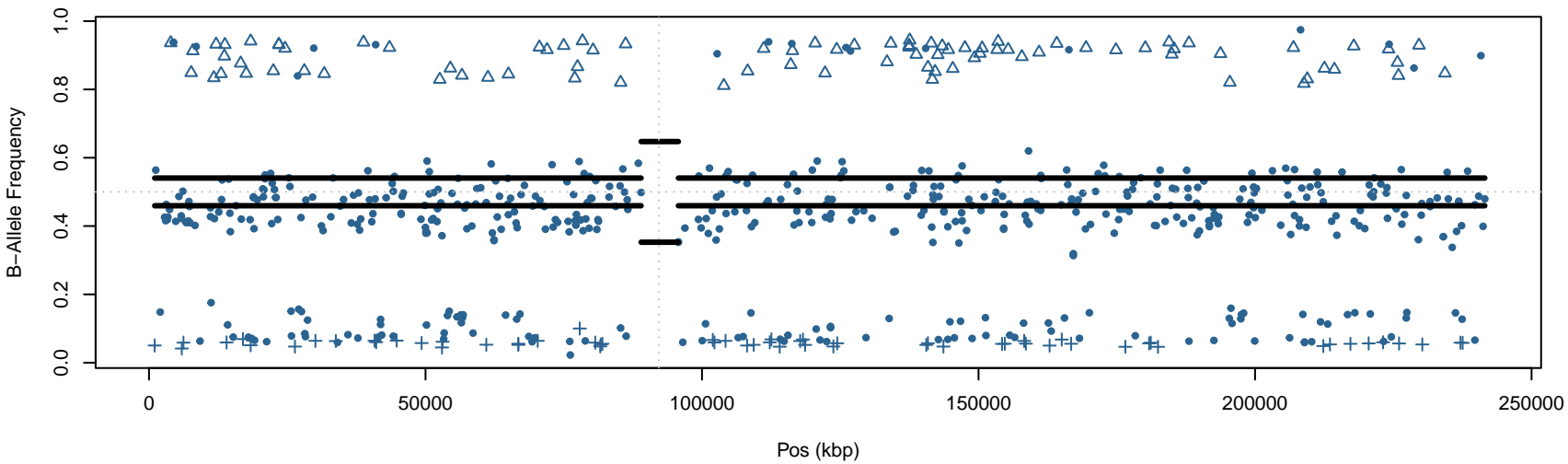


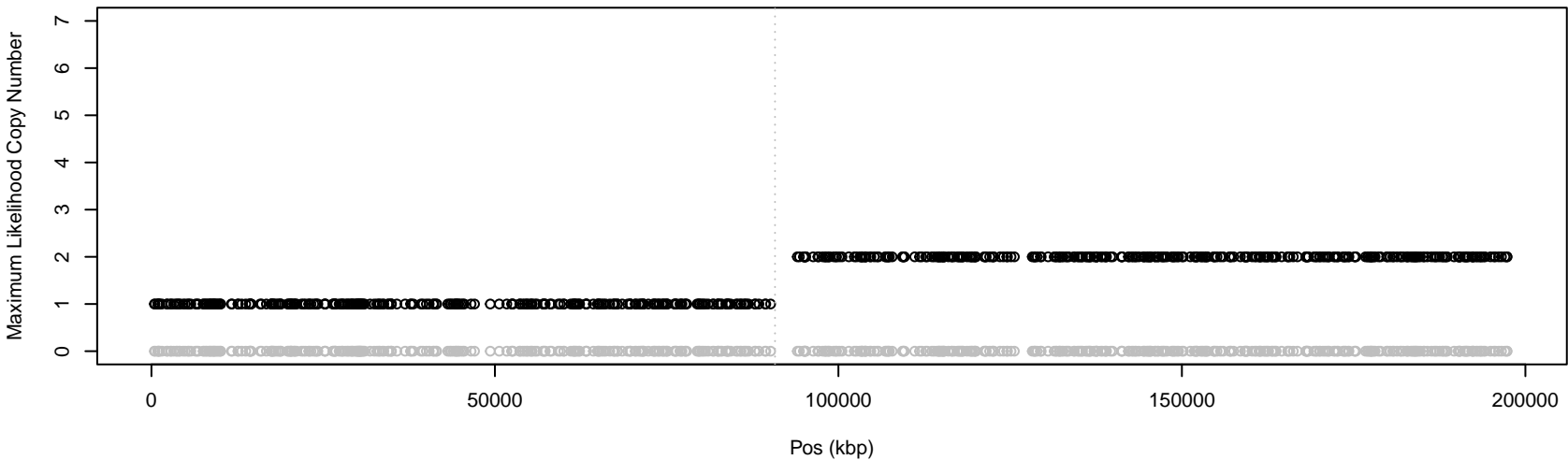
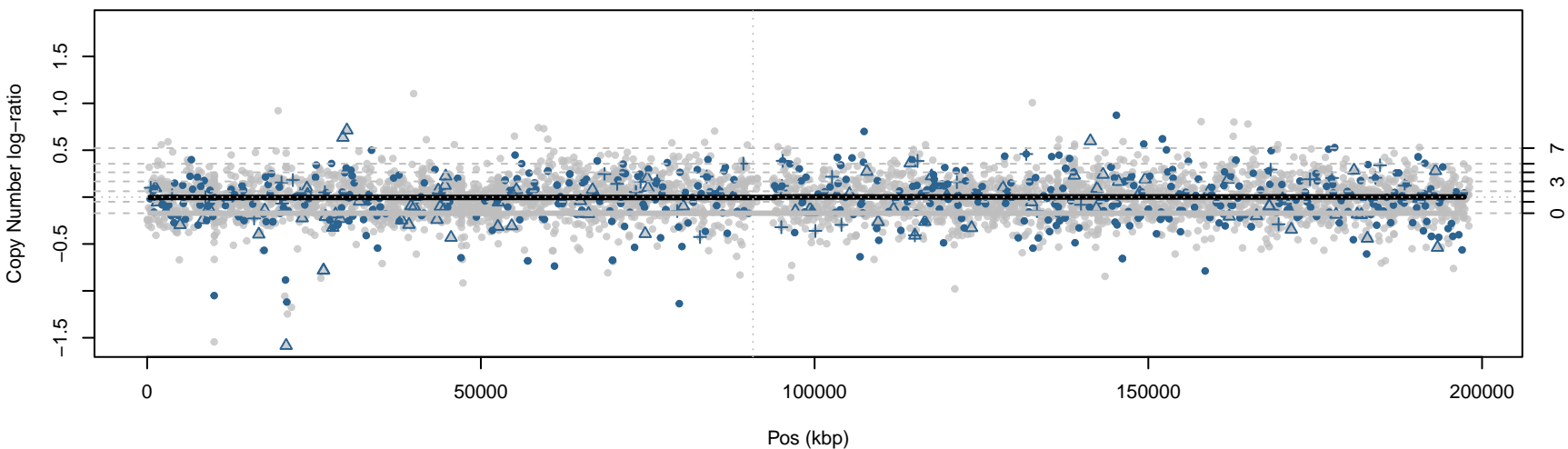
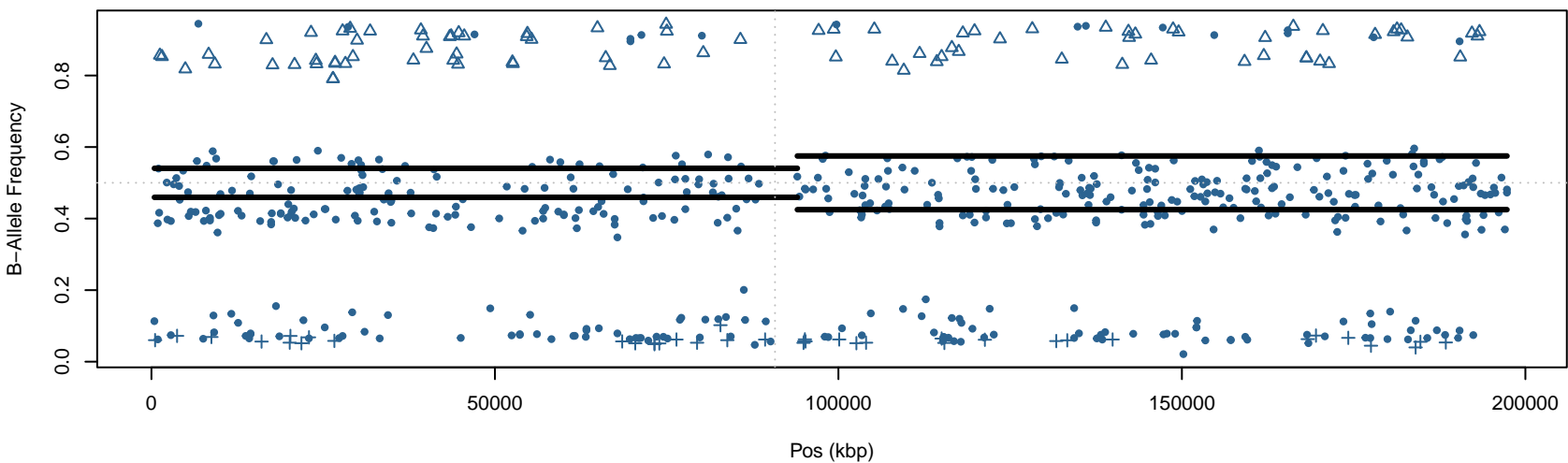
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr1



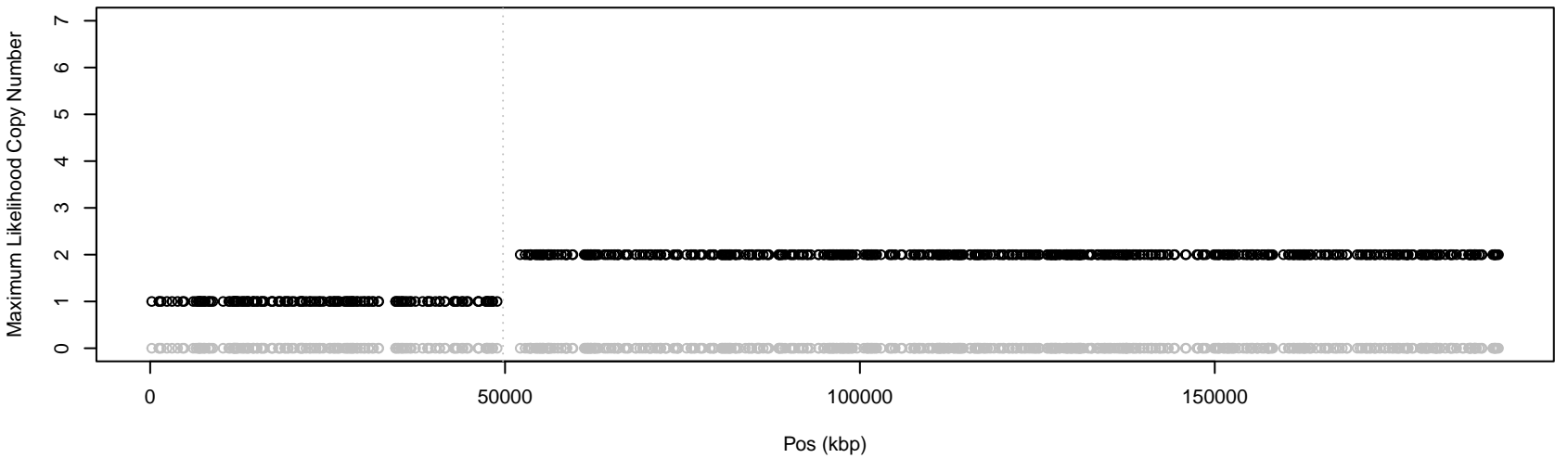
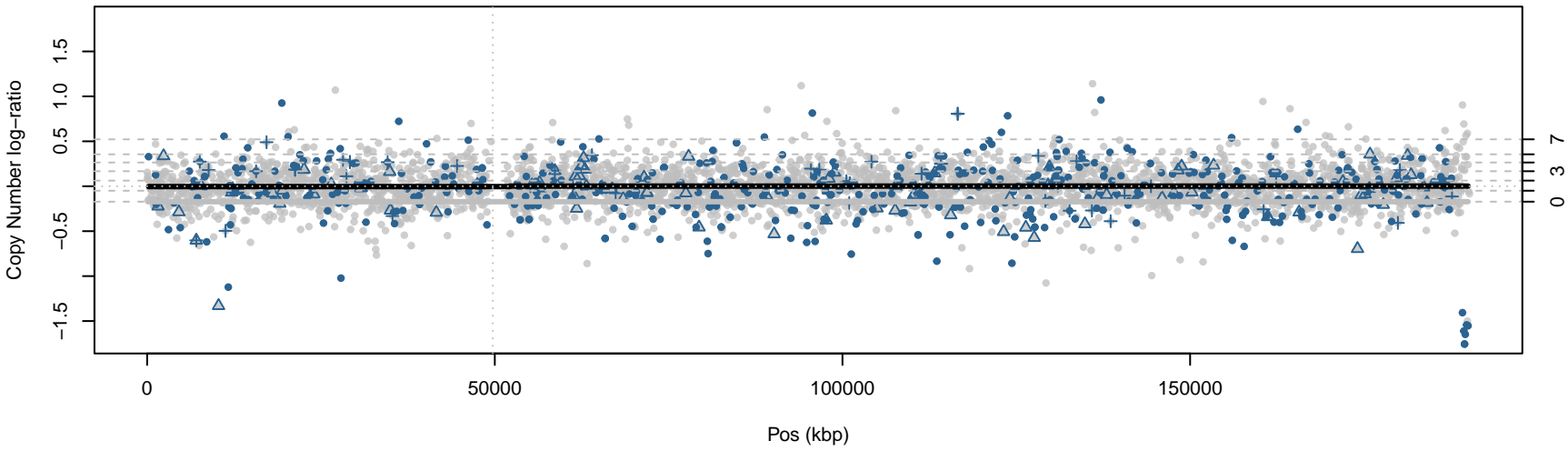
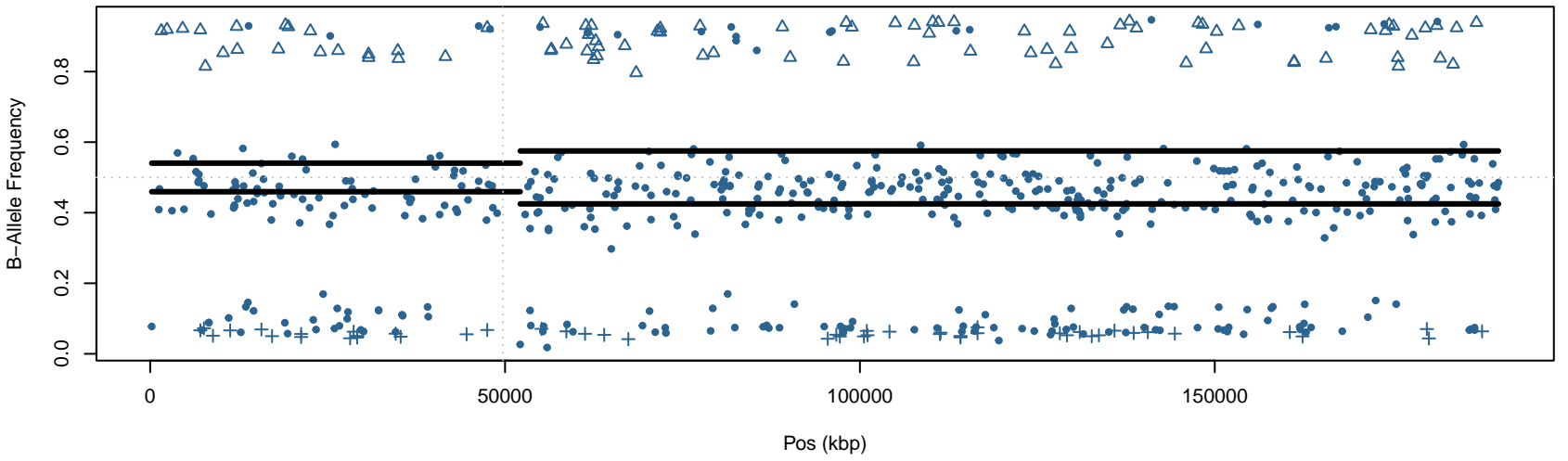
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr2



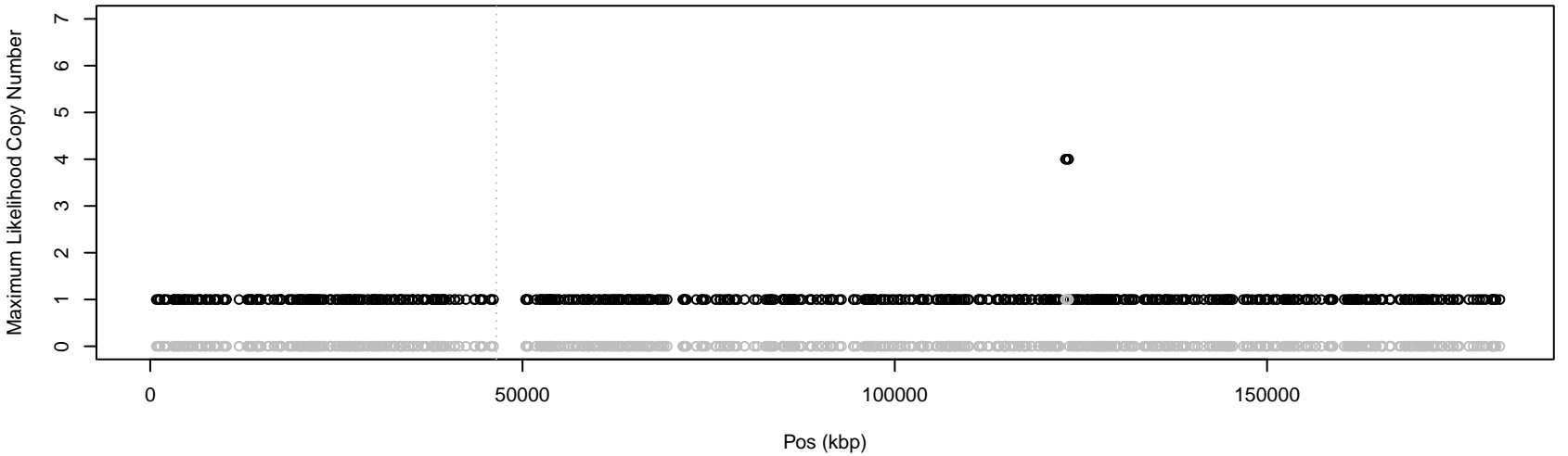
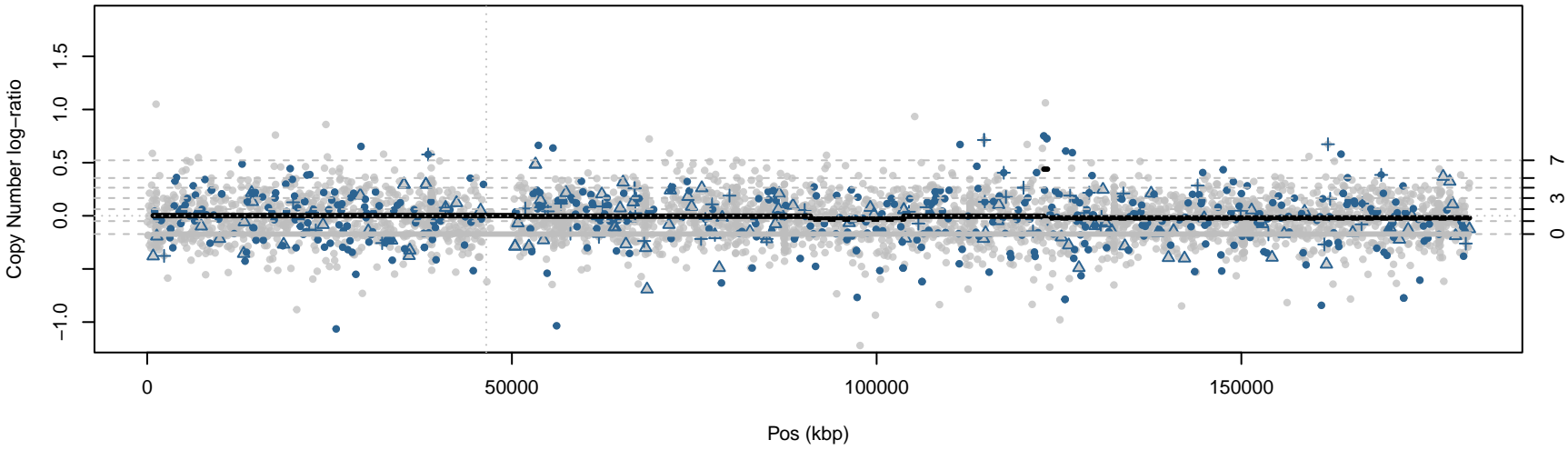
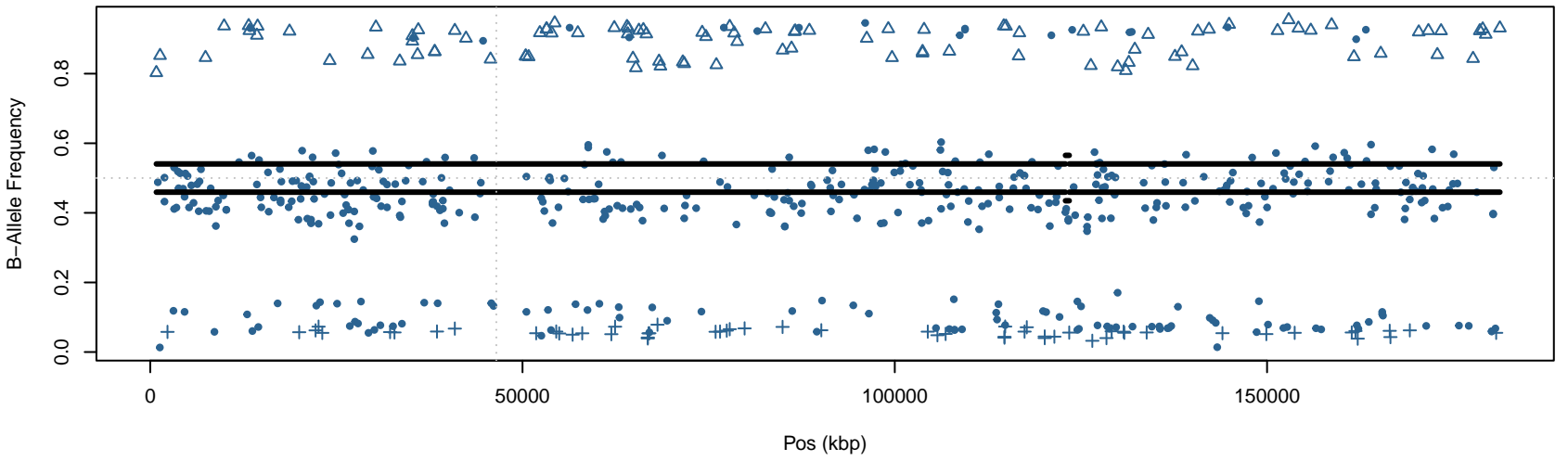
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr3



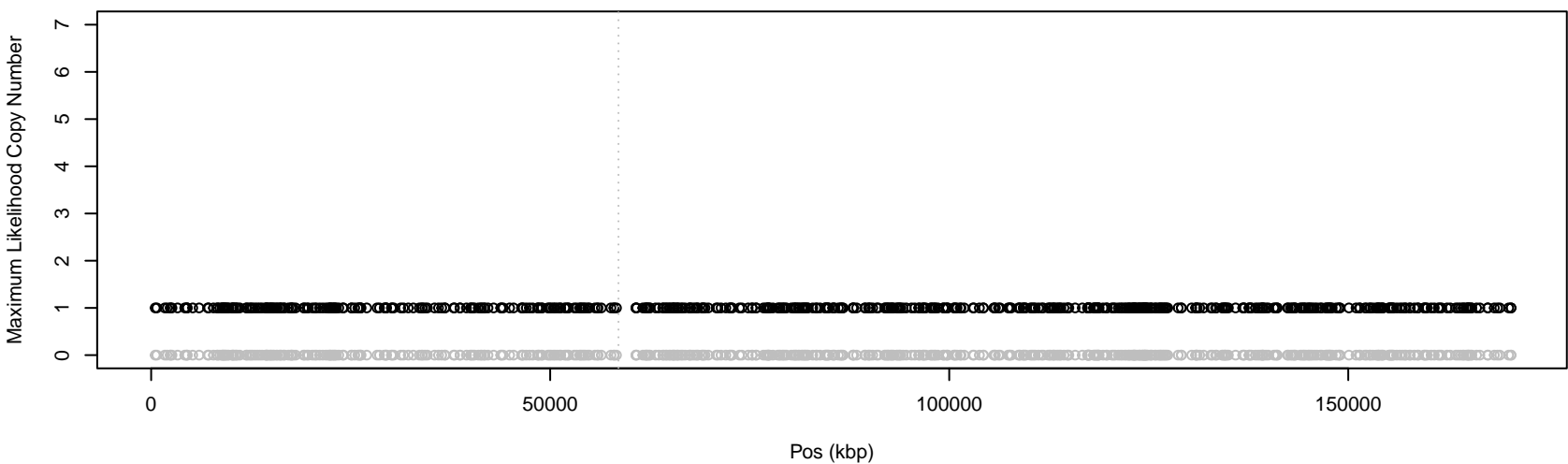
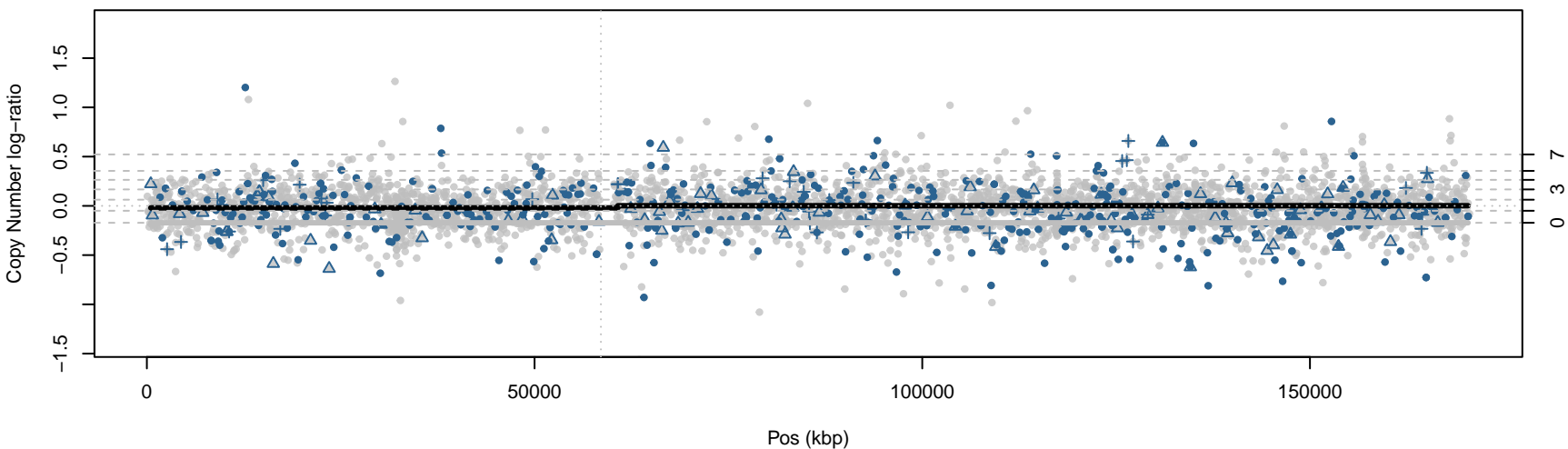
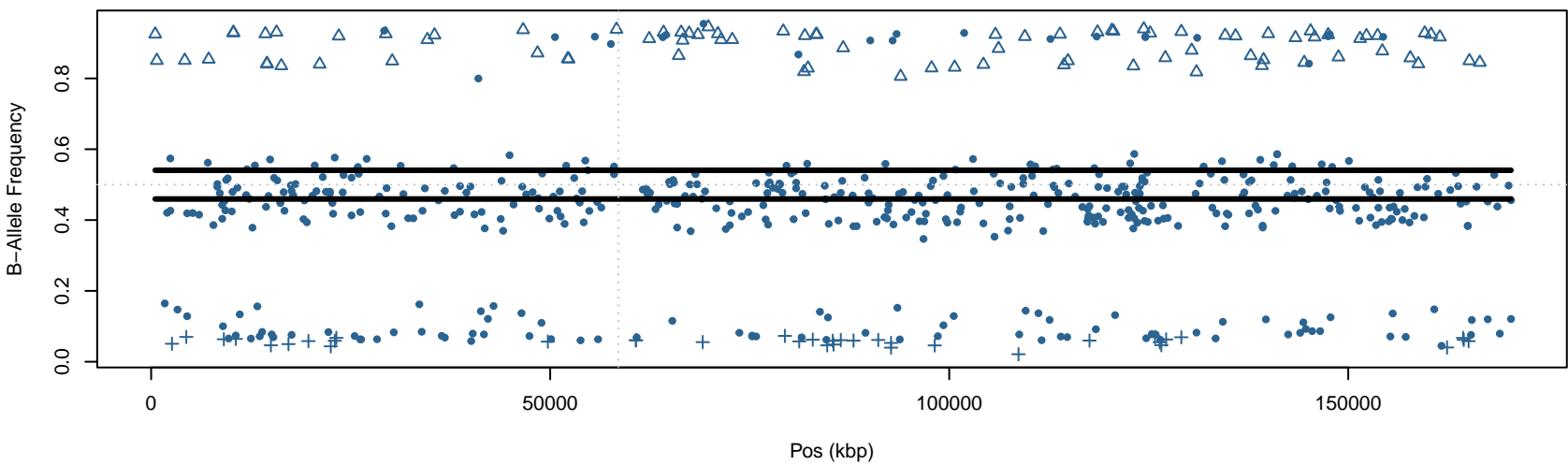
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr4



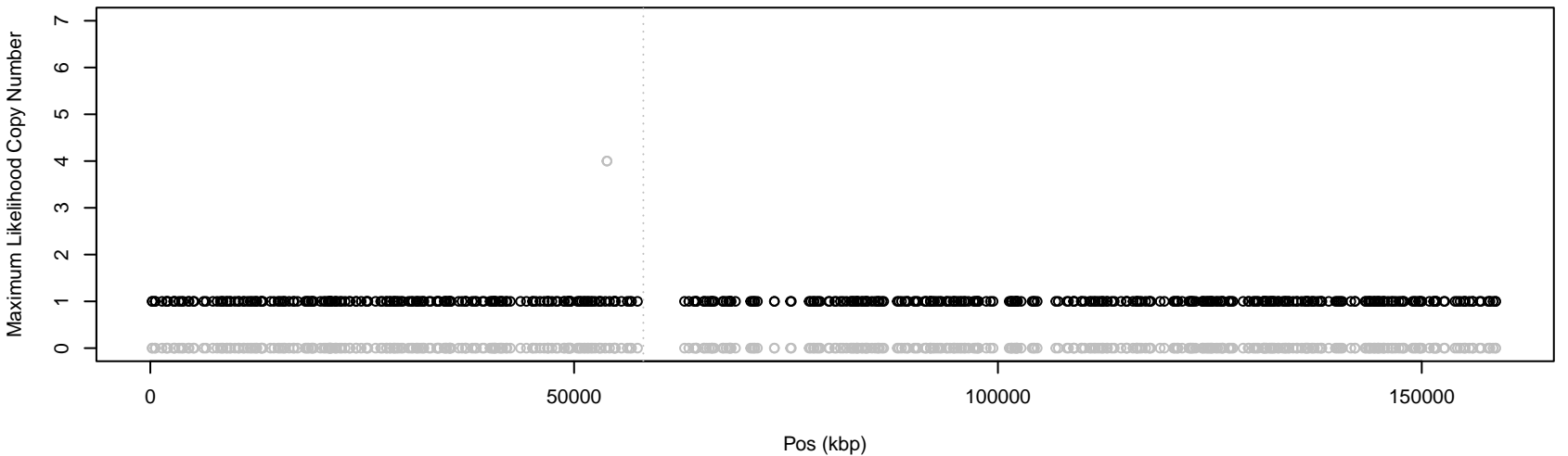
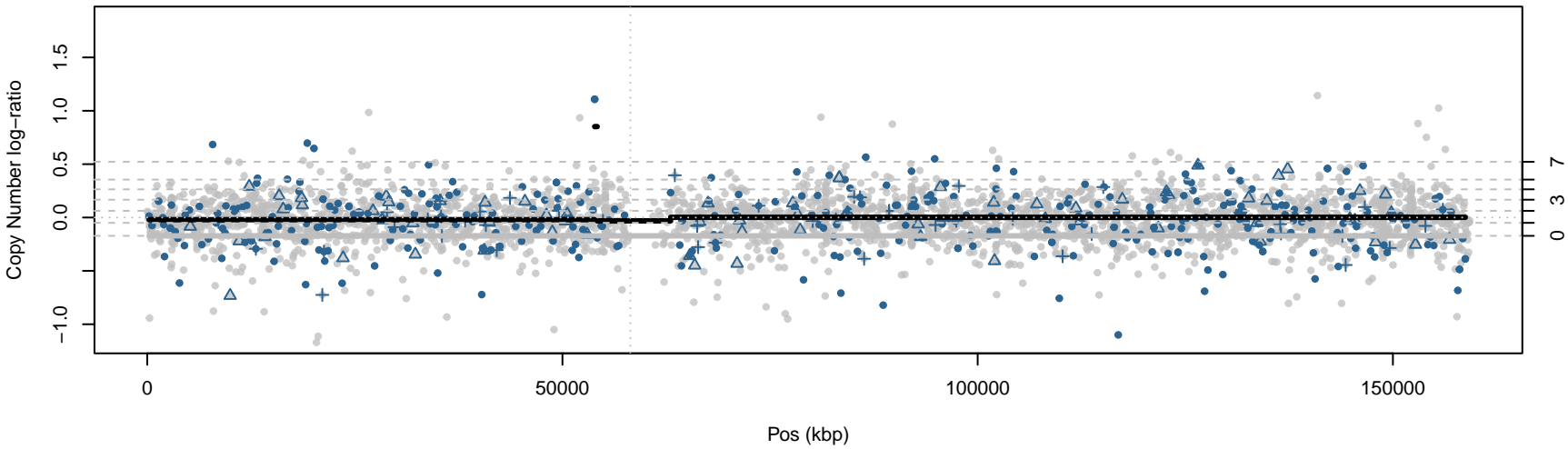
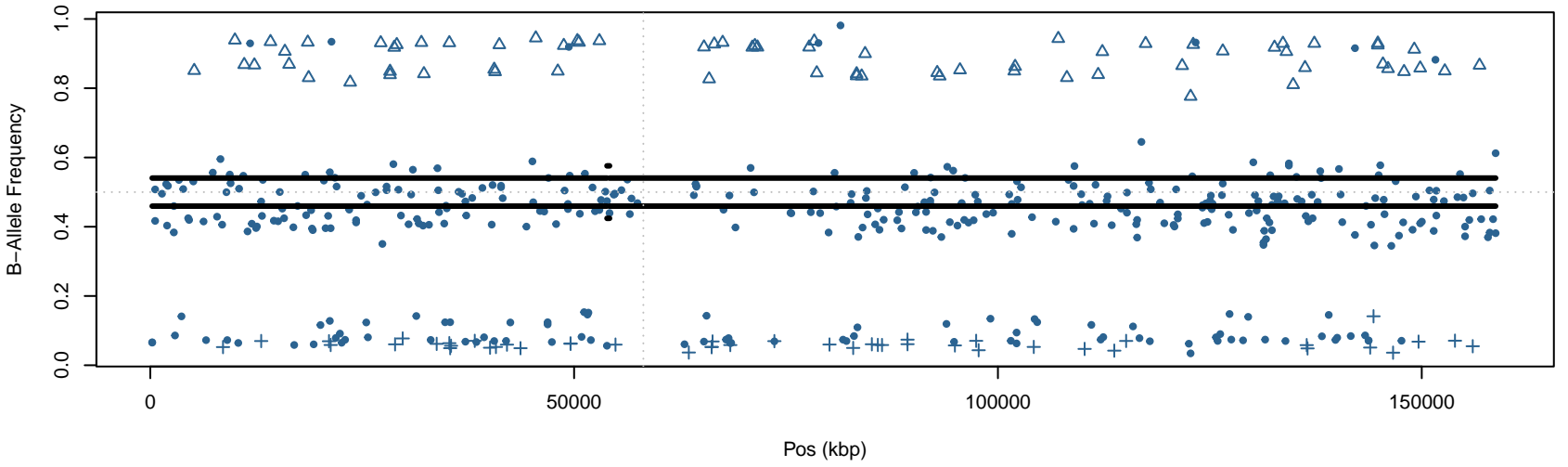
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr5



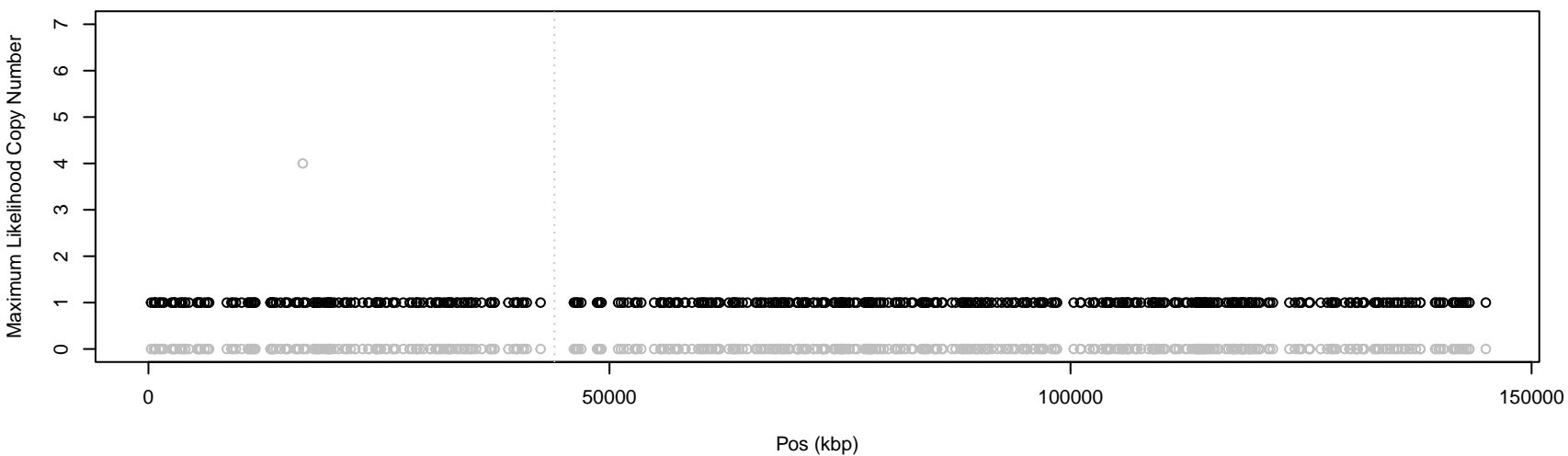
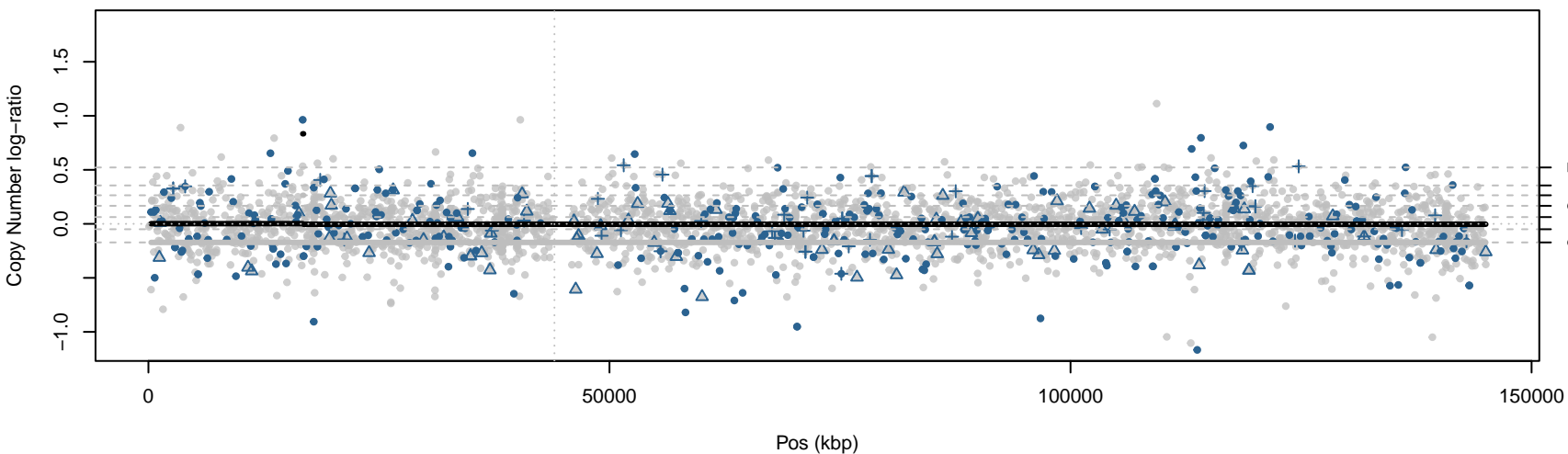
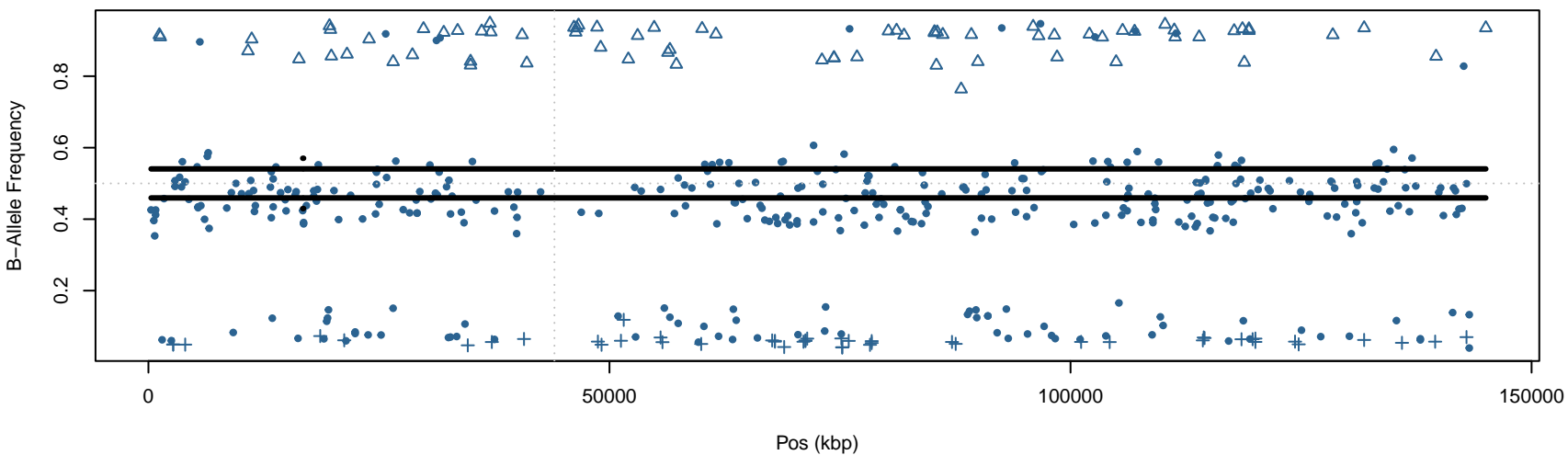
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr6



Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr7

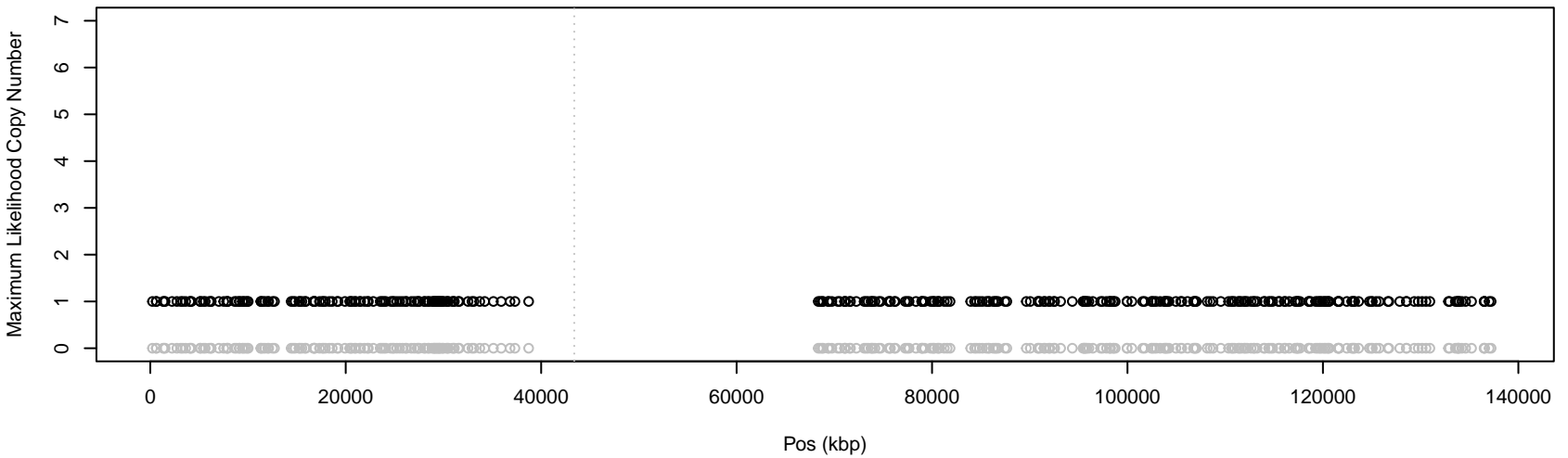
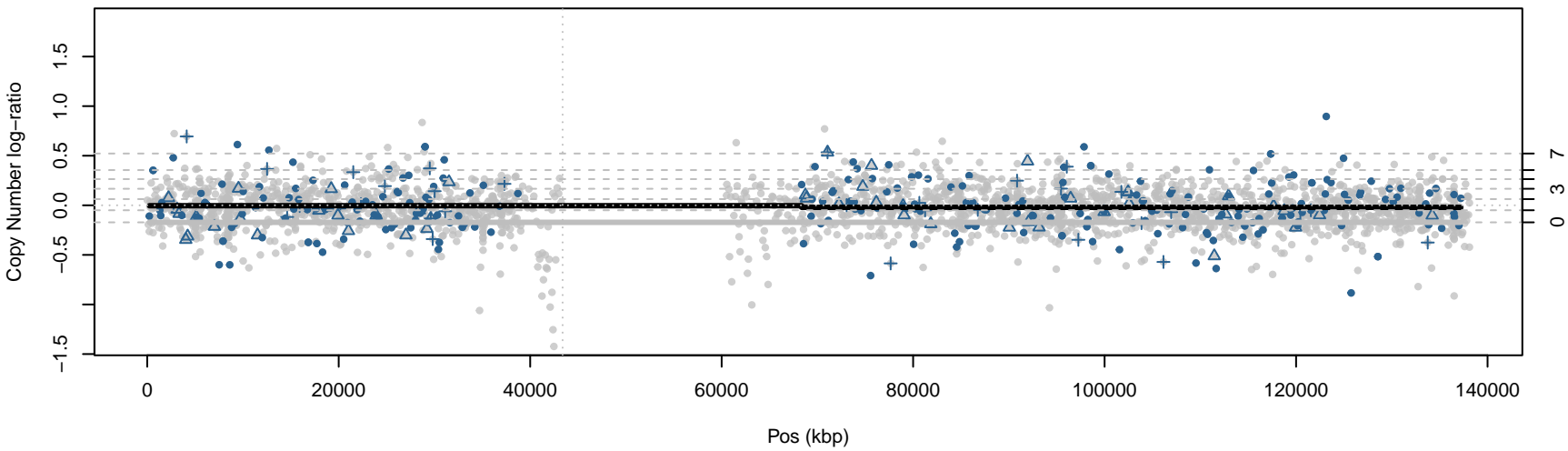
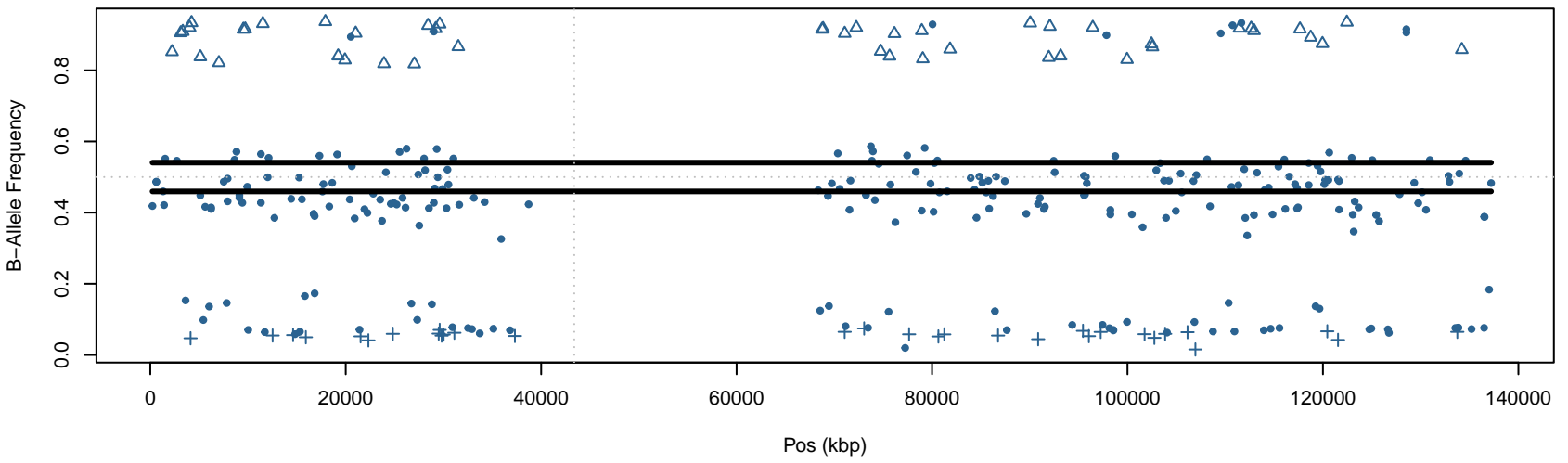


Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr8

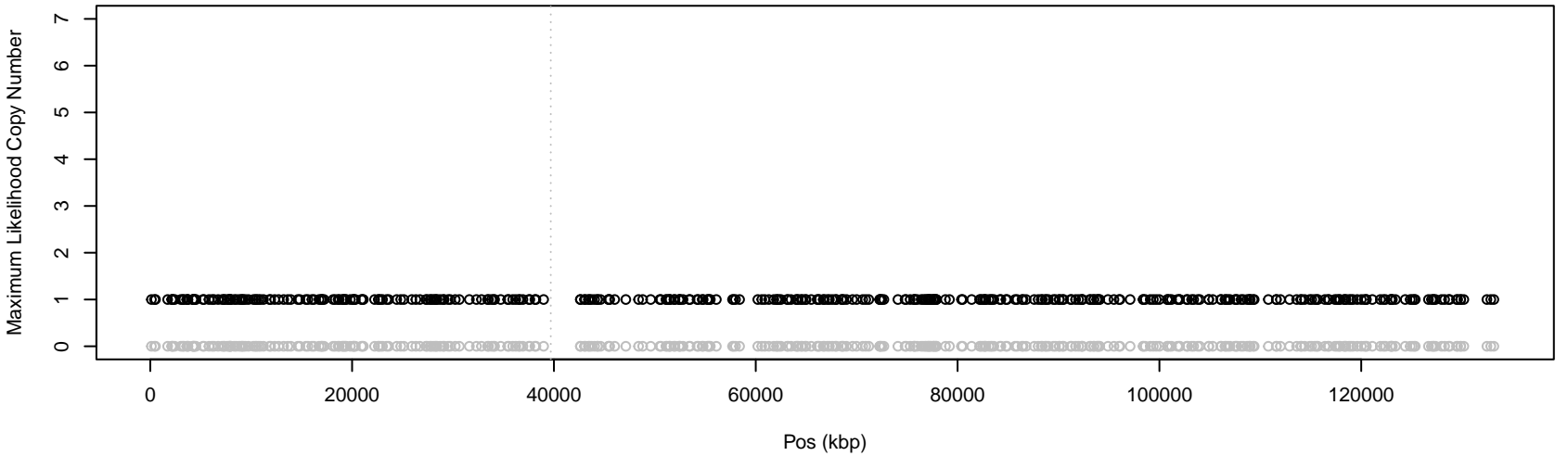
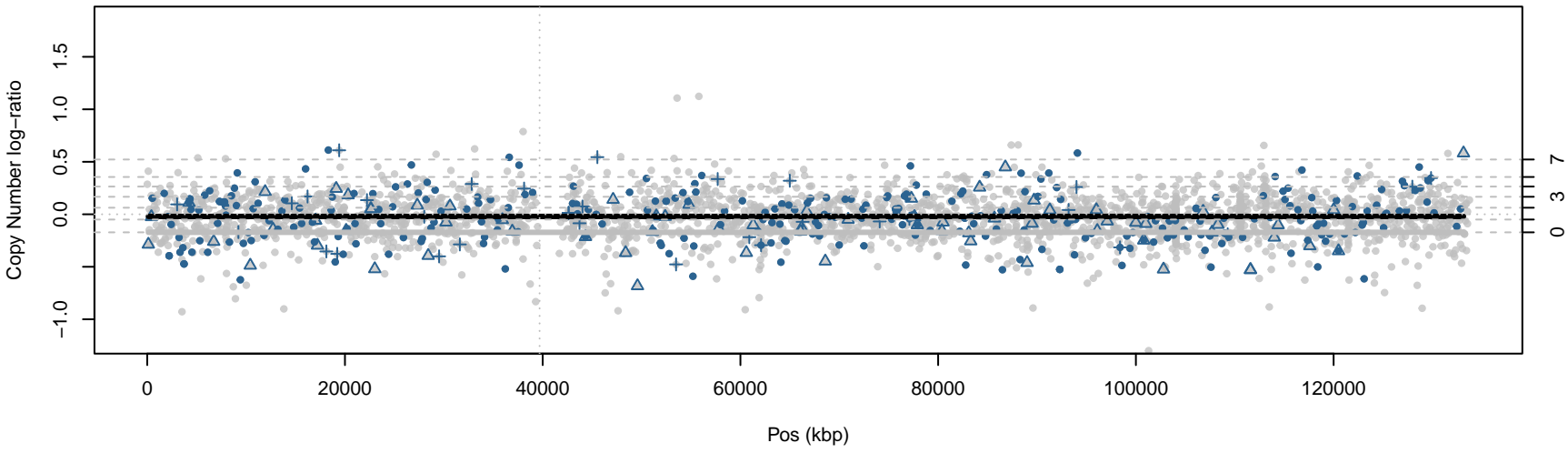
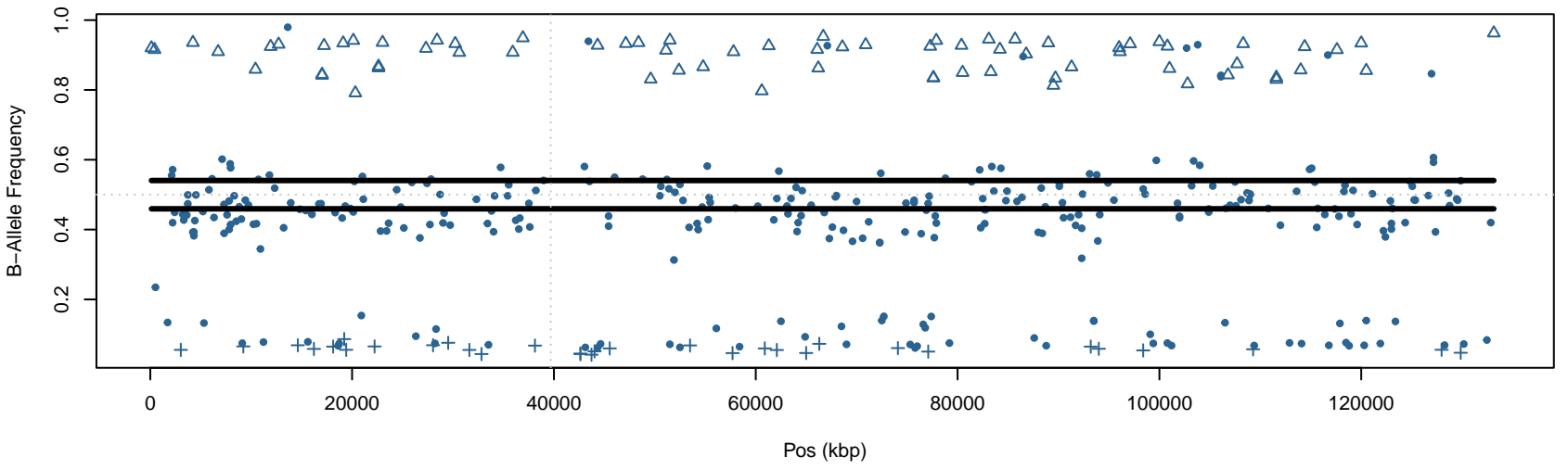




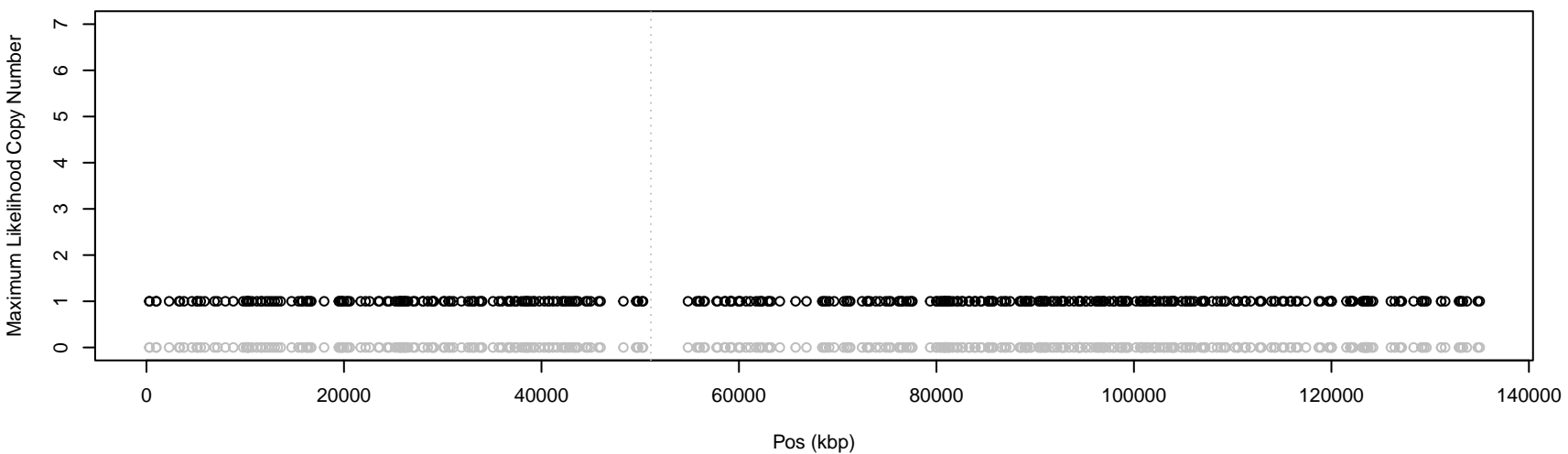
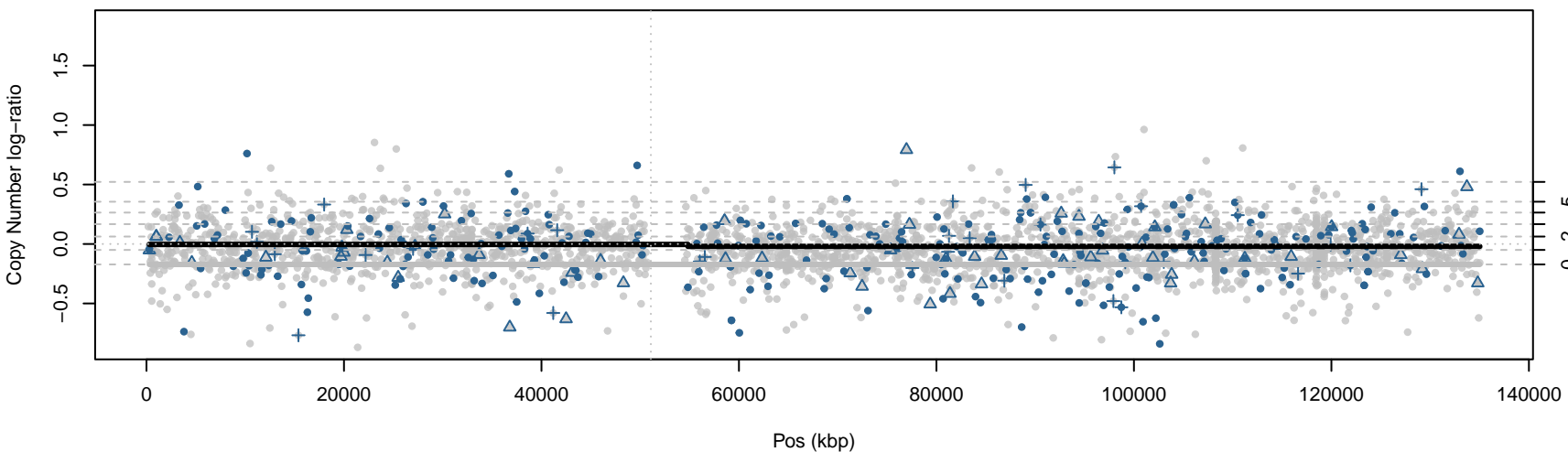
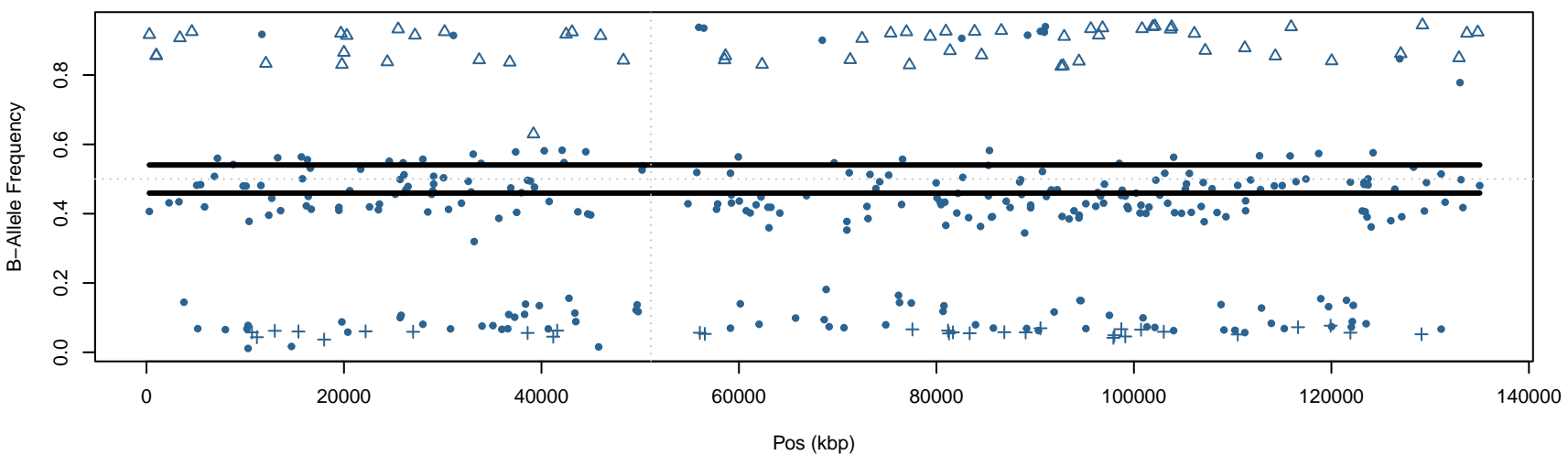
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr9



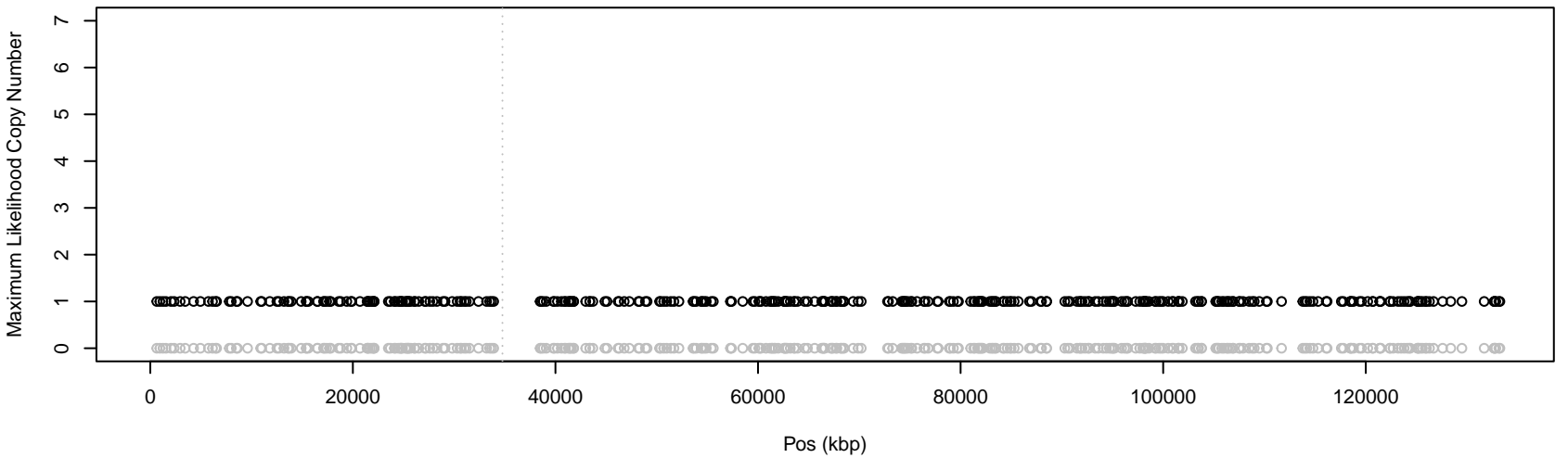
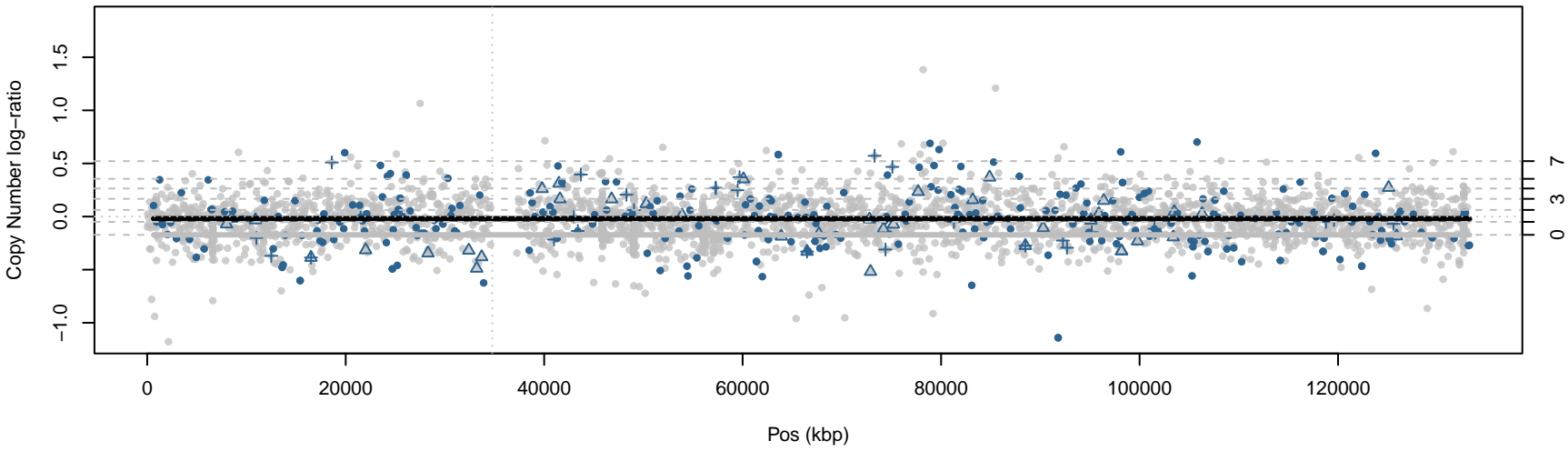
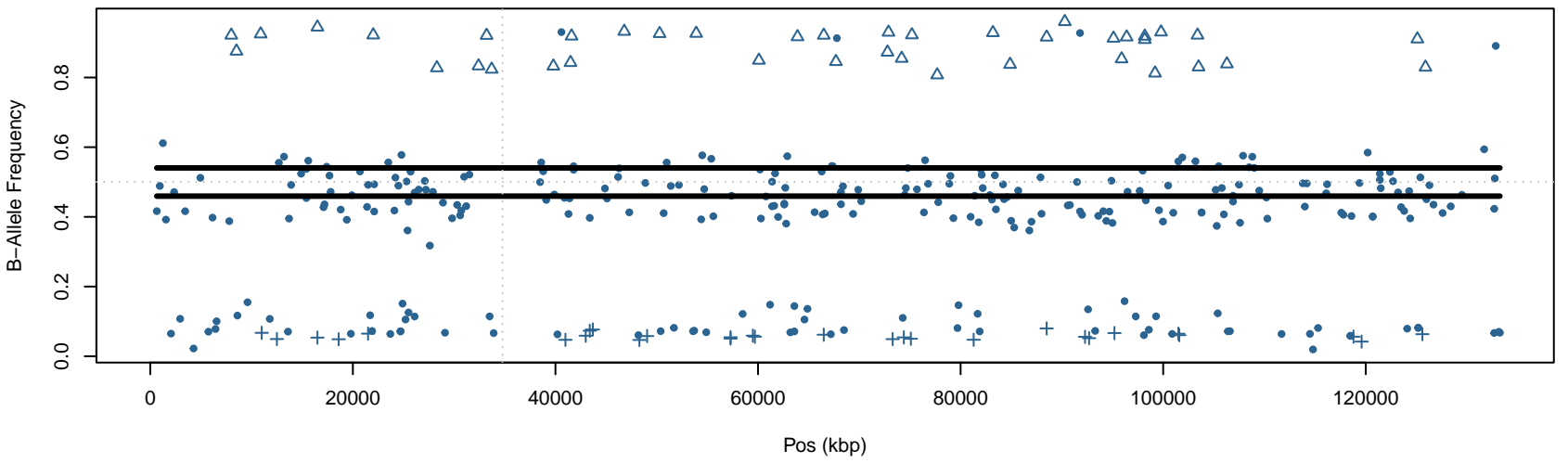
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr10



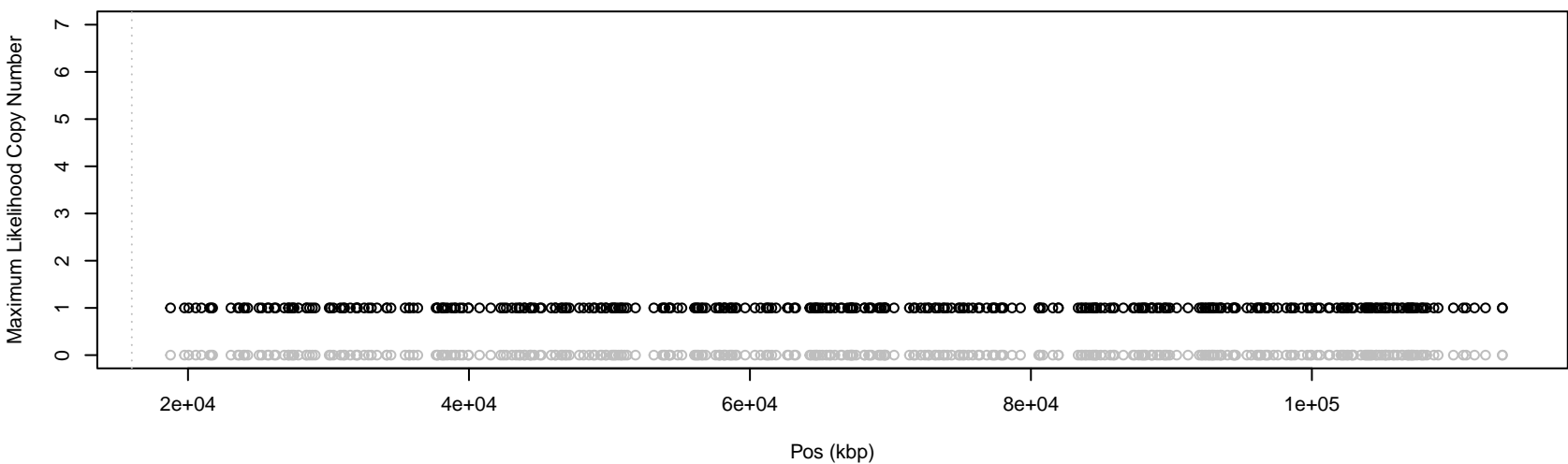
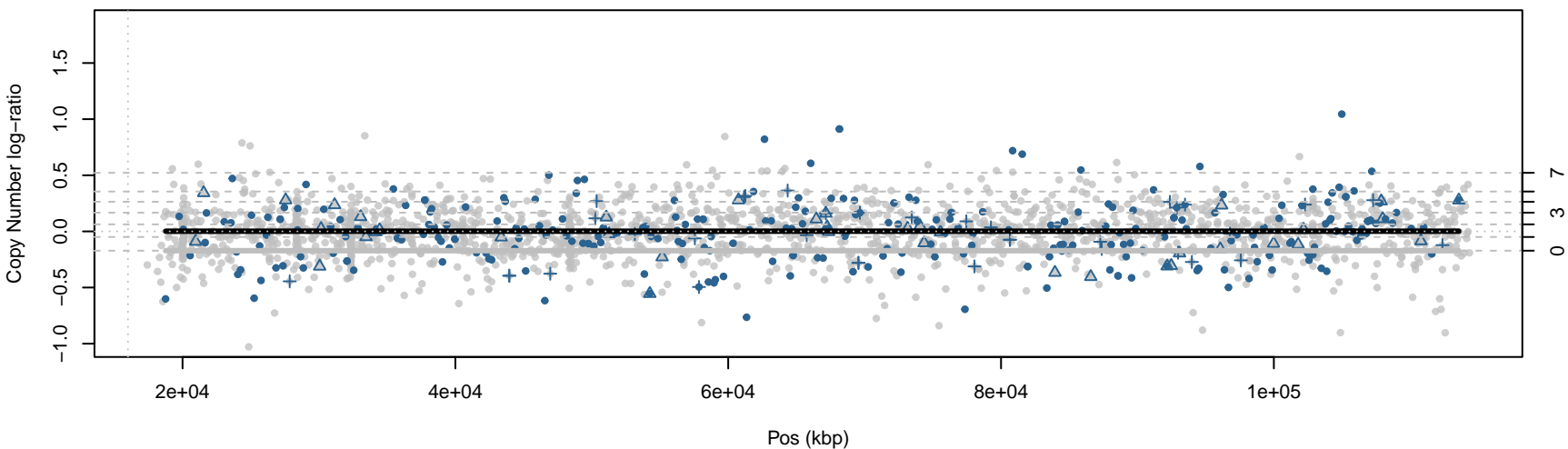
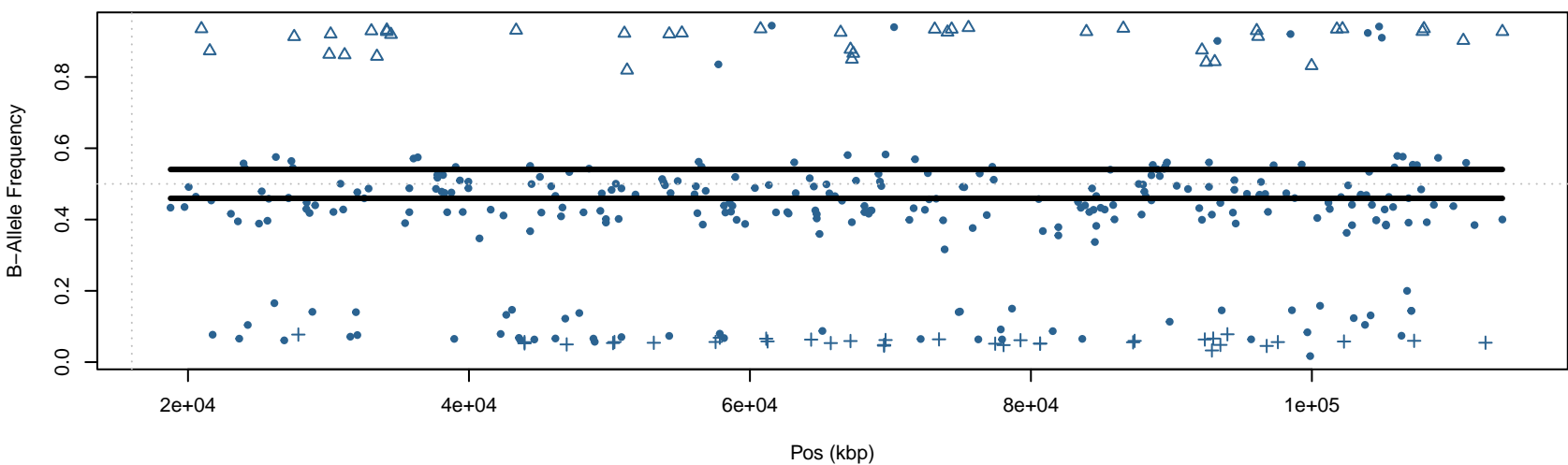
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr11



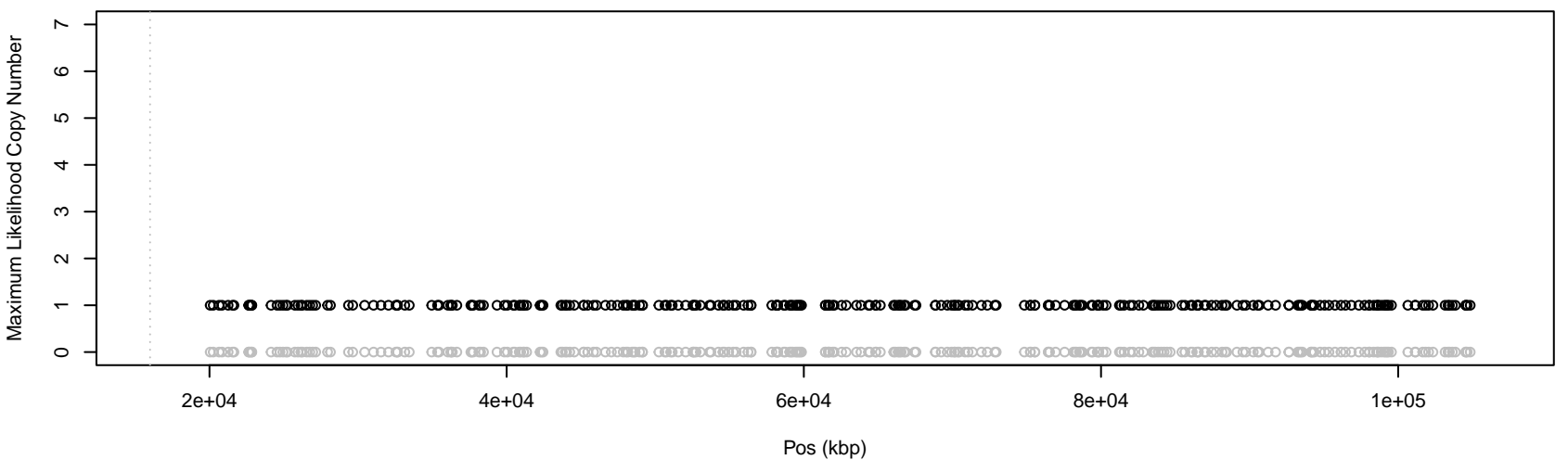
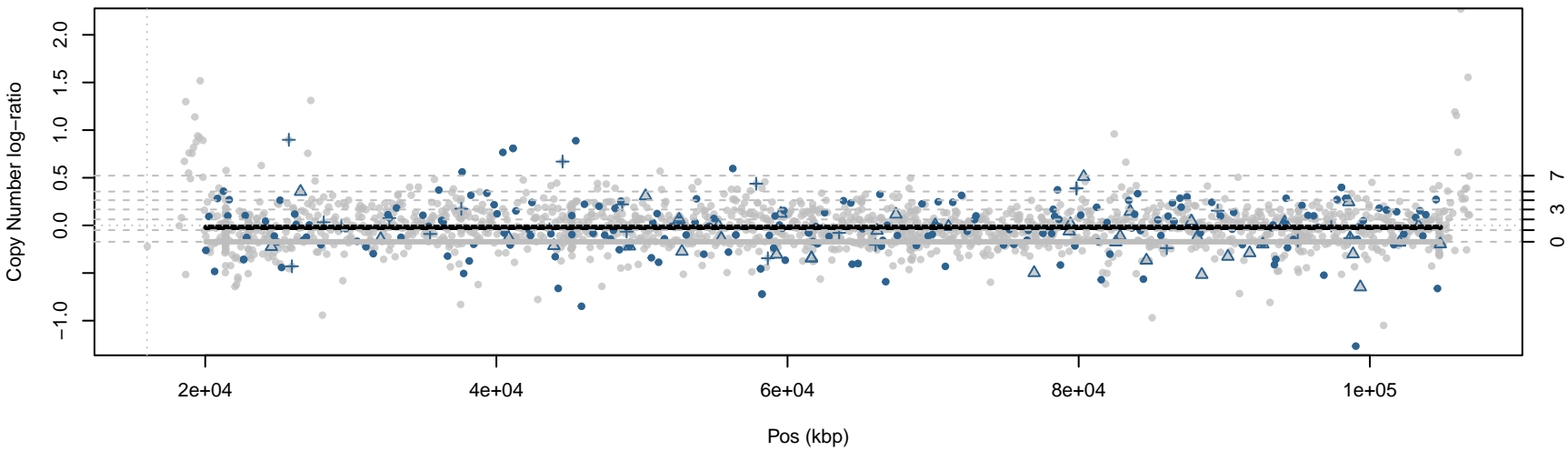
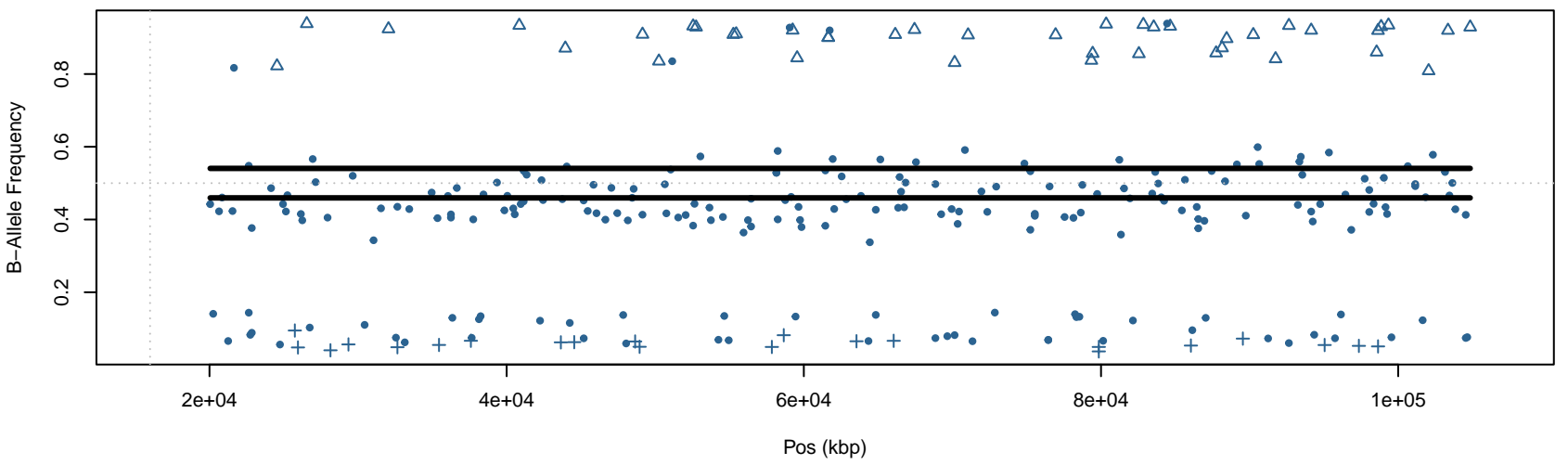
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr12



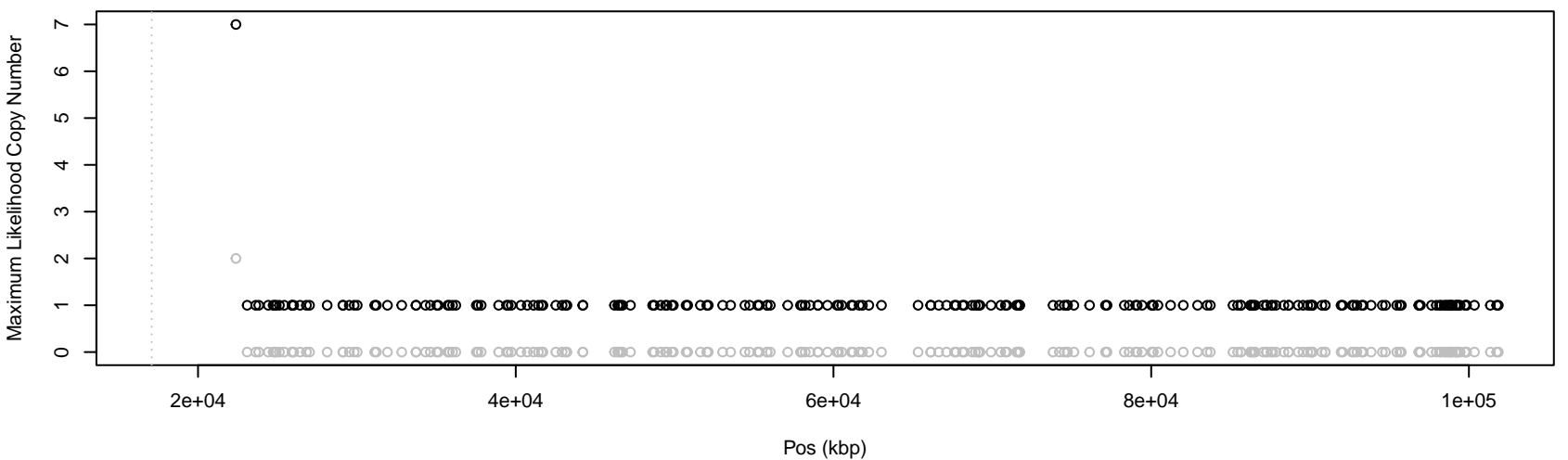
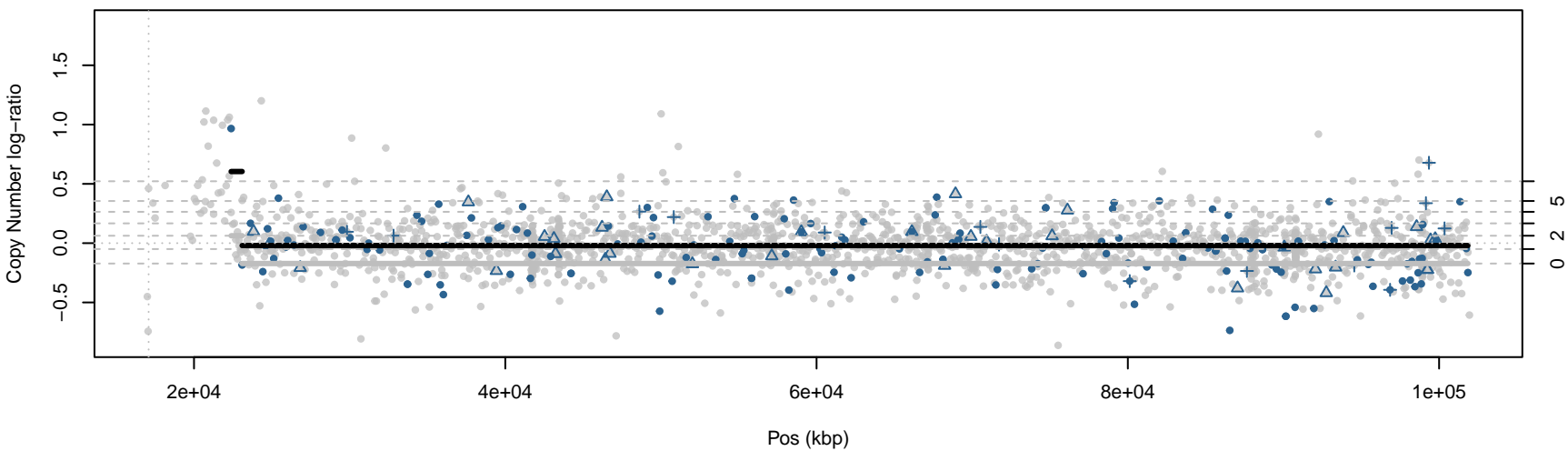
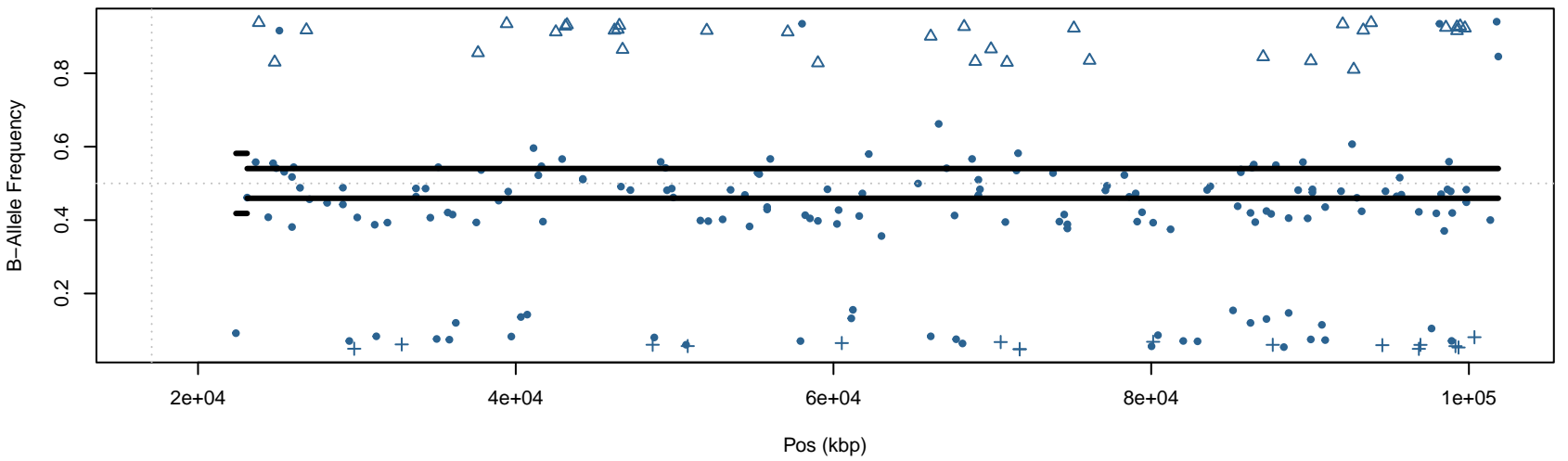
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr13



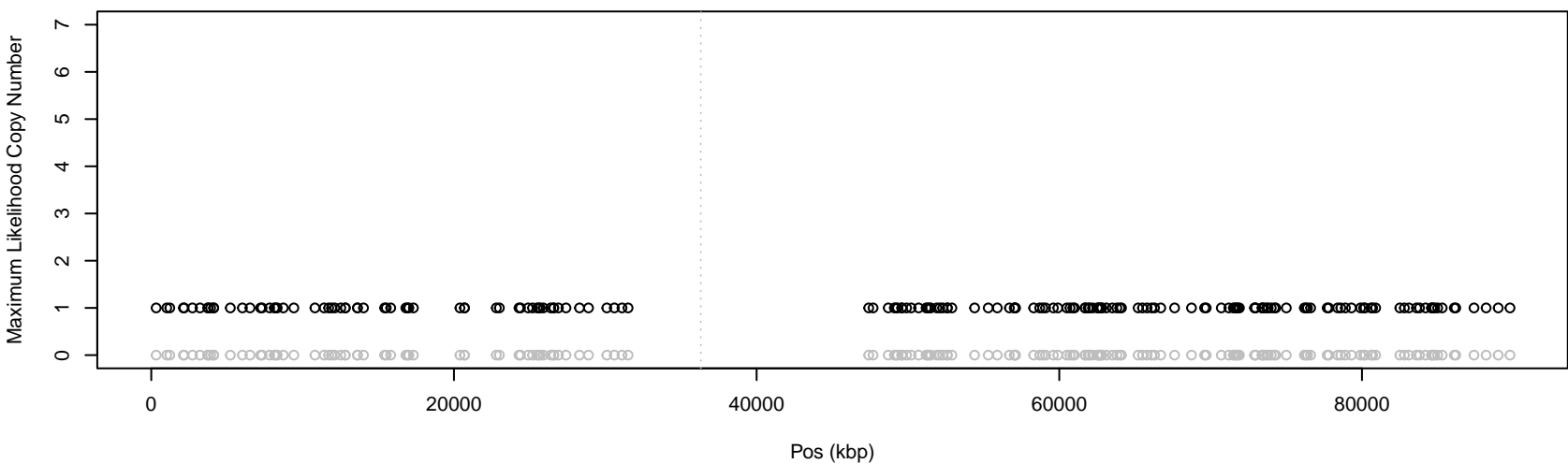
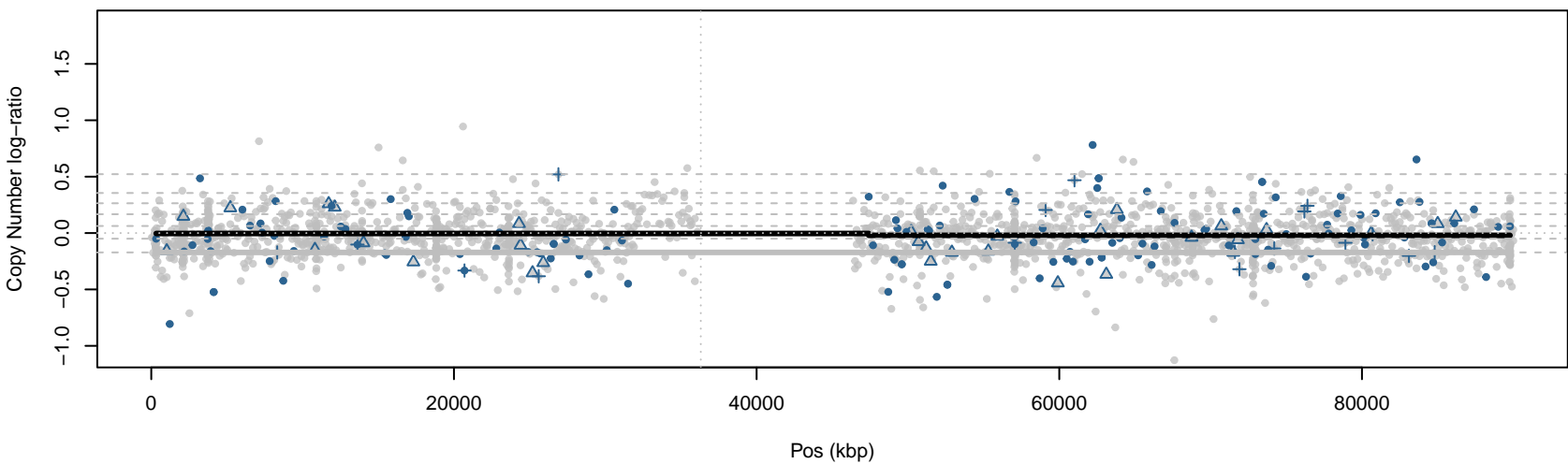
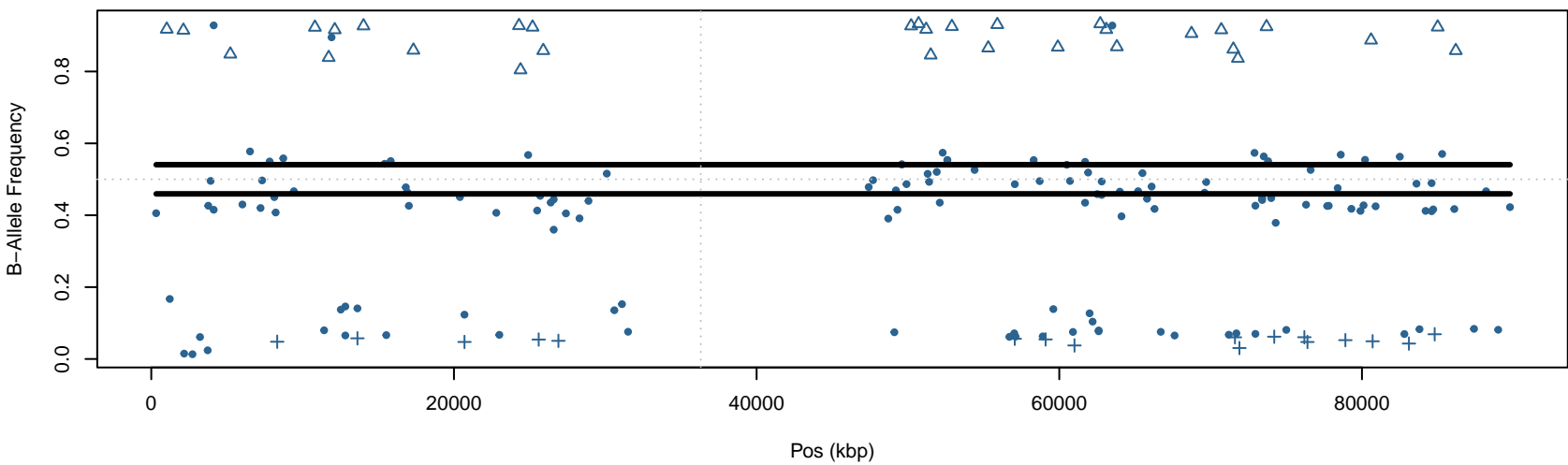
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr14



Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr15

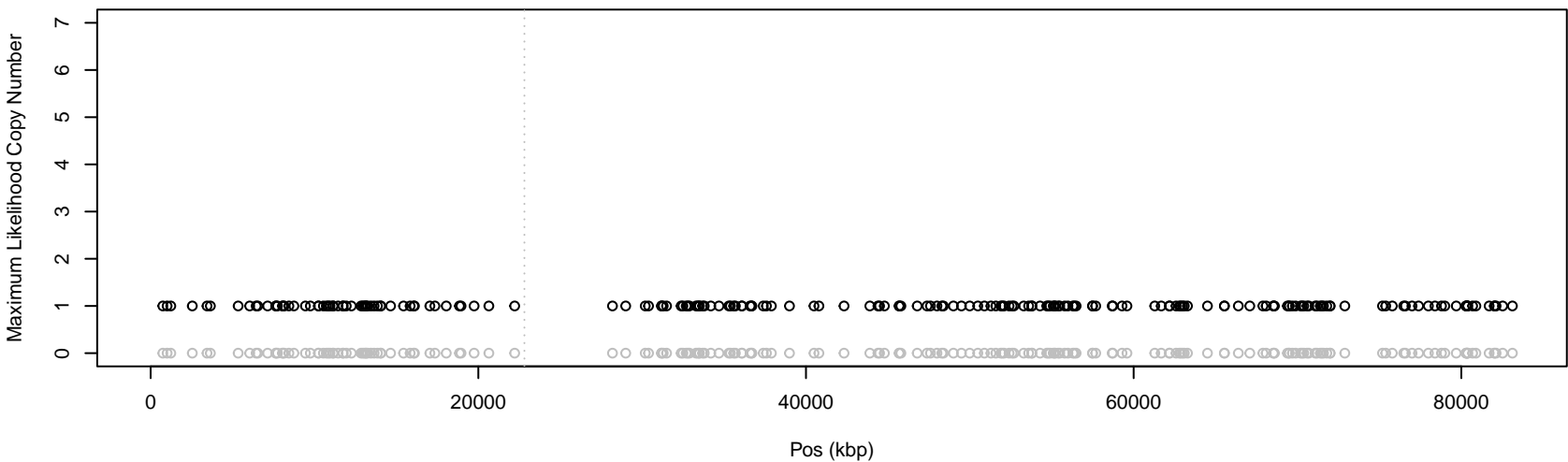
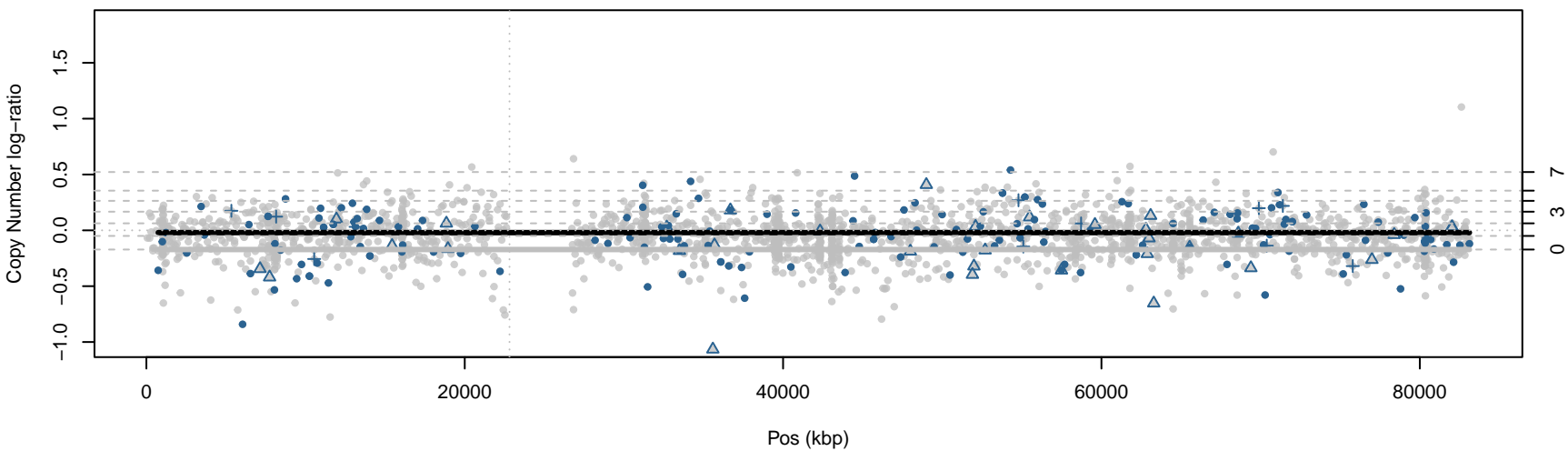
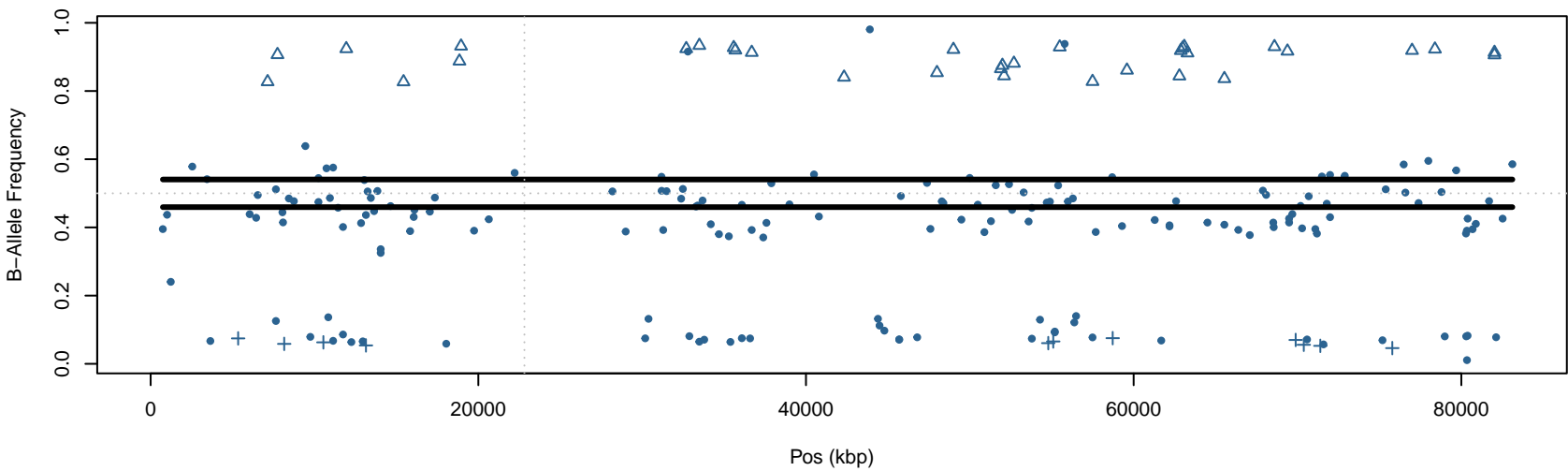


Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr16

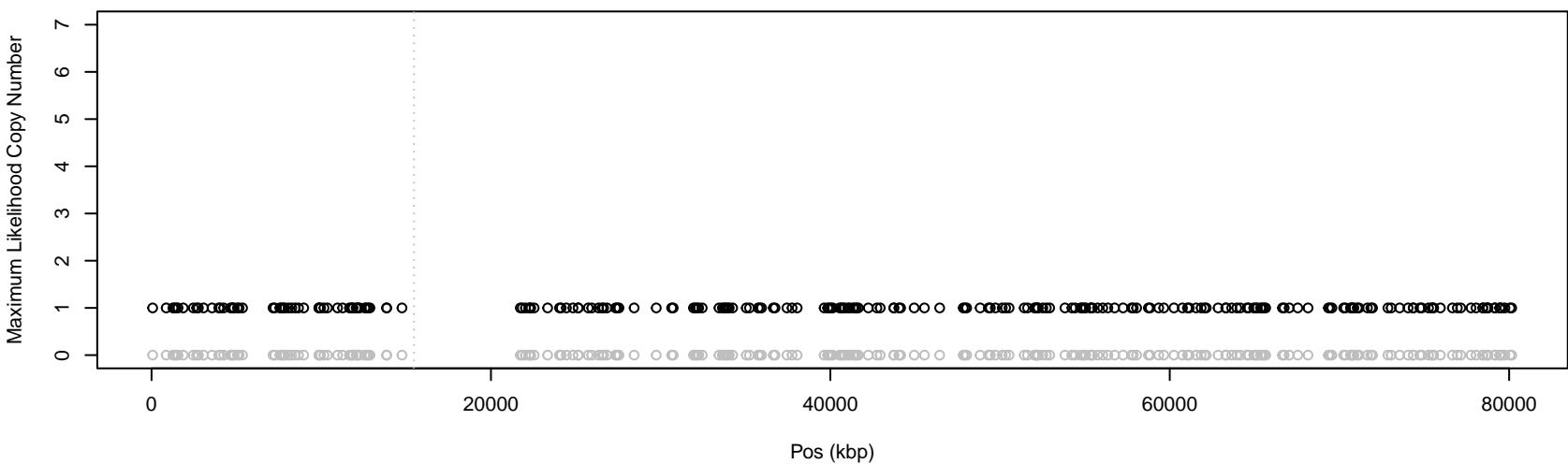
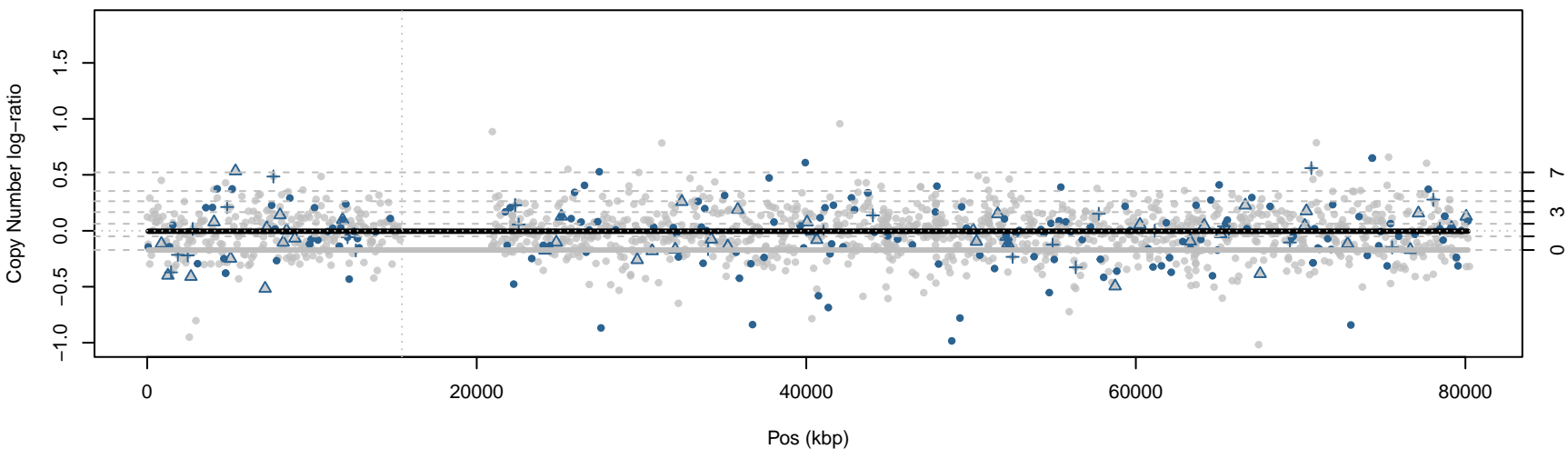
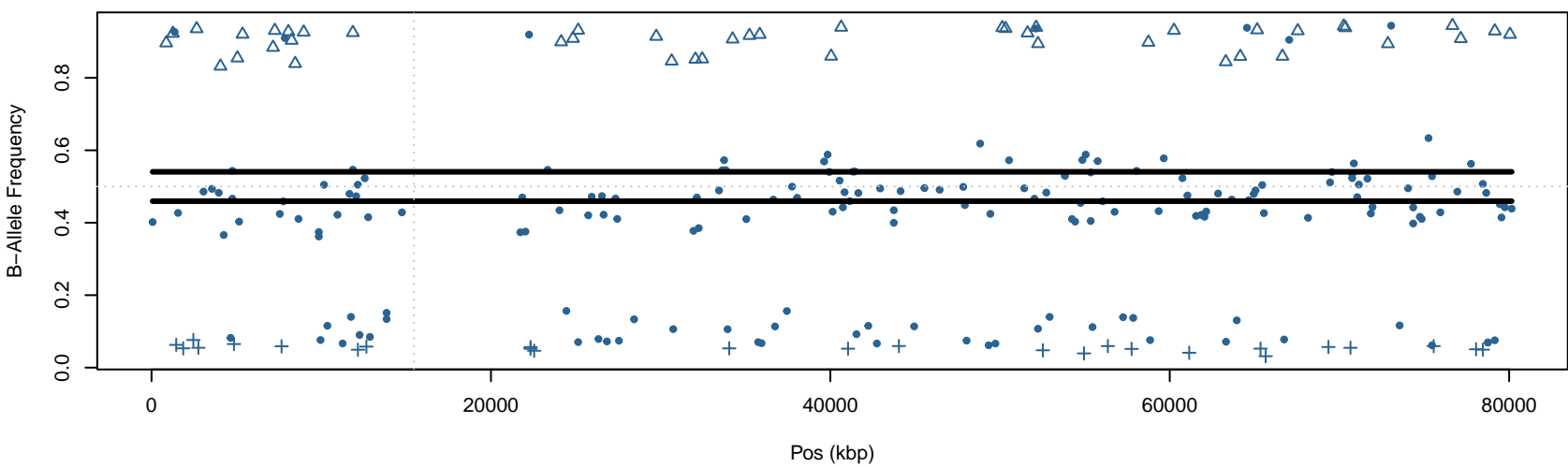




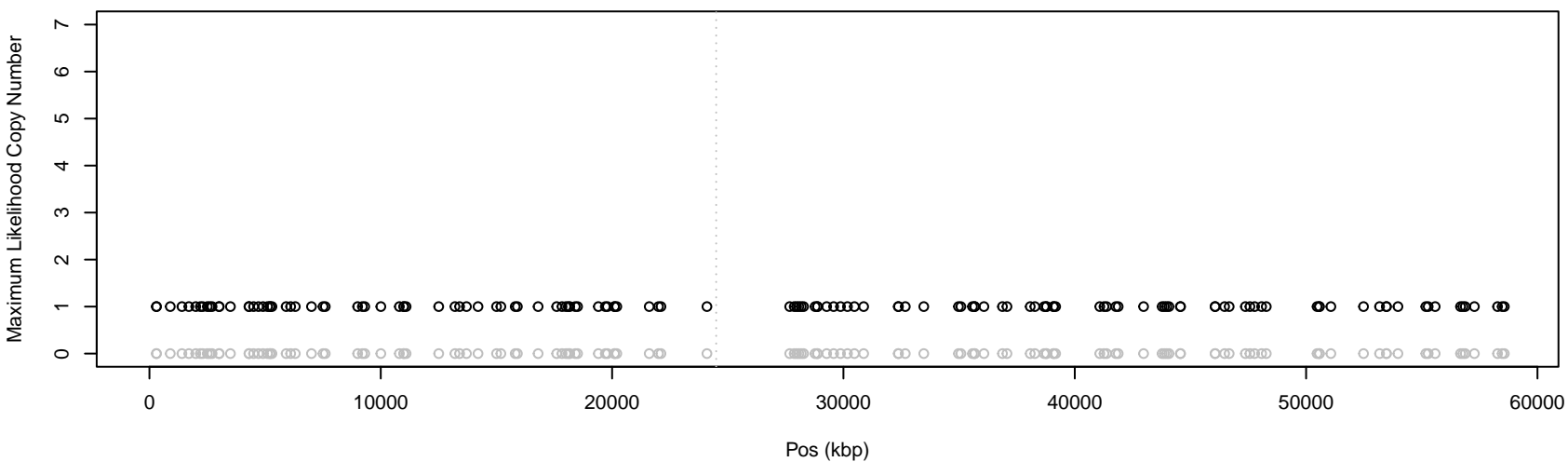
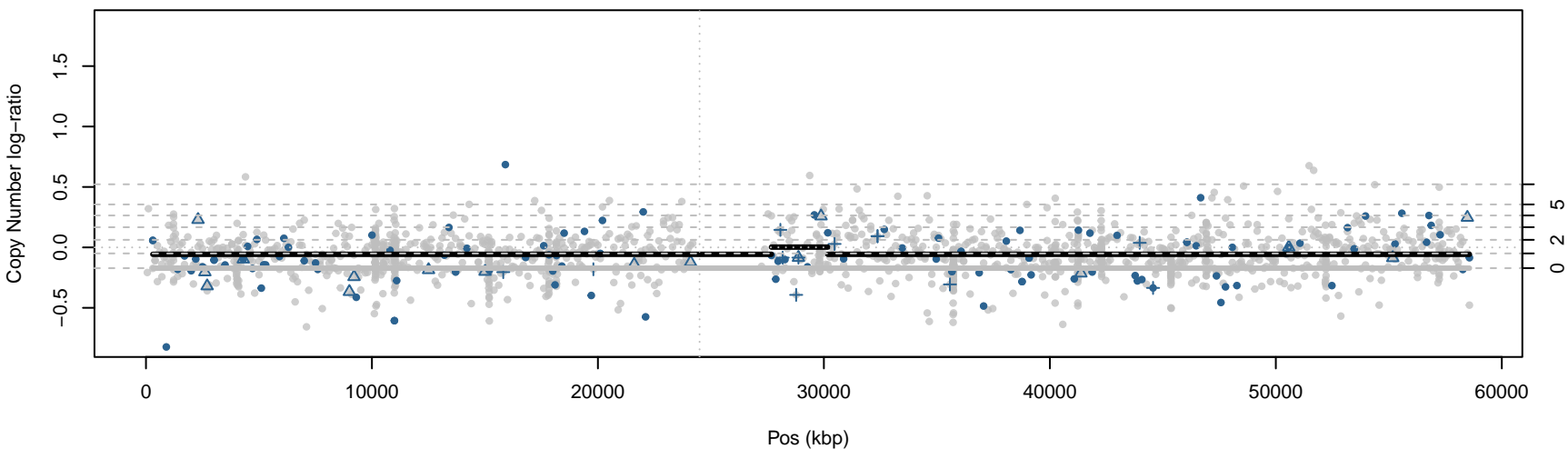
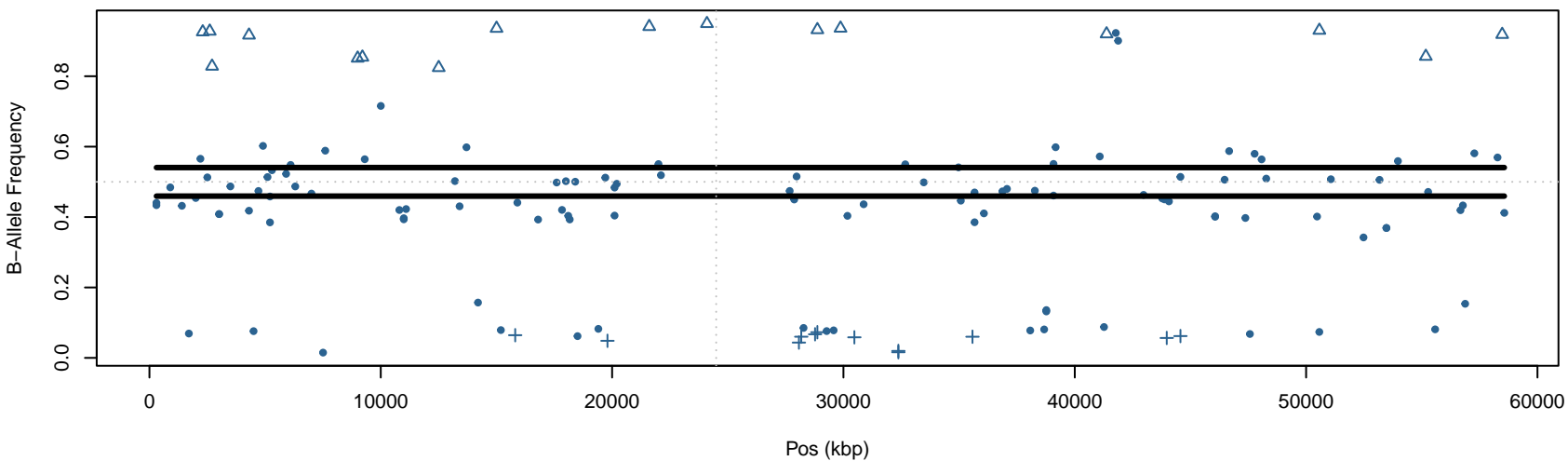
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr17



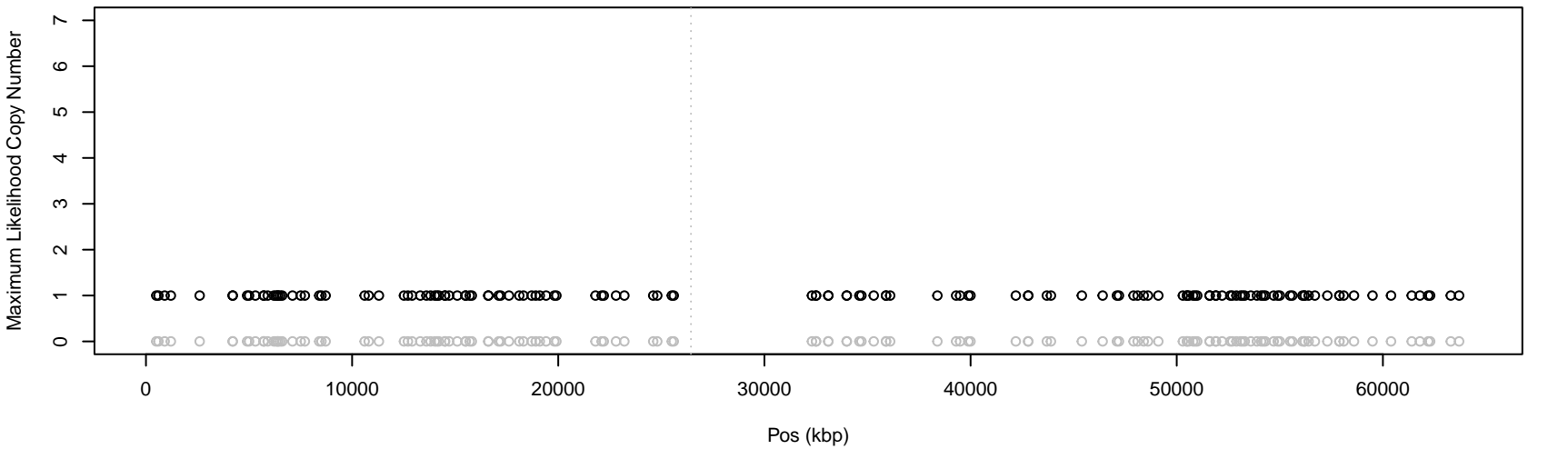
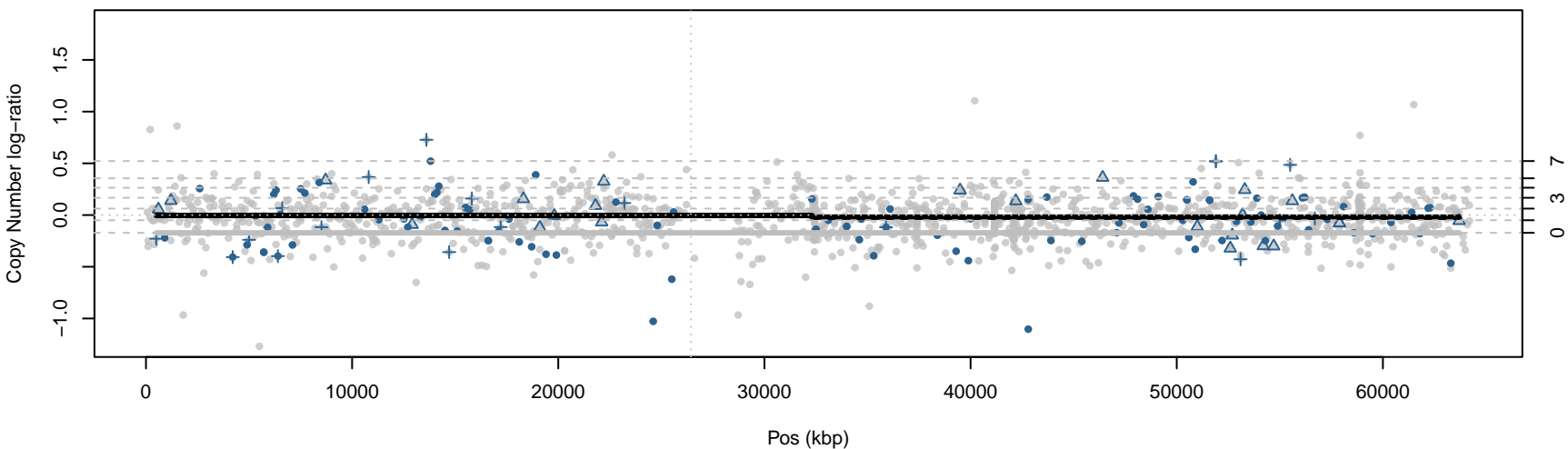
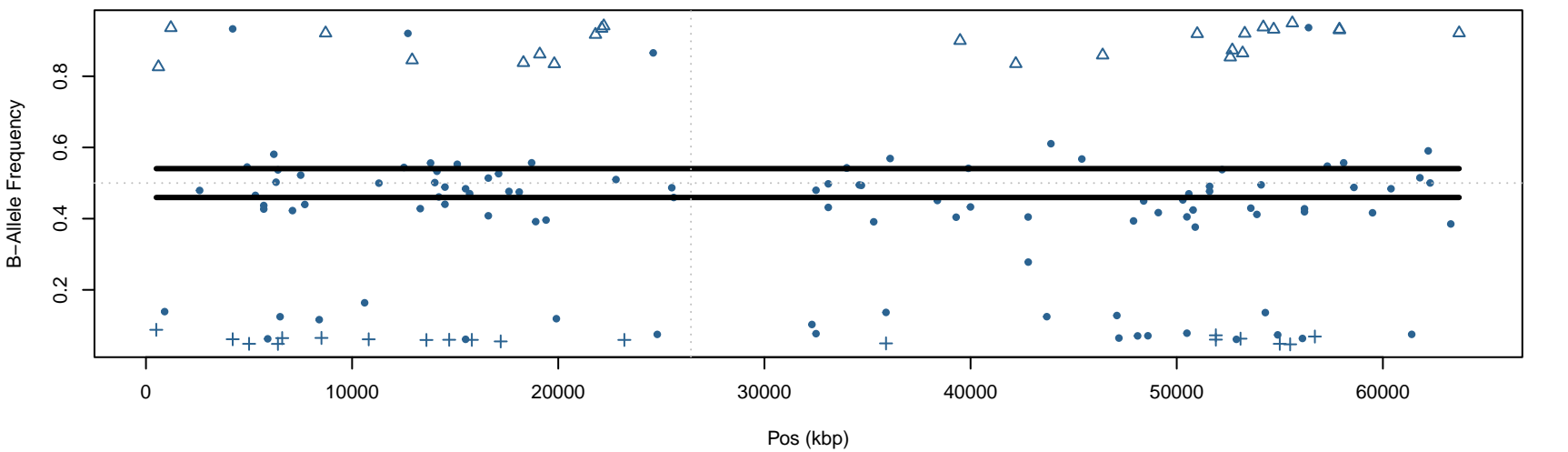
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr18



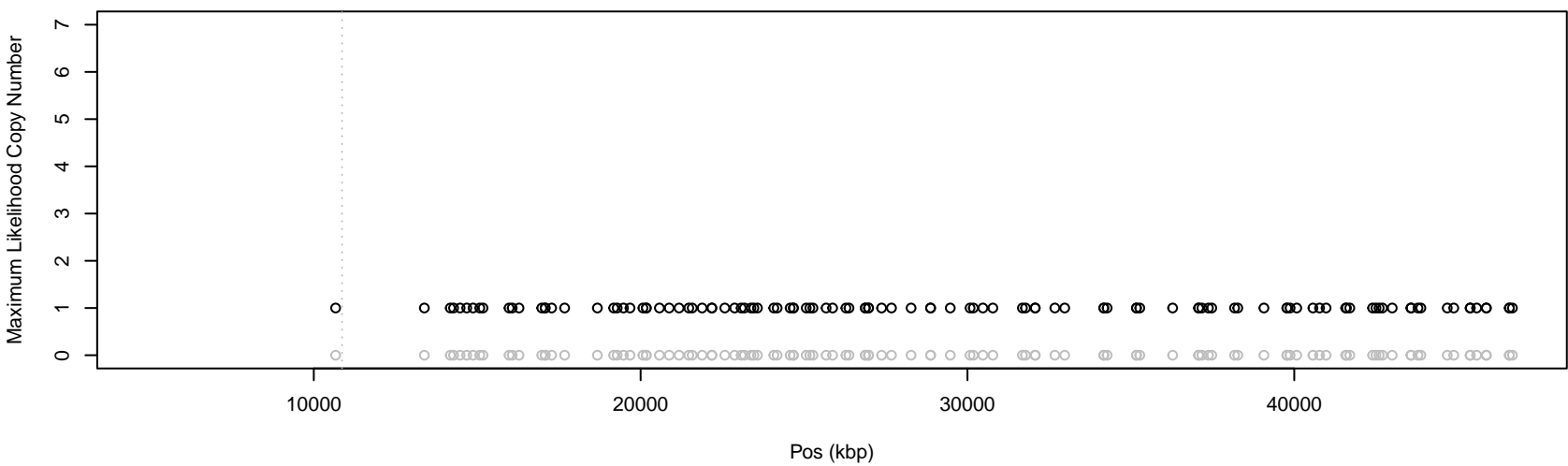
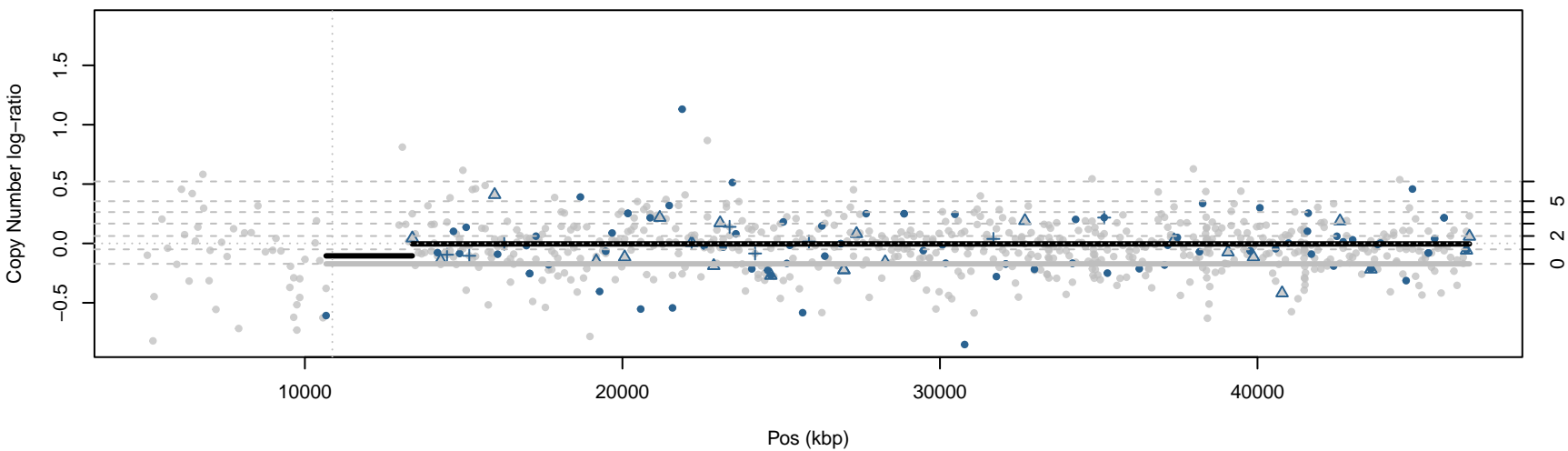
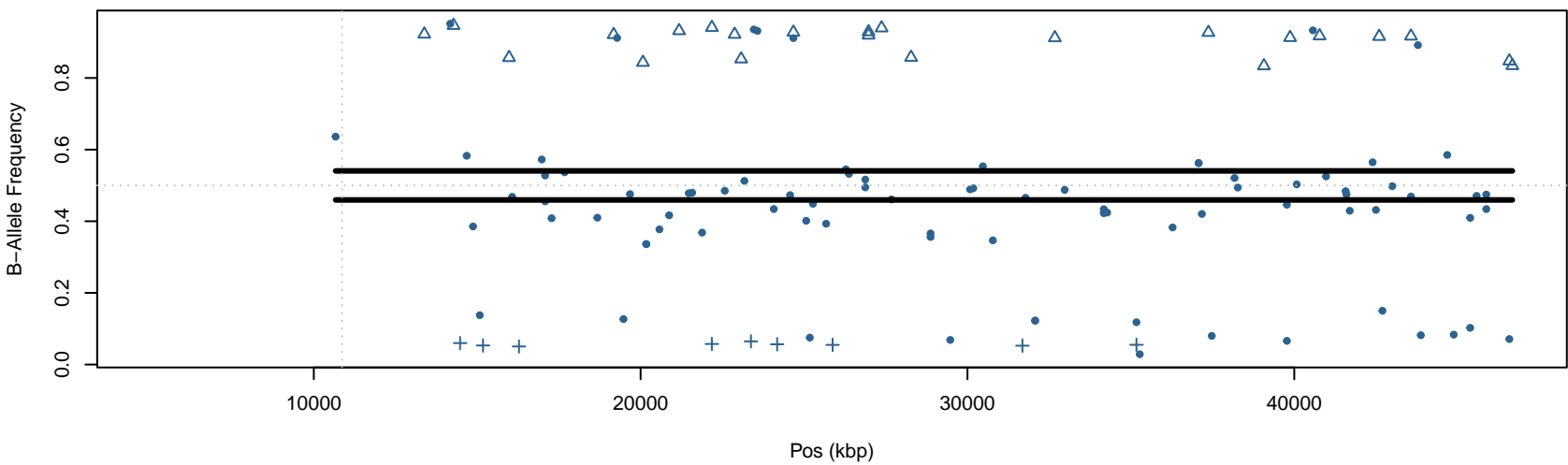
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr19



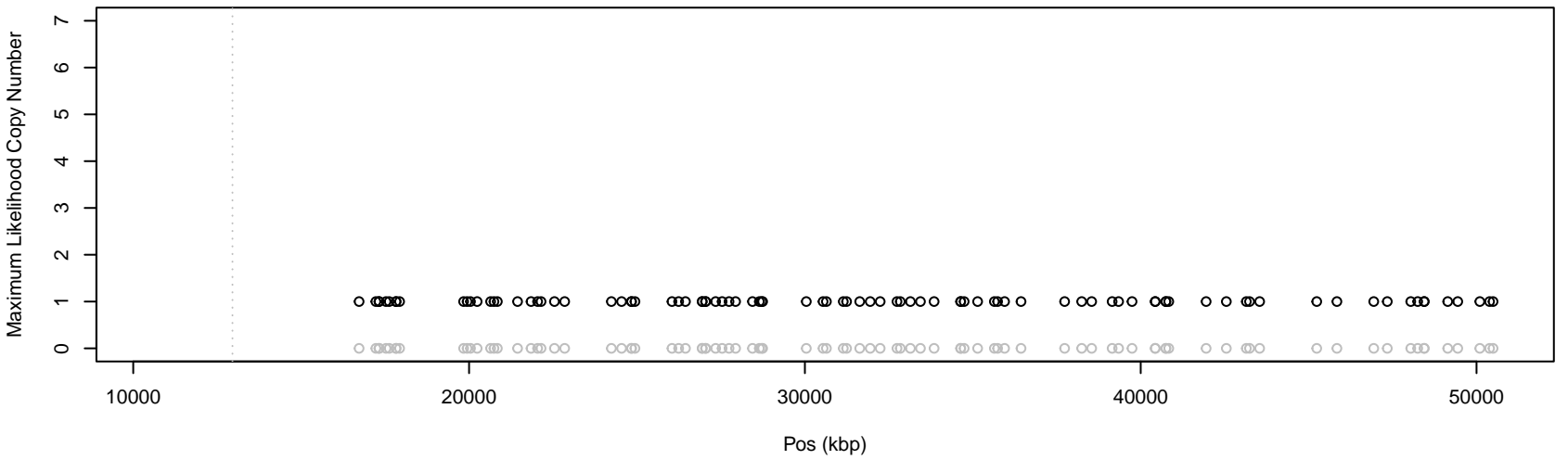
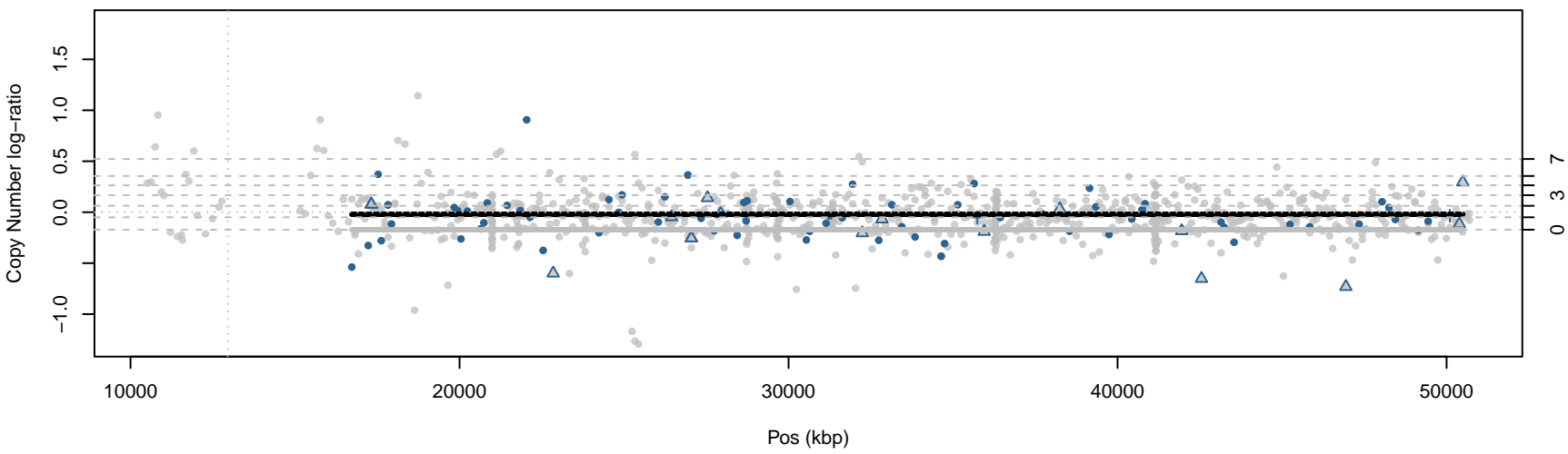
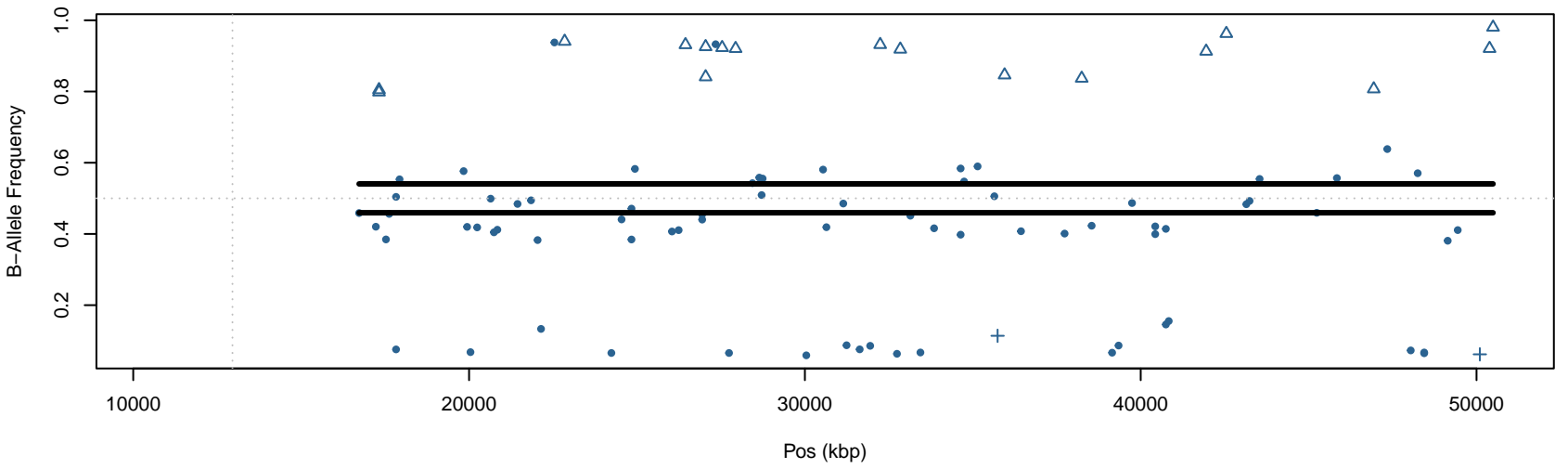
Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr20



Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr21



Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chr22



Purity: 0.15 Tumor ploidy: 1.434 SNV log-likelihood: -122256.27 GoF: 25.5% Mean coverage: 1651 Chromosome: chrX

