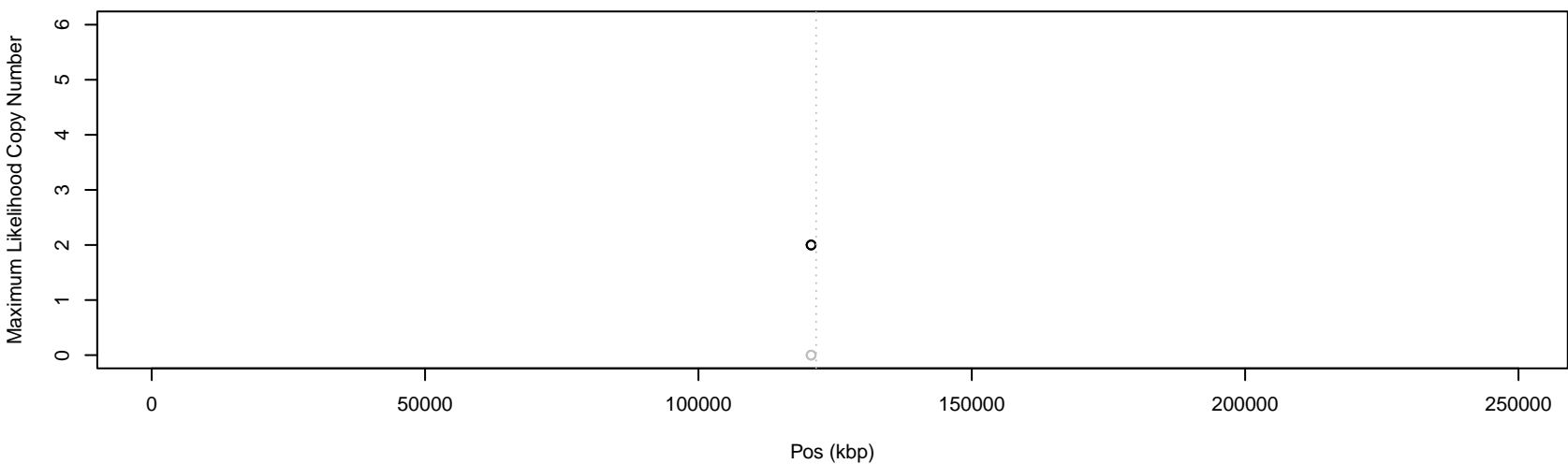
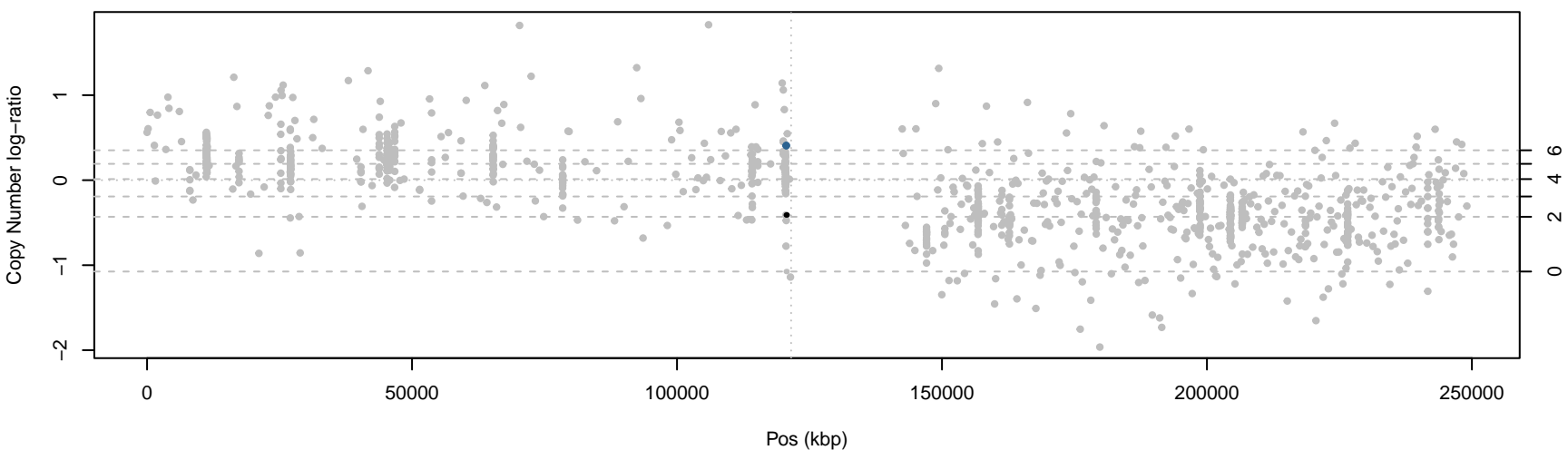
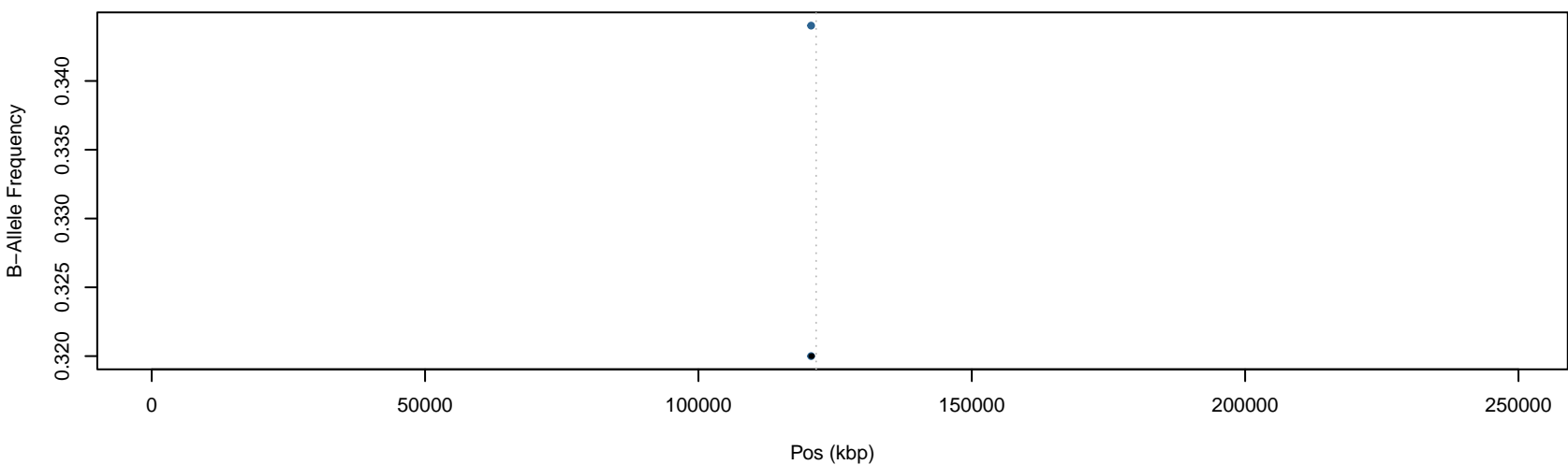
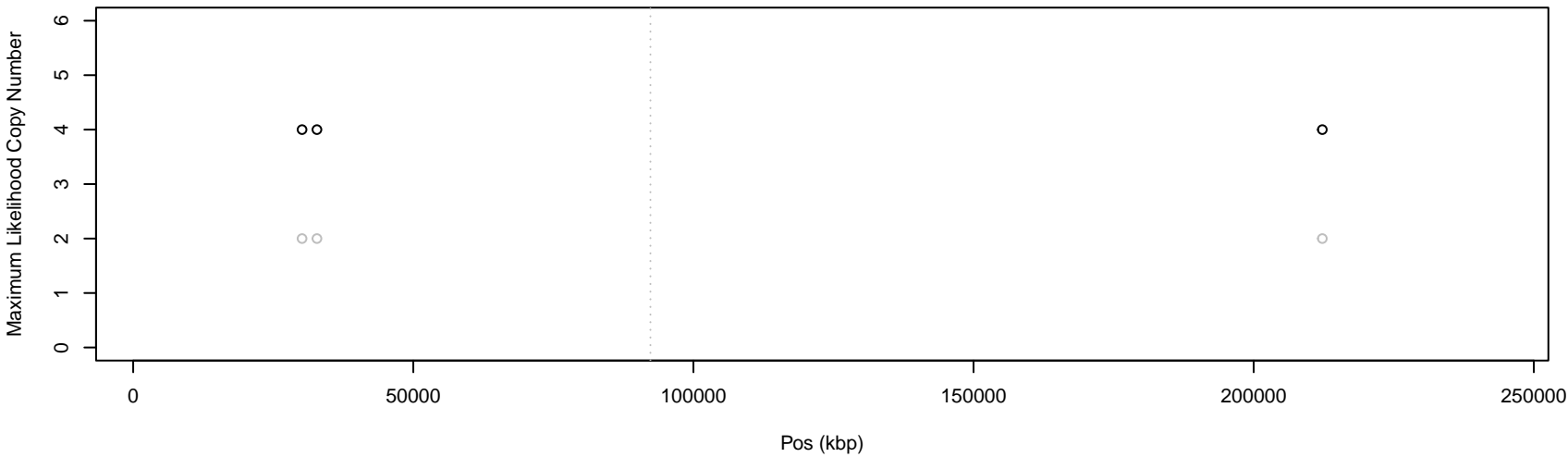
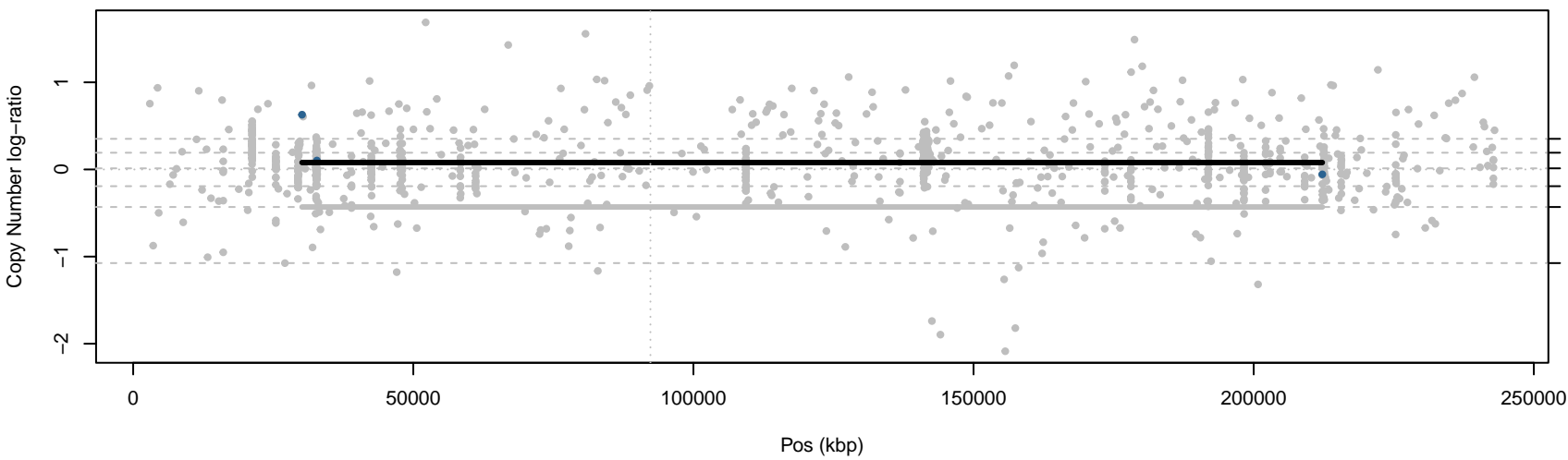
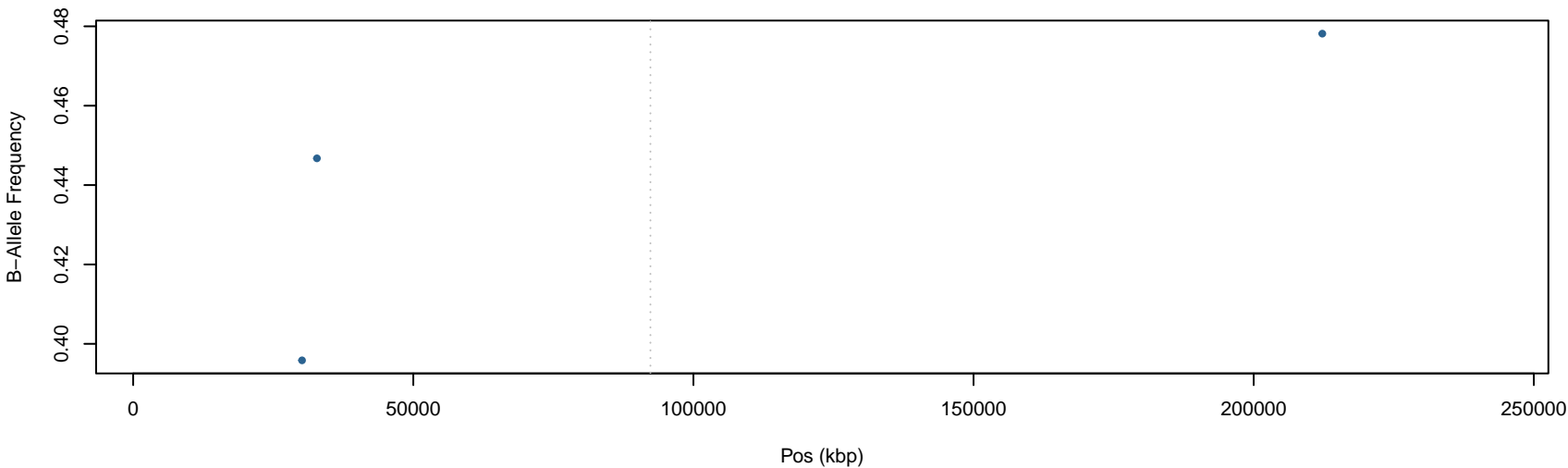


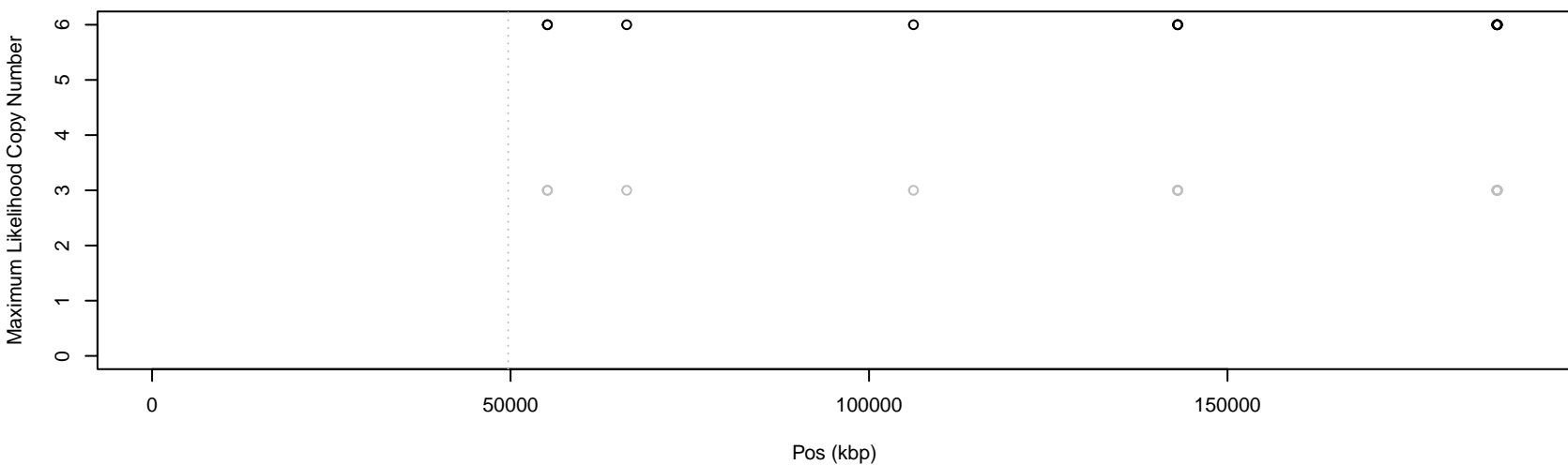
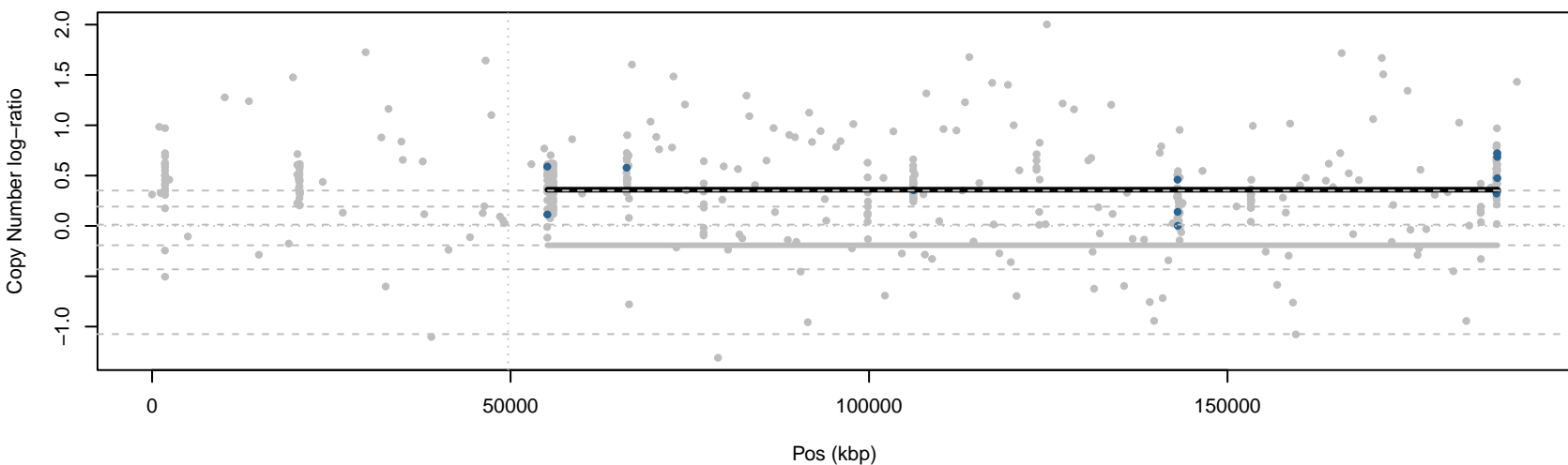
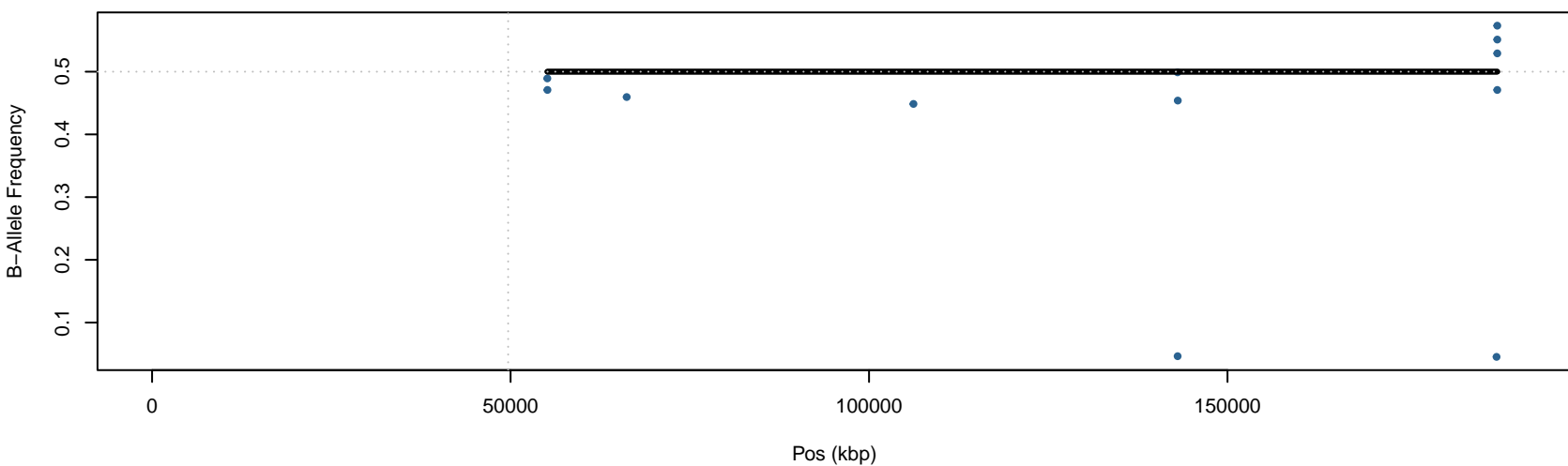
Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 1



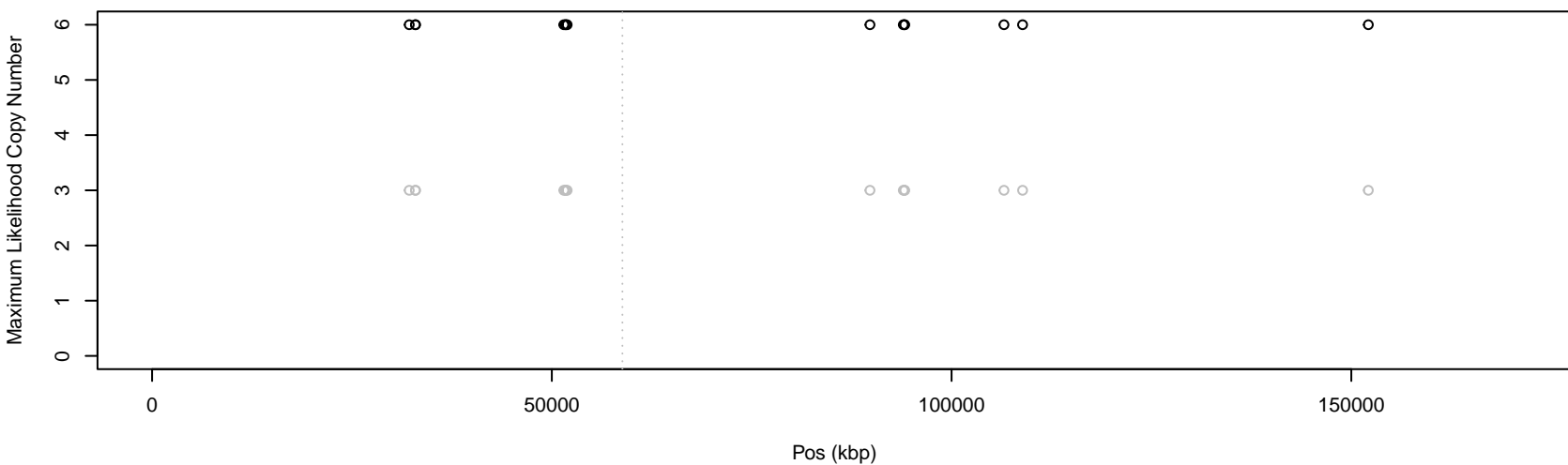
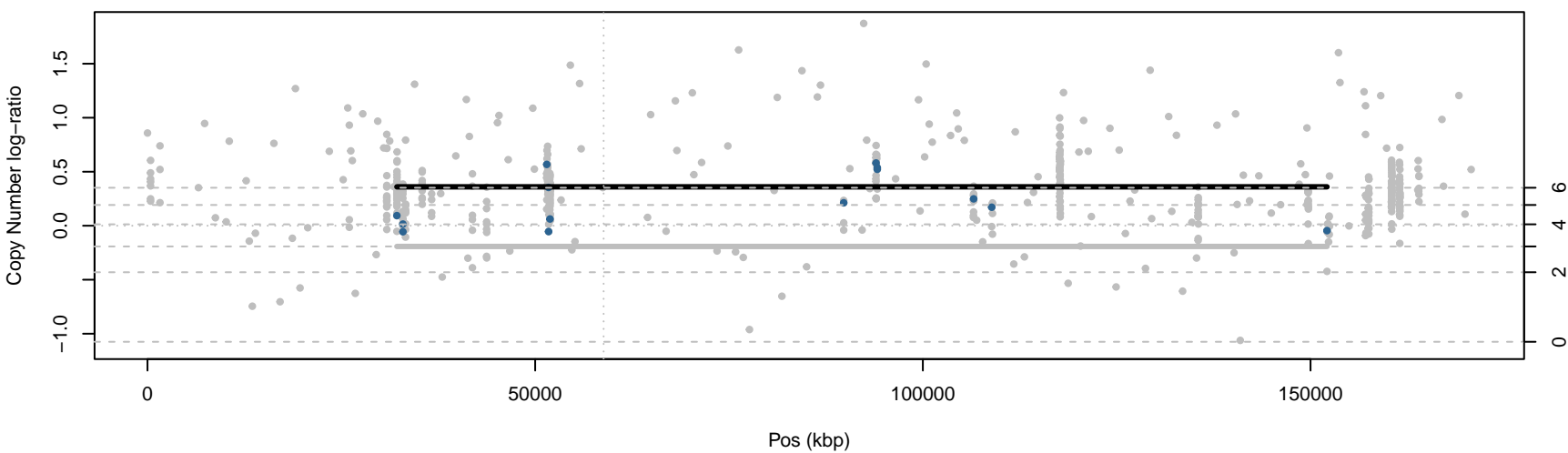
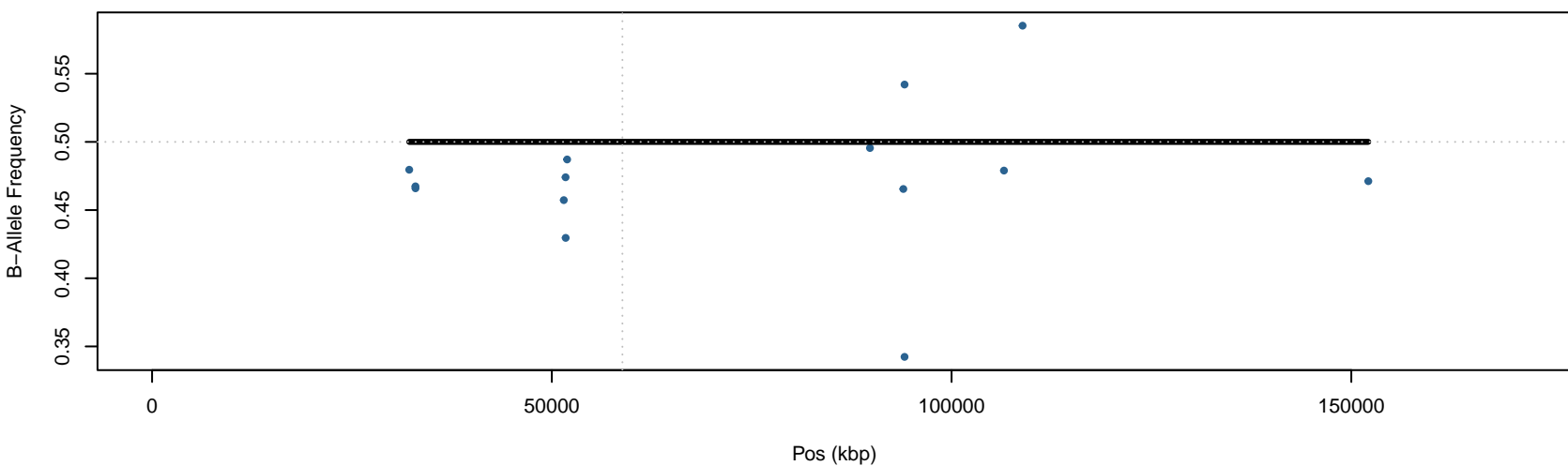
Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 2



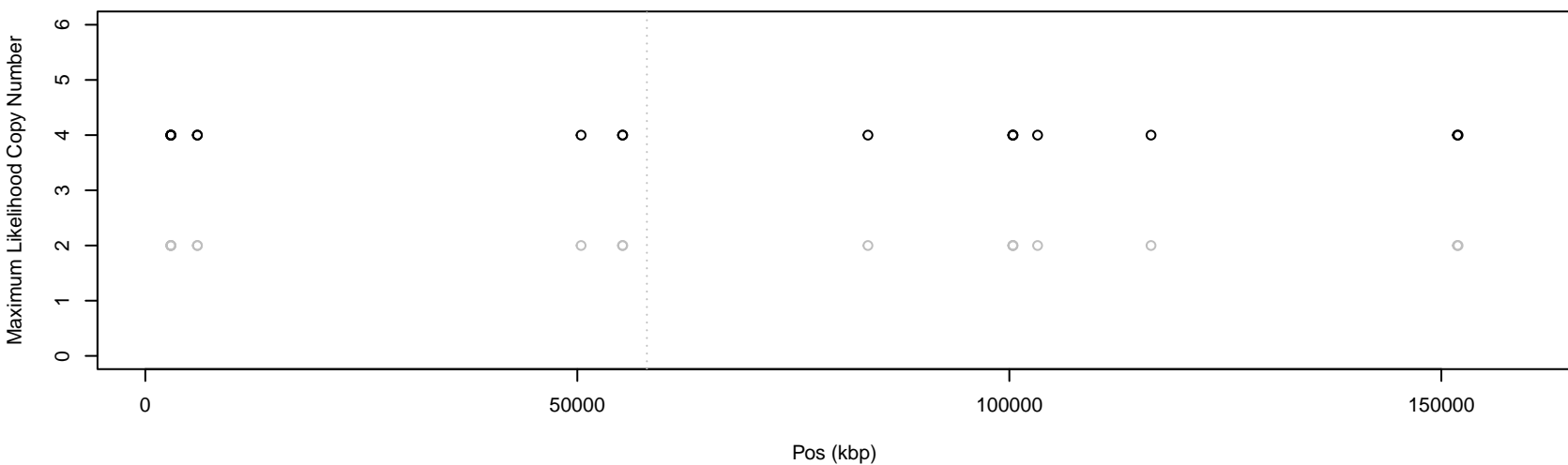
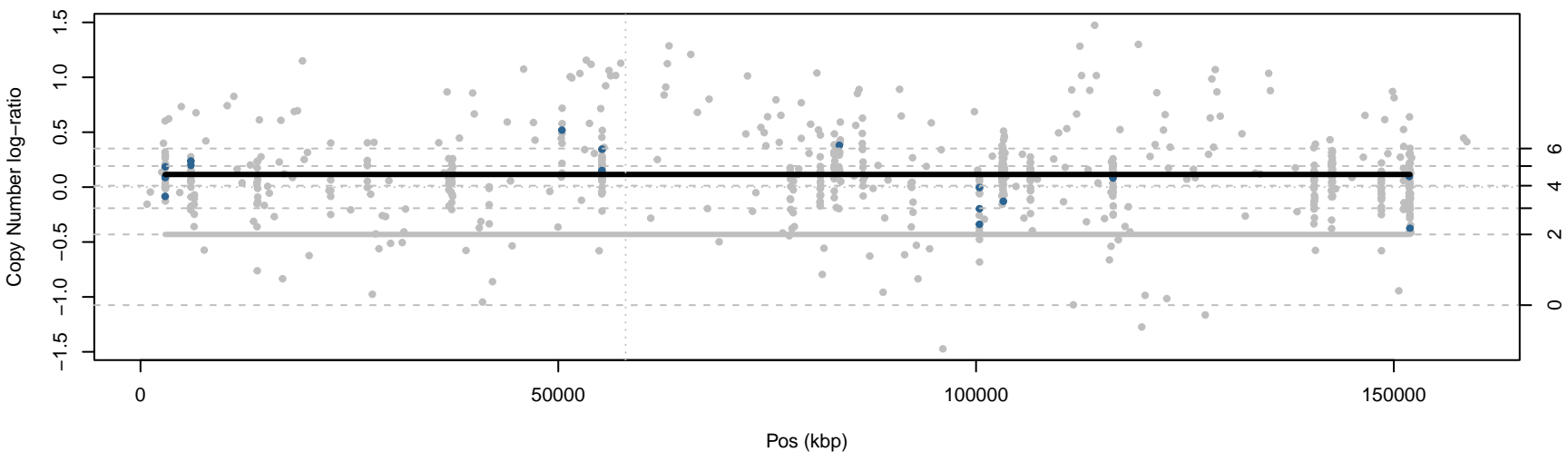
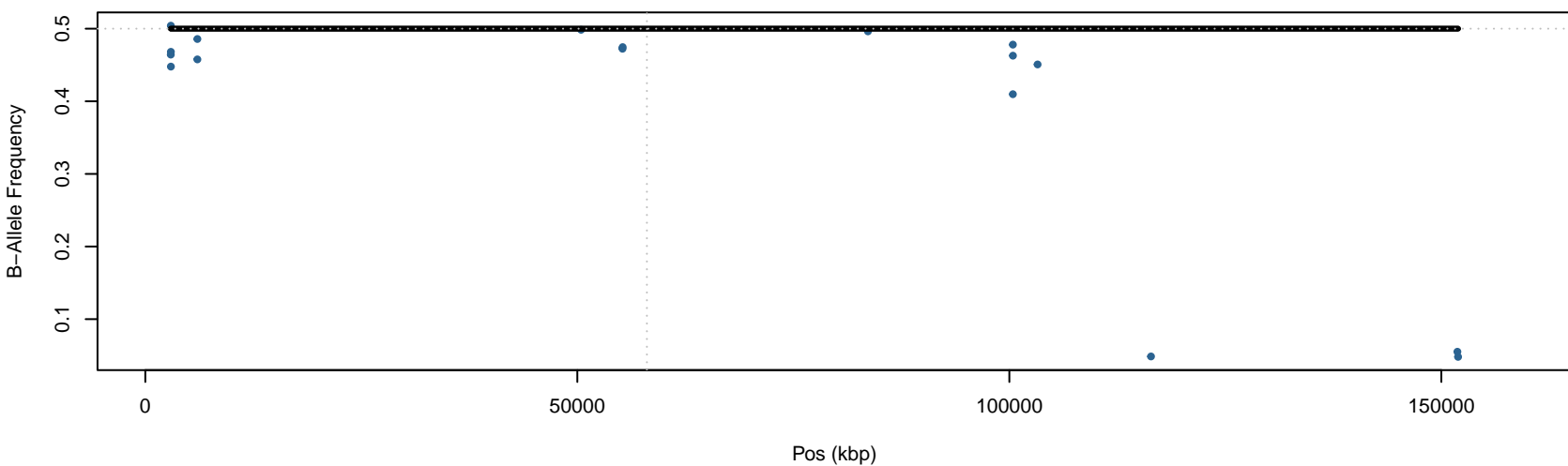
Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 4



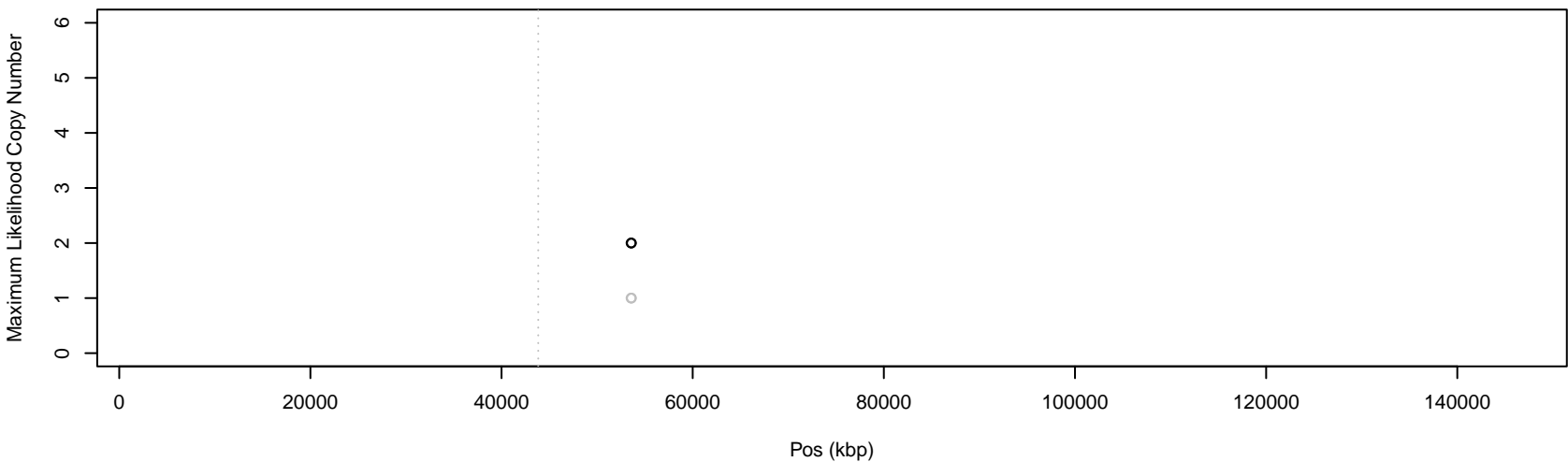
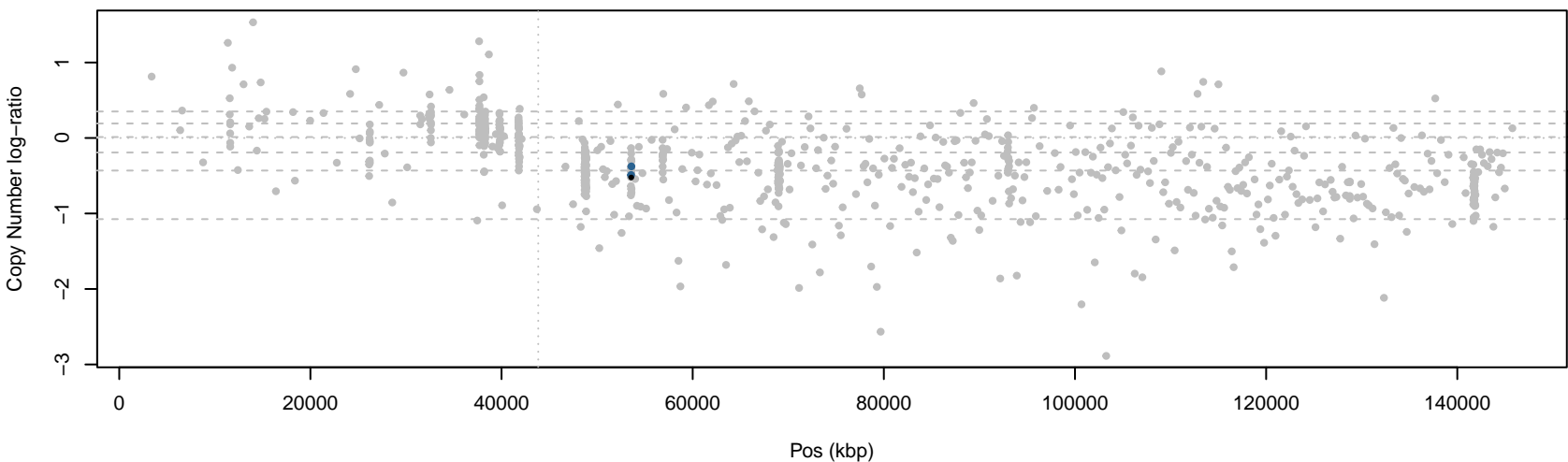
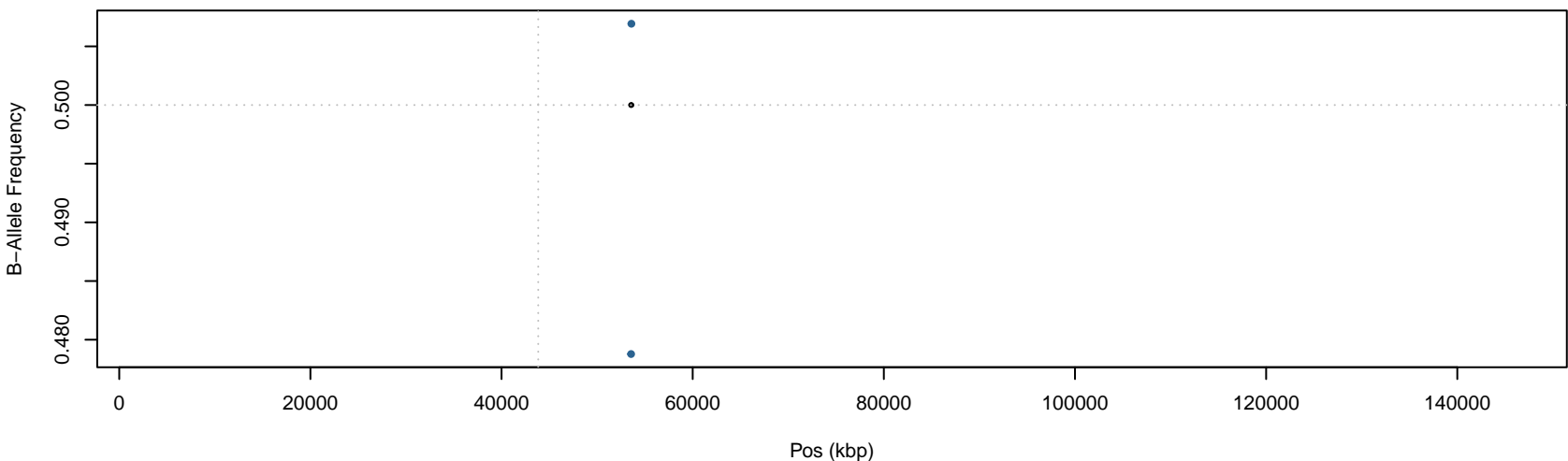
Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 6



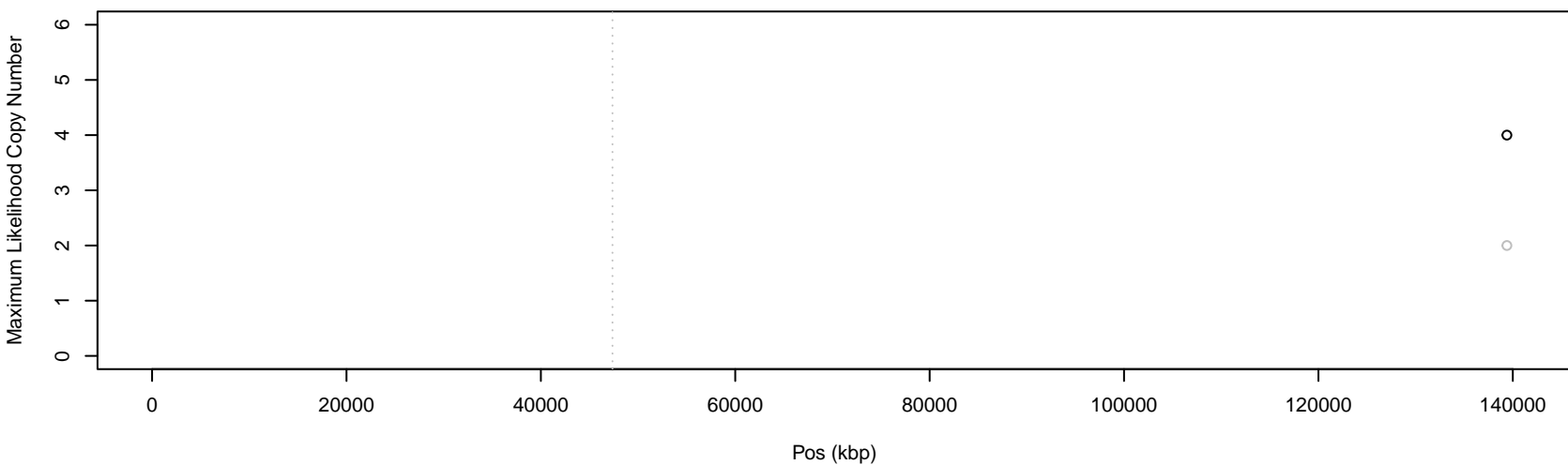
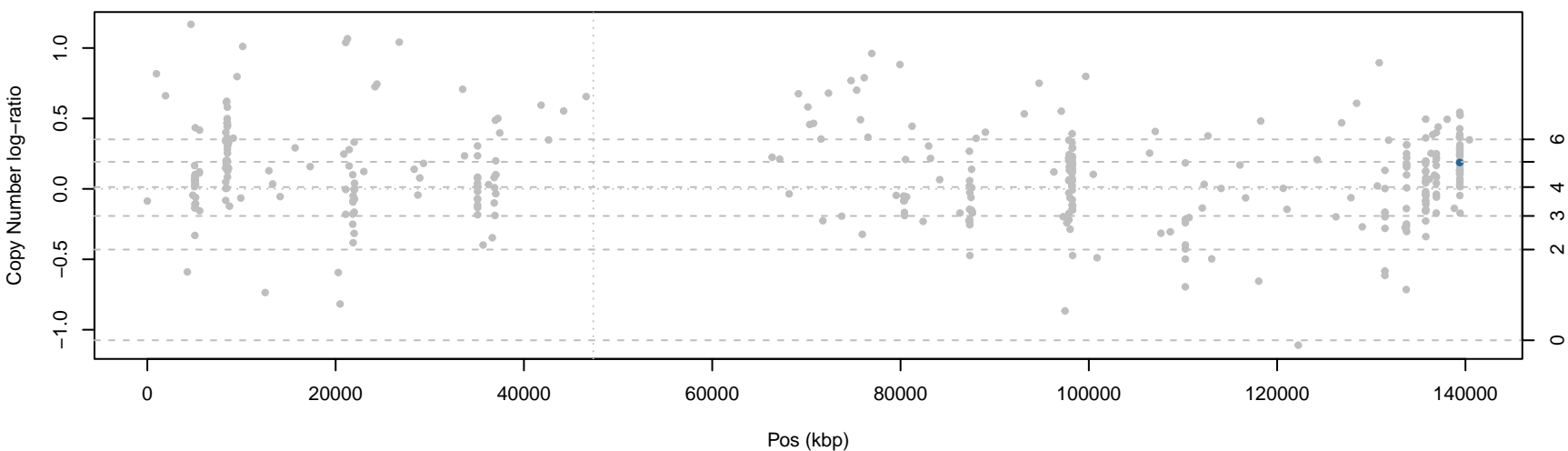
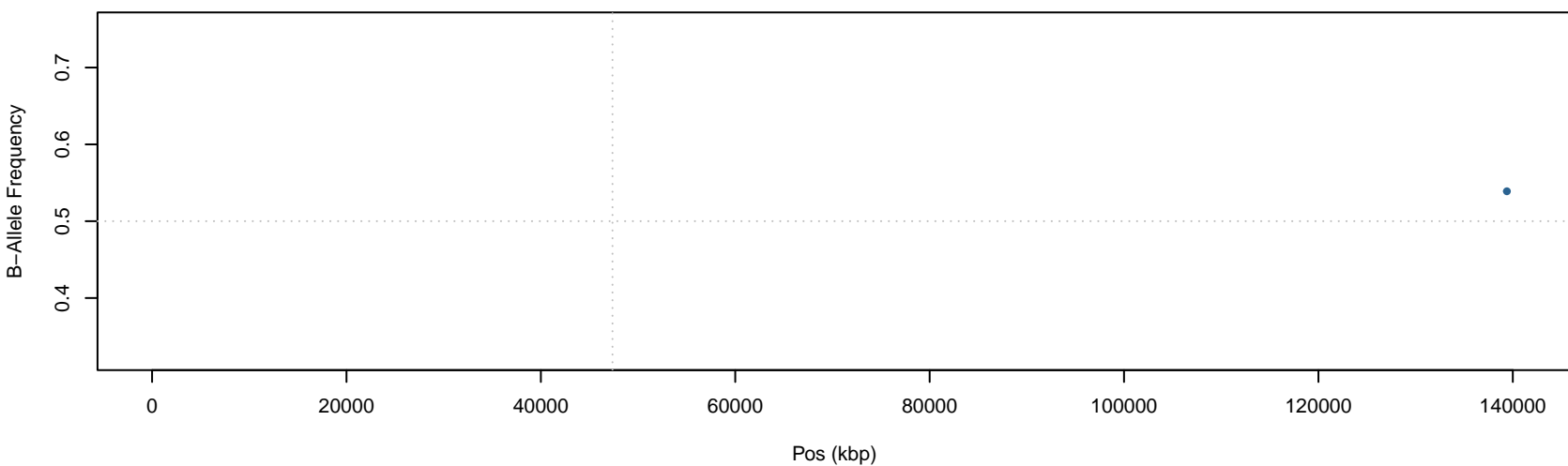
Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 7



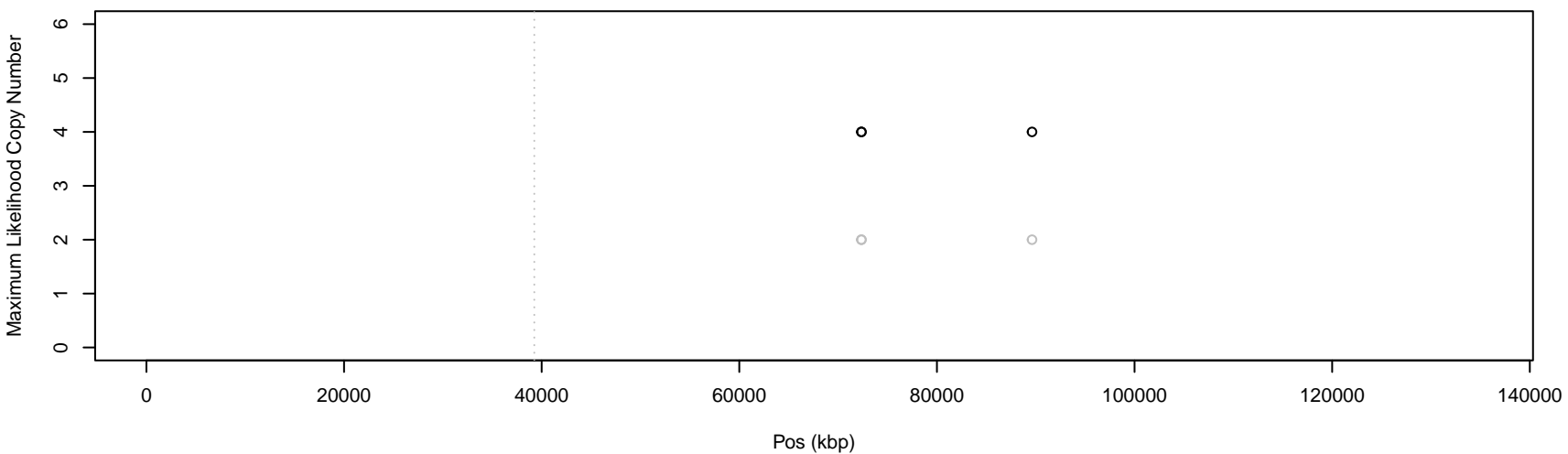
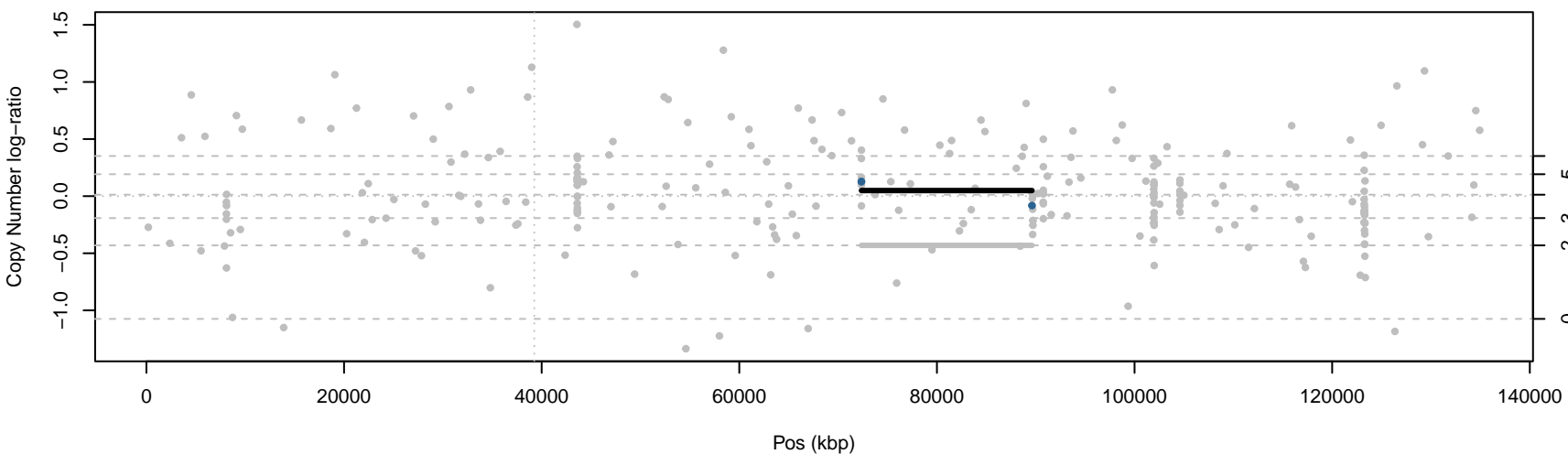
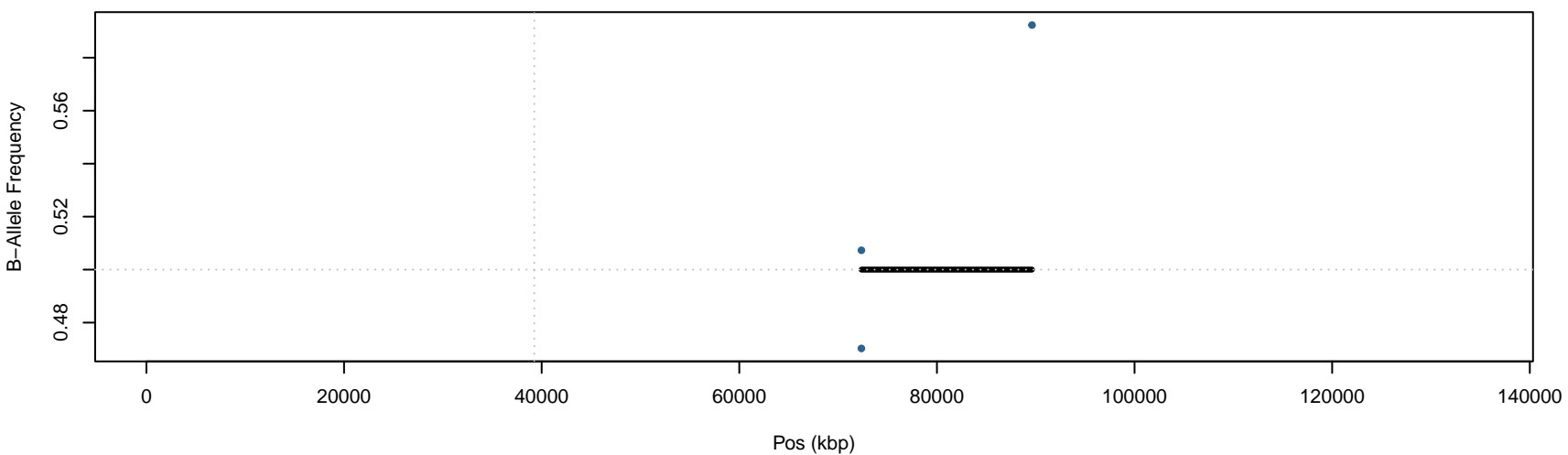
Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 8



Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 9

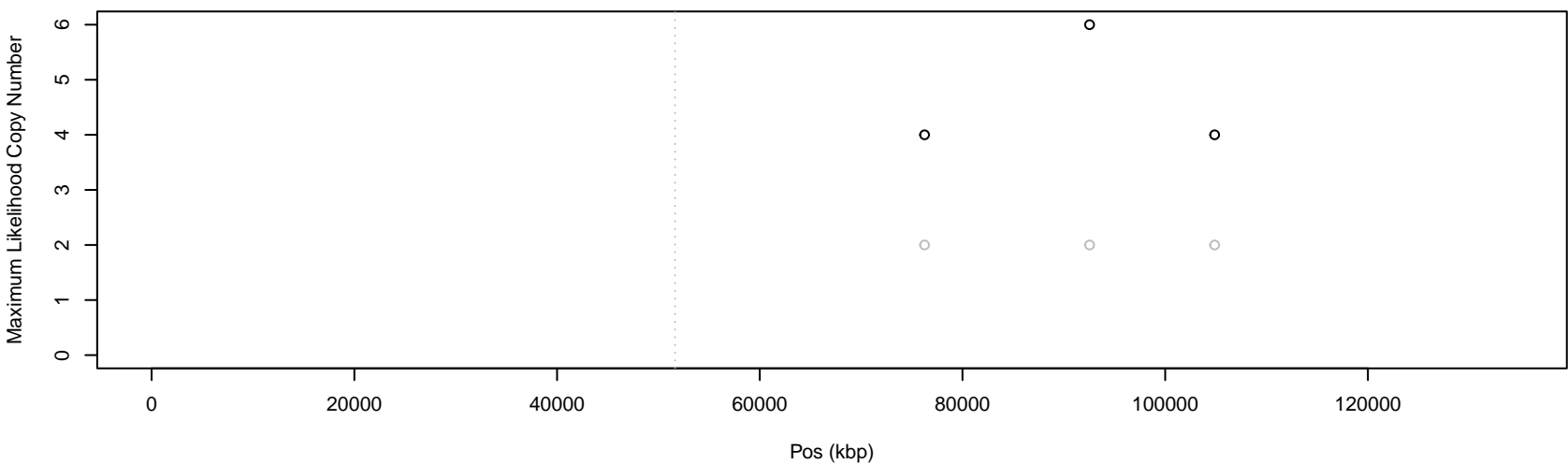
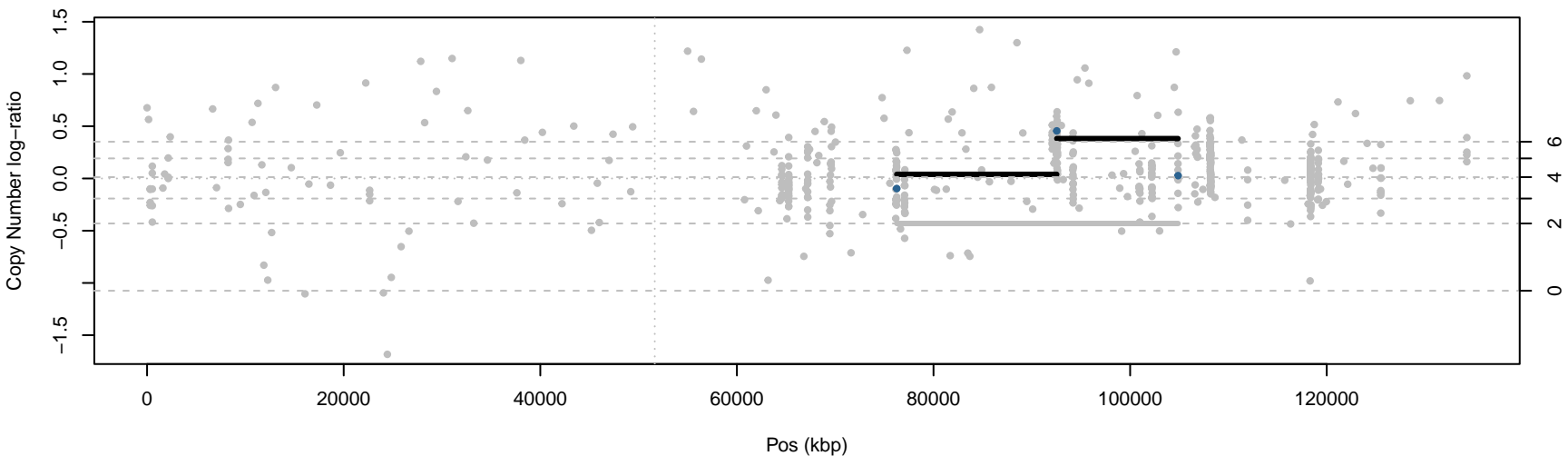
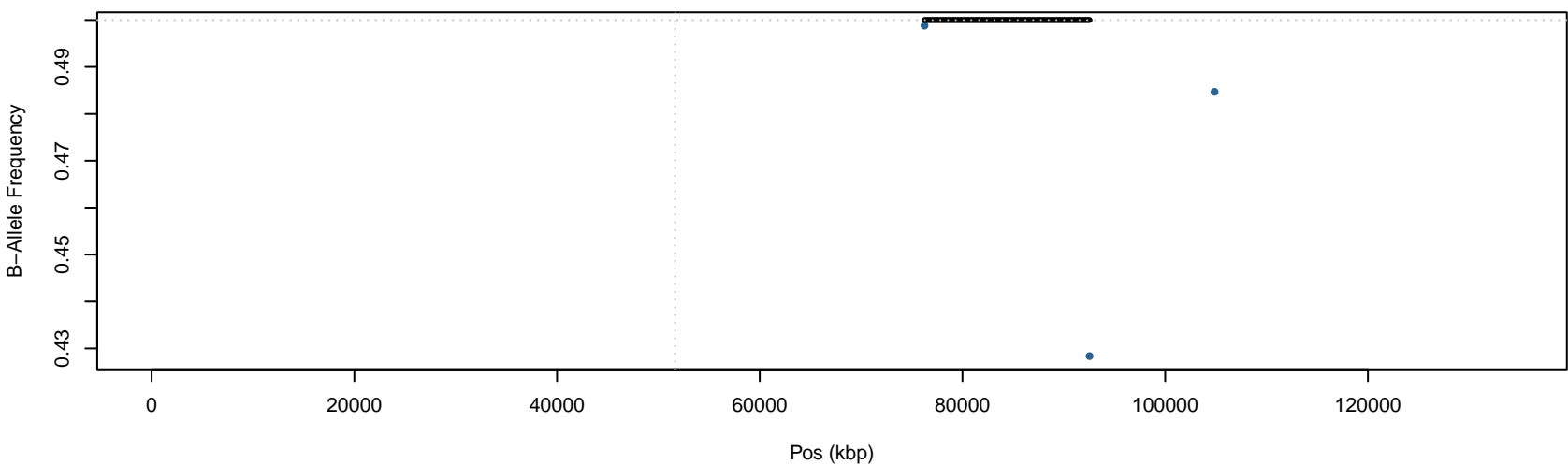


Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 10

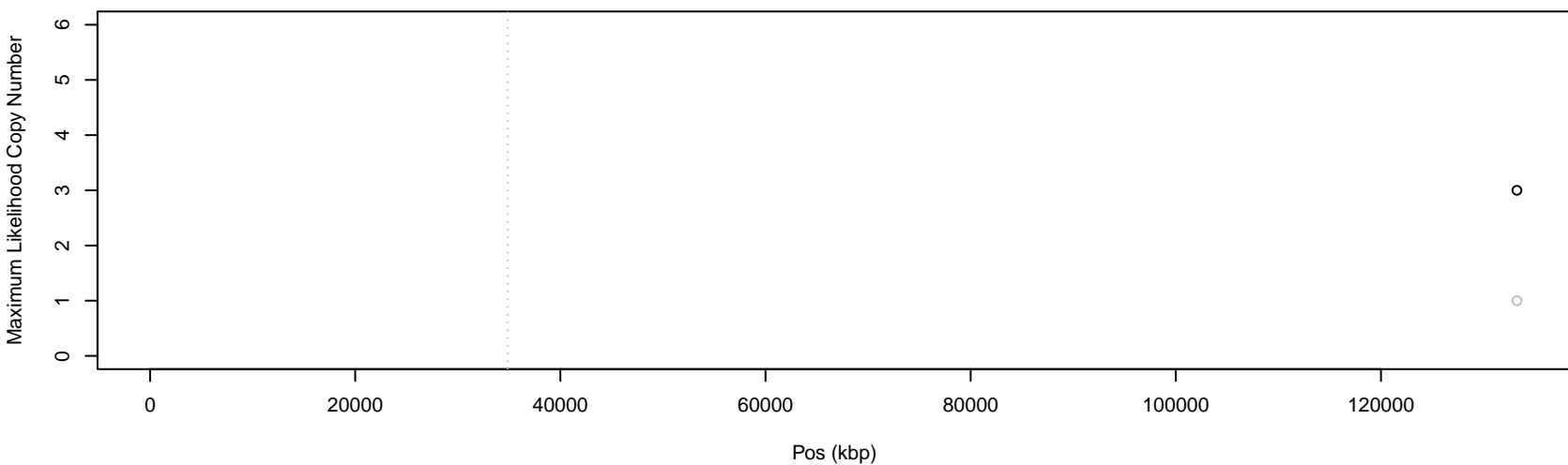
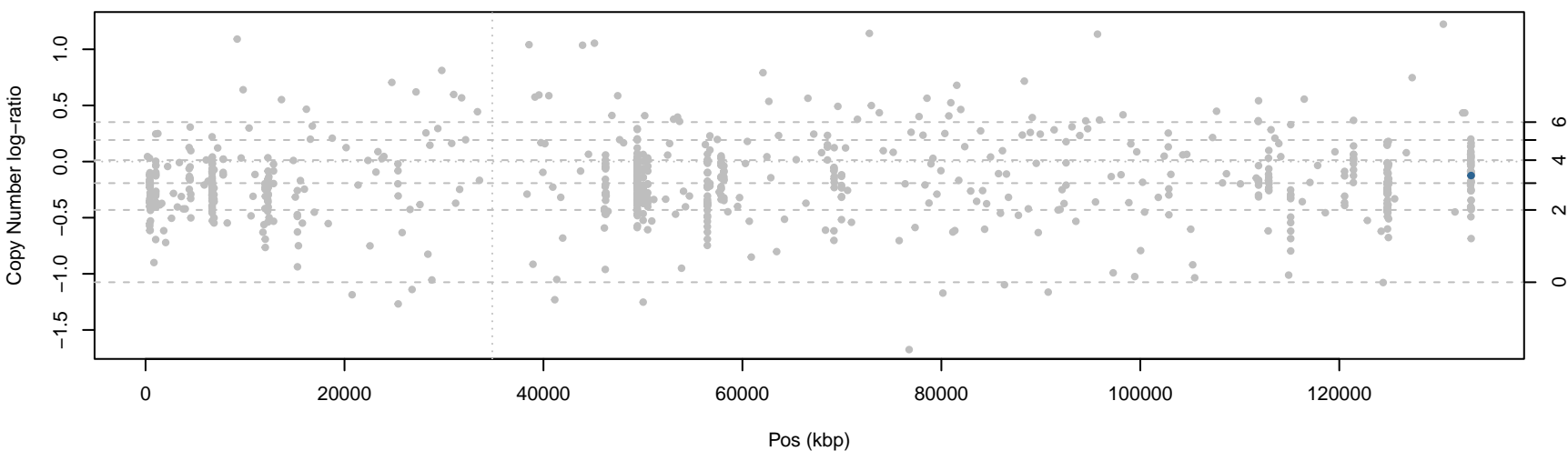
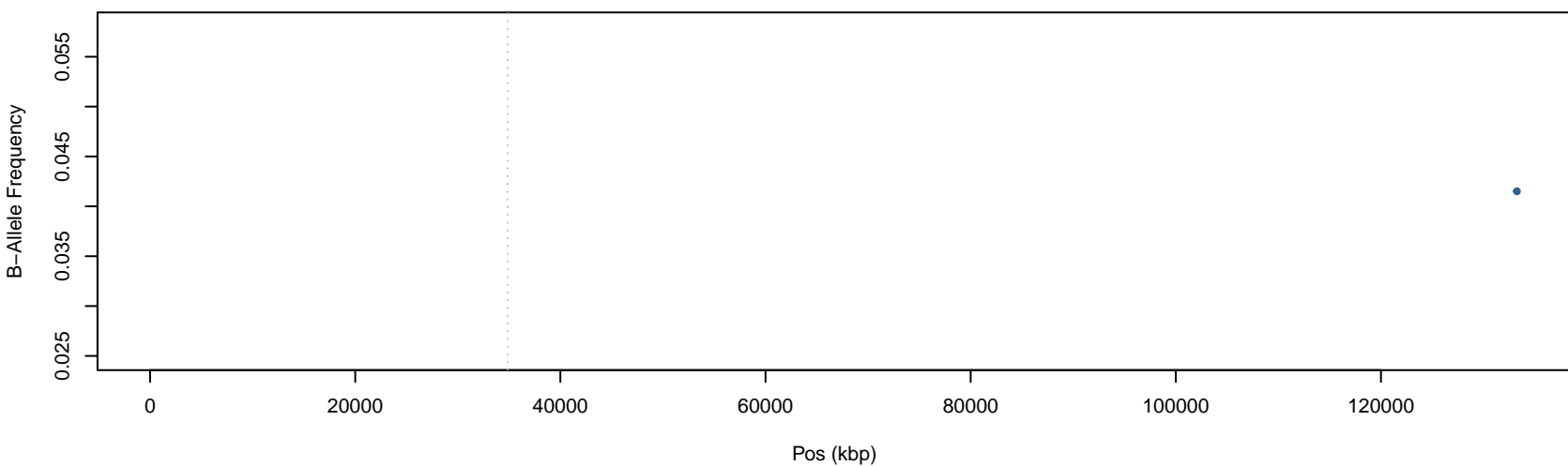




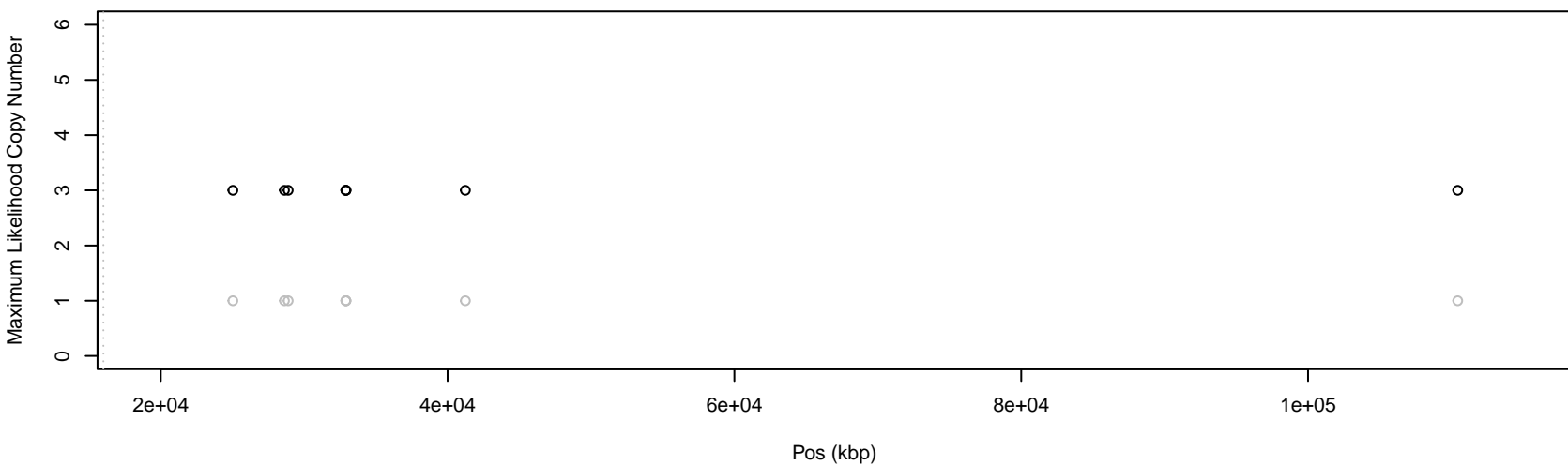
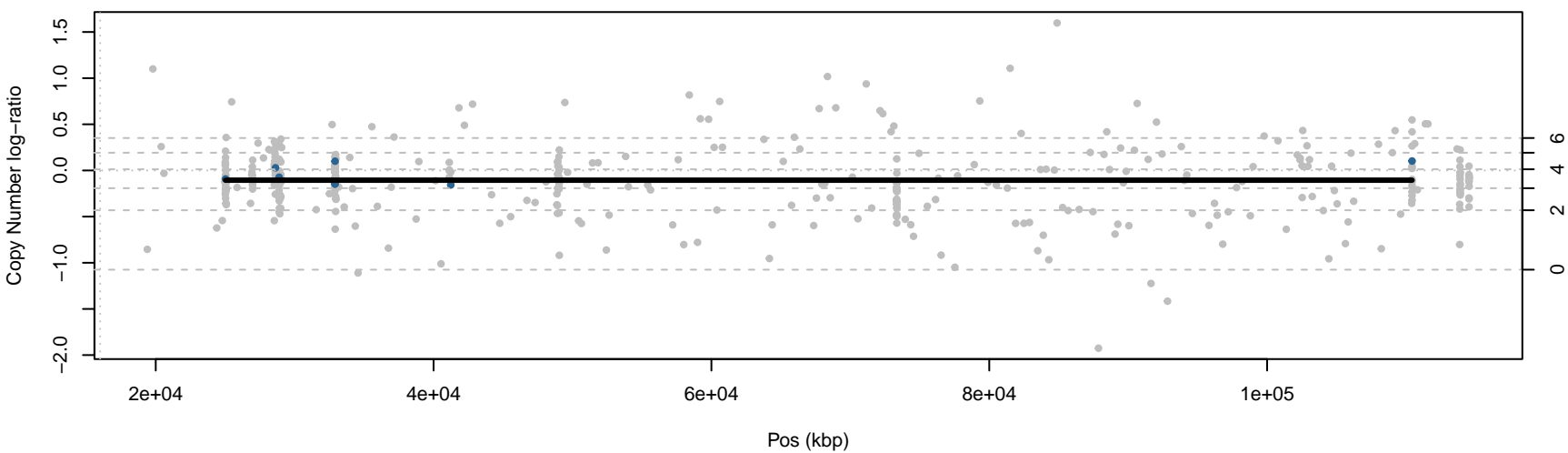
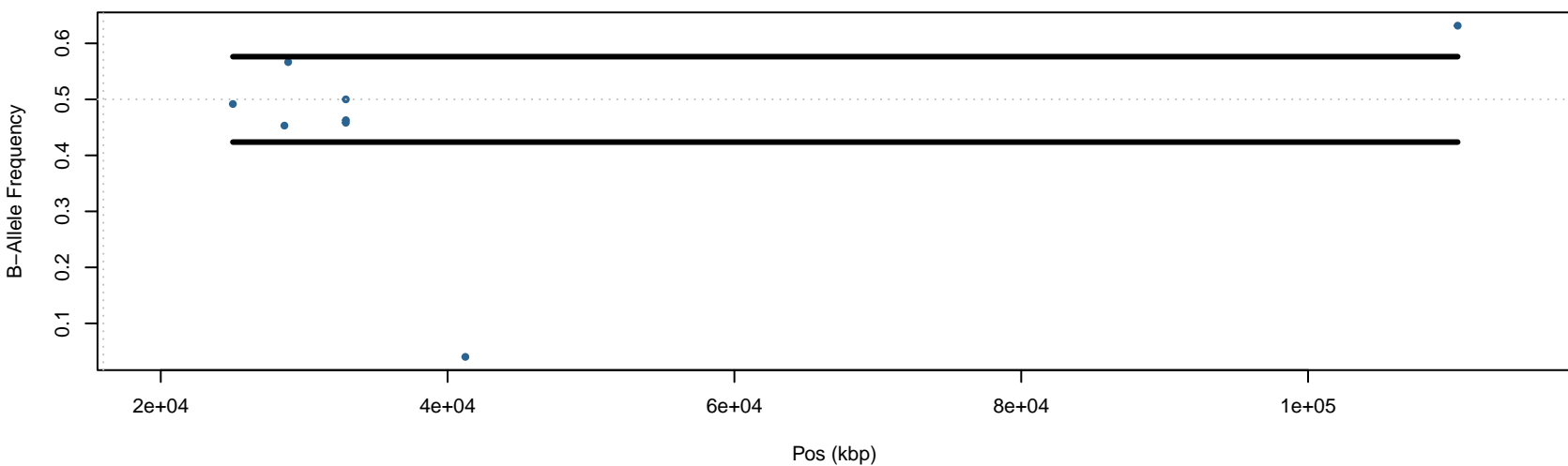
Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 11



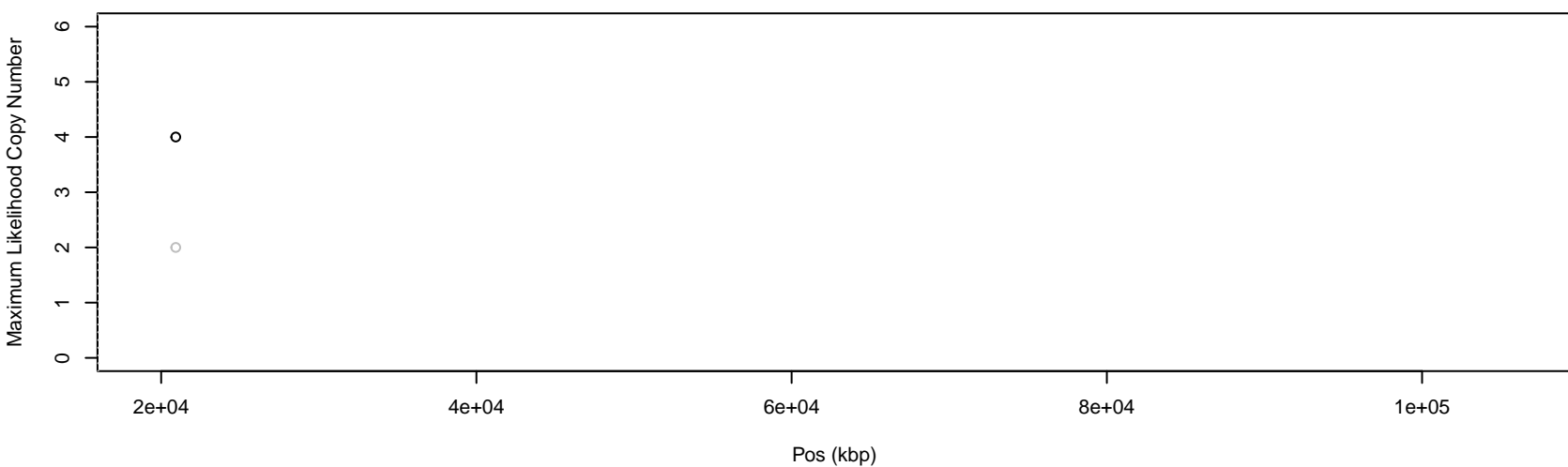
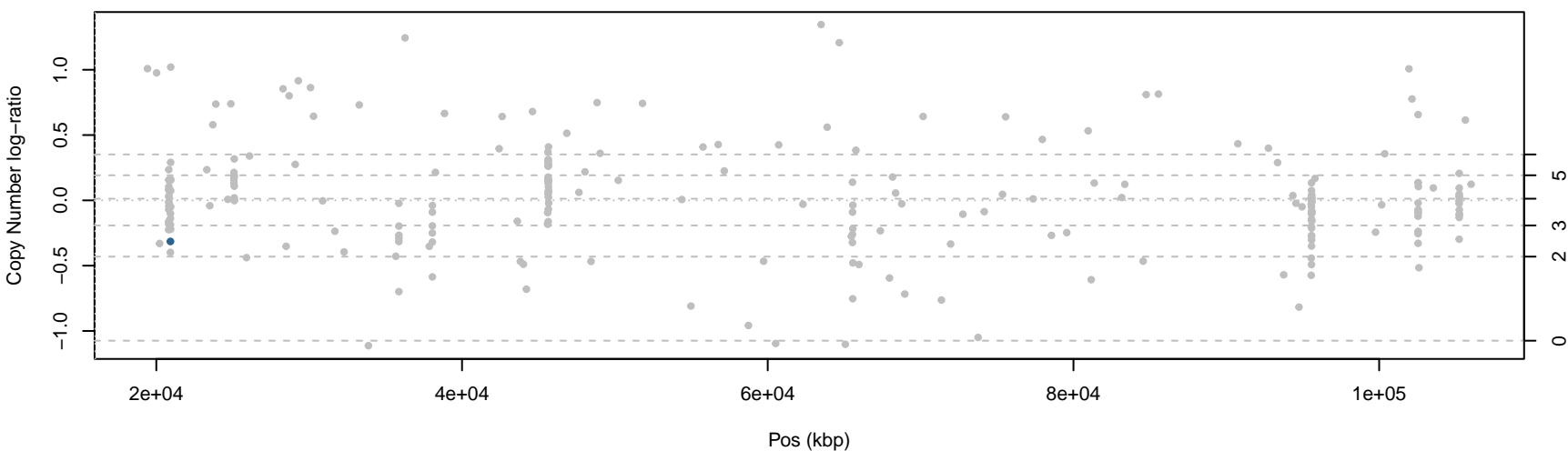
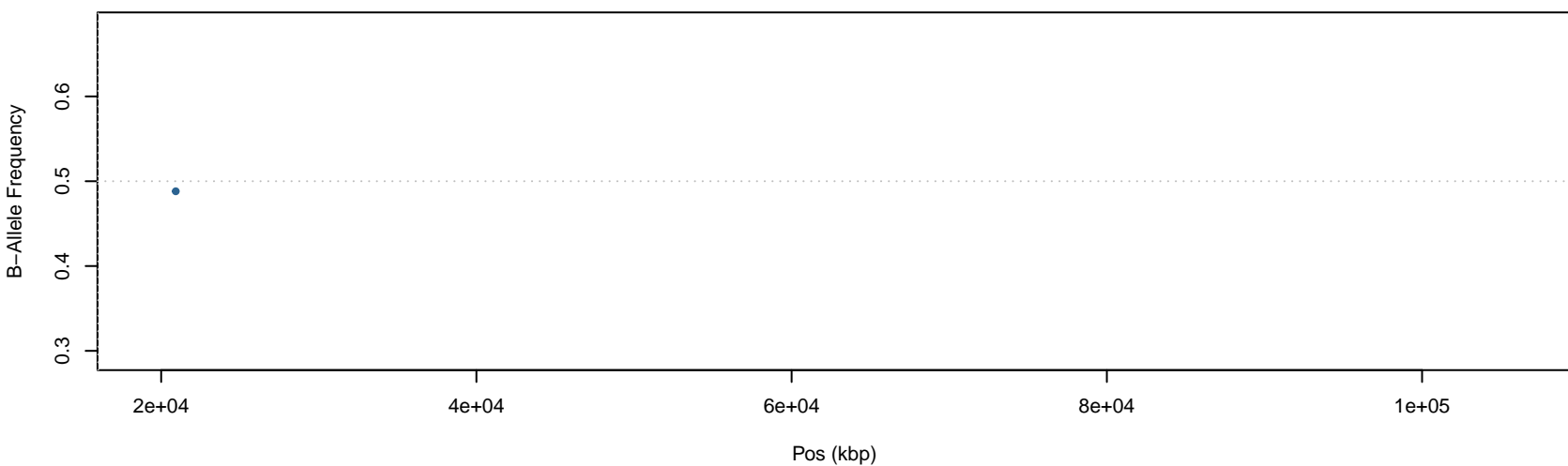
Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 12



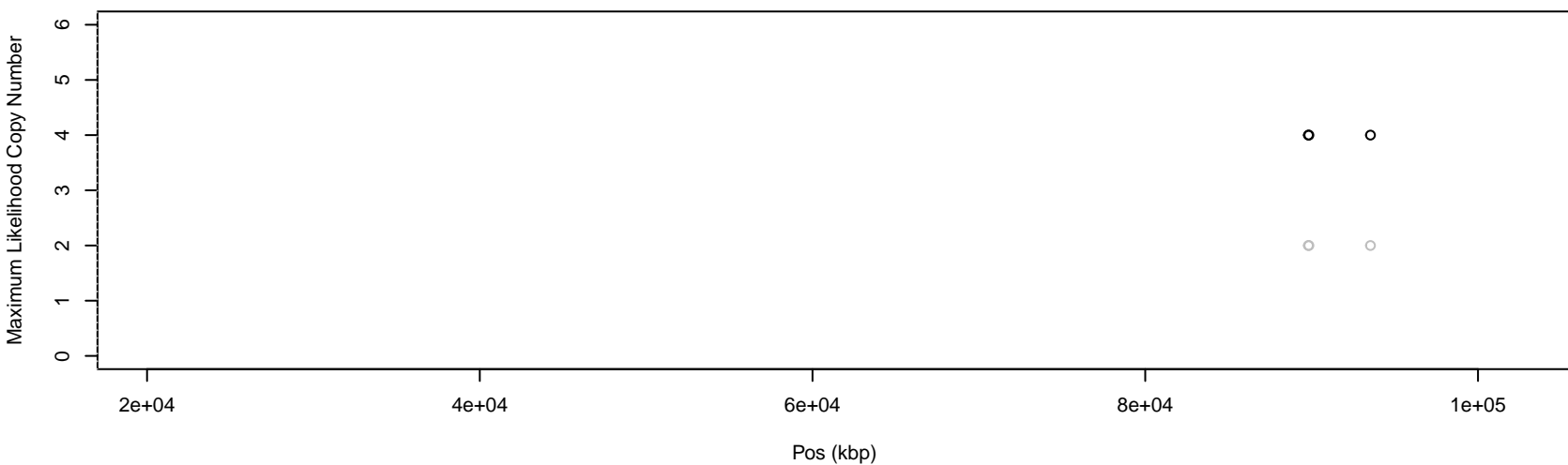
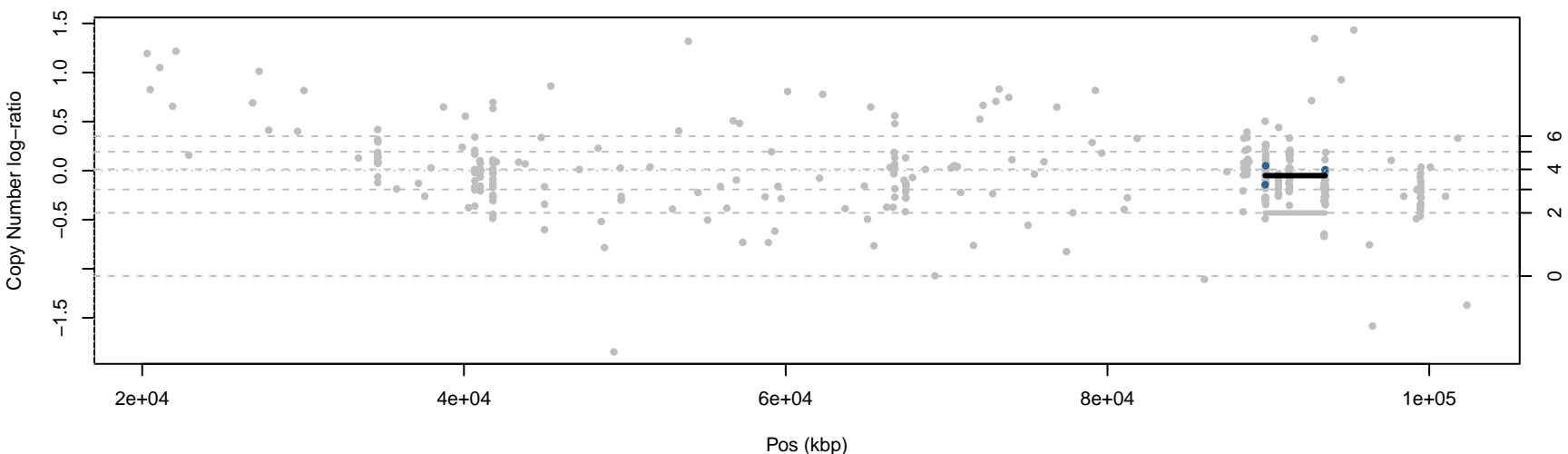
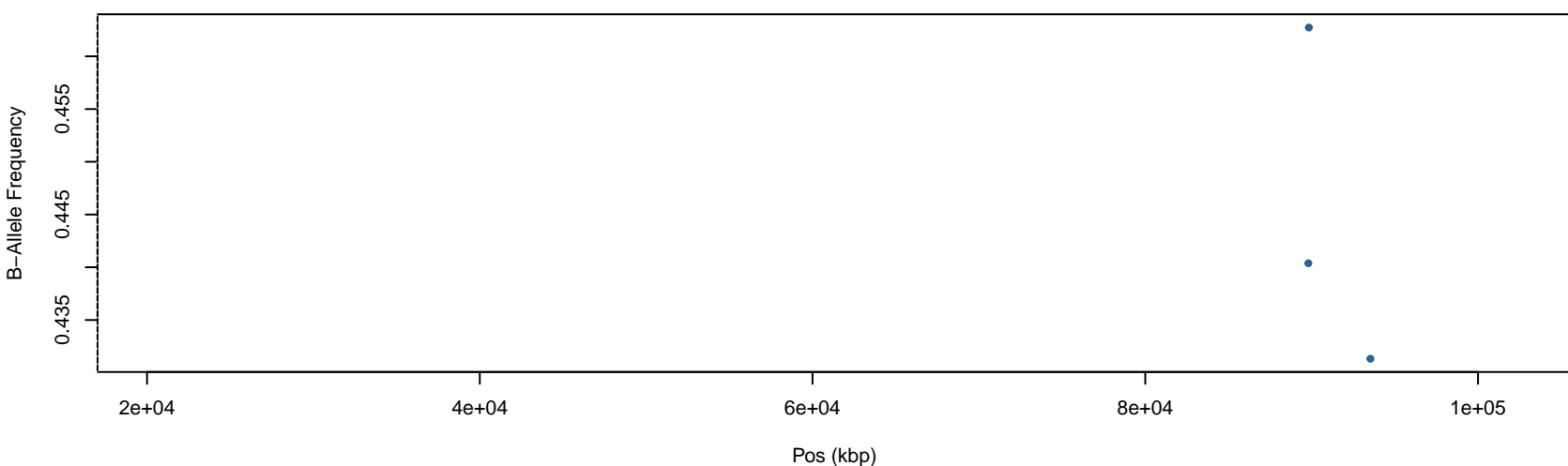
Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 13



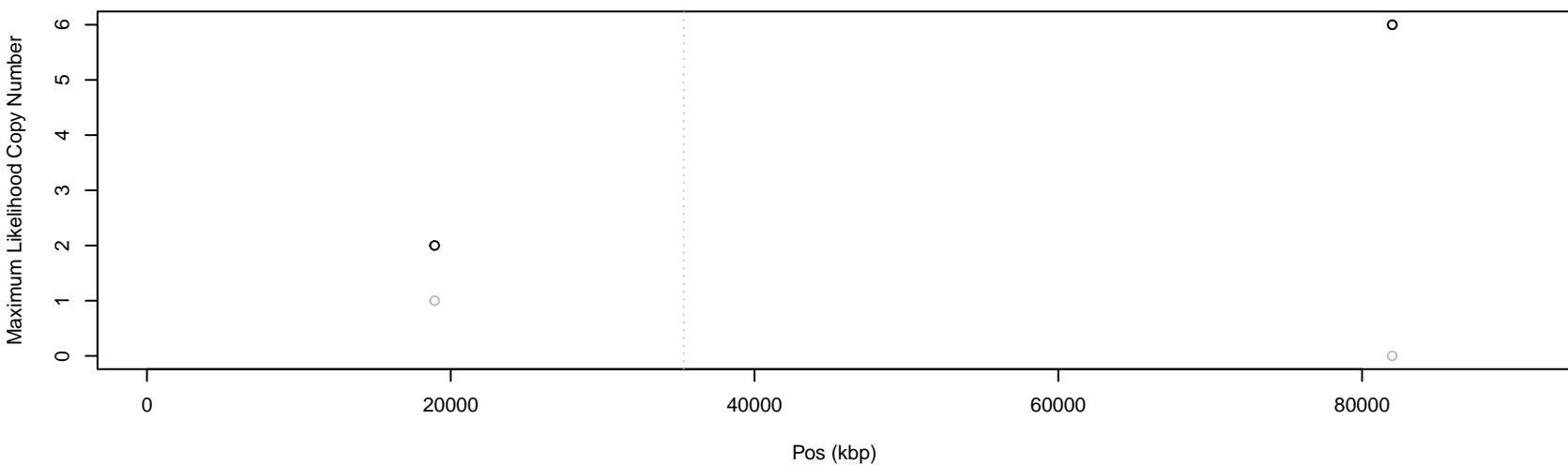
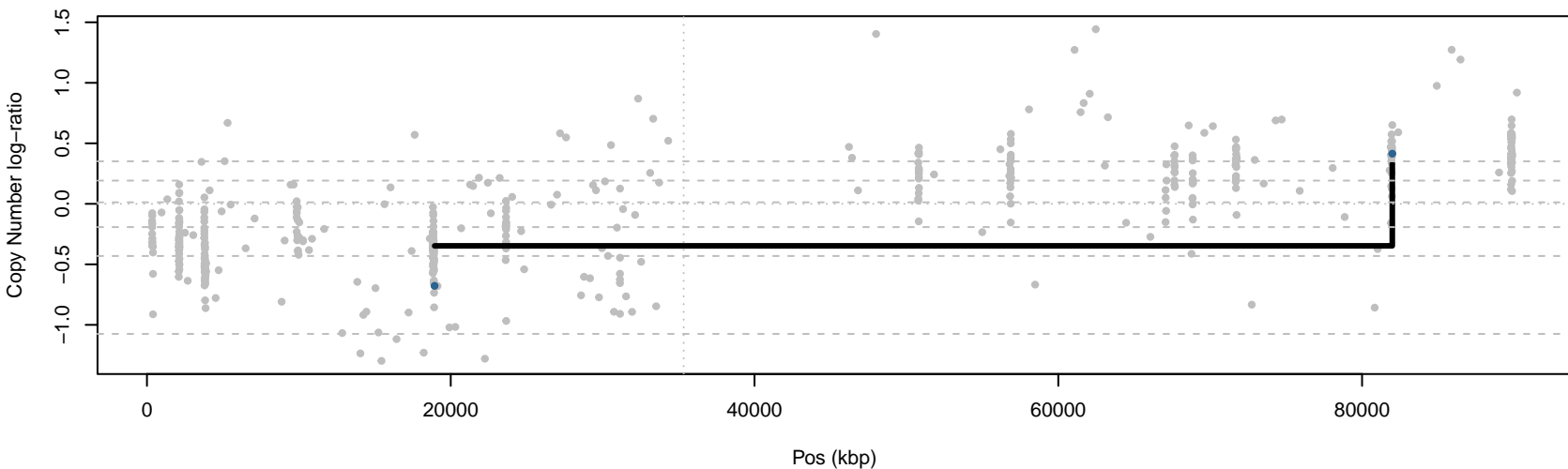
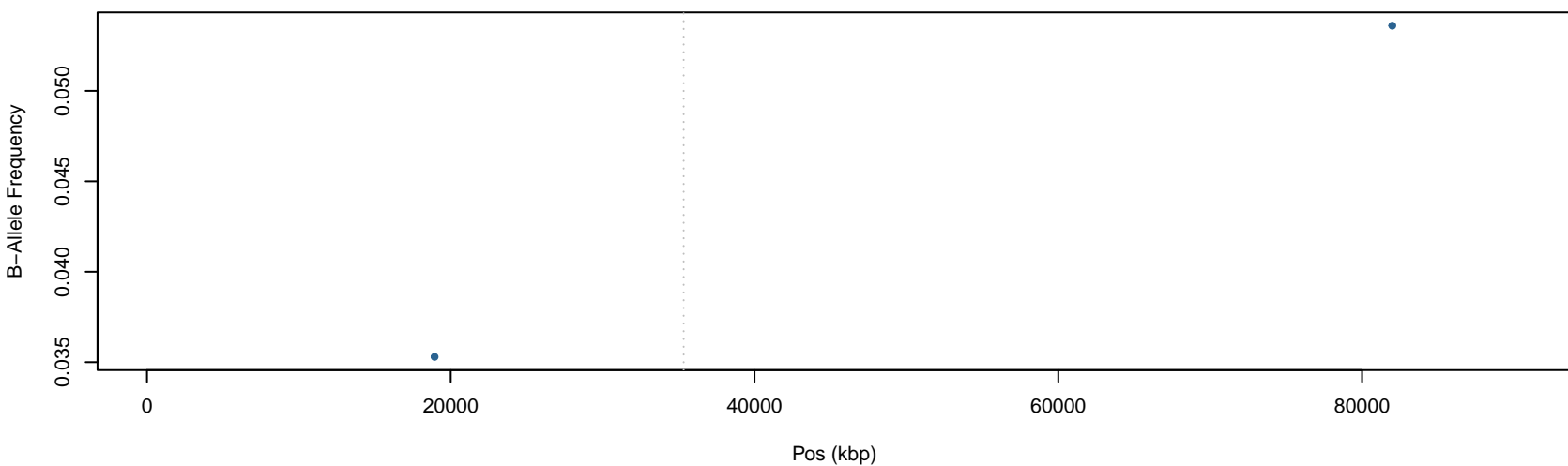
Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 14



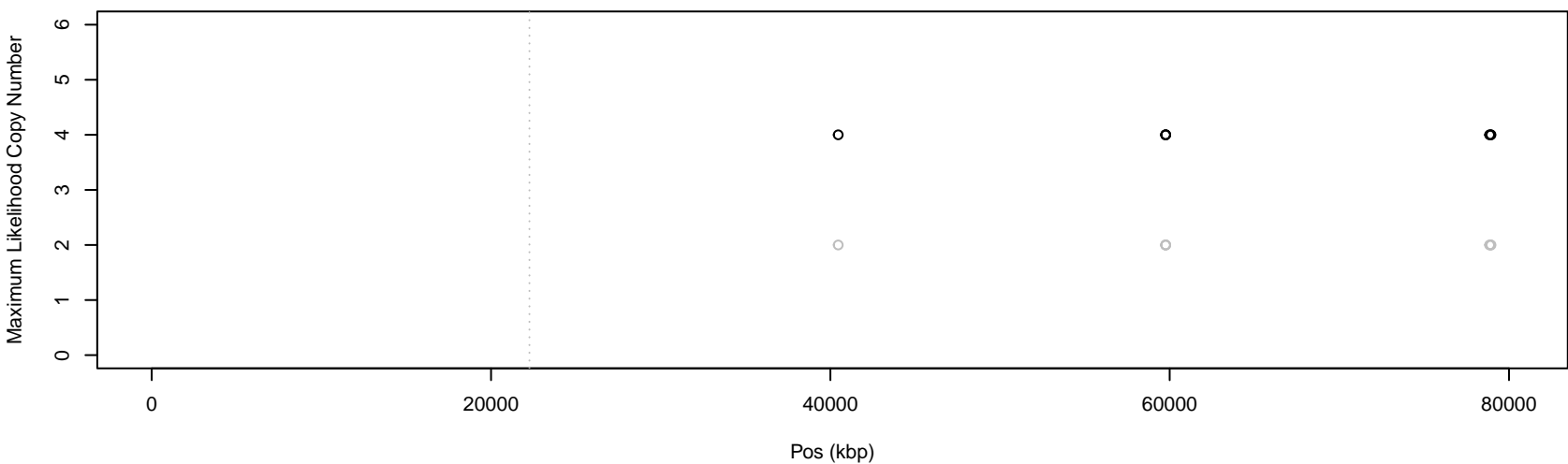
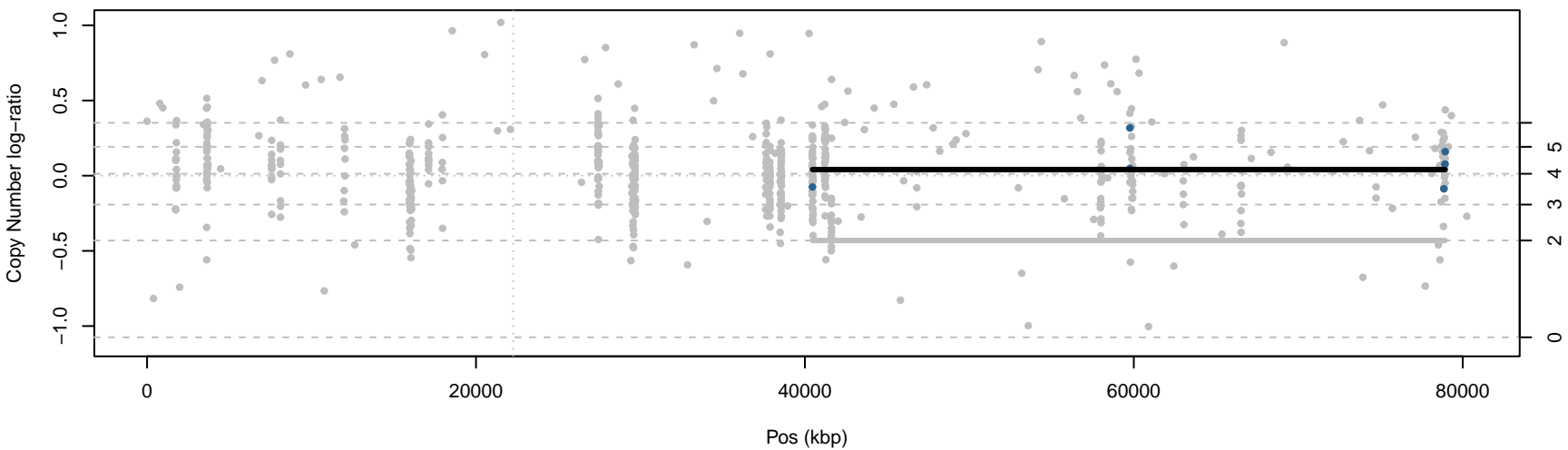
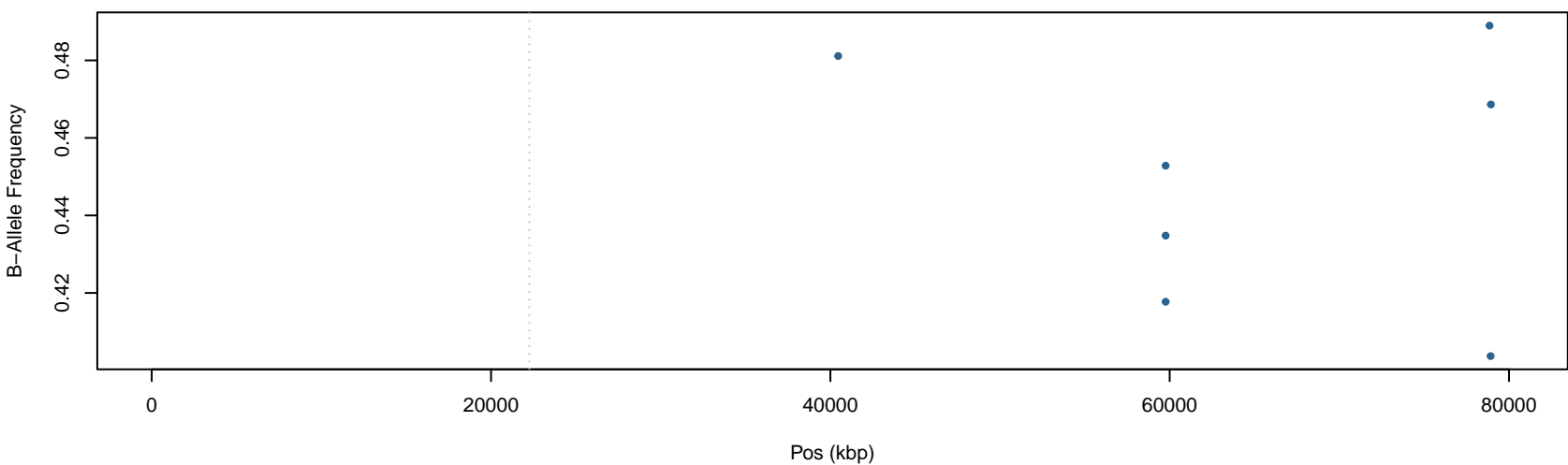
Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 15



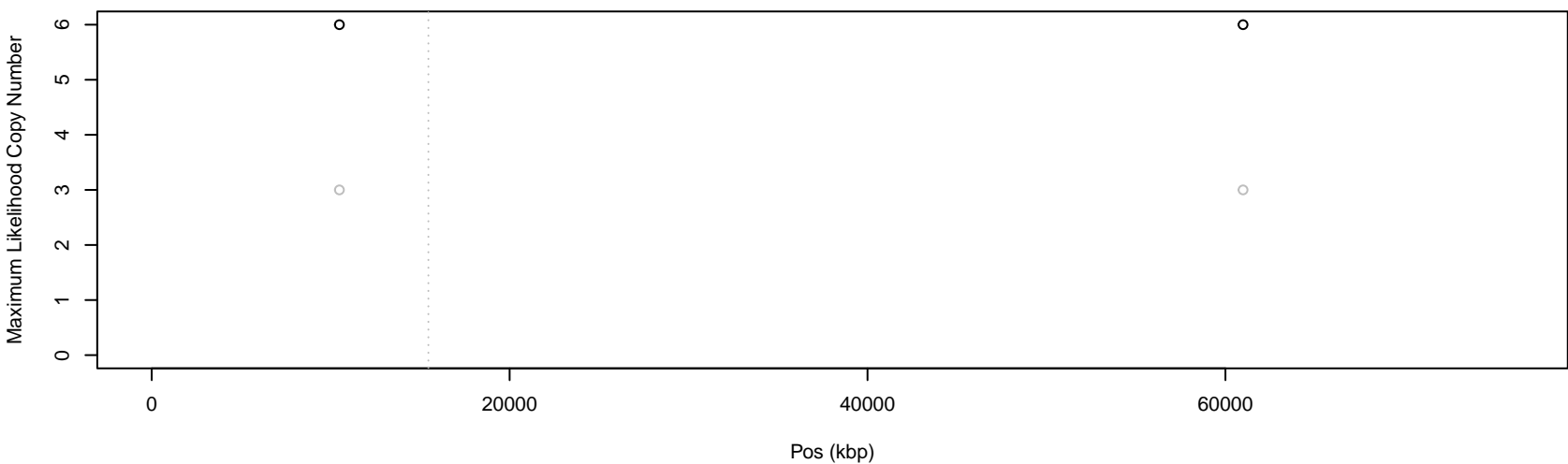
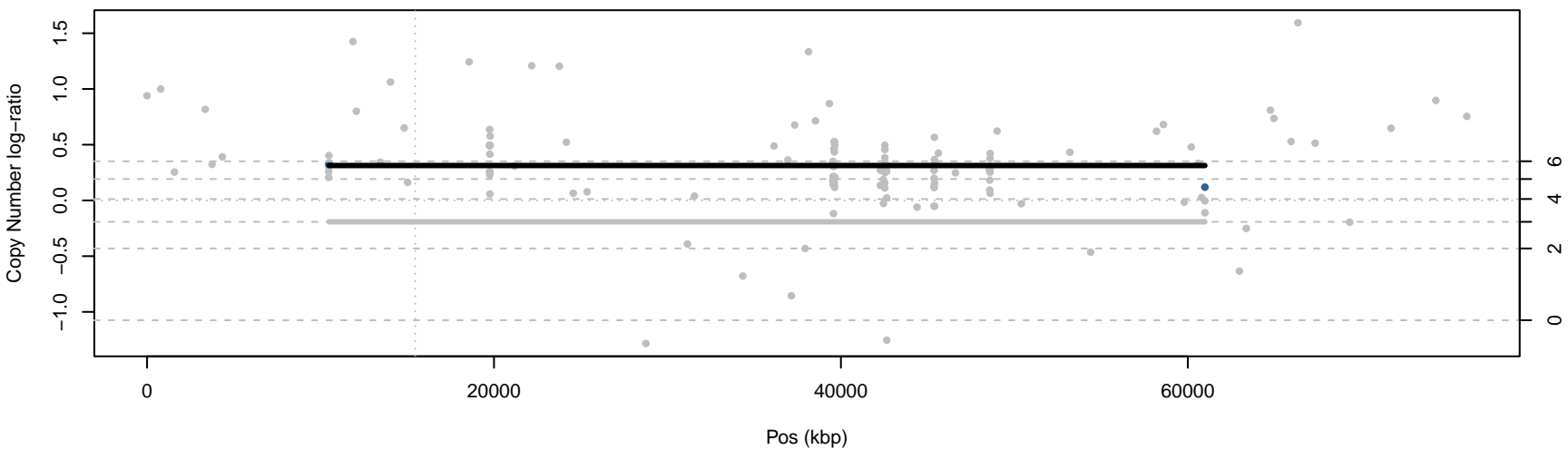
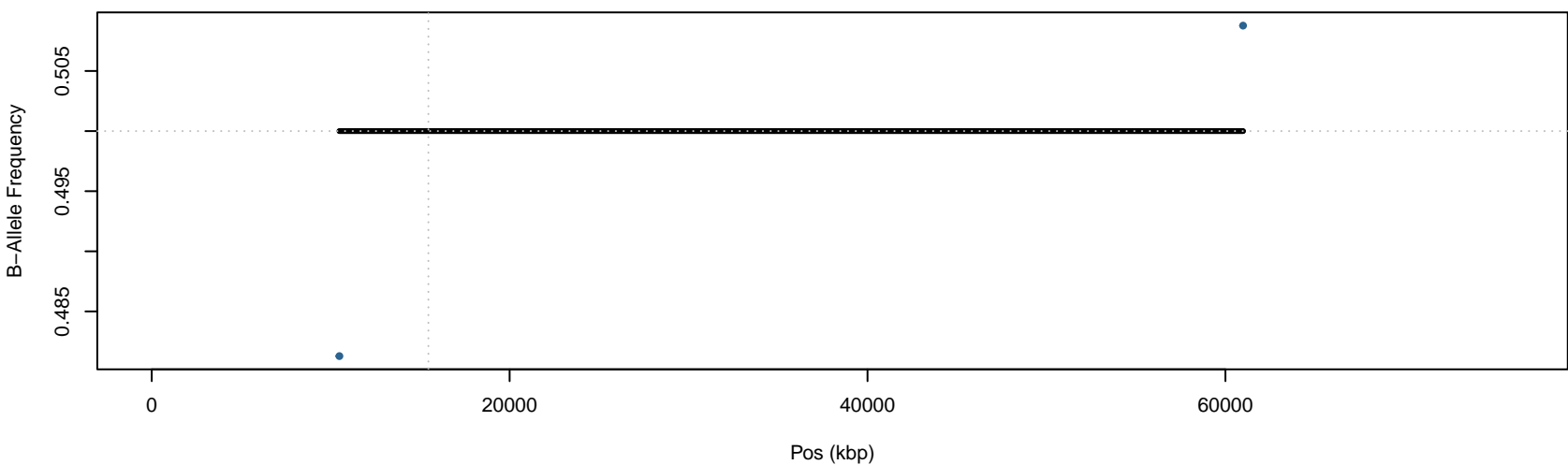
Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 16



Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 17

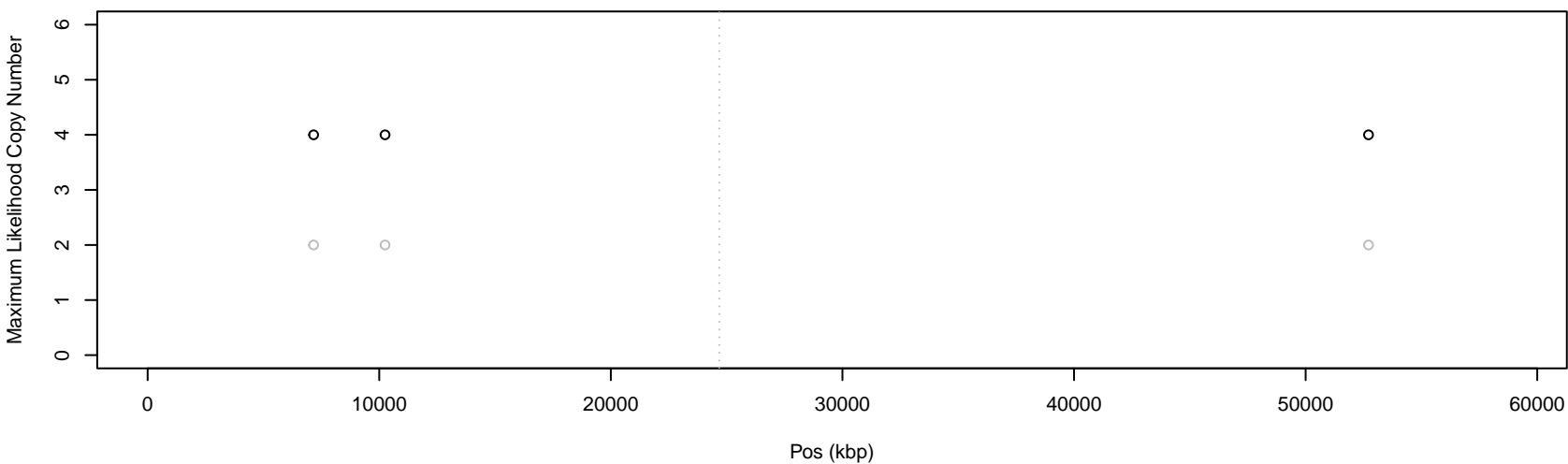
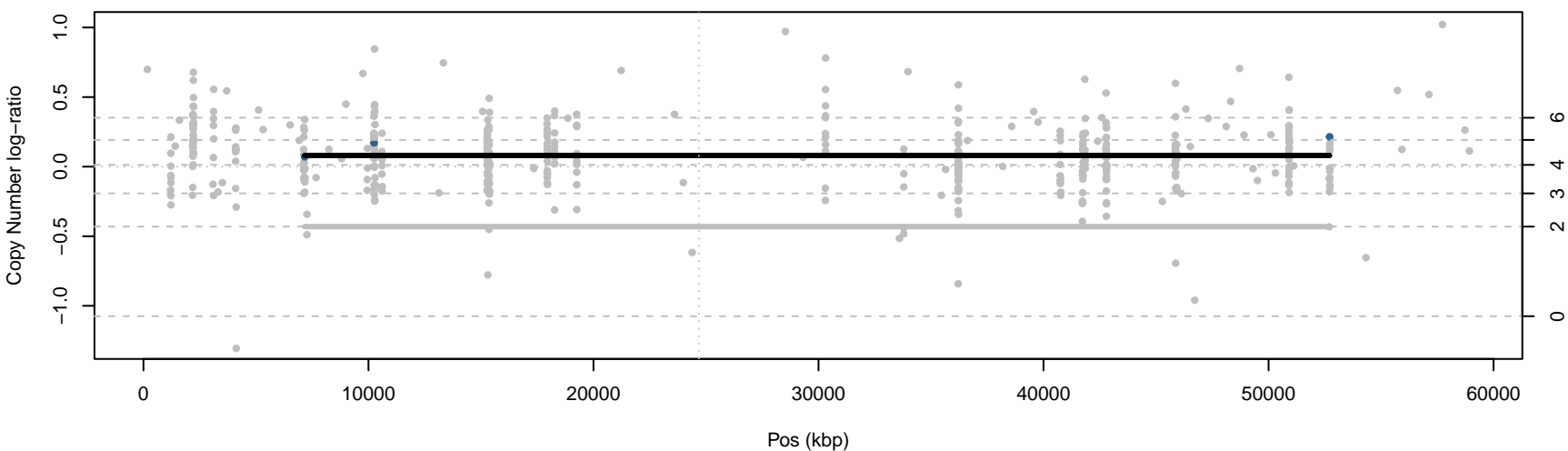
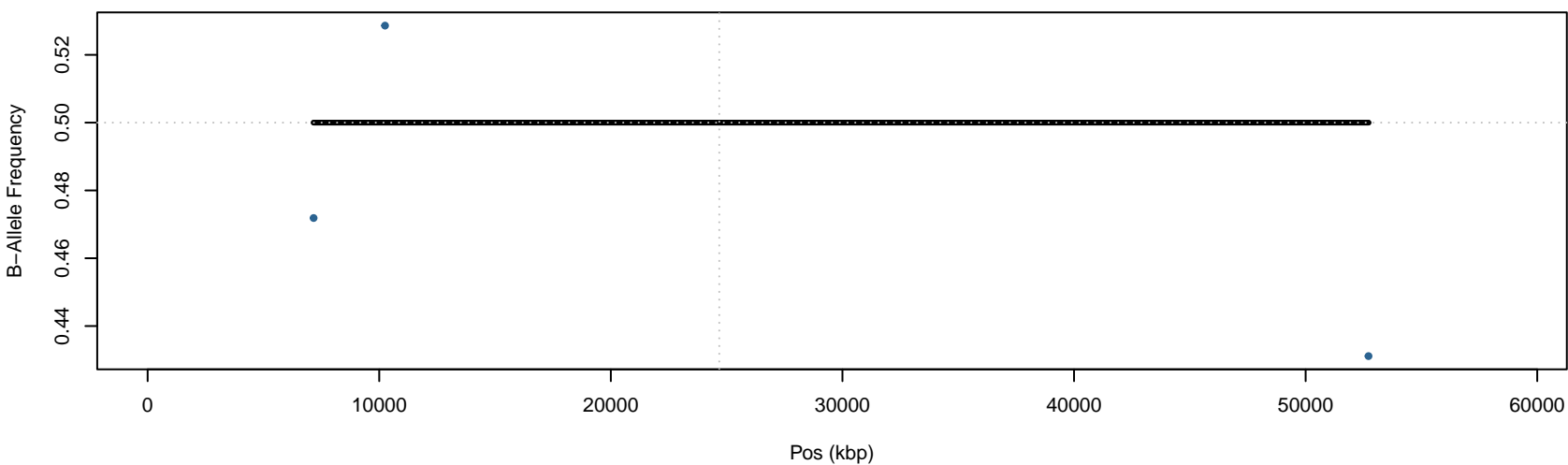


Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 18

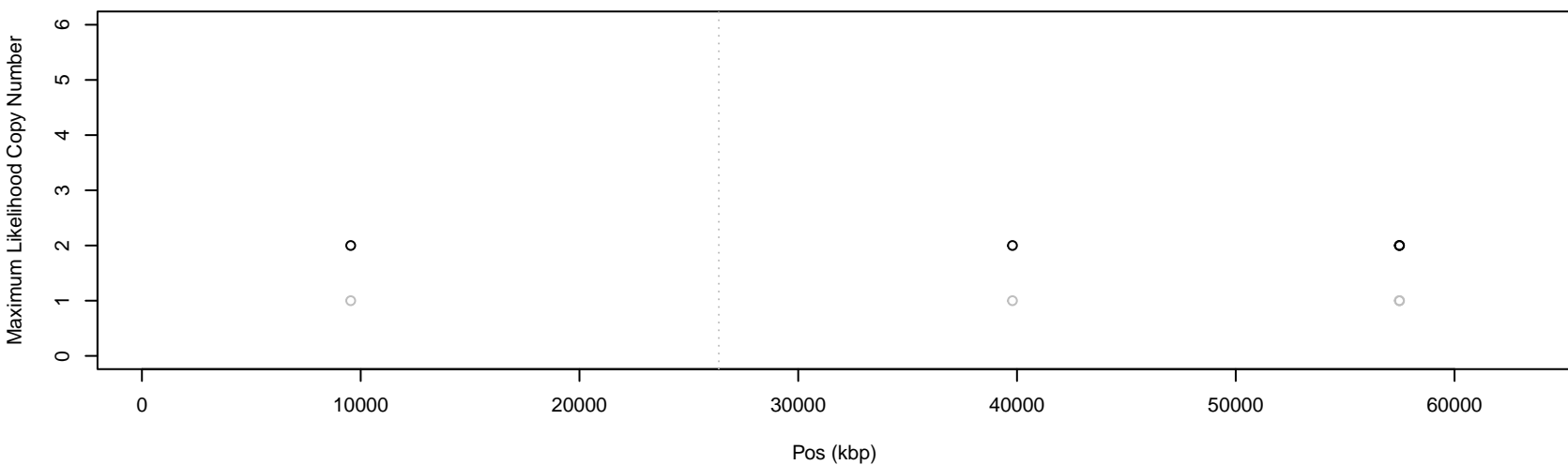
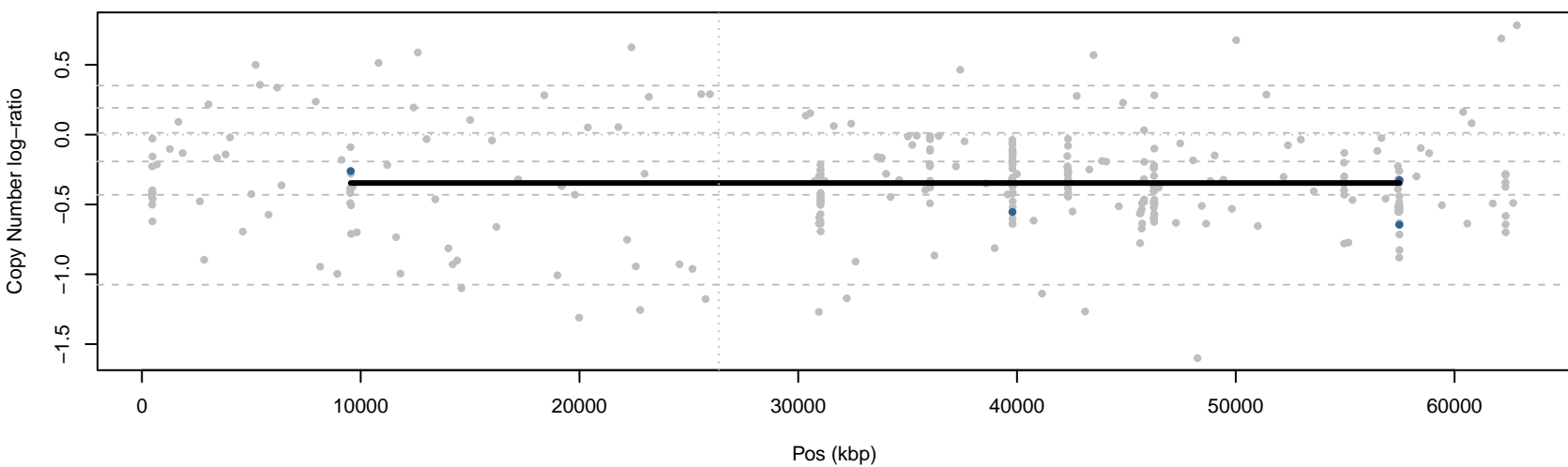
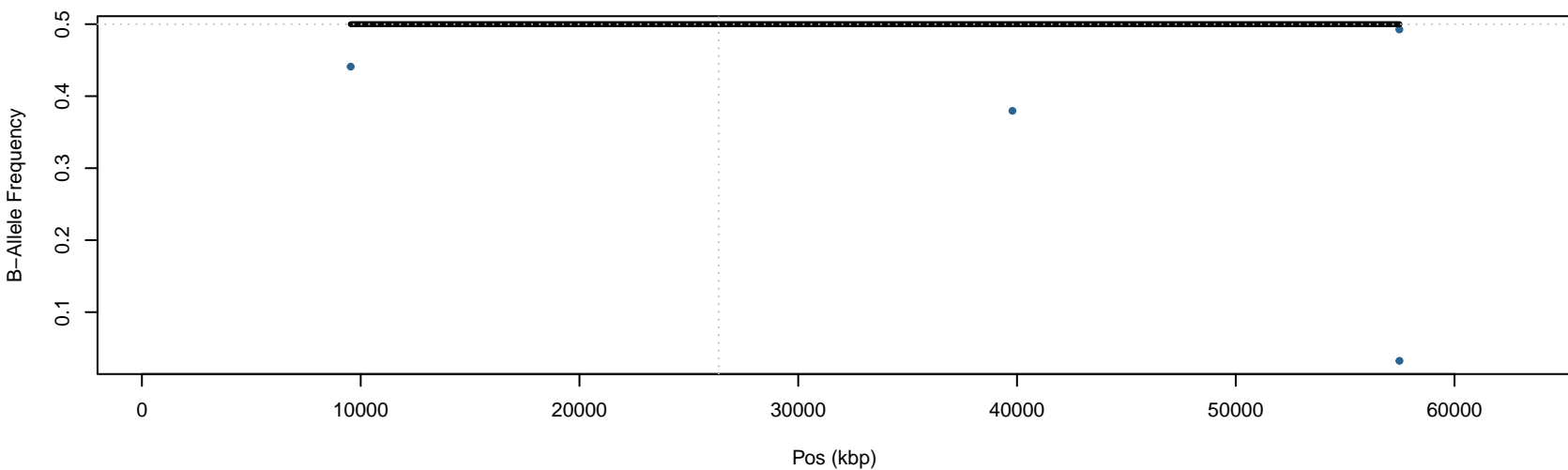




Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 19



Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 20



Purity: 0.36 Tumor ploidy: 3.934 SNV log-likelihood: -27.59 GoF: 90.2% Mean coverage: 454;525 Chromosome: 22

