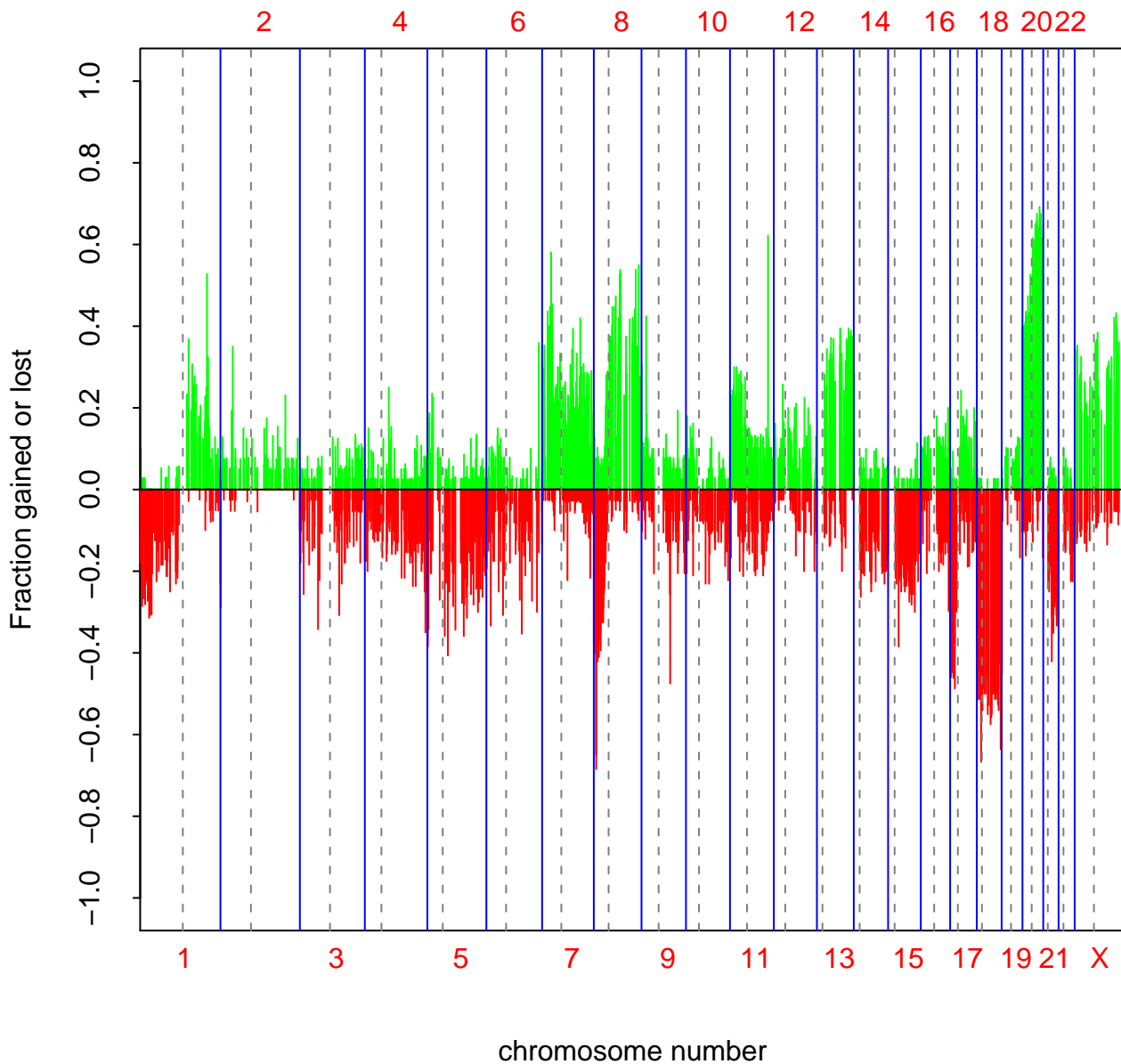
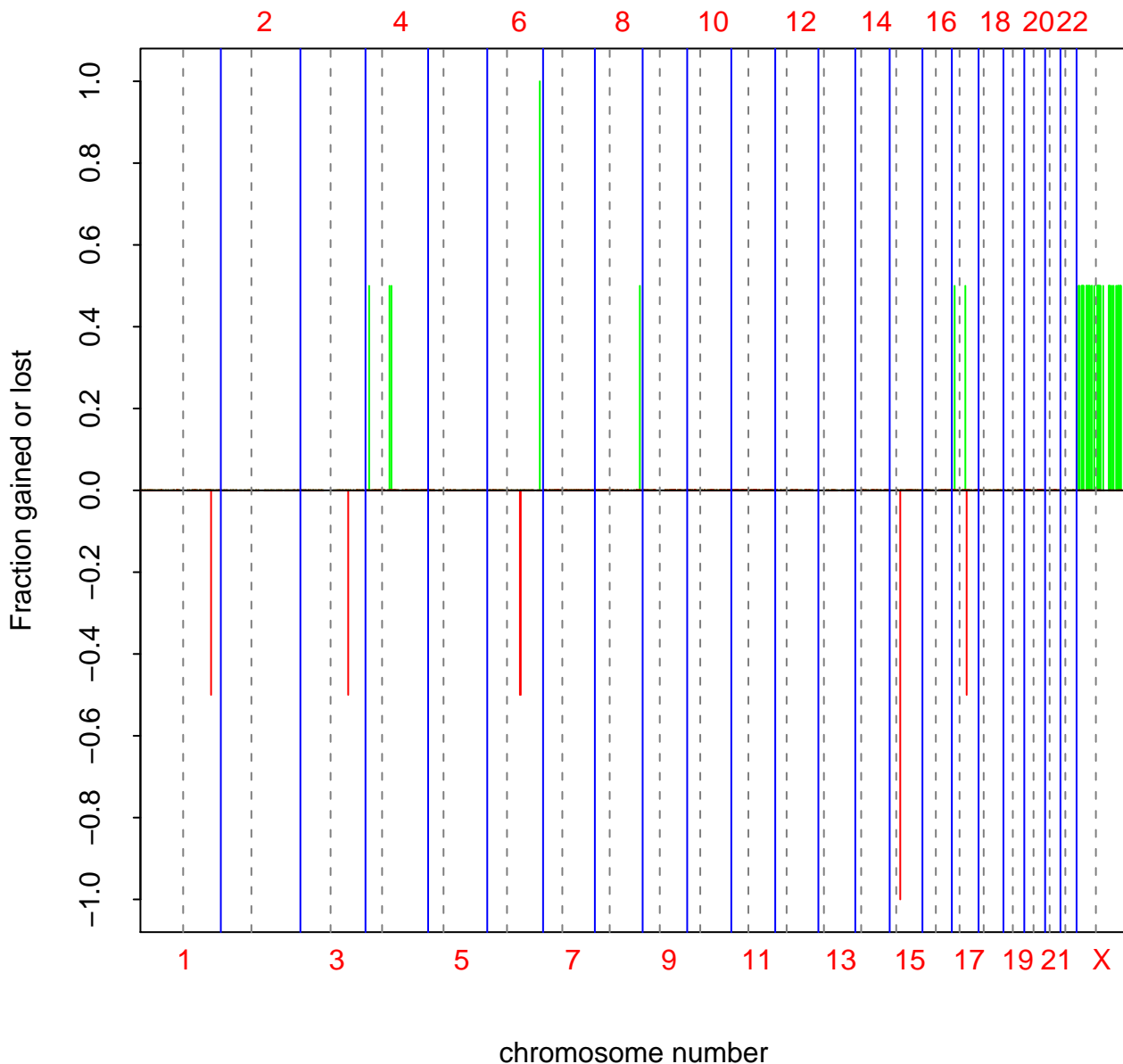


All Samples

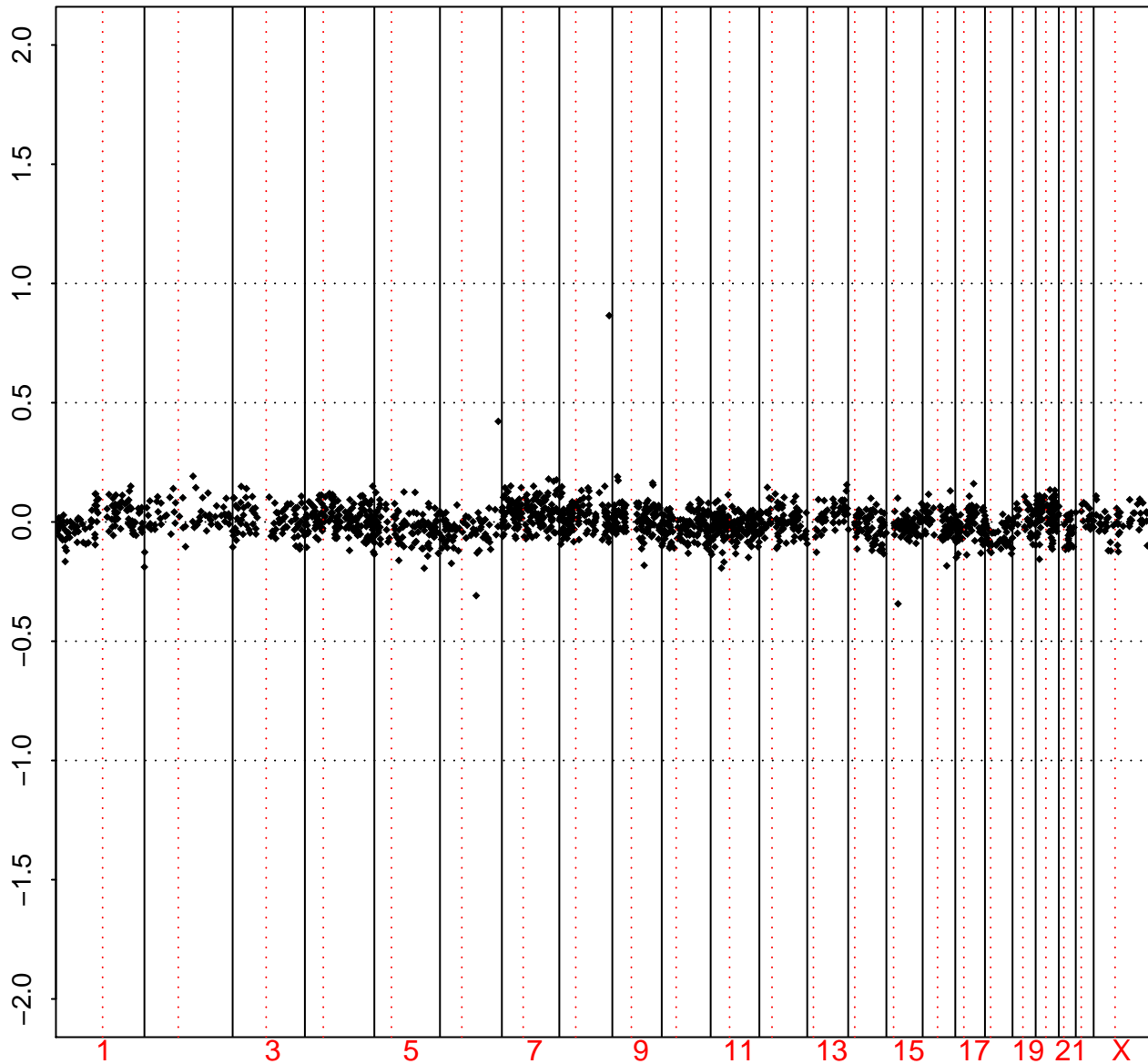


All Samples

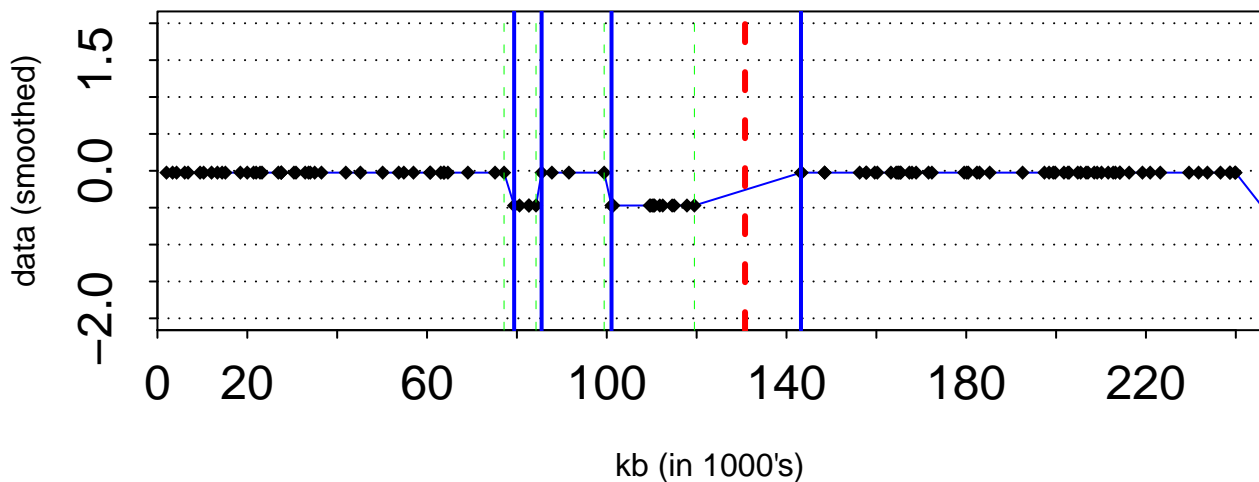
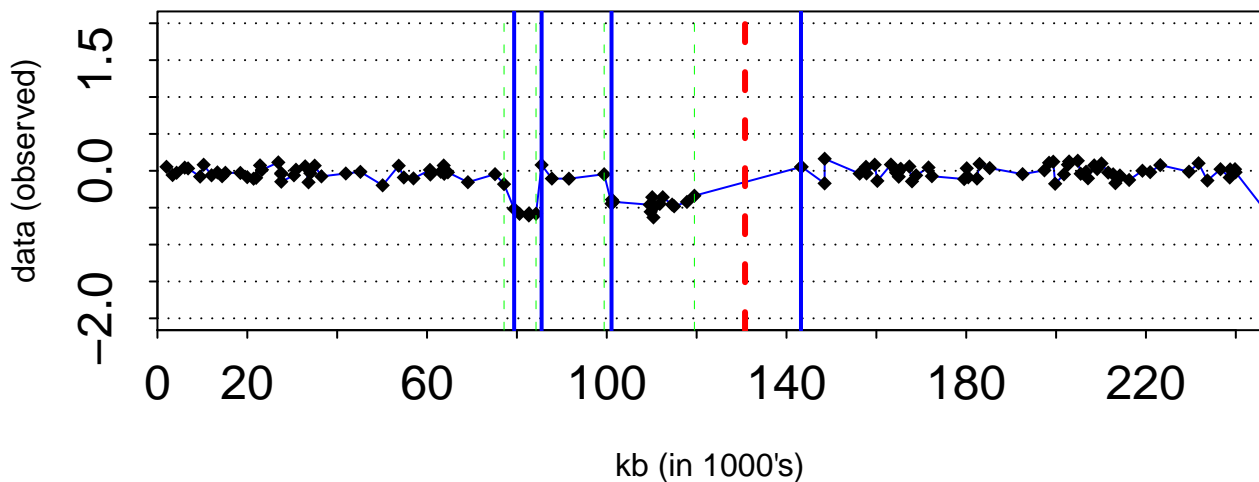


E:/biocbld/bbs-2.4-bioc/tmpdir/Rinst476490618/aCGH/examples/sprocCR4

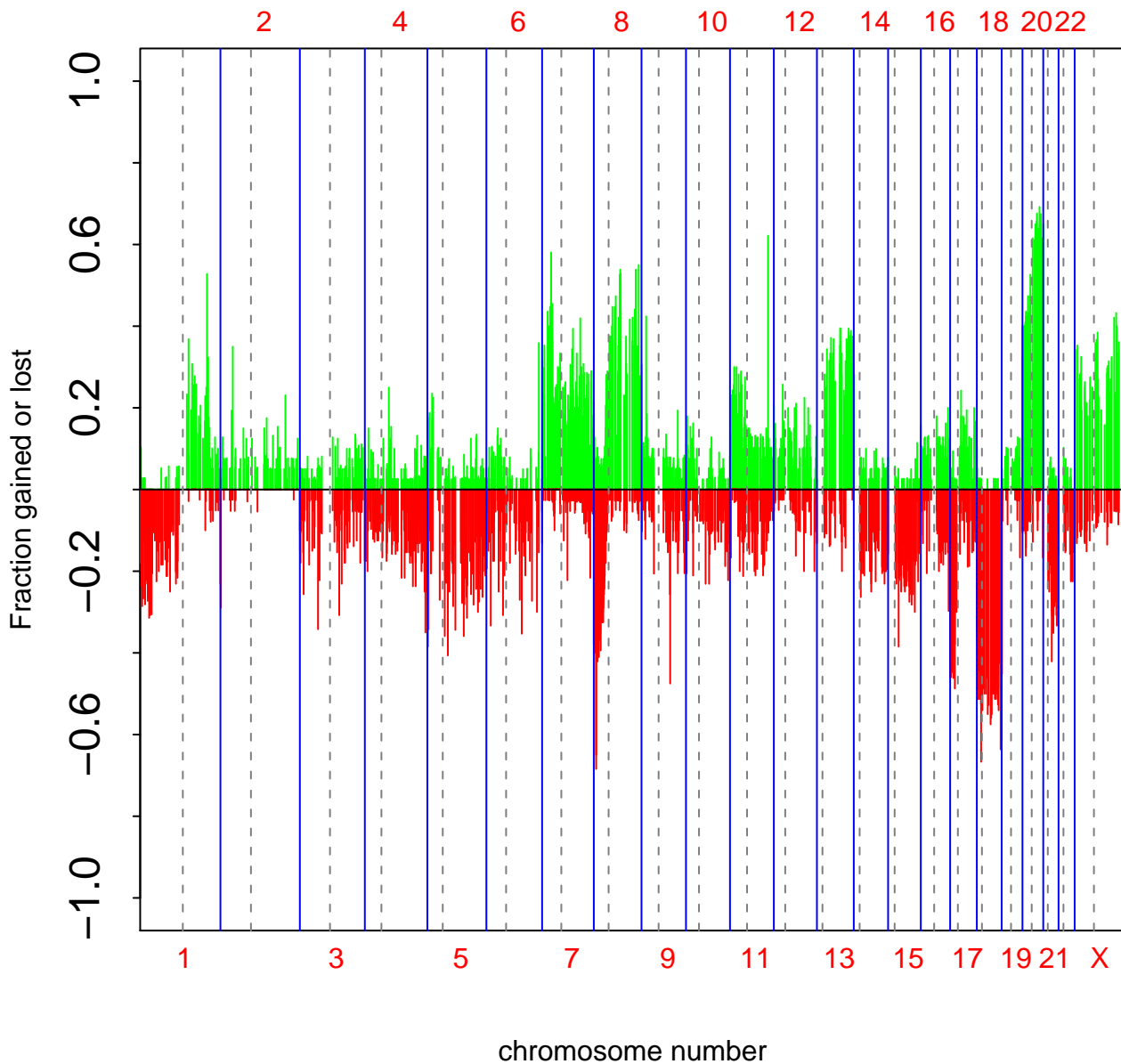
Log2Ratio

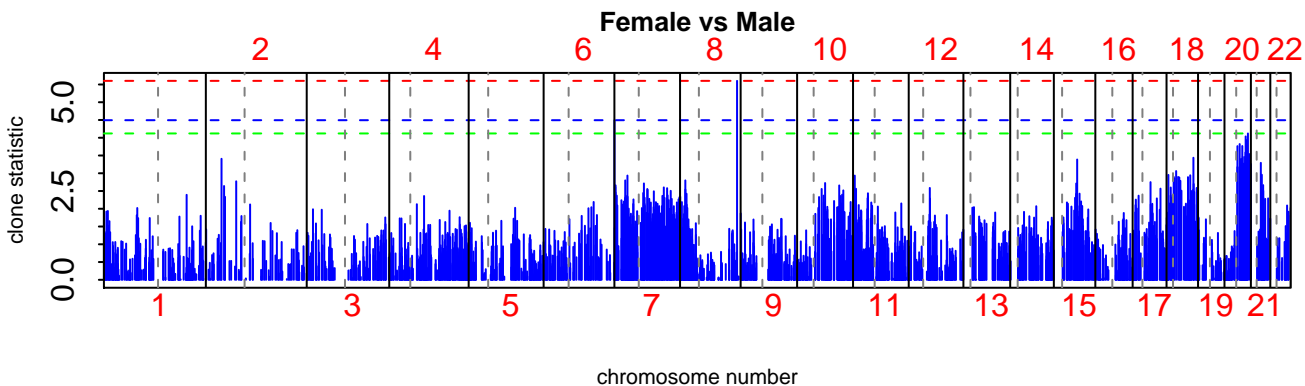
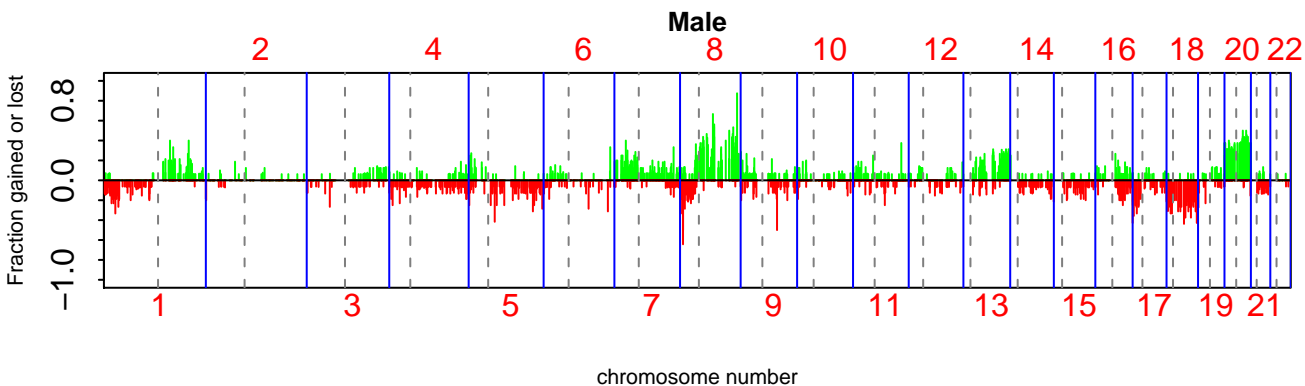
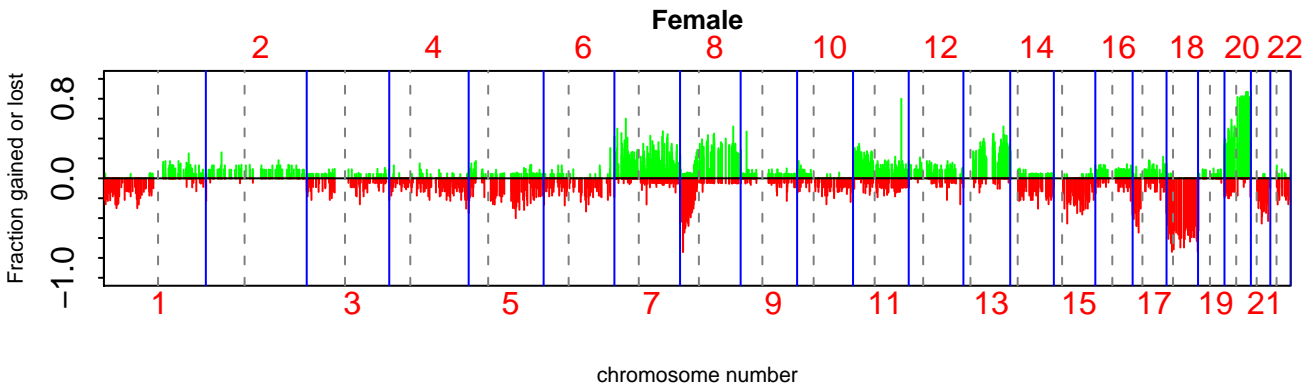


Sample 1 sprocCR31.txt – Chr 1 Number of states 2

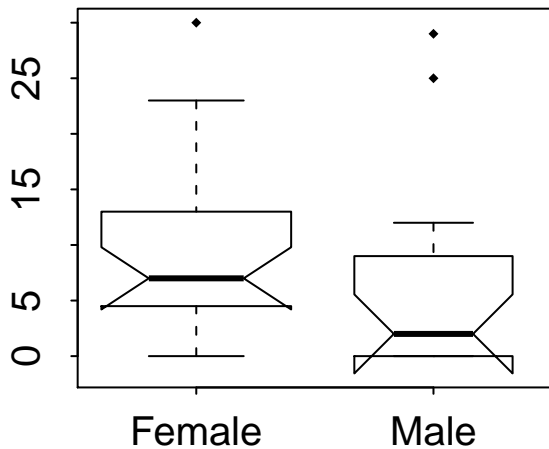


All Samples

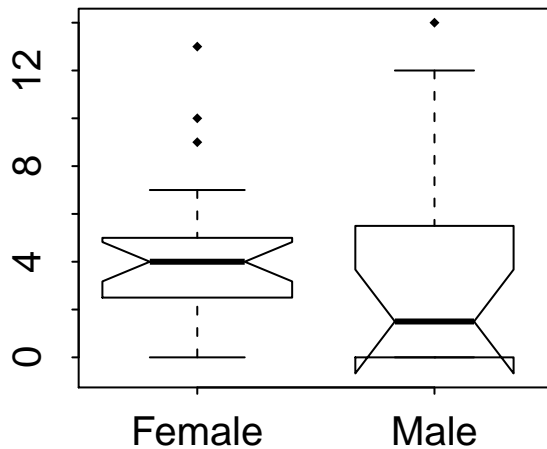




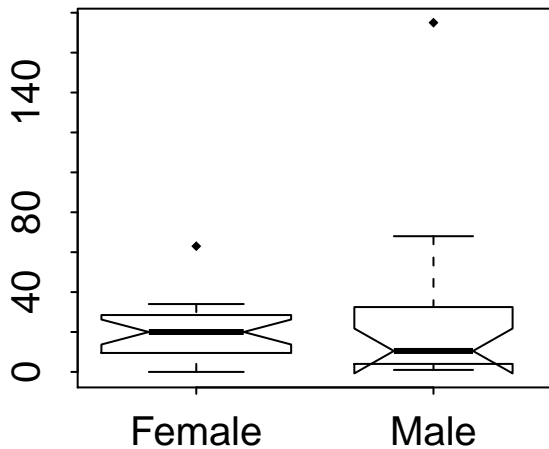
Number of Transitions 0.0950684



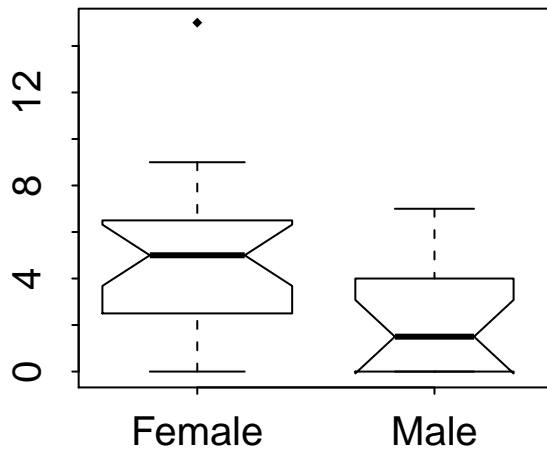
Number of Chrom containing Transitions 0.174915



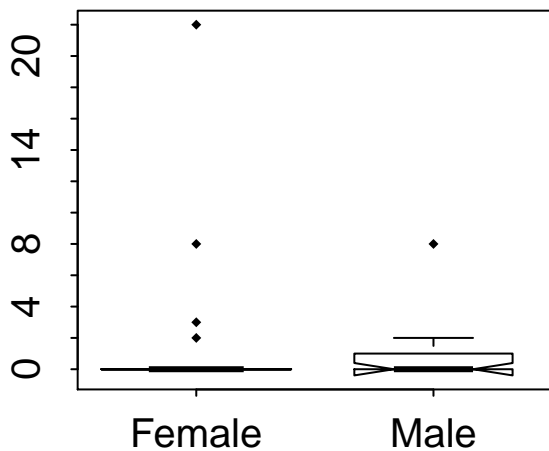
Number of Aberrations 0.841381



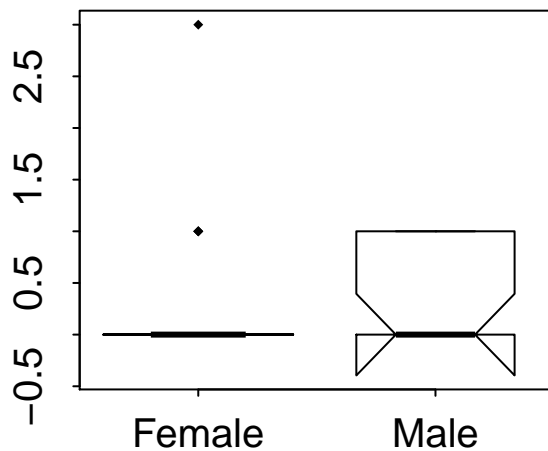
Number of Whole Chrom Changes 0.0174111



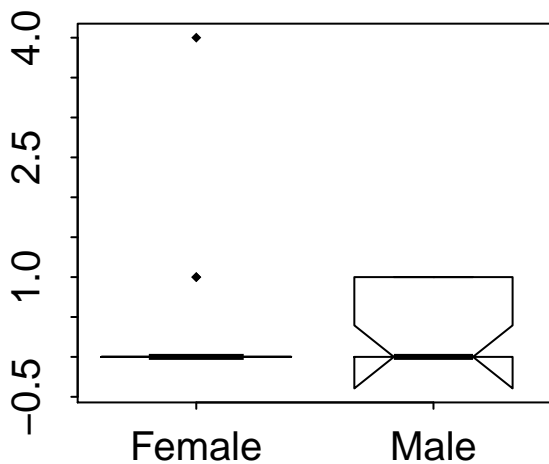
Number of Amplifications 0.462205



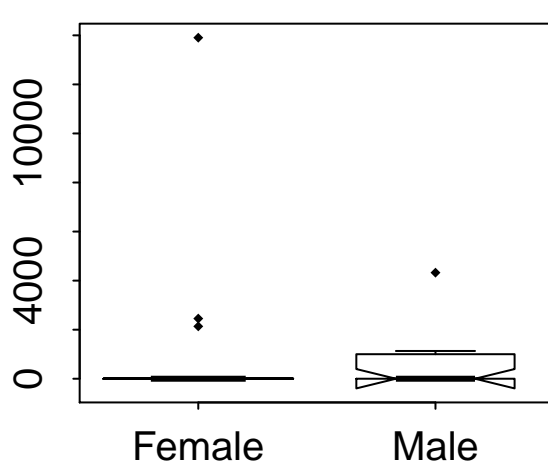
Number of Chrom containing Amplifications 0.369971



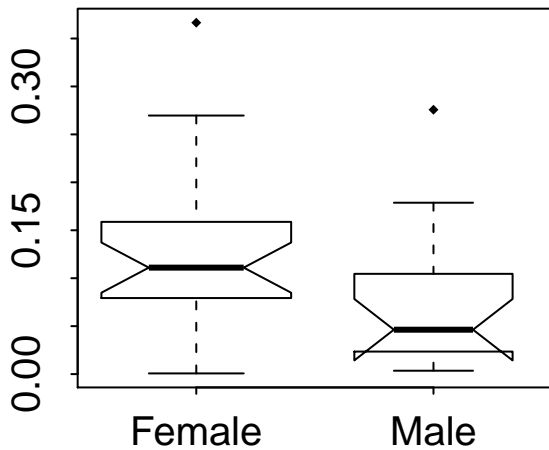
Number of Amplicons 0.207136



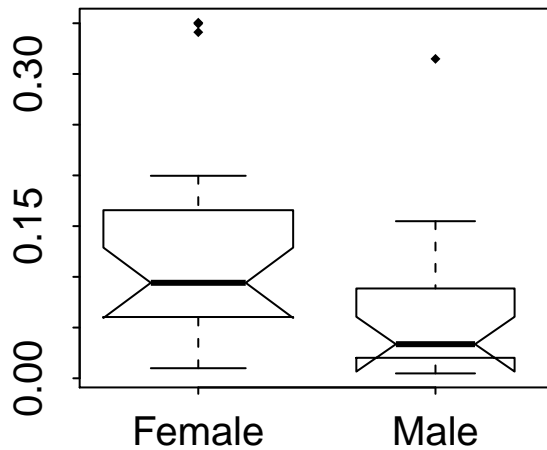
Amount of Genome Amplified 0.257209



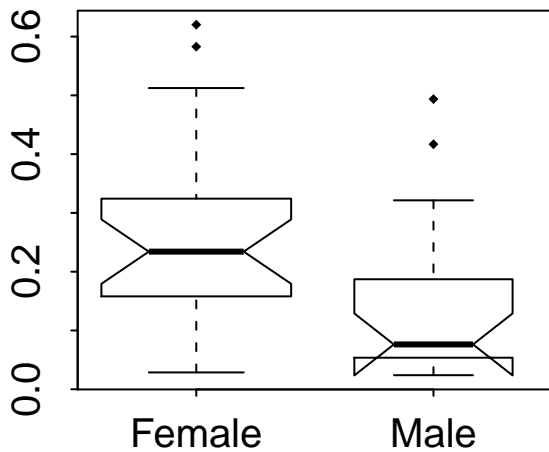
Fraction of Genome Gained 0.0300104

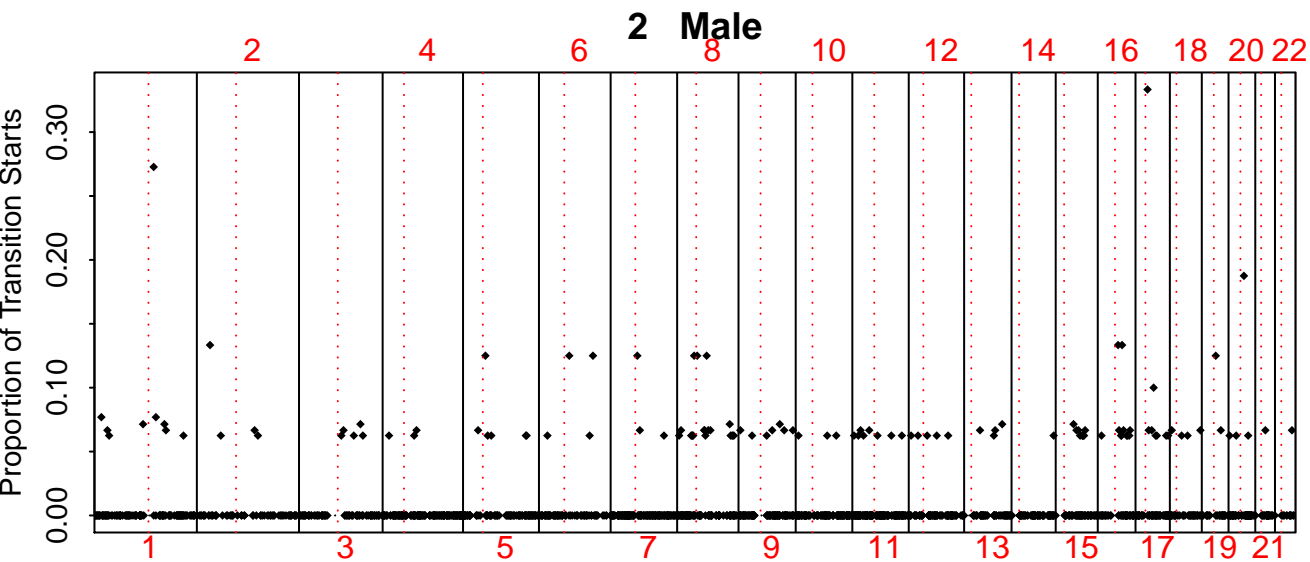
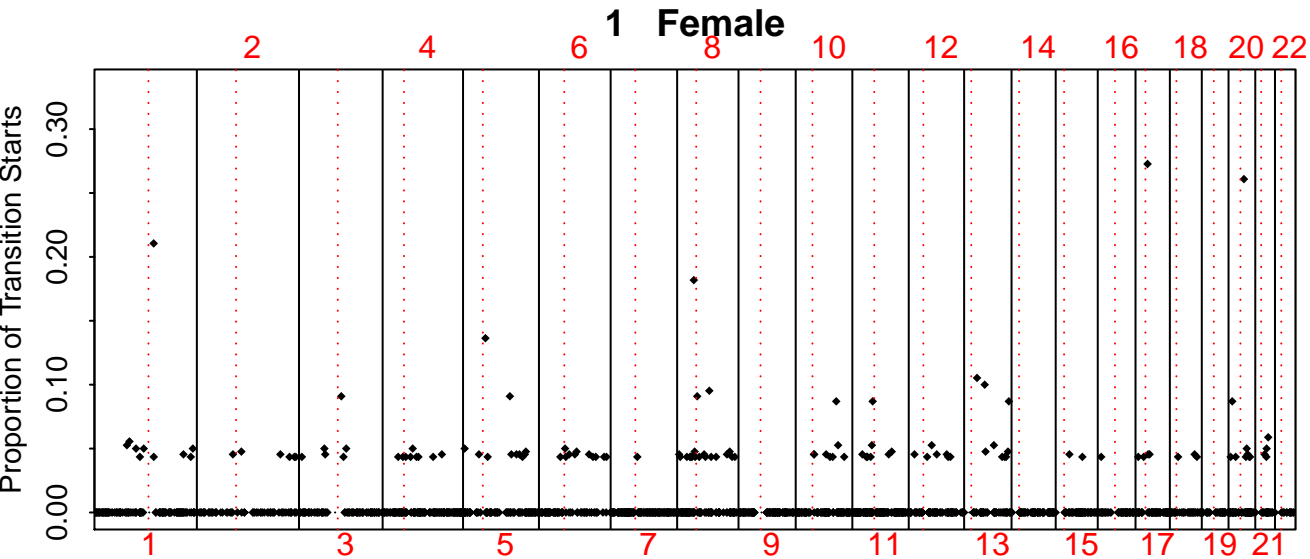


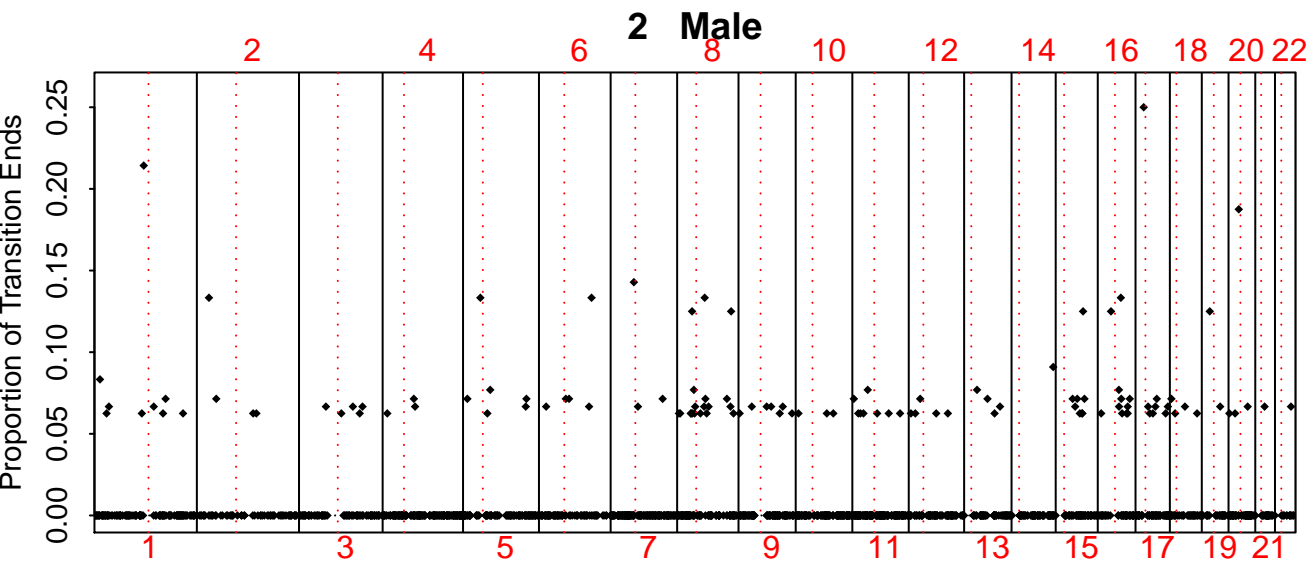
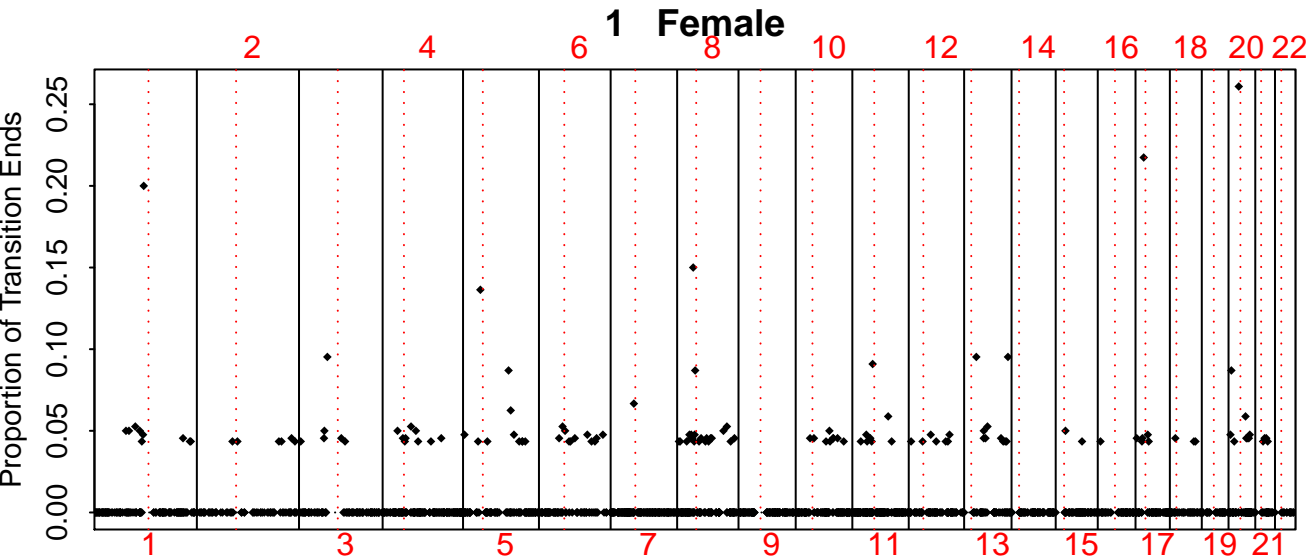
Fraction of Genome Lost 0.0300104



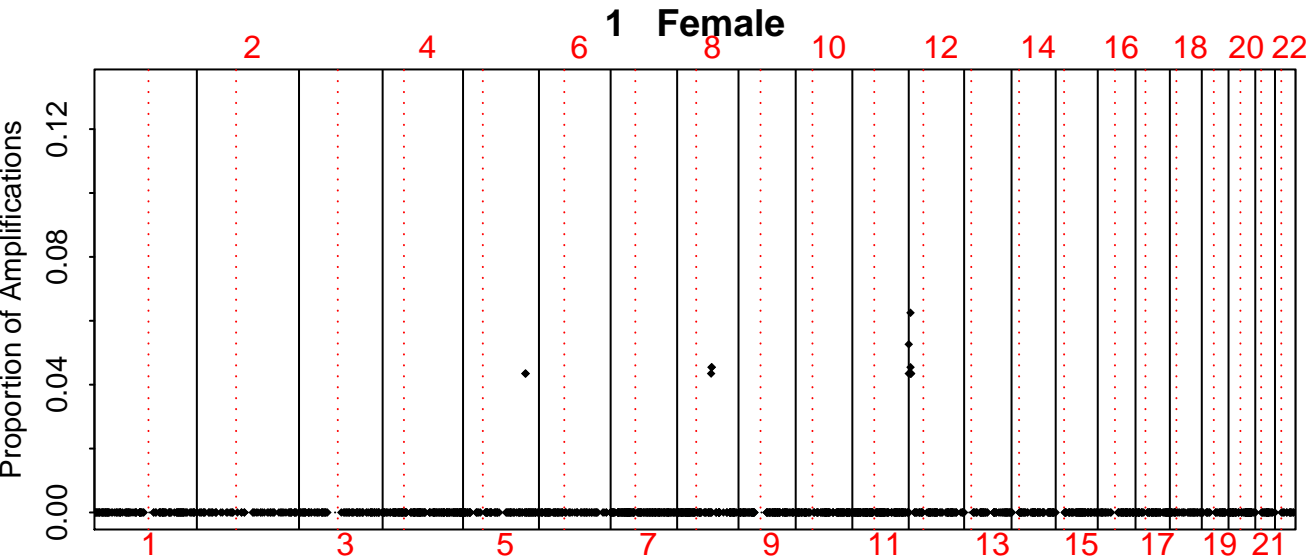
Fraction of Genome Altered 0.0240951



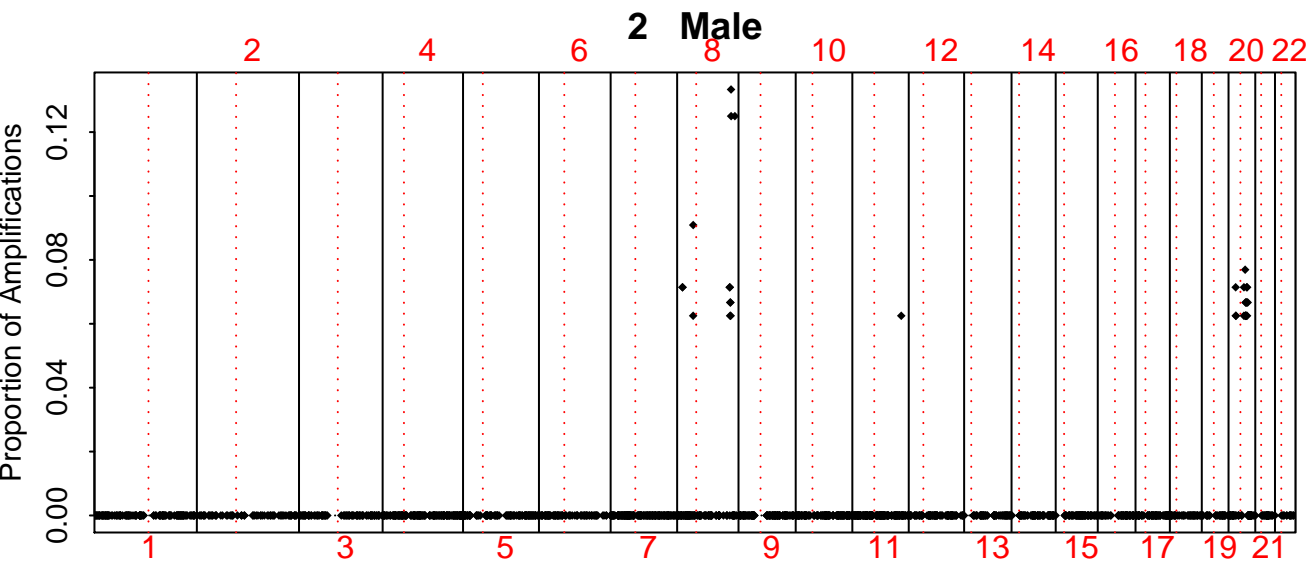




Proportion of Amplifications

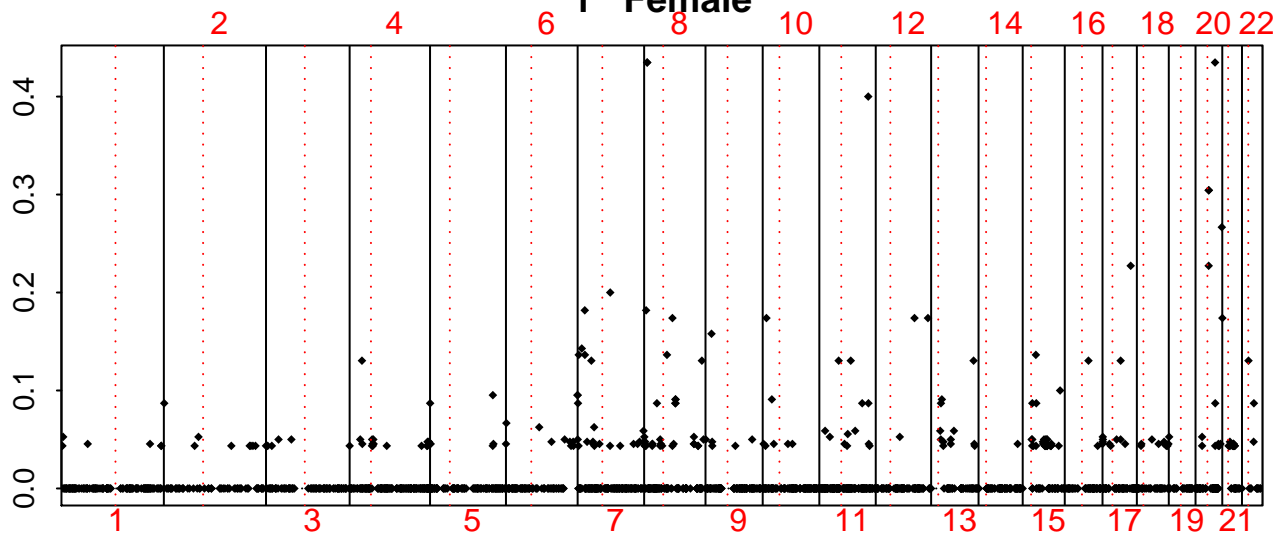


Proportion of Amplifications



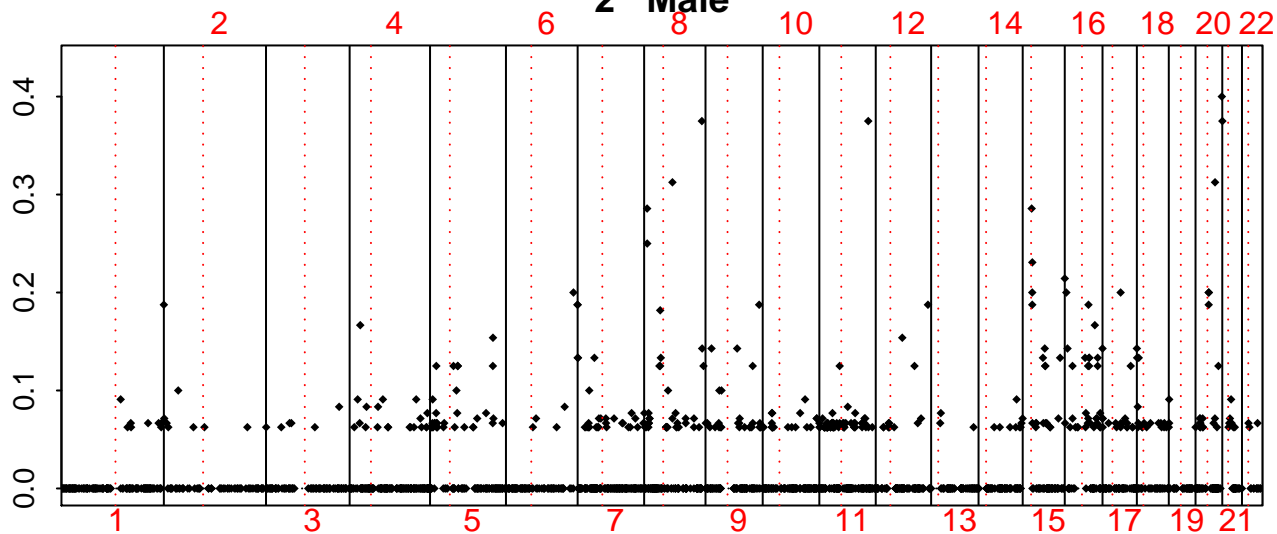
Proportion of Aberrations

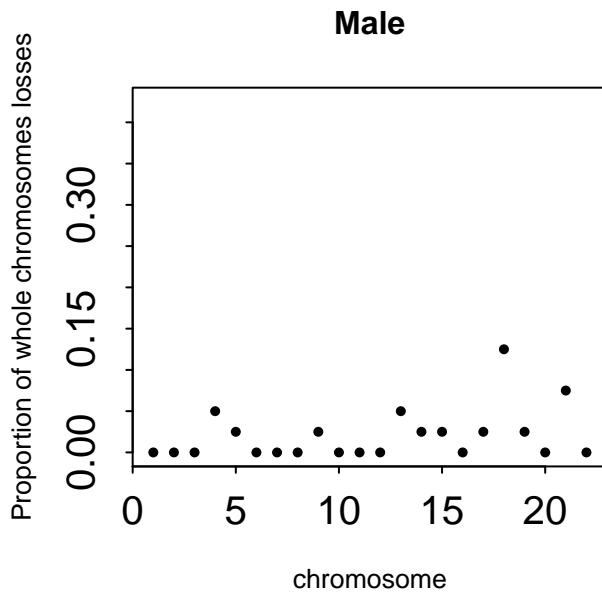
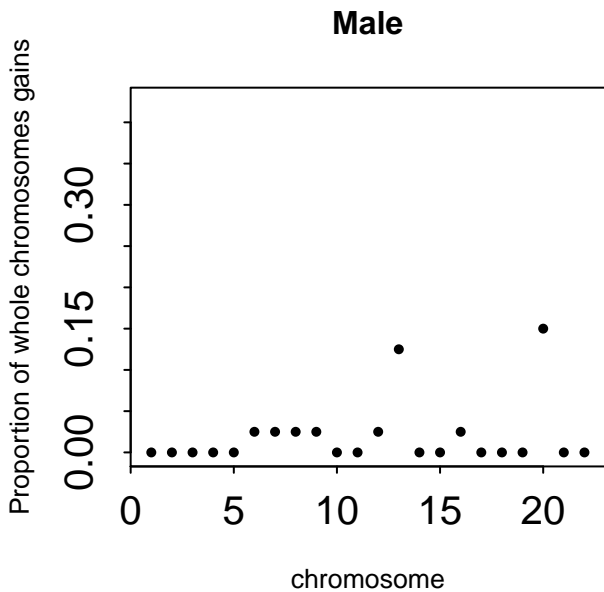
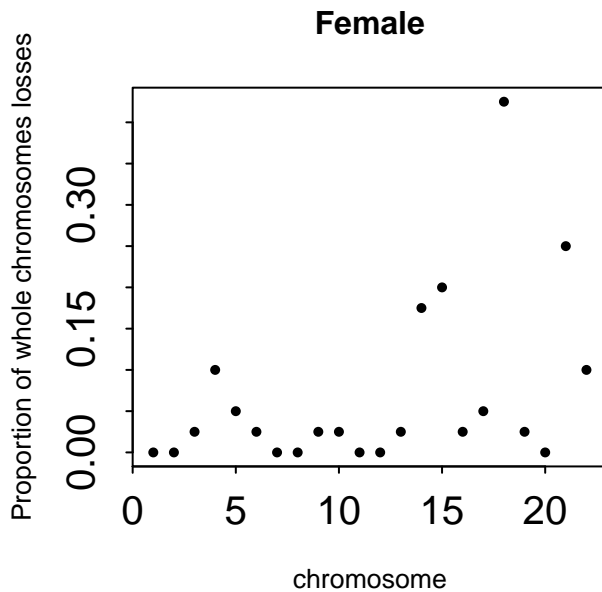
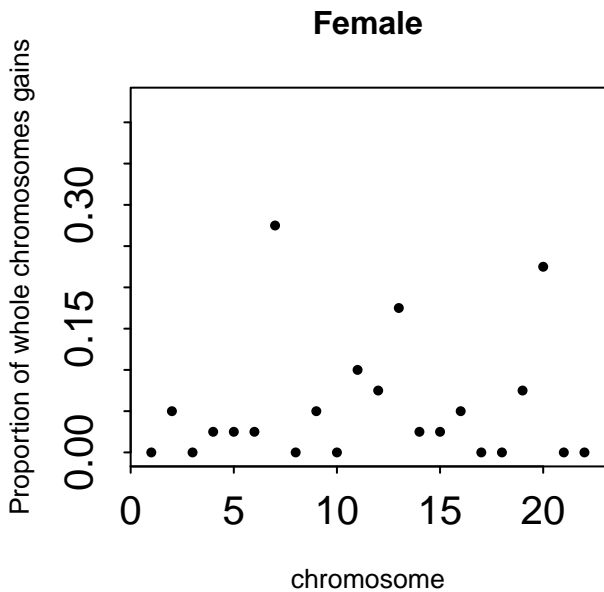
1 Female

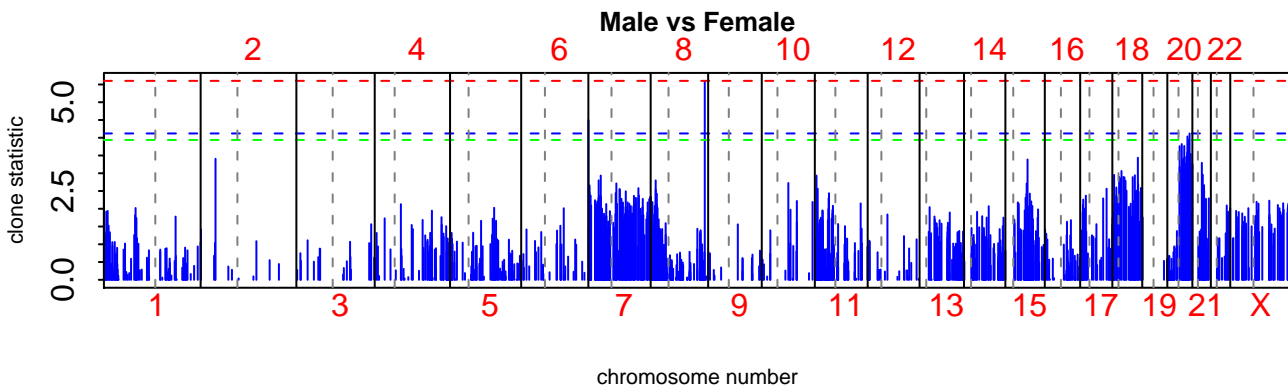
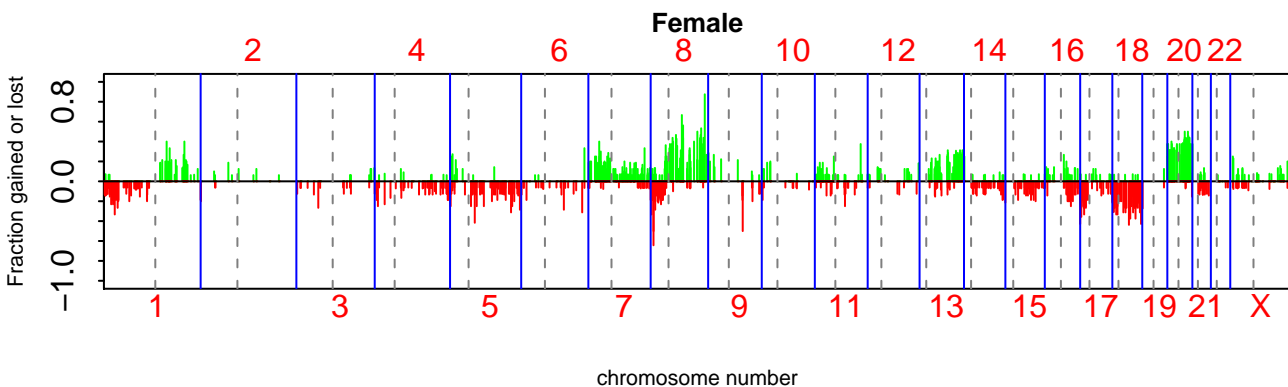
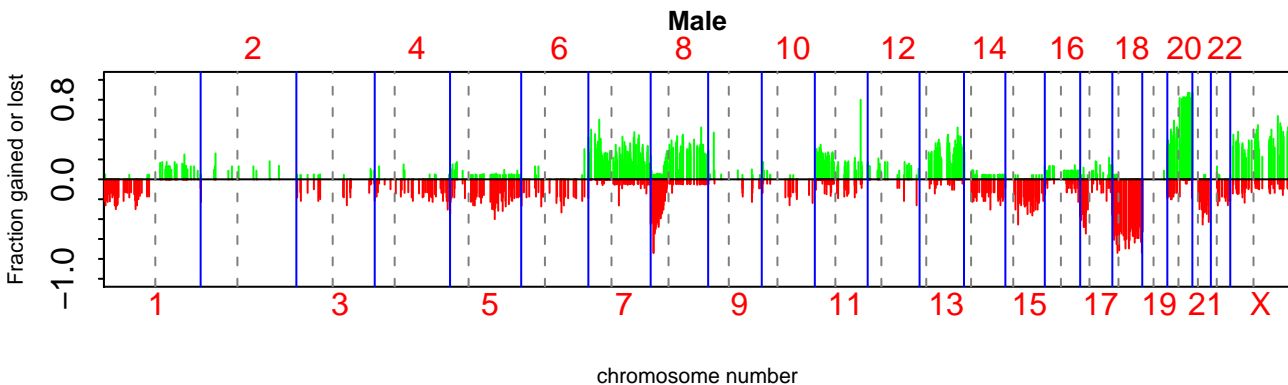


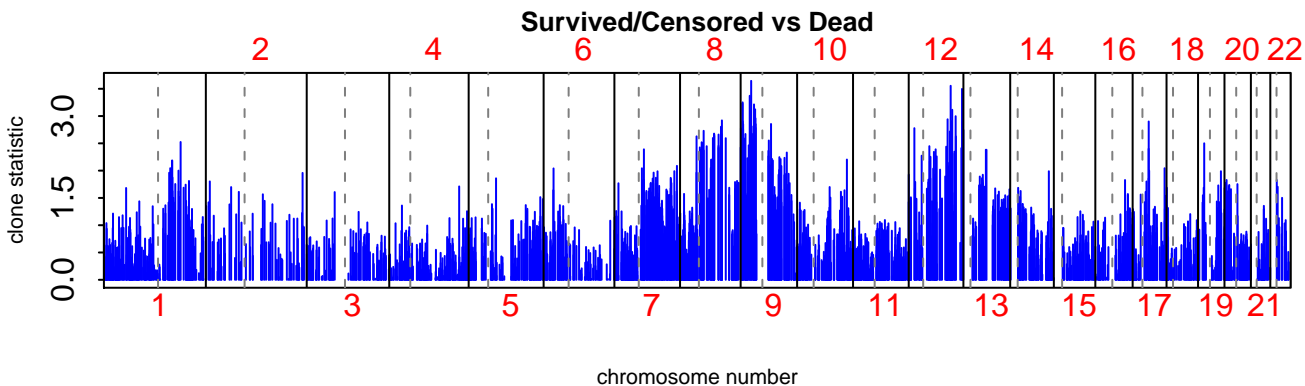
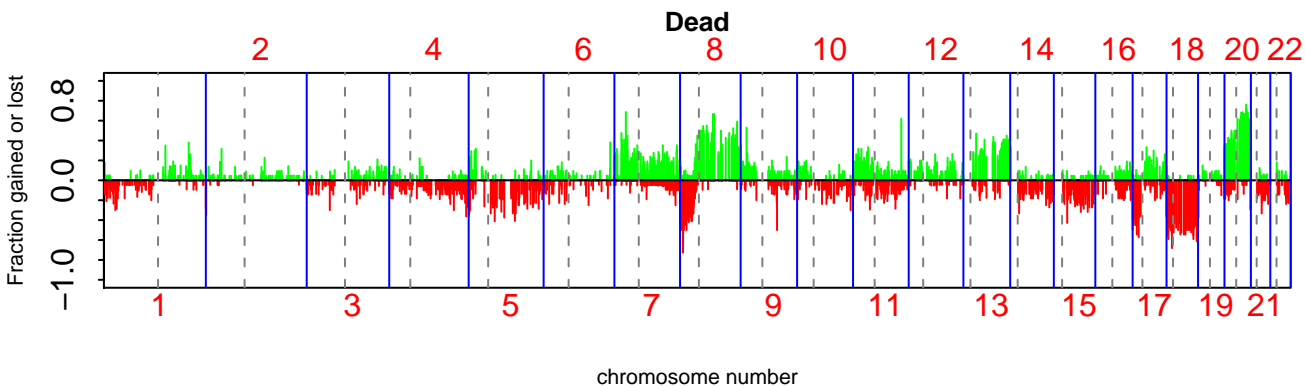
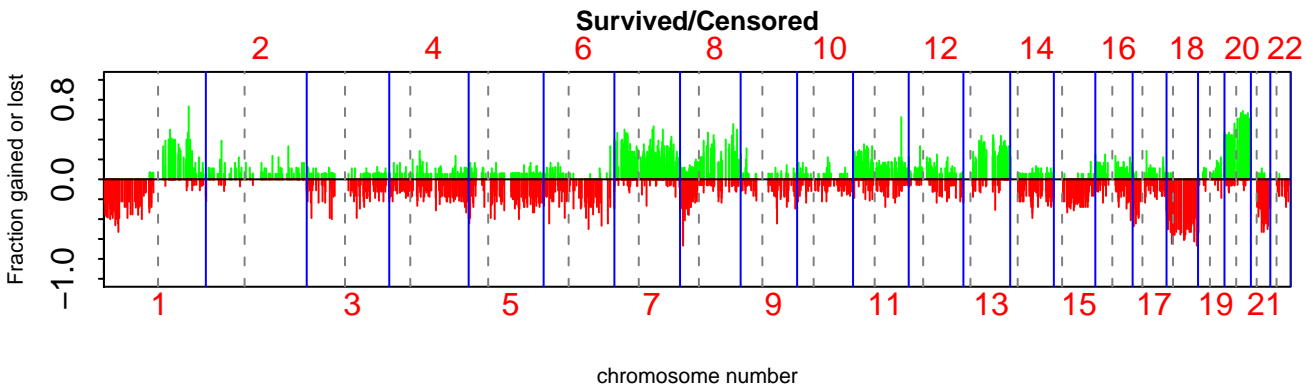
Proportion of Aberrations

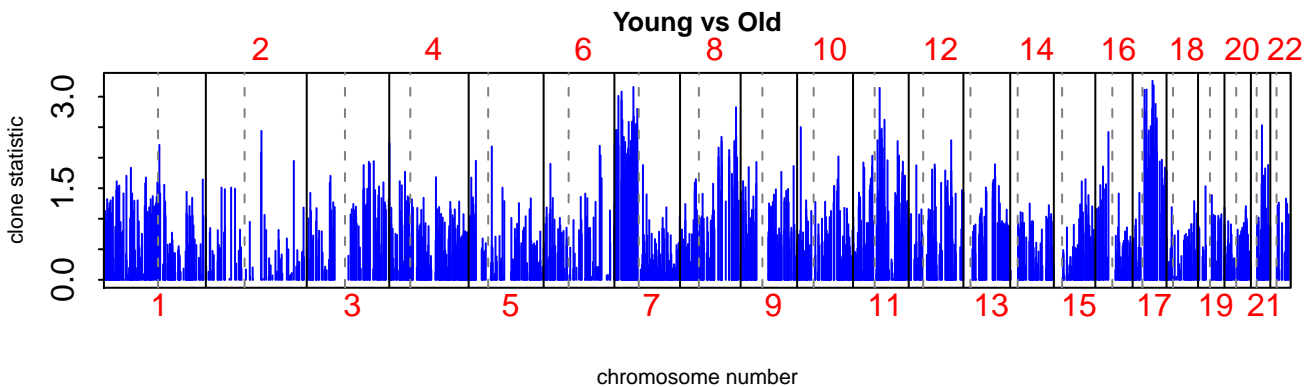
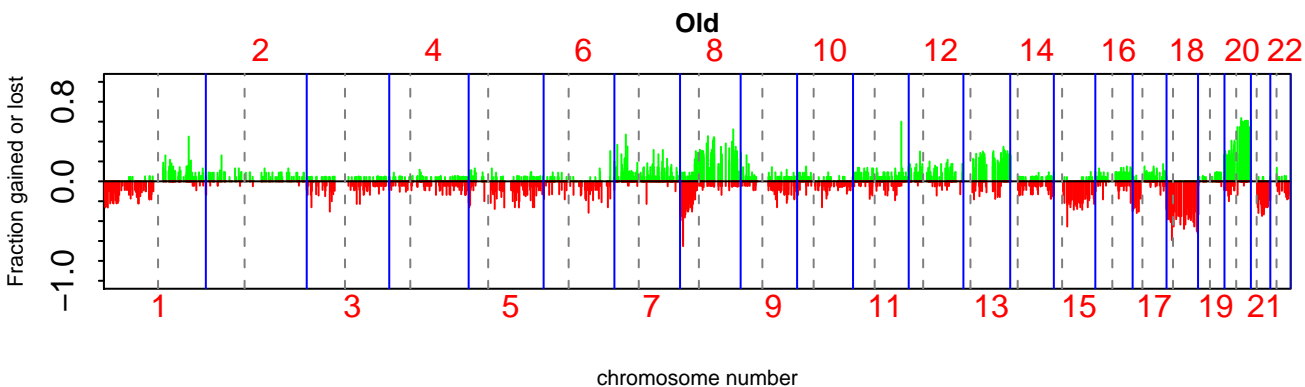
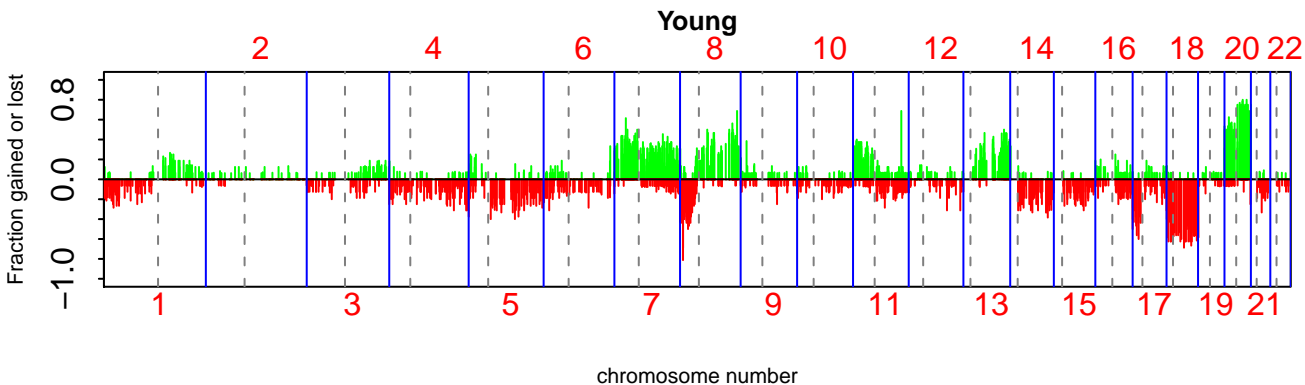
2 Male











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