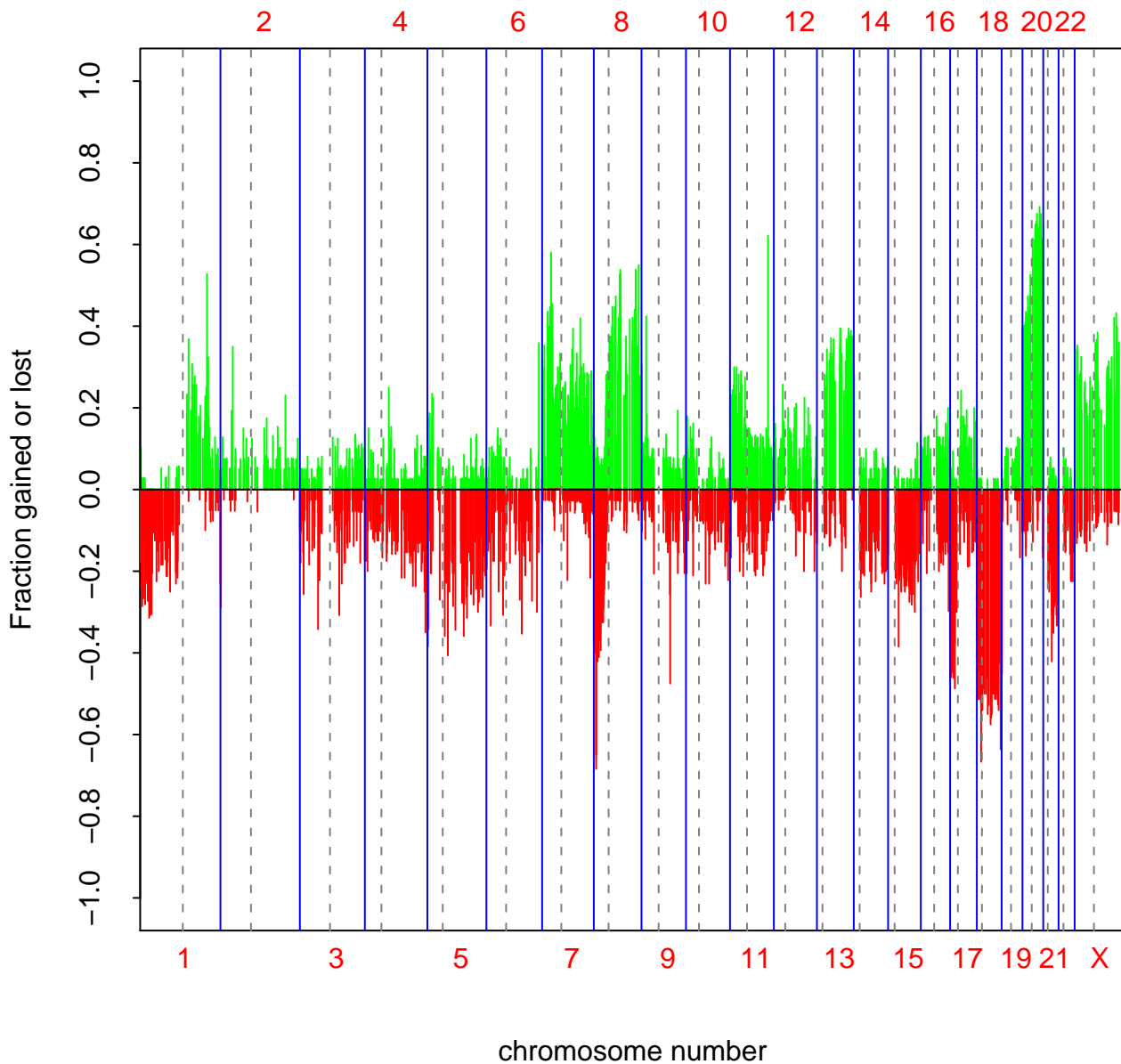


All Samples



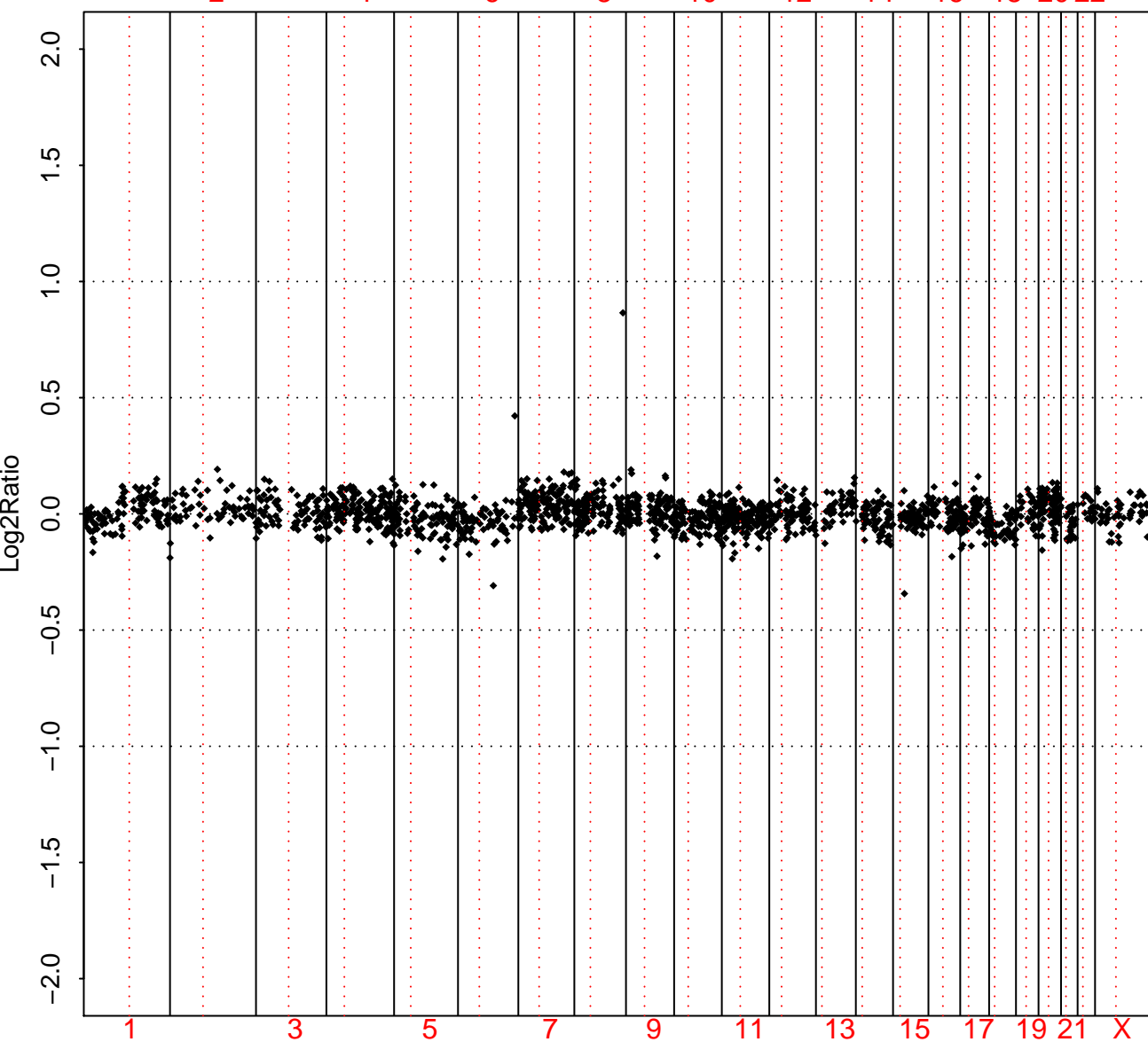
All Samples

22

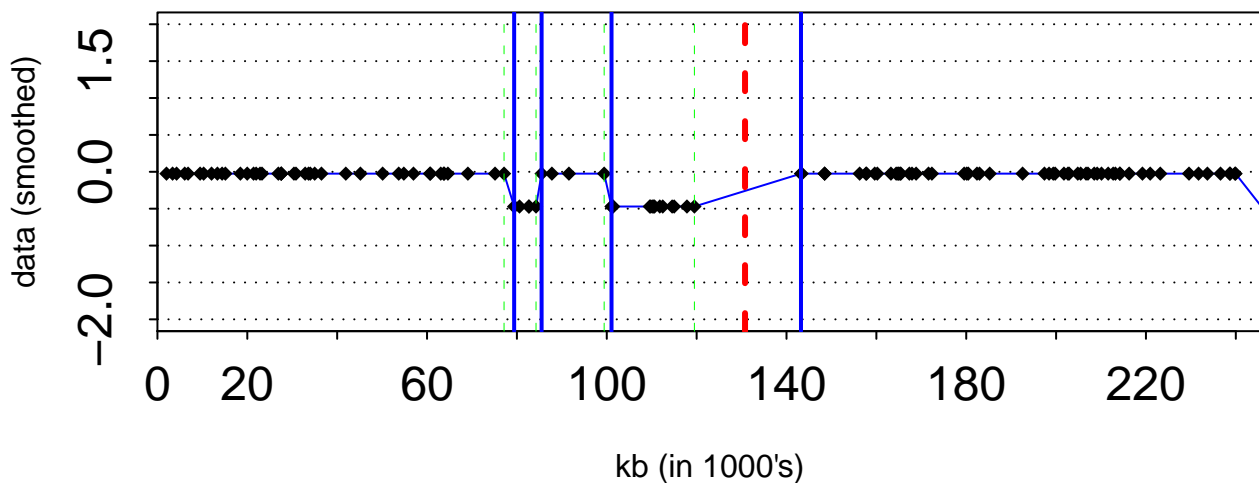
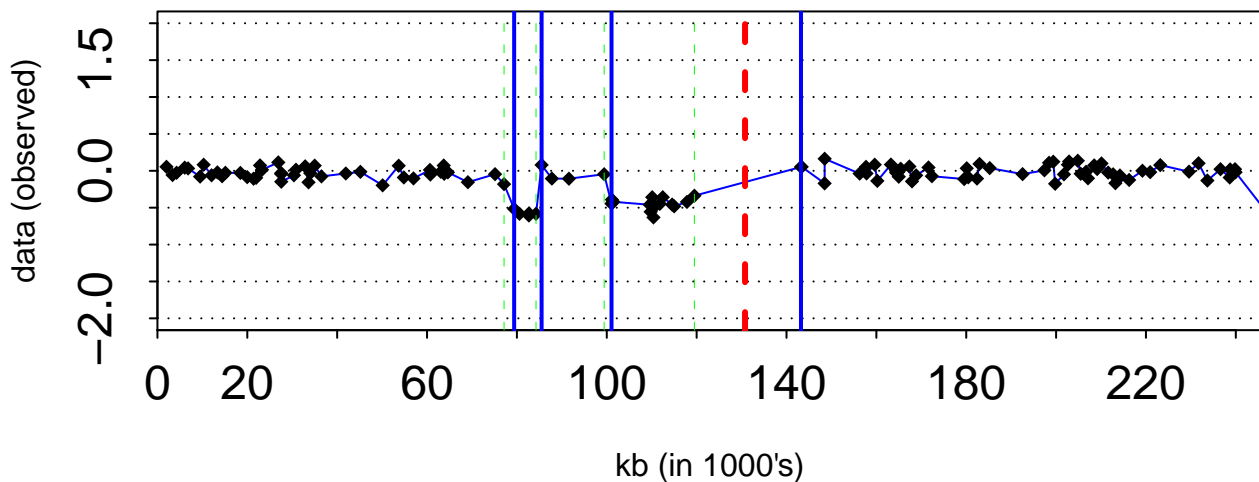
Fraction gained or lost

chromosome number

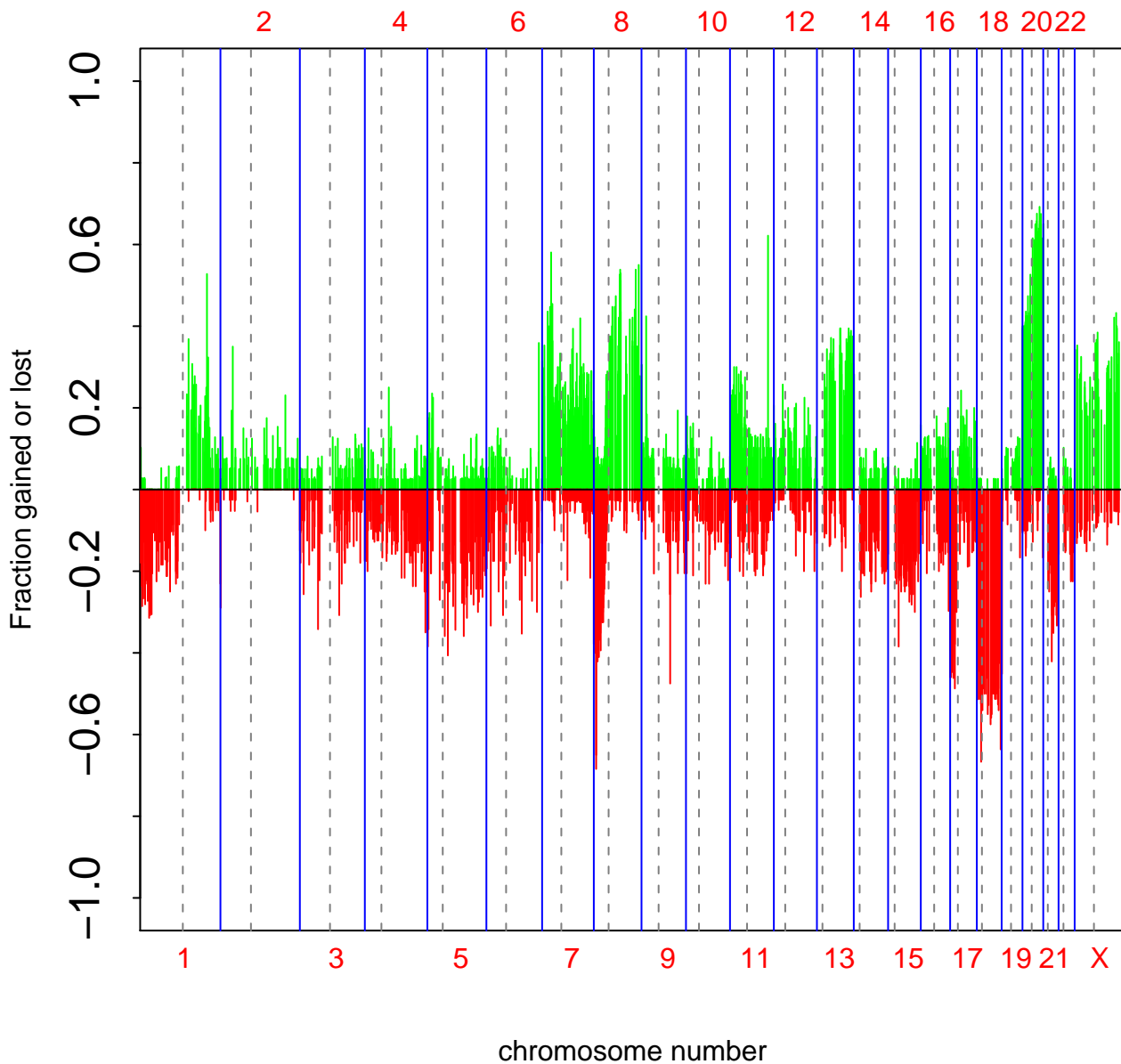
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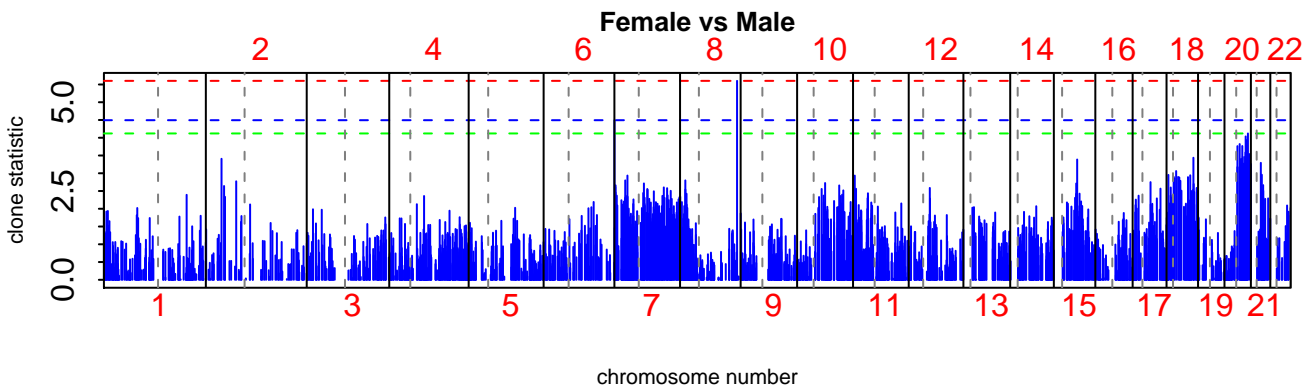
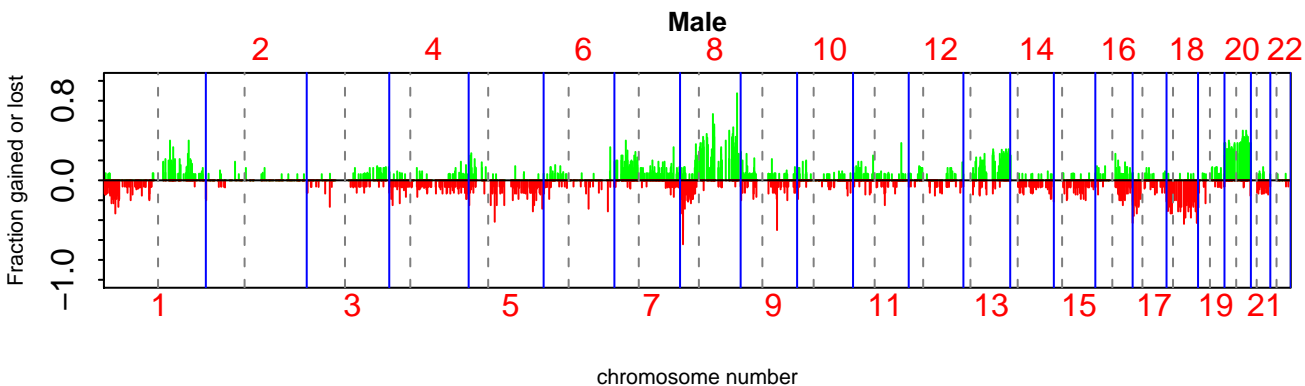
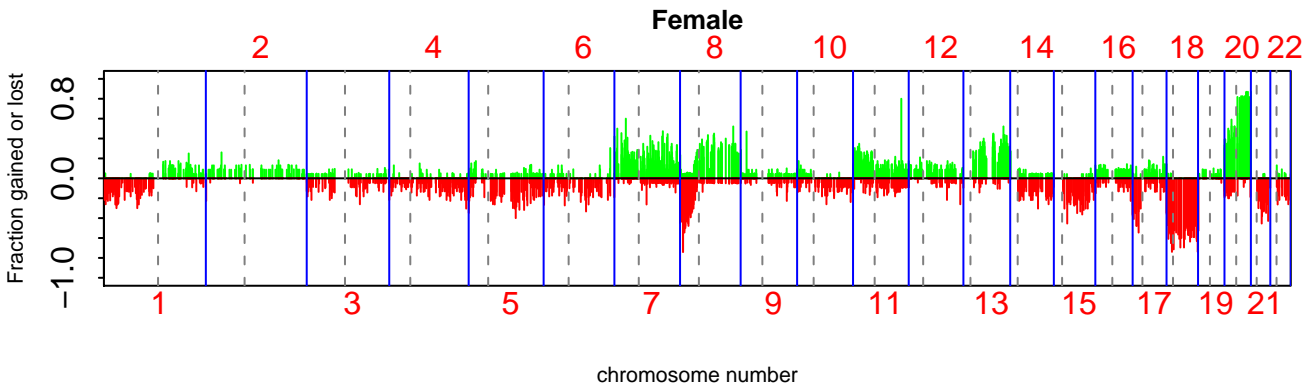


Sample 1 sprocCR31.txt – Chr 1 Number of states 2

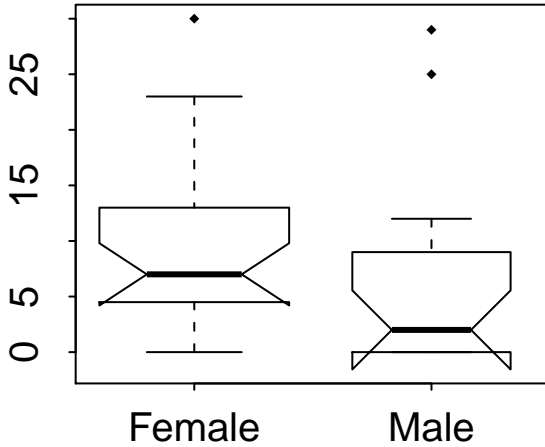


All Samples

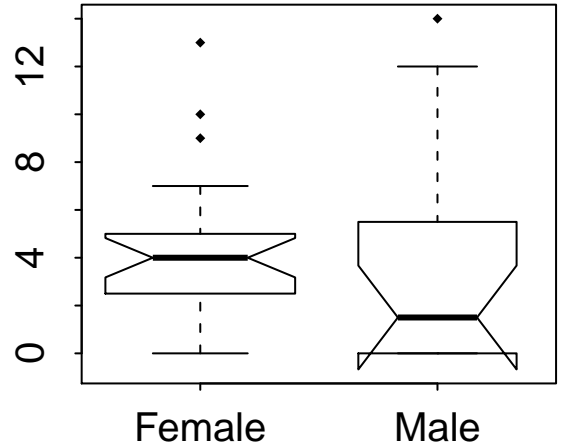




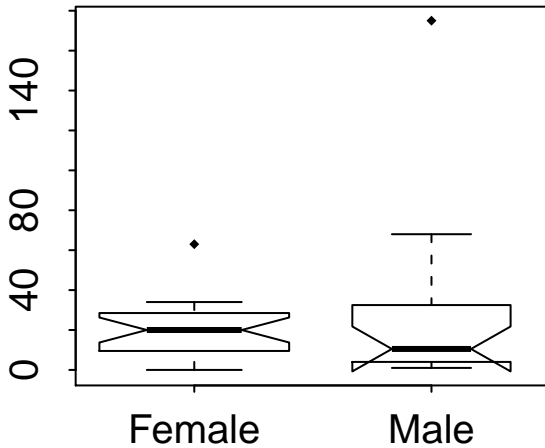
Number of Transitions 0.0950684



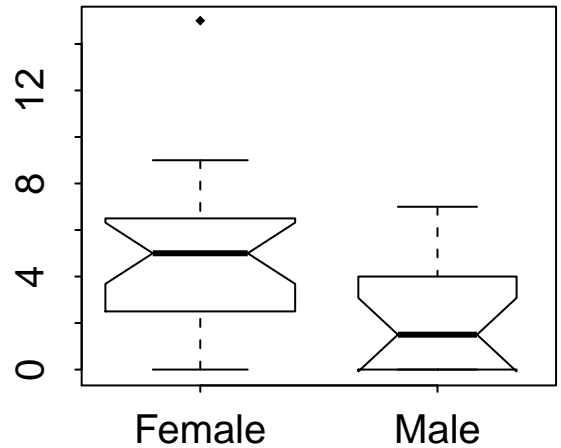
Number of Chrom containing Transitions 0.174915



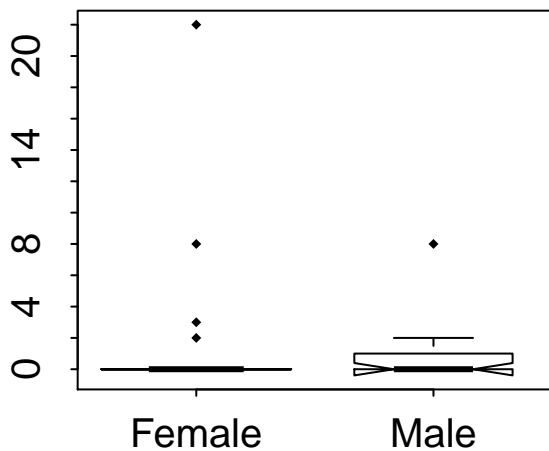
Number of Aberrations 0.841381



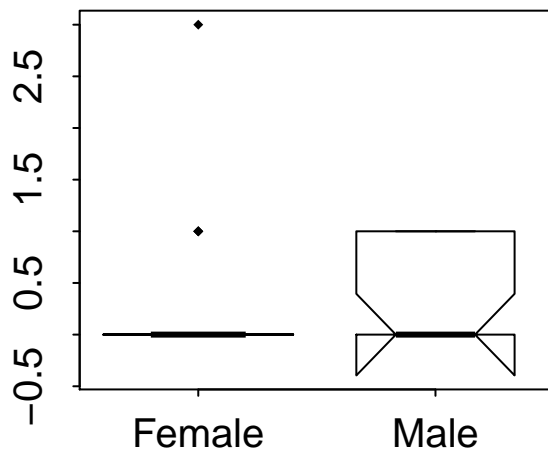
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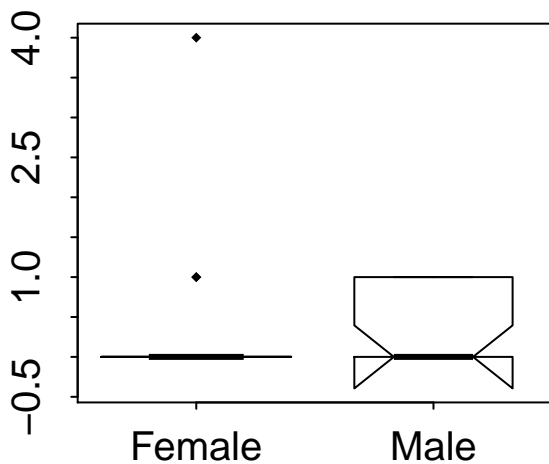
Number of Amplifications 0.462205



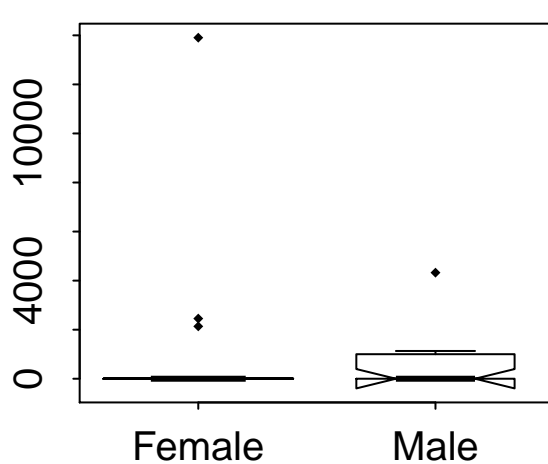
Number of Chrom containing Amplifications 0.369971



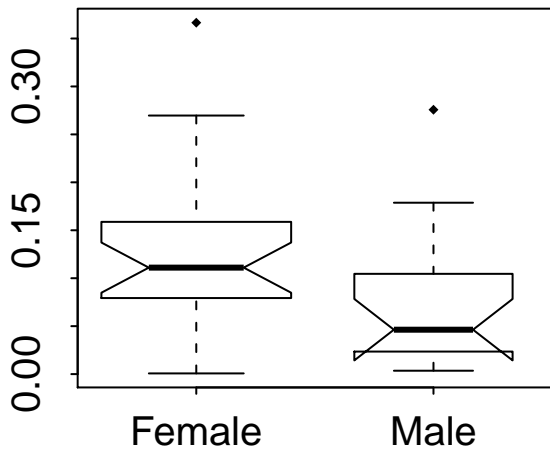
Number of Amplicons 0.207136



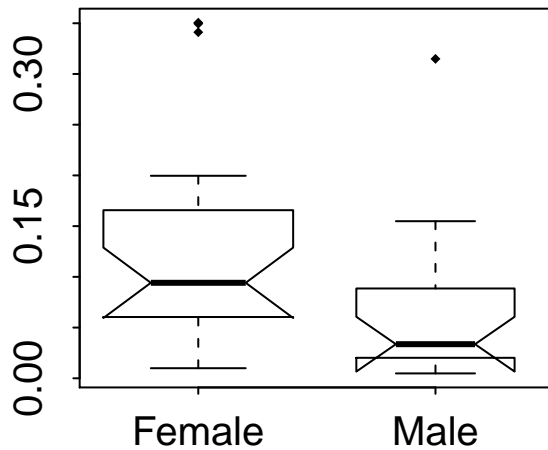
Amount of Genome Amplified 0.257209



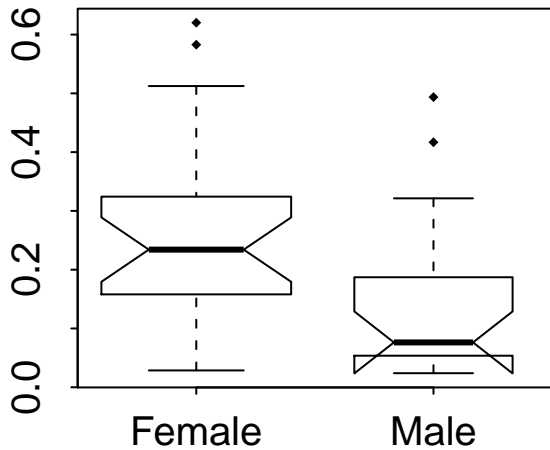
Fraction of Genome Gained 0.0300104

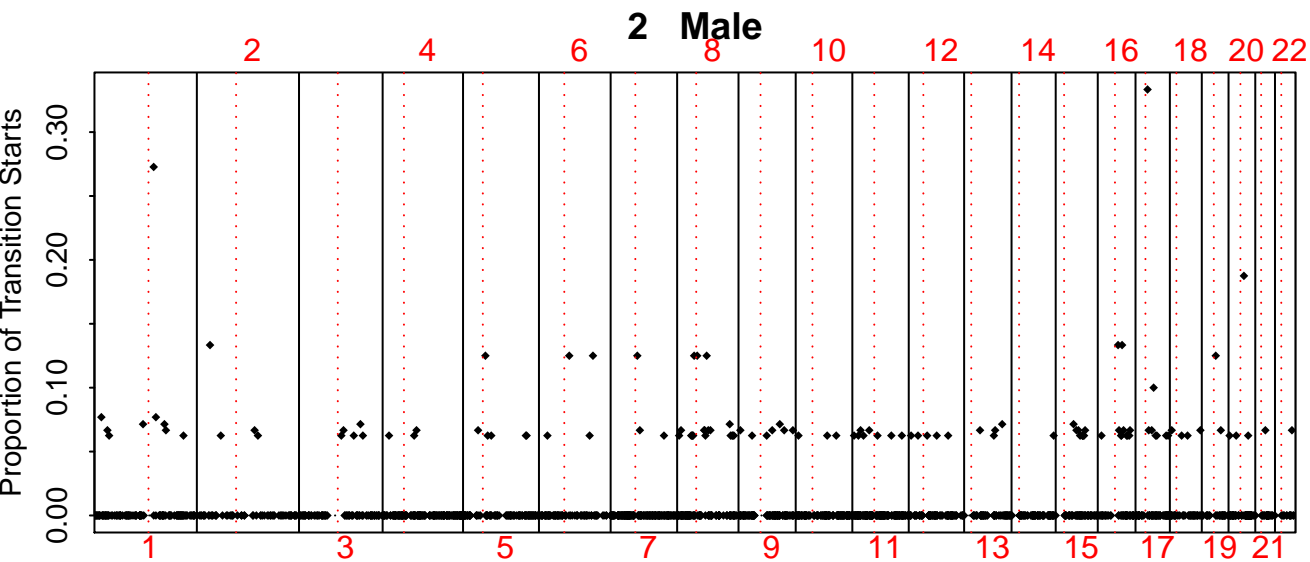
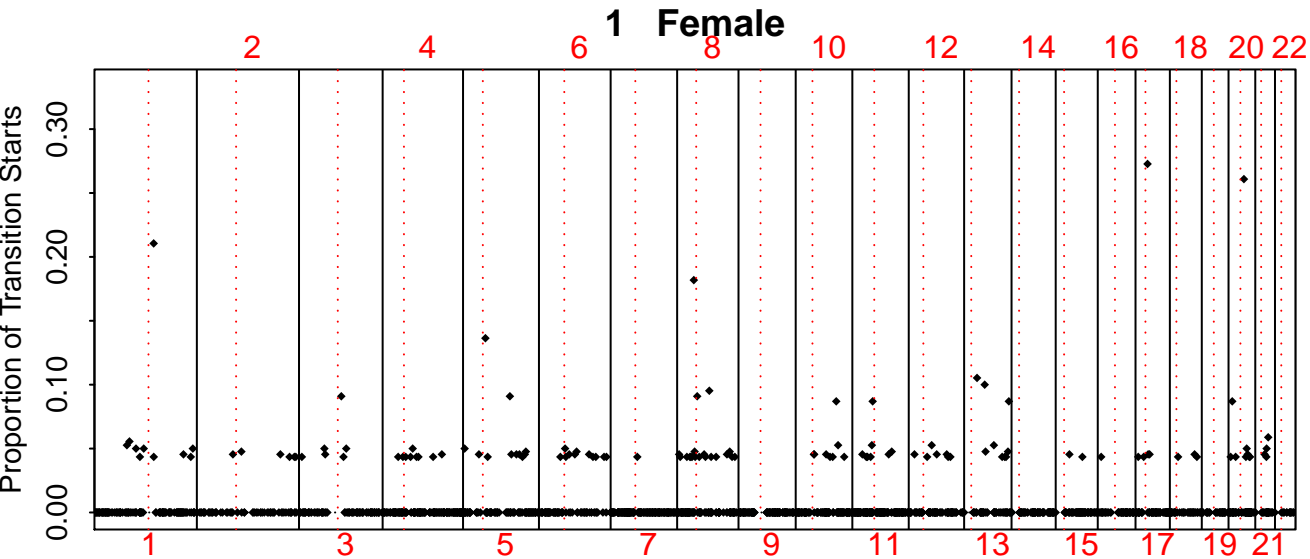


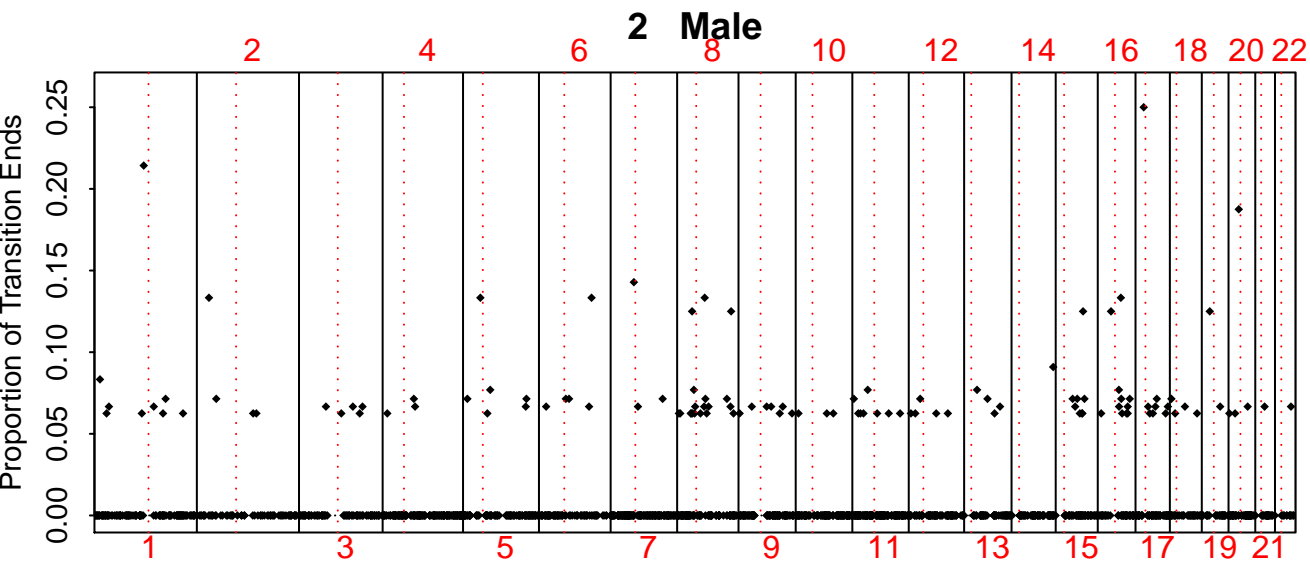
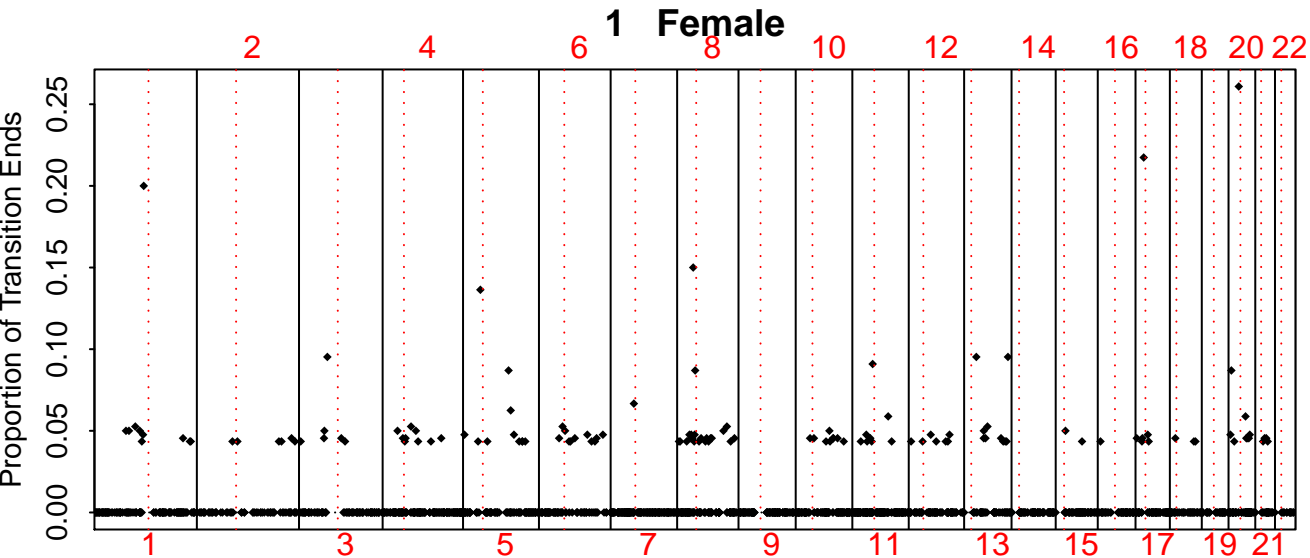
Fraction of Genome Lost 0.0300104



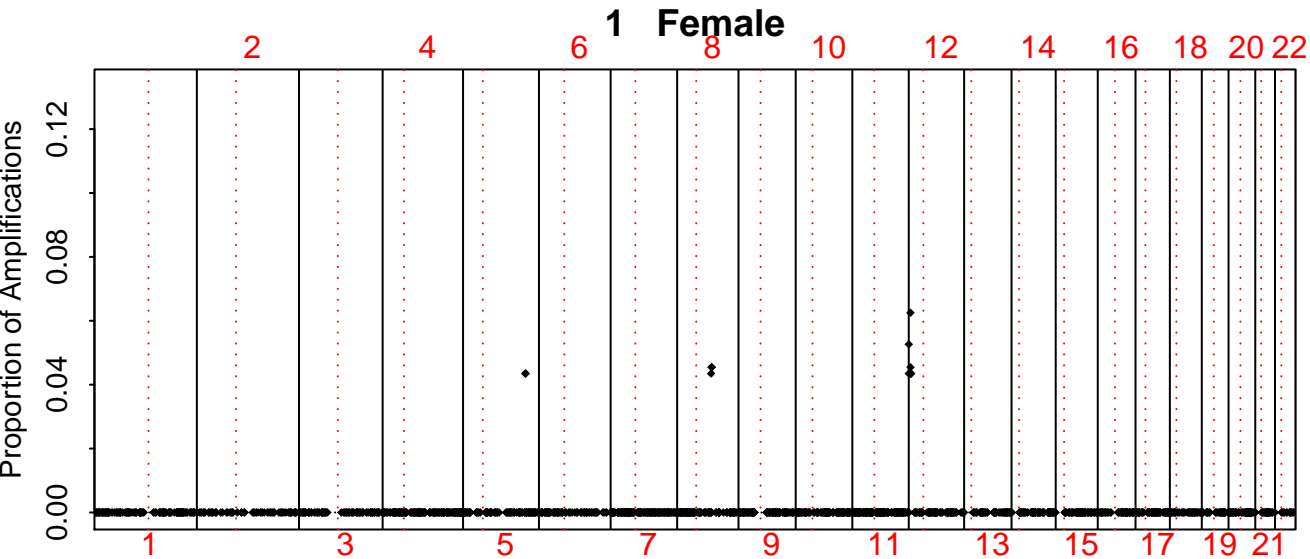
Fraction of Genome Altered 0.0240951



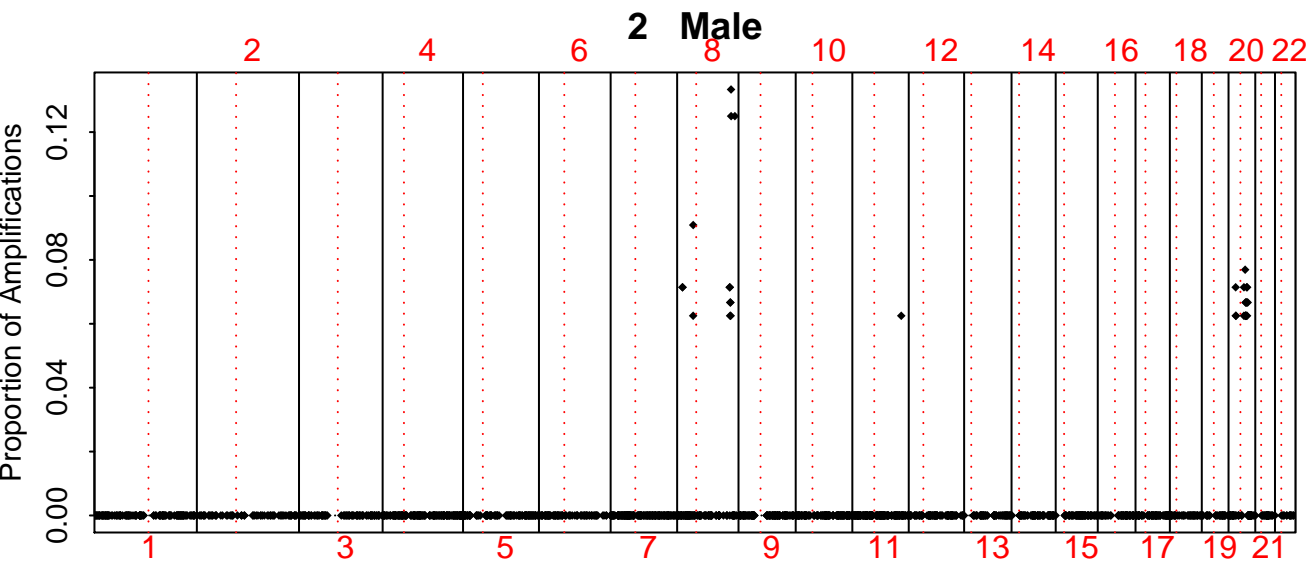




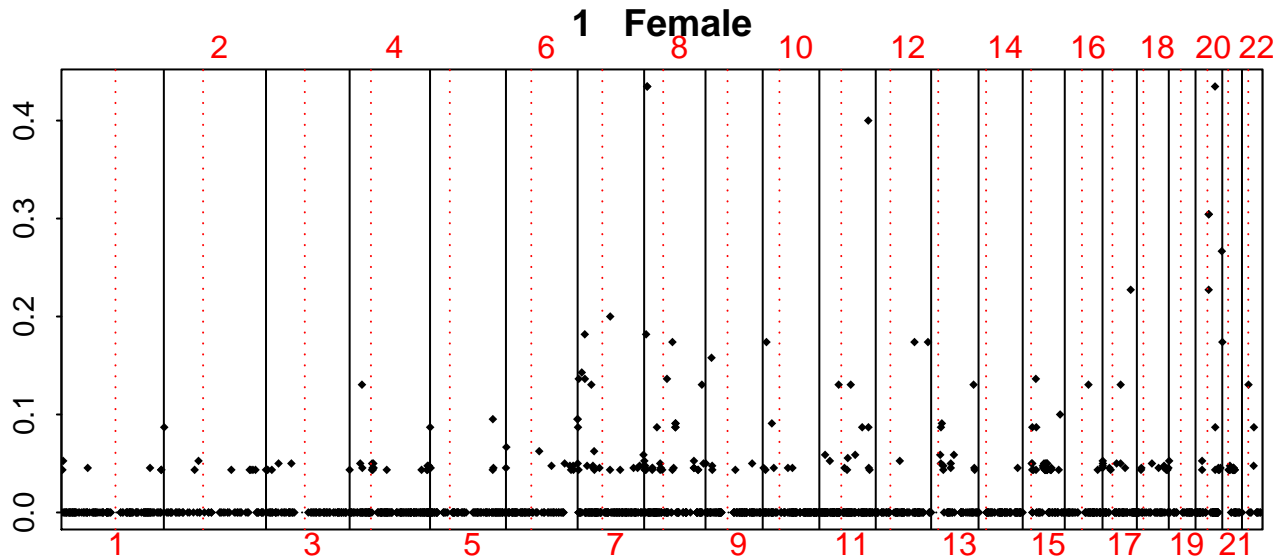
Proportion of Amplifications



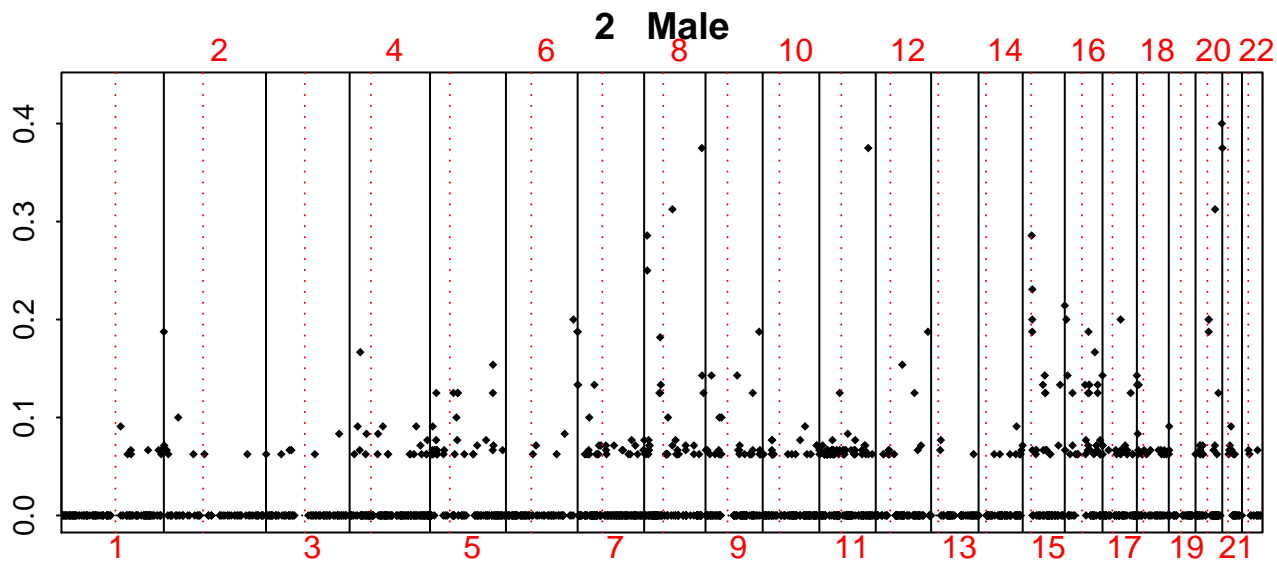
Proportion of Amplifications

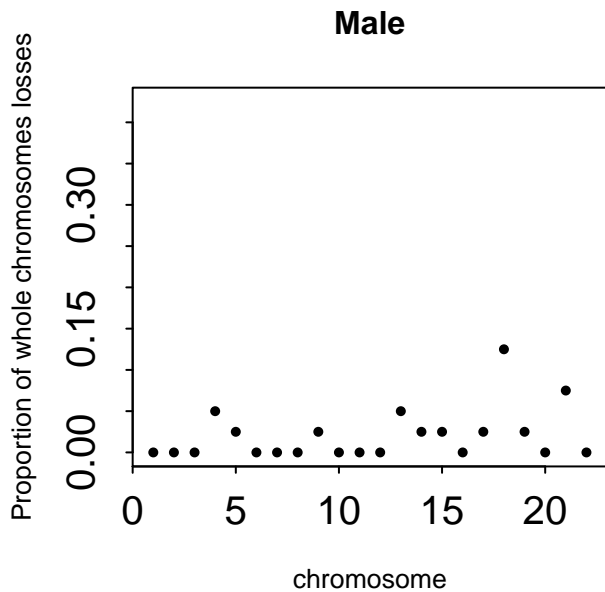
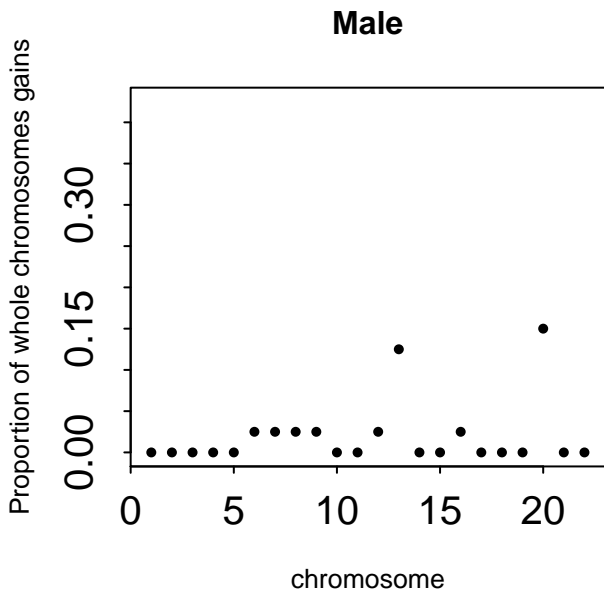
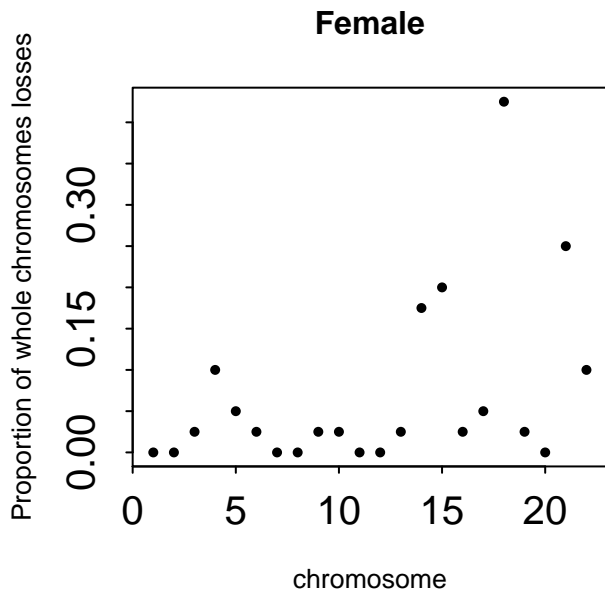
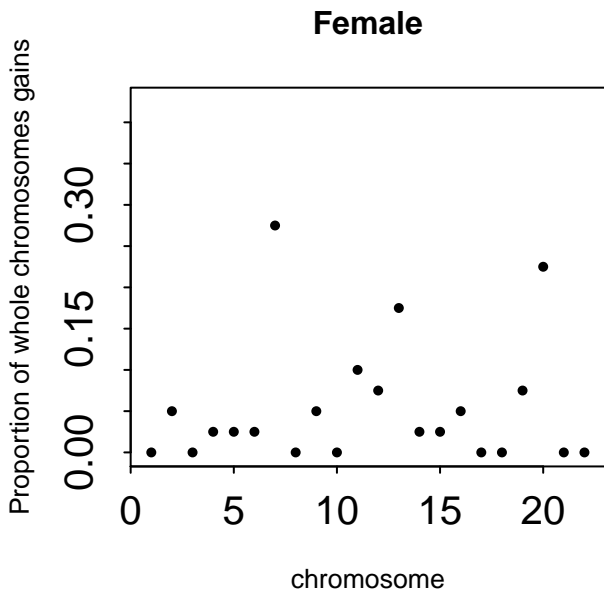


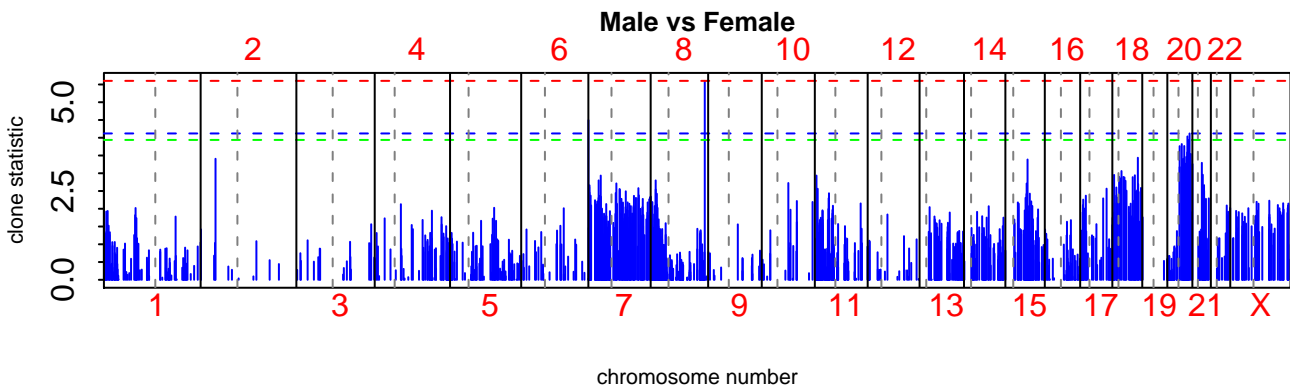
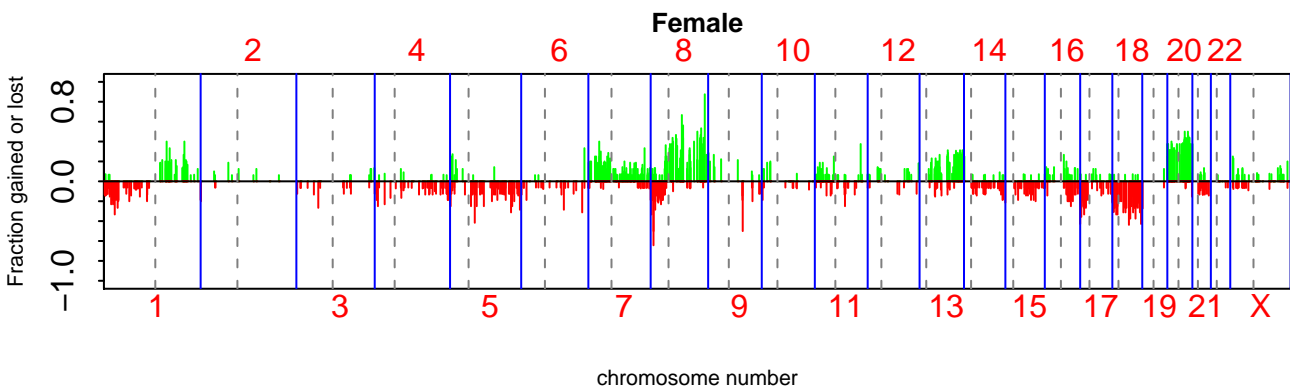
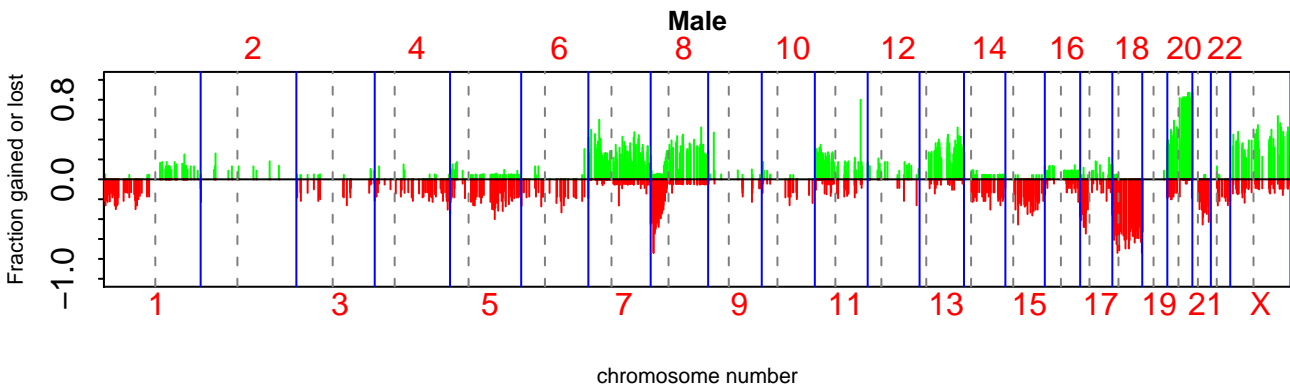
Proportion of Aberrations

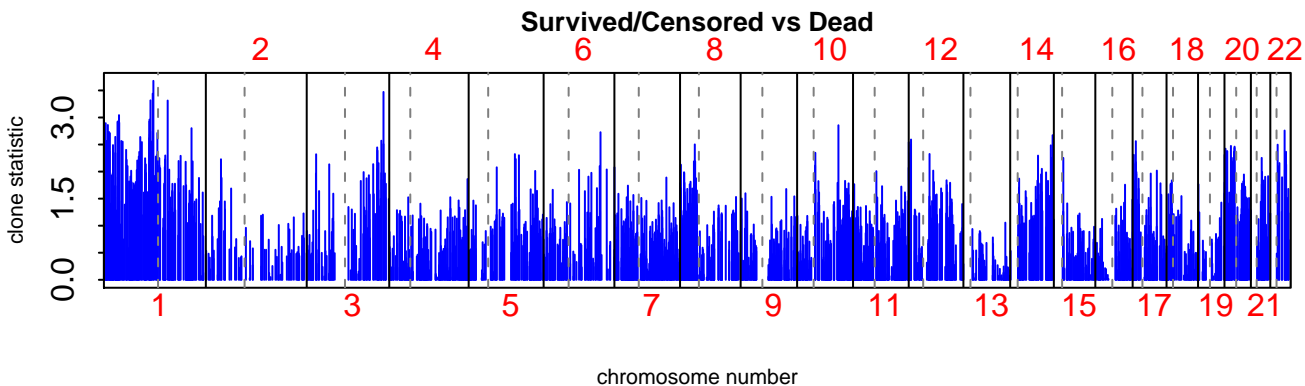
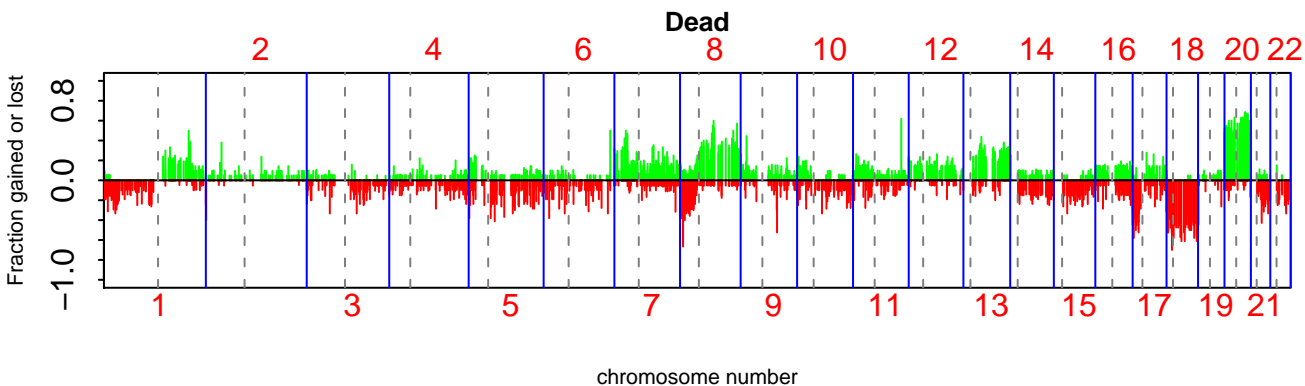
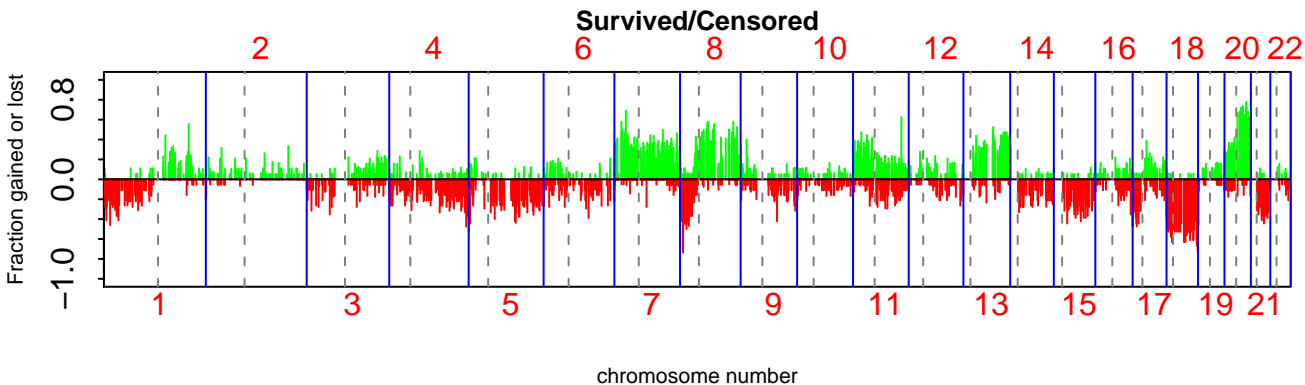


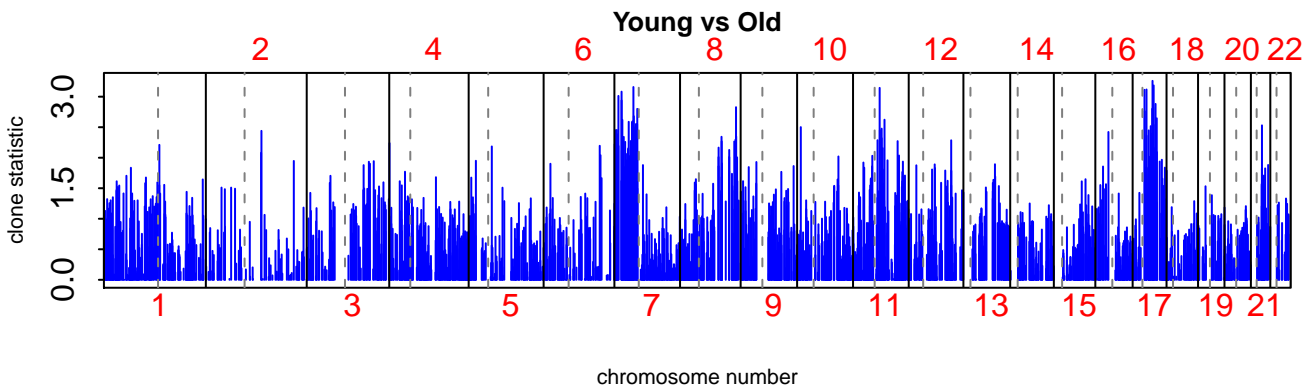
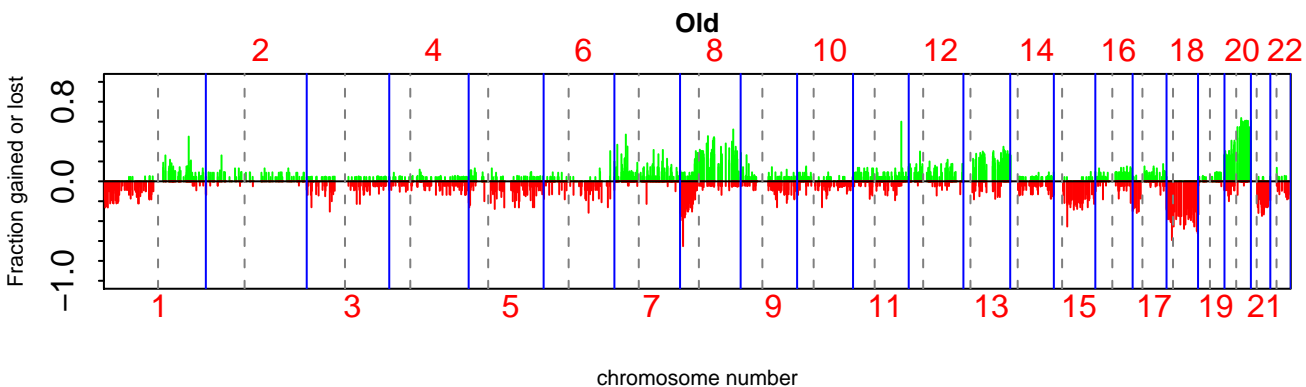
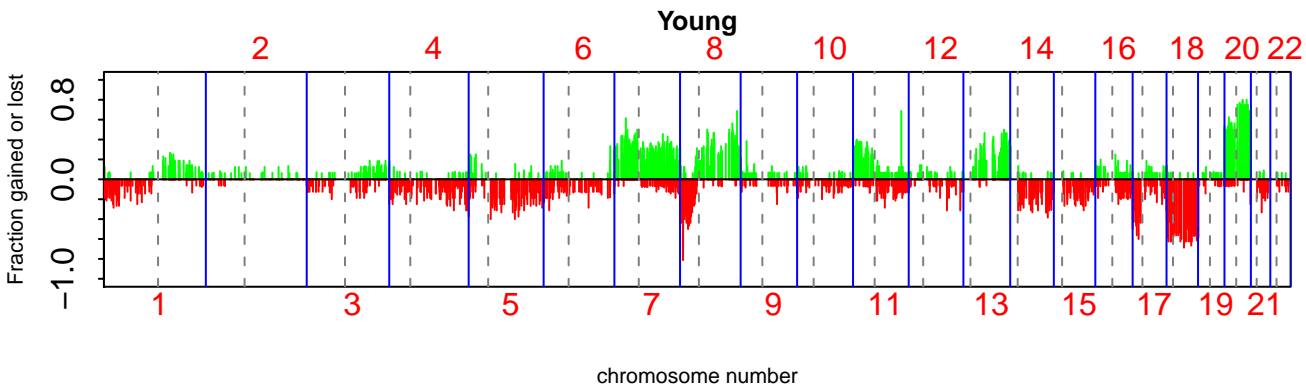
Proportion of Aberrations







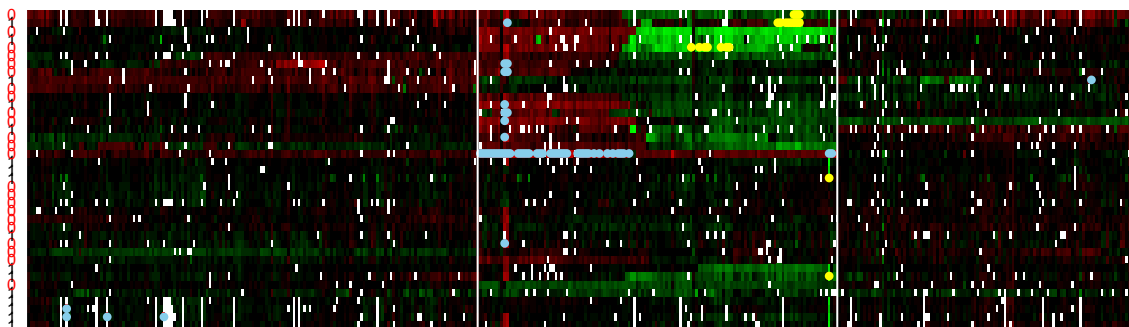




sample

Female
Male

8

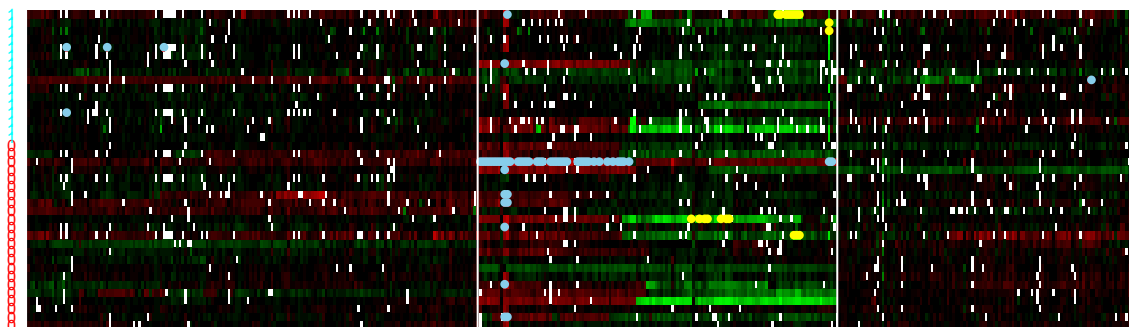


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sprocR170.0
sprocR136.0
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sprocR169.0
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sprocR152.0
sprocR154.0
sprocR161.0
sprocR395.0
sprocR340.0
sprocR112.0
sprocR177.0
sprocR196.0
sprocR100.0
sprocR359.0
sprocR193.0
sprocR174.0
sprocR218.0
sprocR212.0
sprocR217.0
sprocR232.0

clone

Female
Male

8



sprocR210.0
sprocR124.0
sprocR395.0
sprocR210.0
sprocR217.0
sprocR353.0
sprocR174.0
sprocR174.0
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sprocR112.0
sprocR77.0

4

9

clone