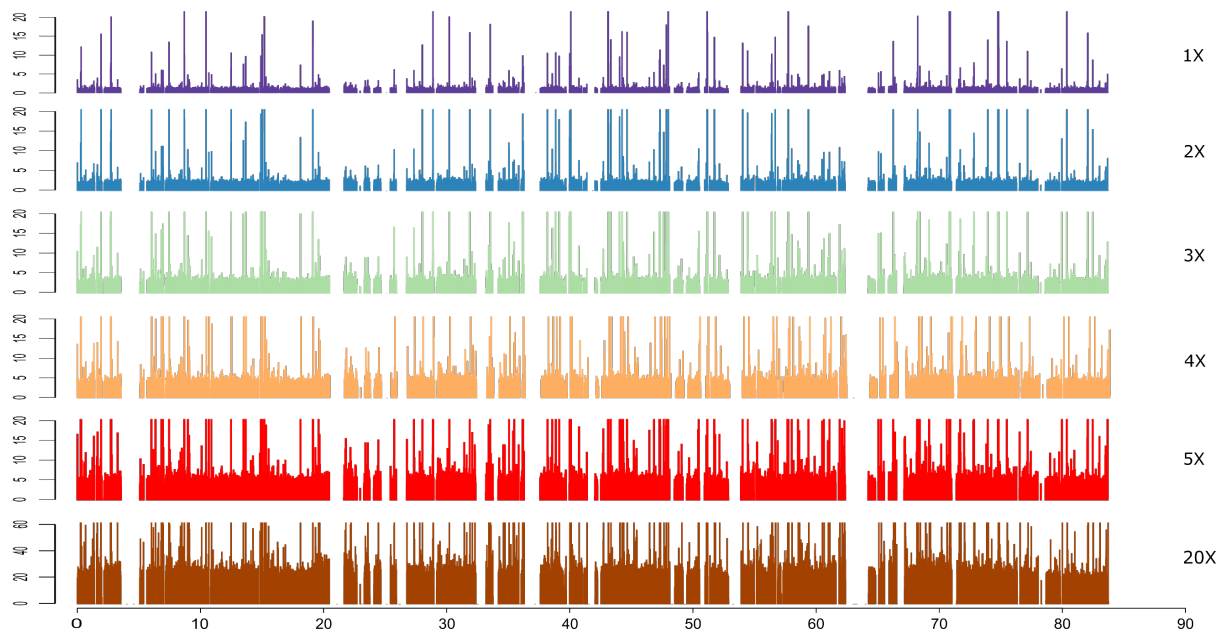
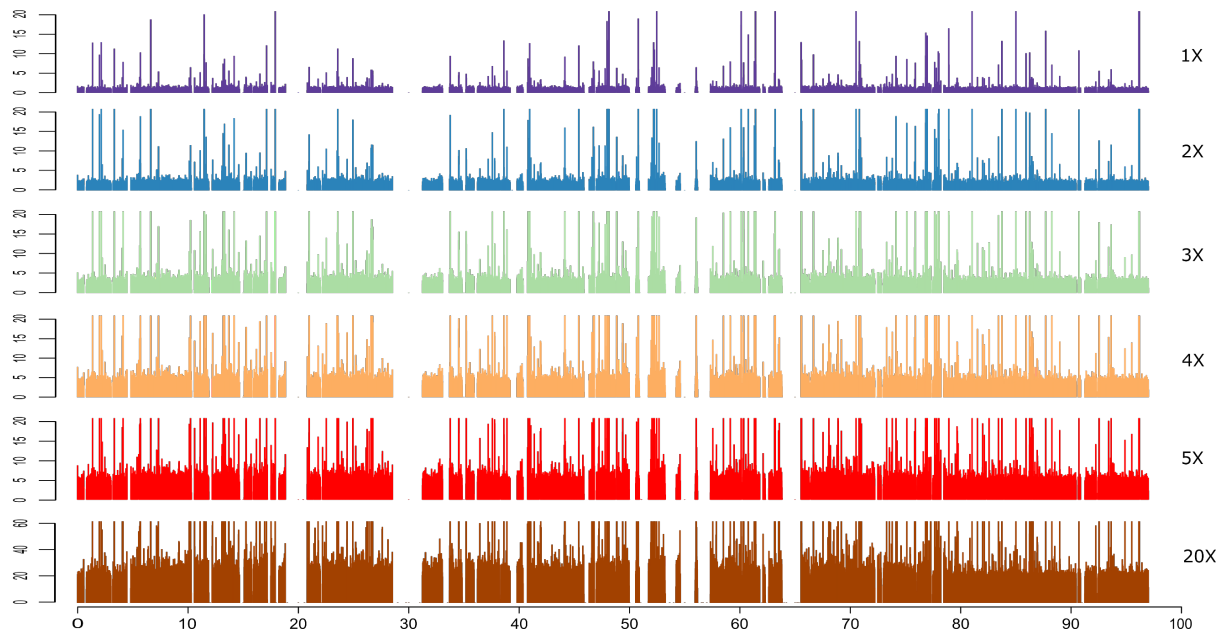


Additional file 4. Distribution of mapped read across the founder accession A by sequencing coverage. The maximum cutoff is set at DP 20 for low coverages and DP 60 for 20X data. Peaks represent regions of high sequencing within 10 Kbp windows.

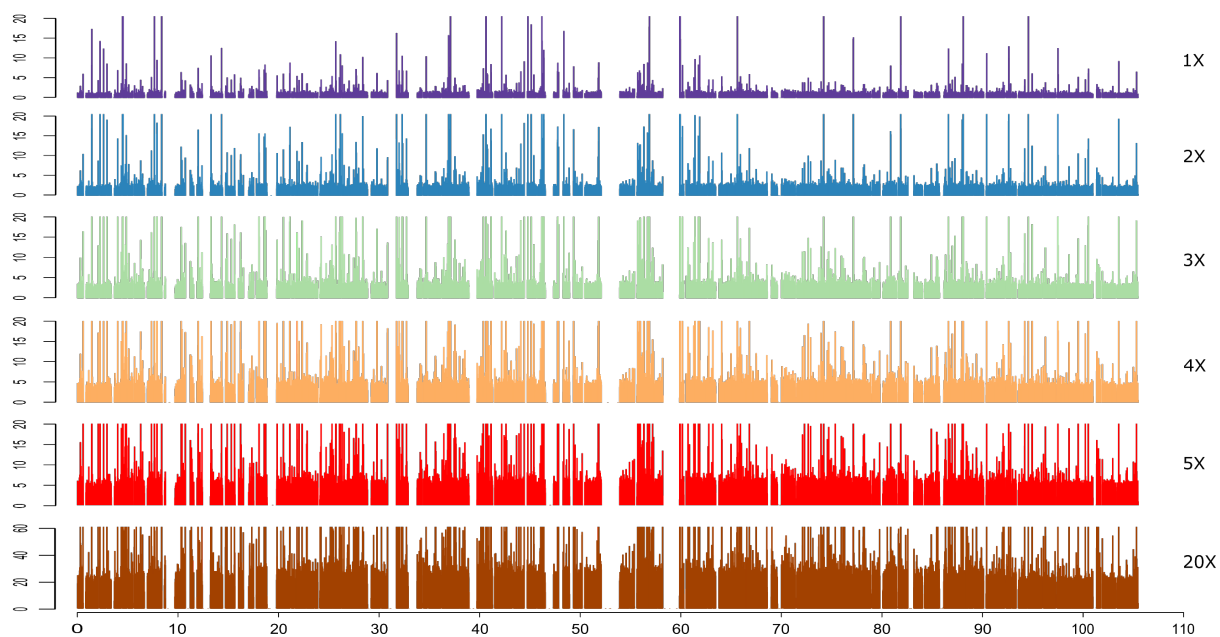
Chromosome 2



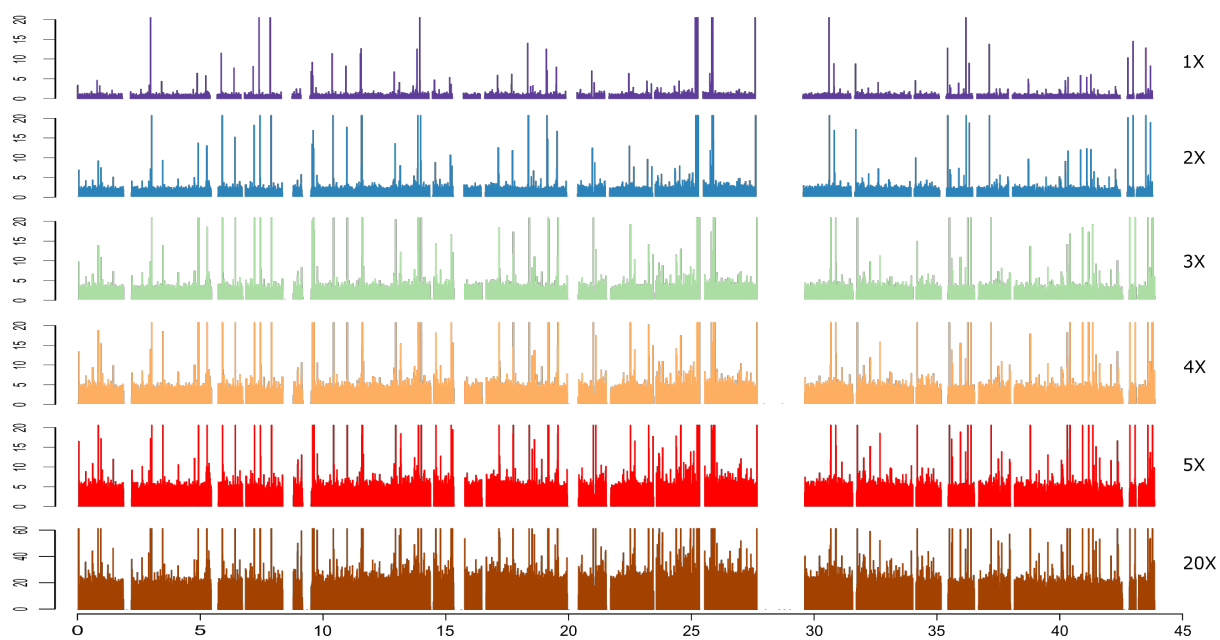
Chromosome 3



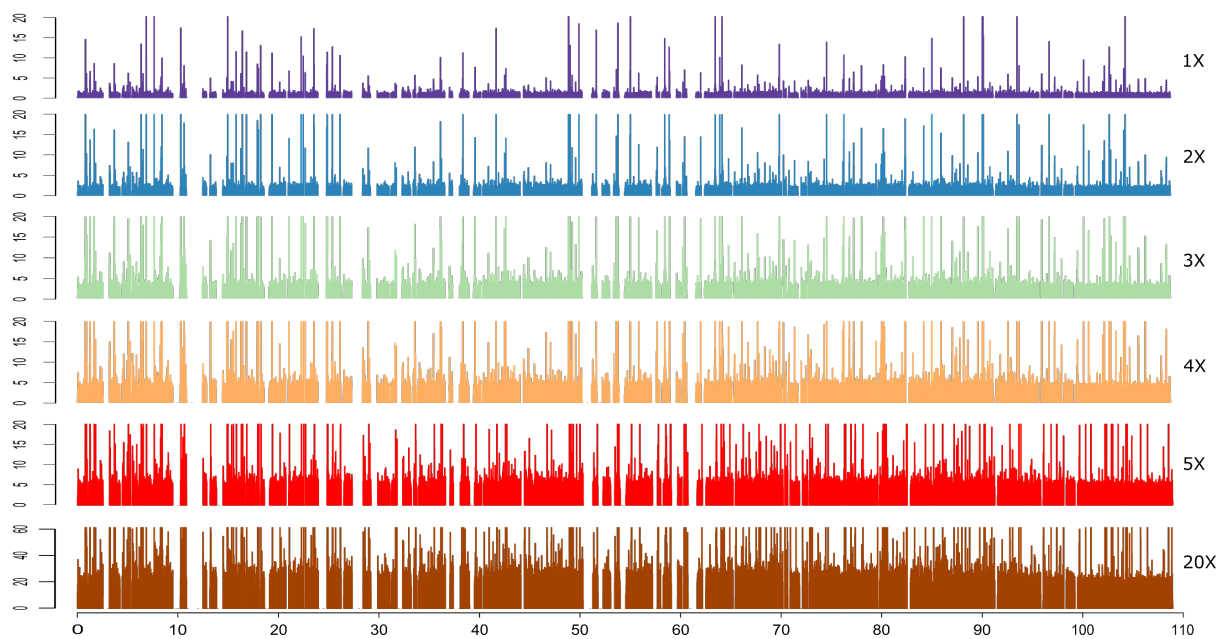
Chromosome 4



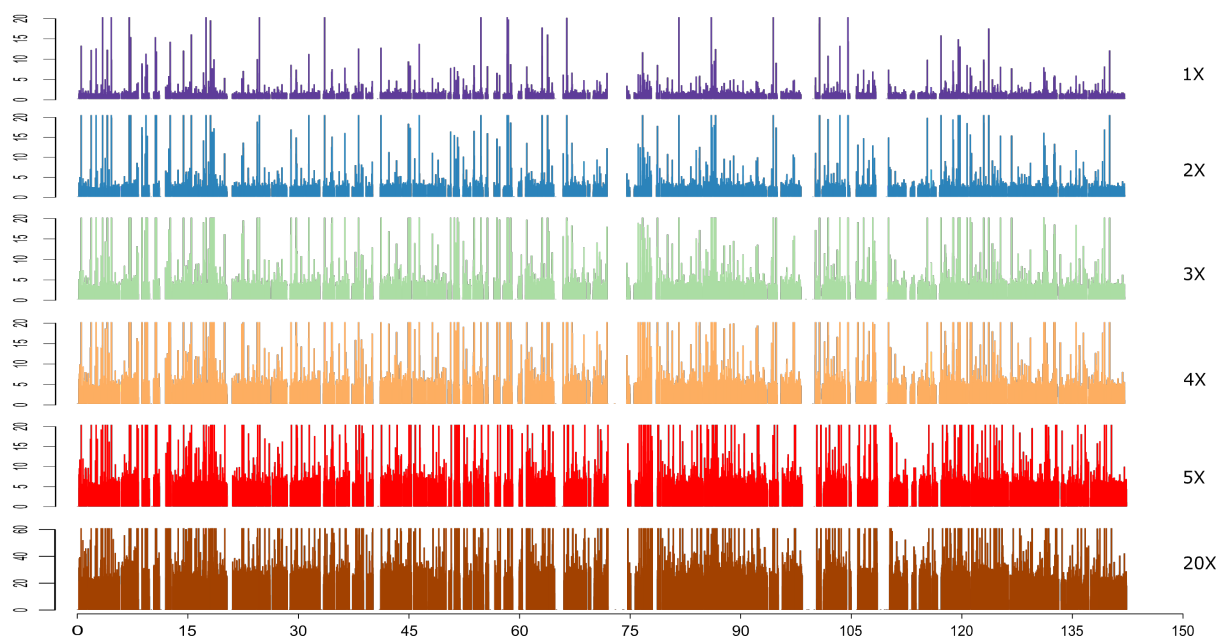
Chromosome 5



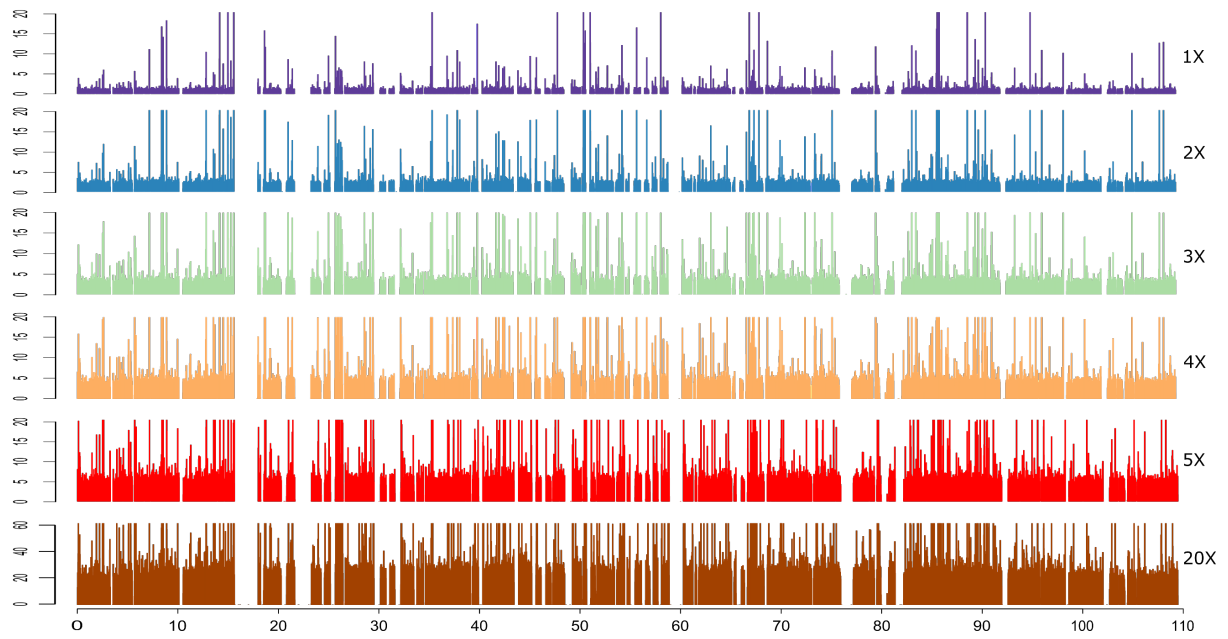
Chromosome 6



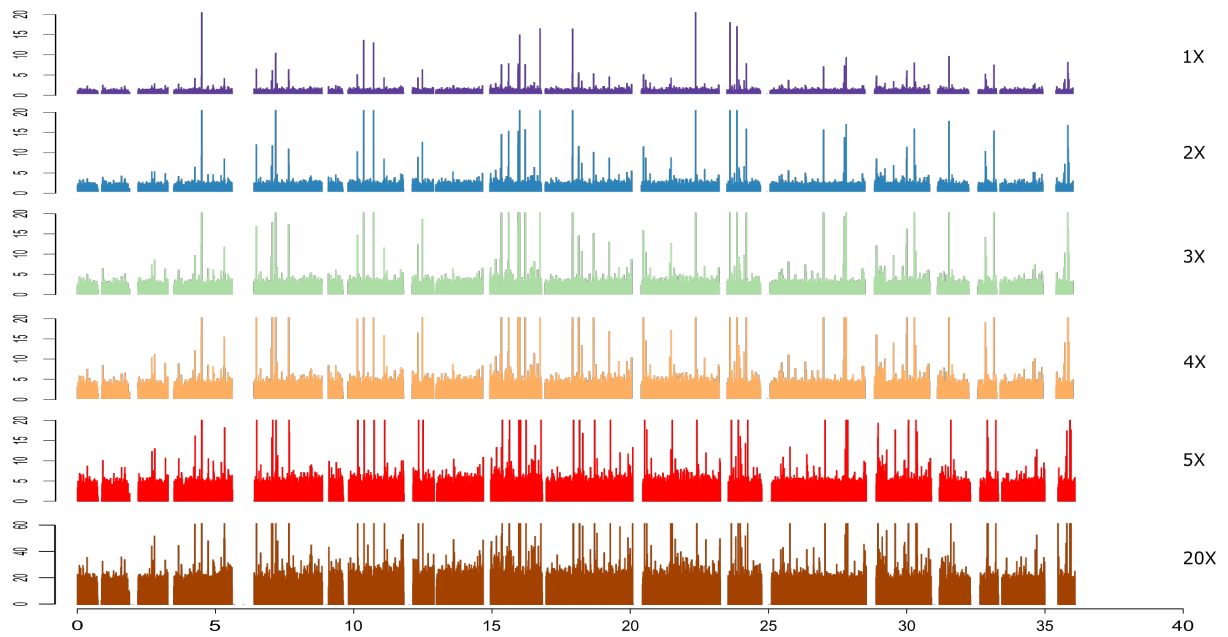
Chromosome 7



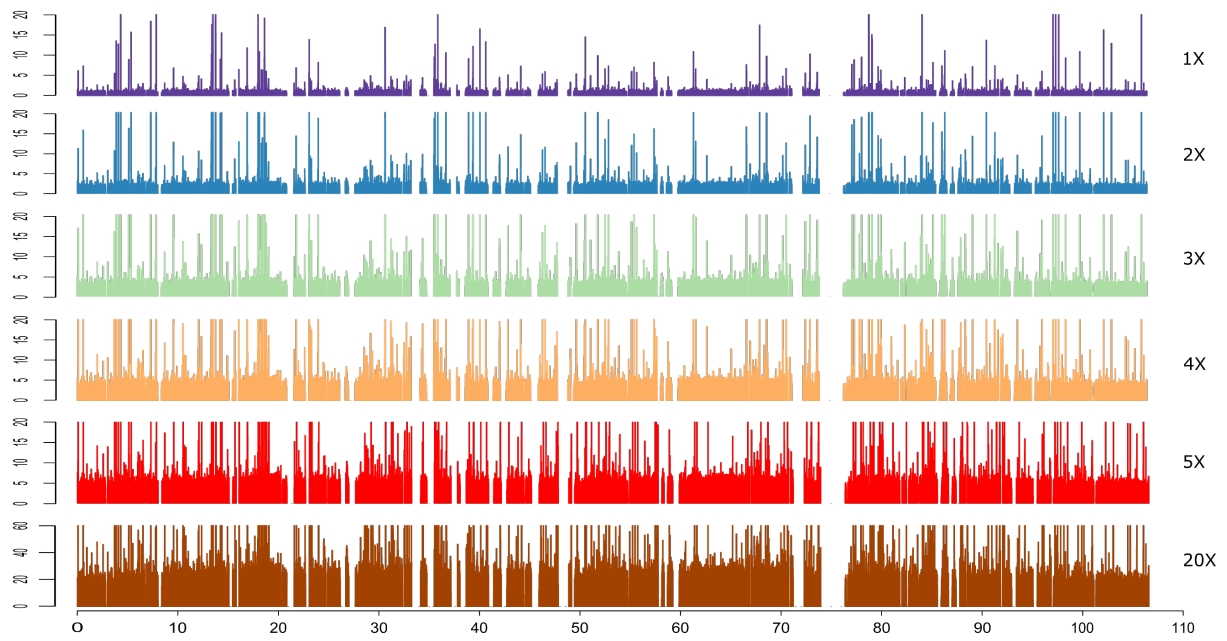
Chromosome 8



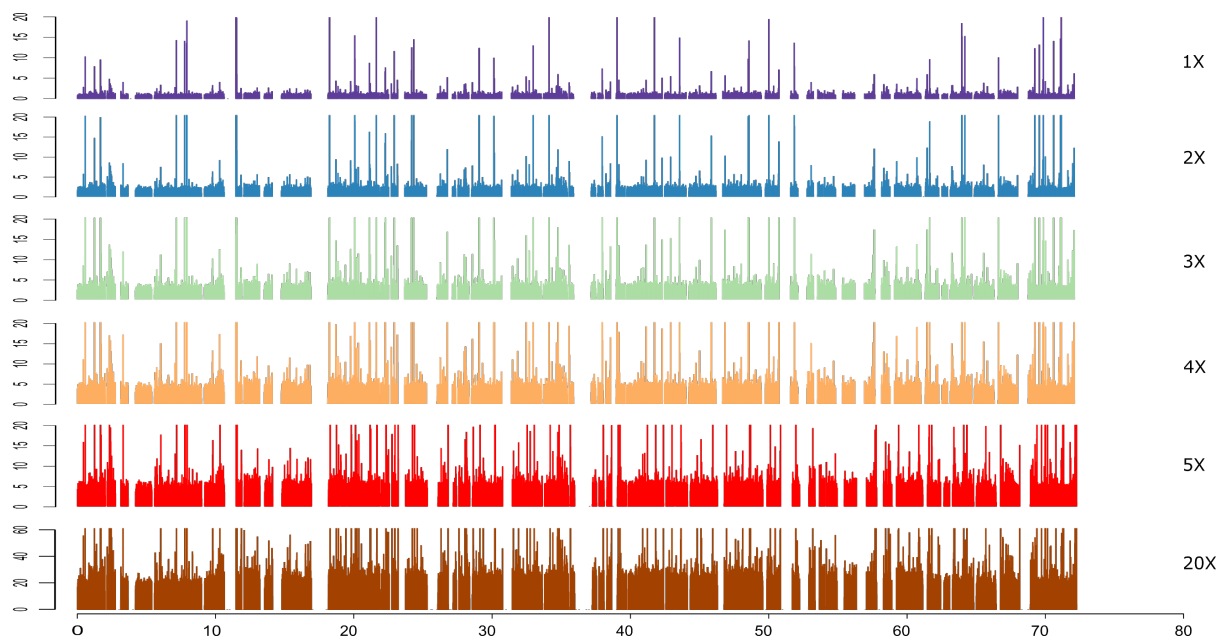
Chromosome 9



Chromosome 10



Chromosome 11



Chromosome 12

