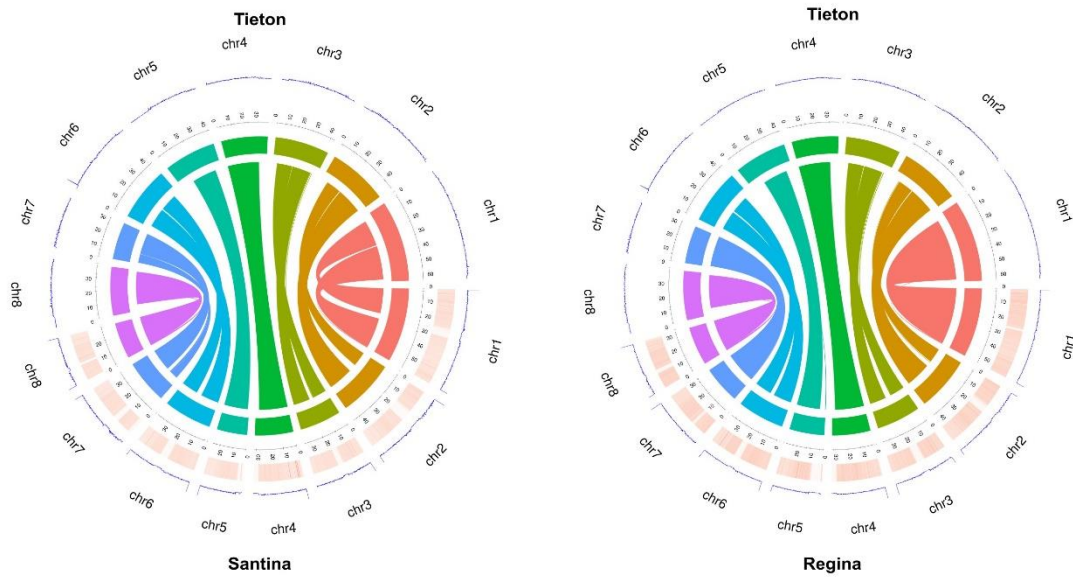




Supplementary figure 2. Santina and Regina synteny with Tieton genome assembly. In left panel is represented the synteny between Tieton (top) and Santina (bottom) genome assemblies. Blue histograms show the frequency of telomeric repeat sequences, while the red heatmap indicates the frequency of repetitive elements along the chromosomes. Colored bars correspond to chromosomes, and connecting lines represent syntenic relationships. In right panel is represented the synteny between Tieton (top) and Regina (bottom) genome assemblies.