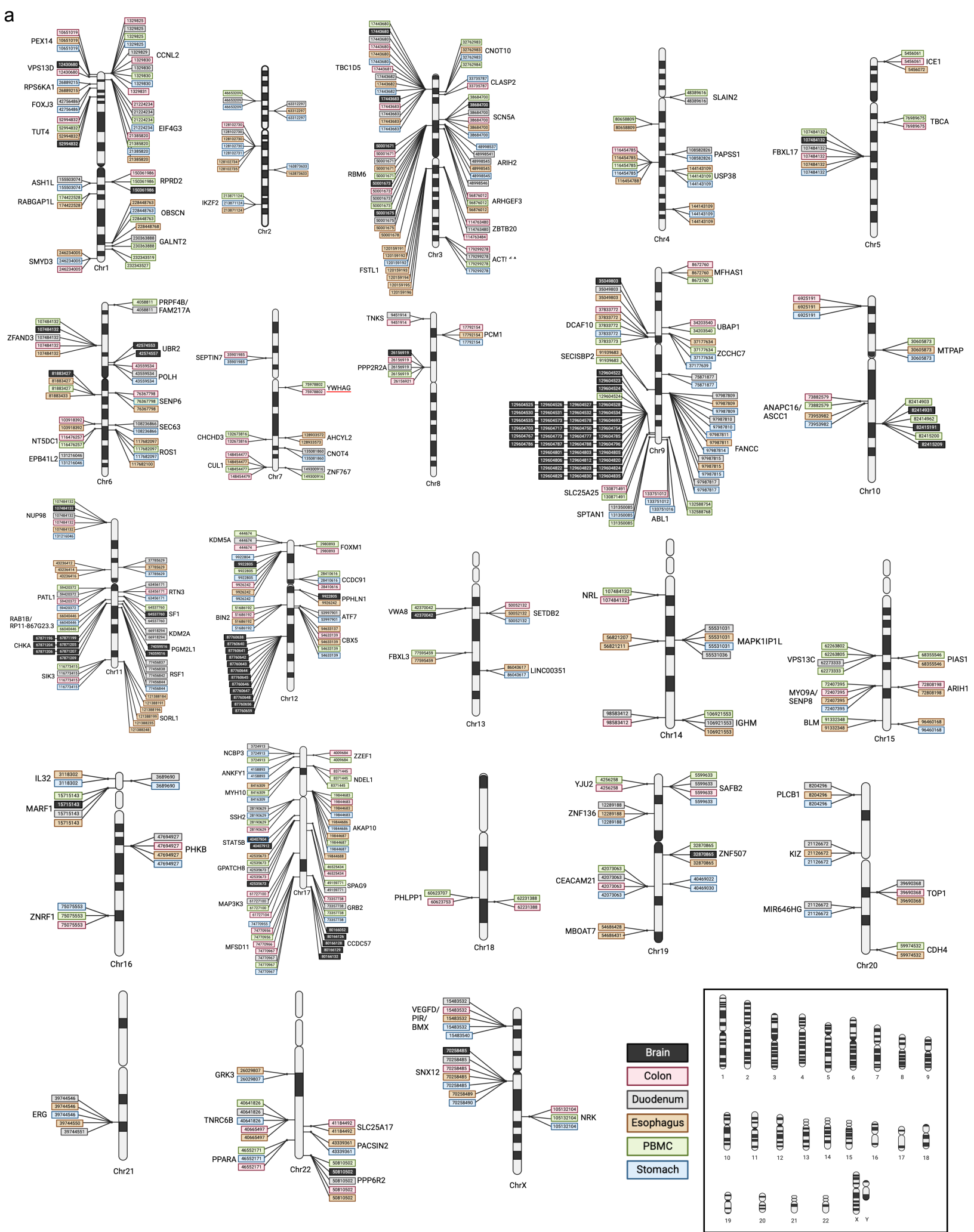
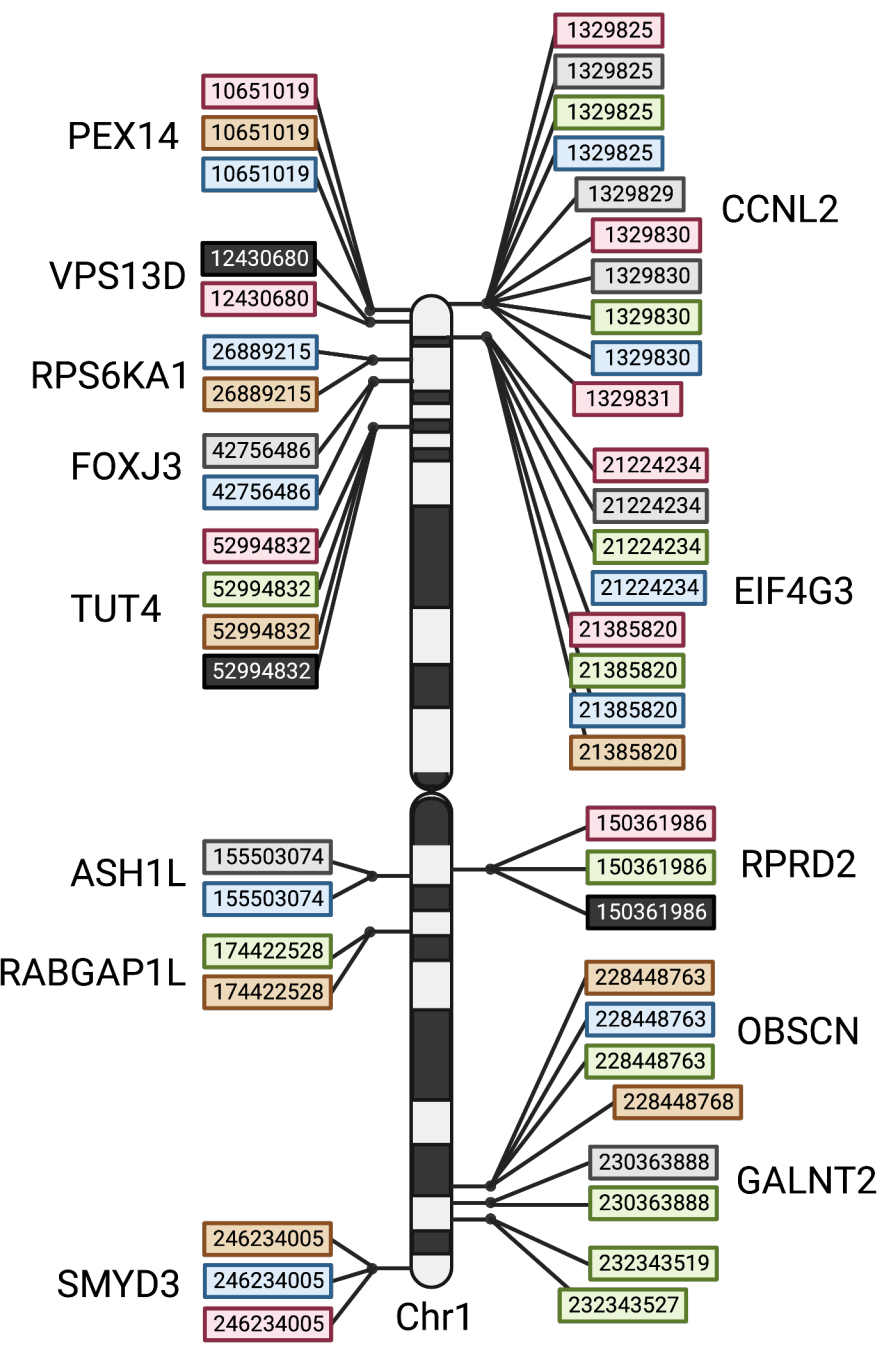


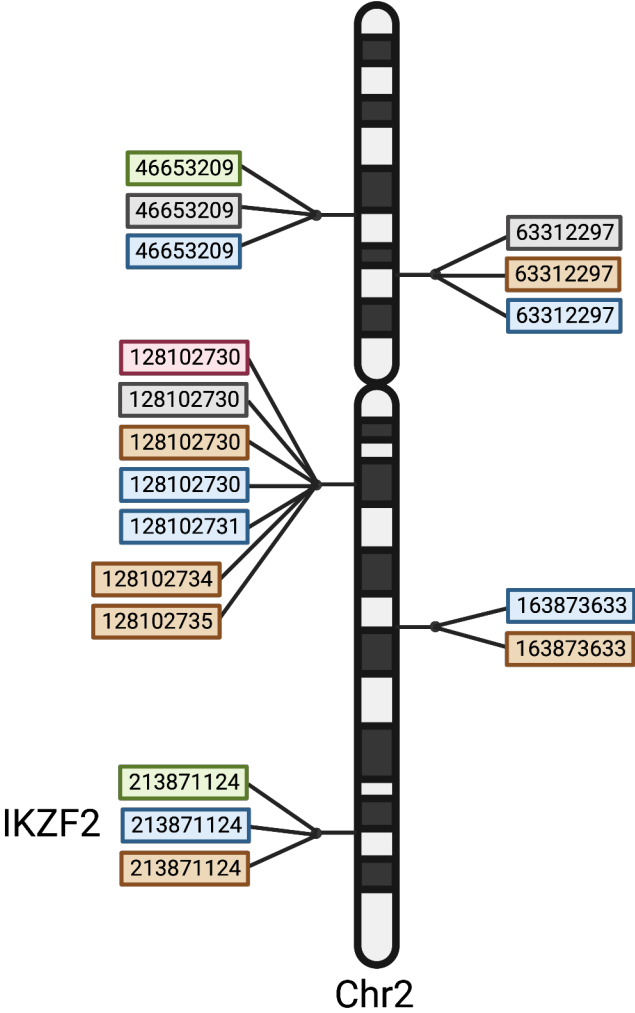
Supplementary Figure 1: Chromosomal distribution of HIV-1 integration sites across tissues.

(a) Overview map of HIV-1 integration site placements across the human genome, showing locations of tissue-specific integration site hotspots. Each cluster represents integration events identified from different anatomical sites. If a hotspot is located within a gene, the corresponding gene name is labeled adjacent to the cluster. Numbers in colored boxes indicate chromosomal coordinates. (b–x) Enlarged views of HIV-1 integration site placements for each individual chromosome.

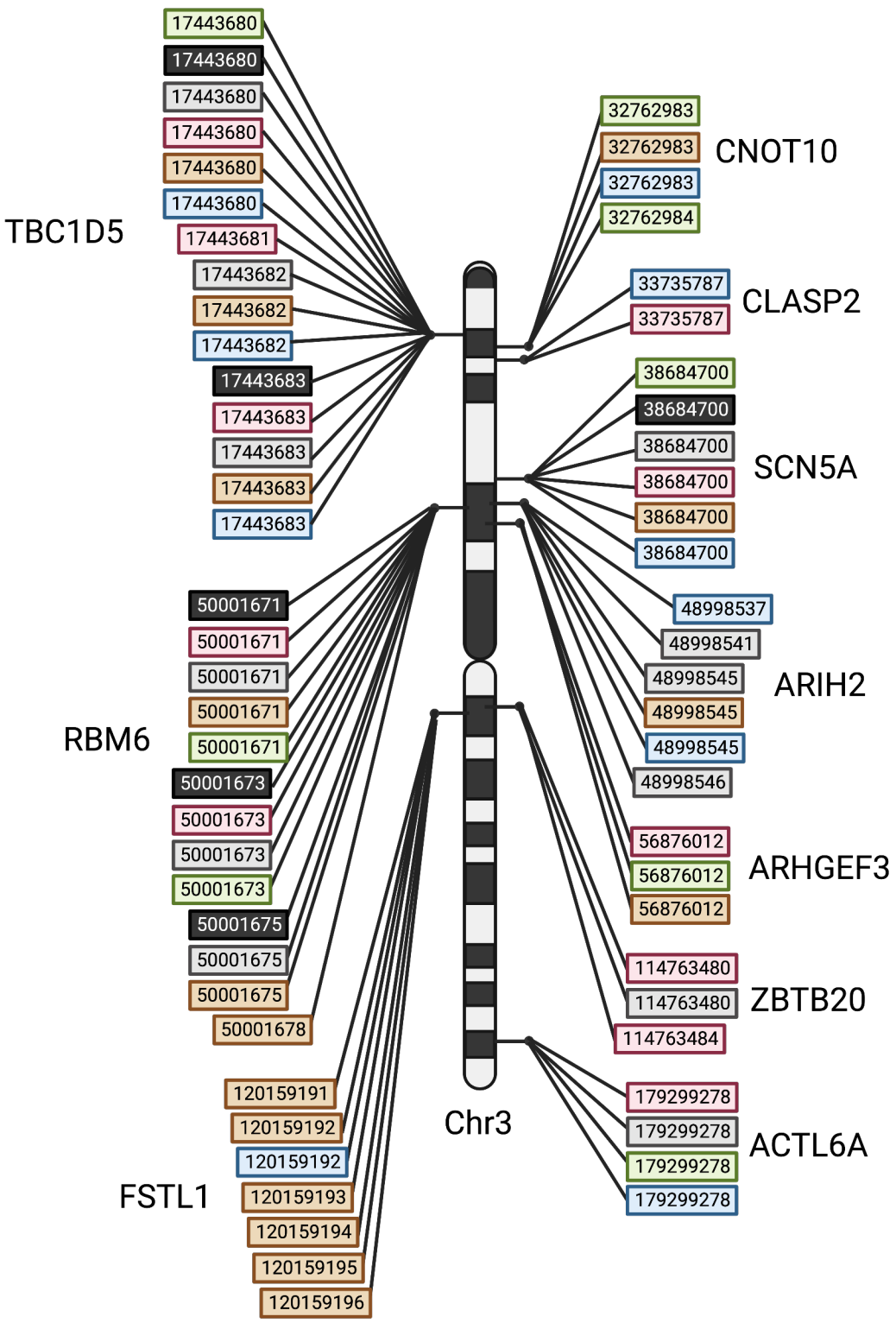


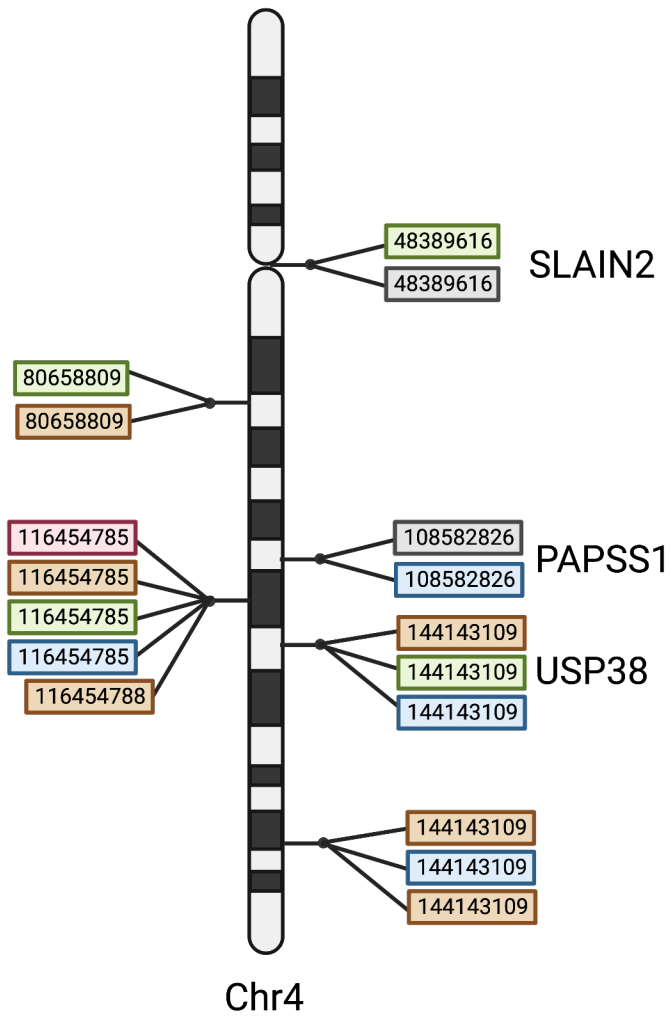
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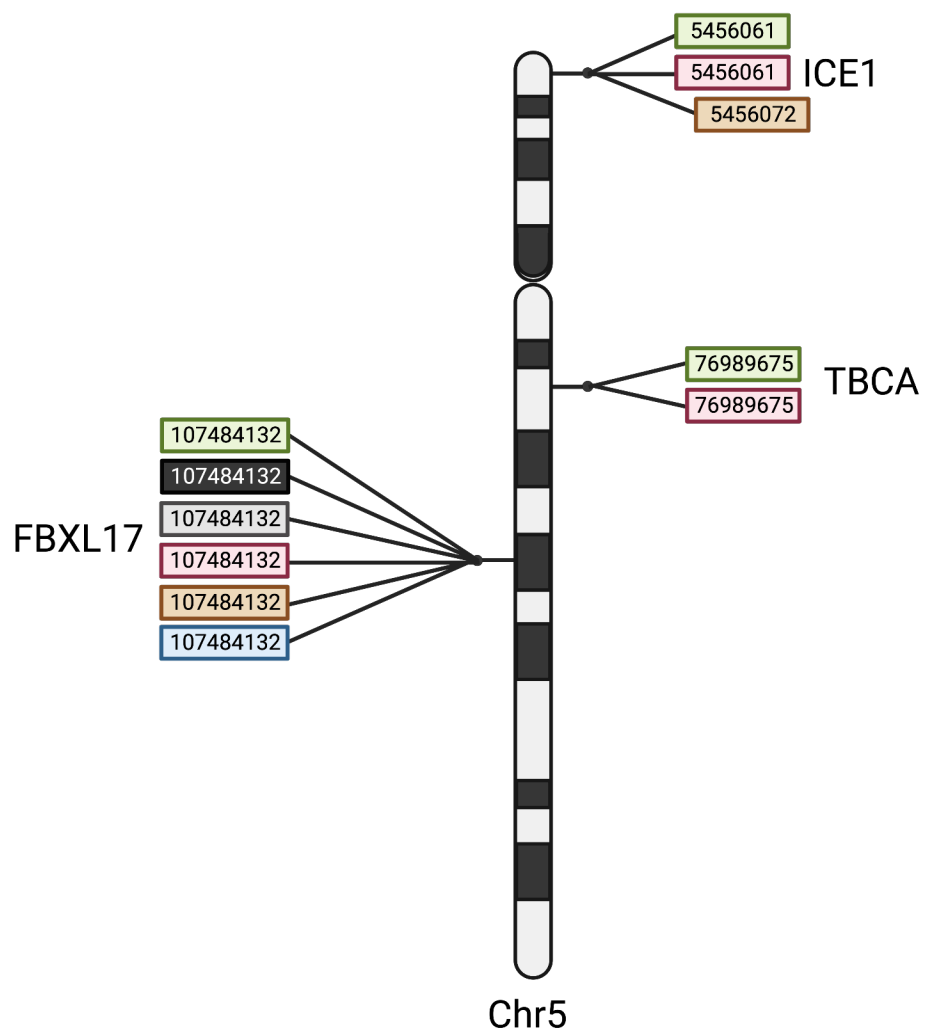


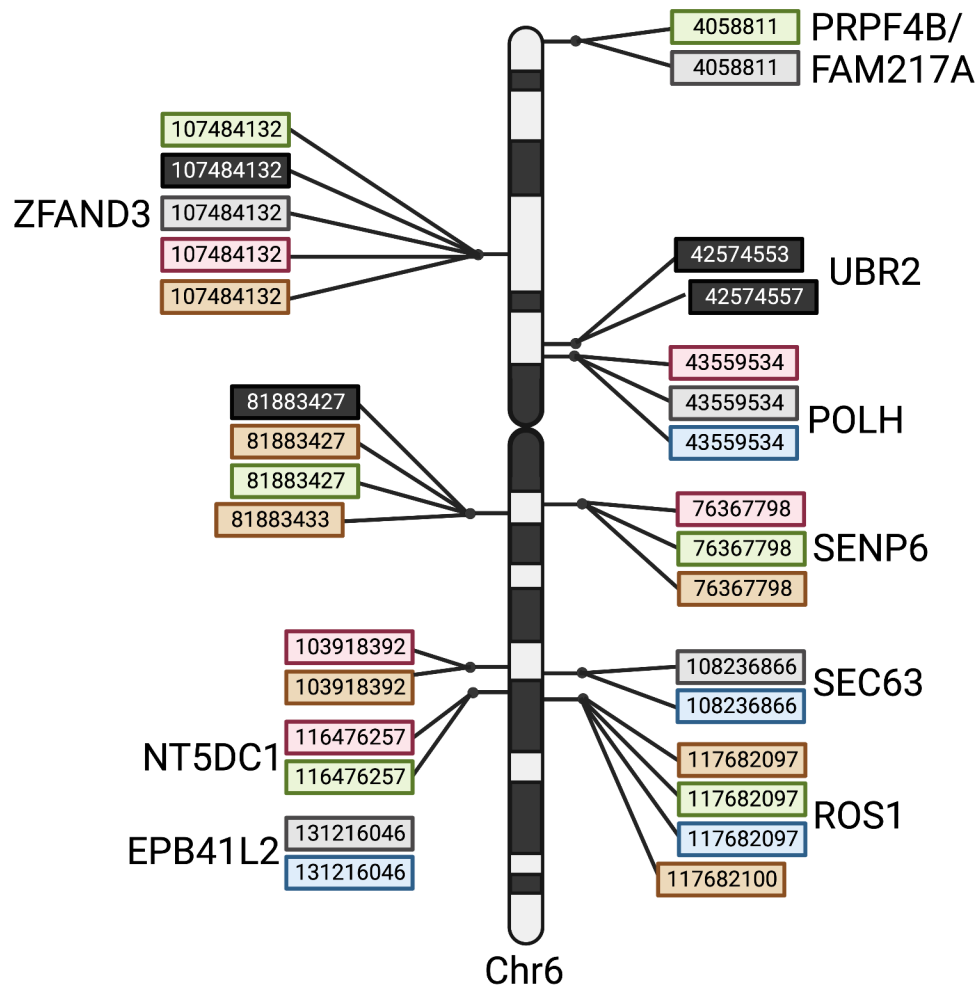


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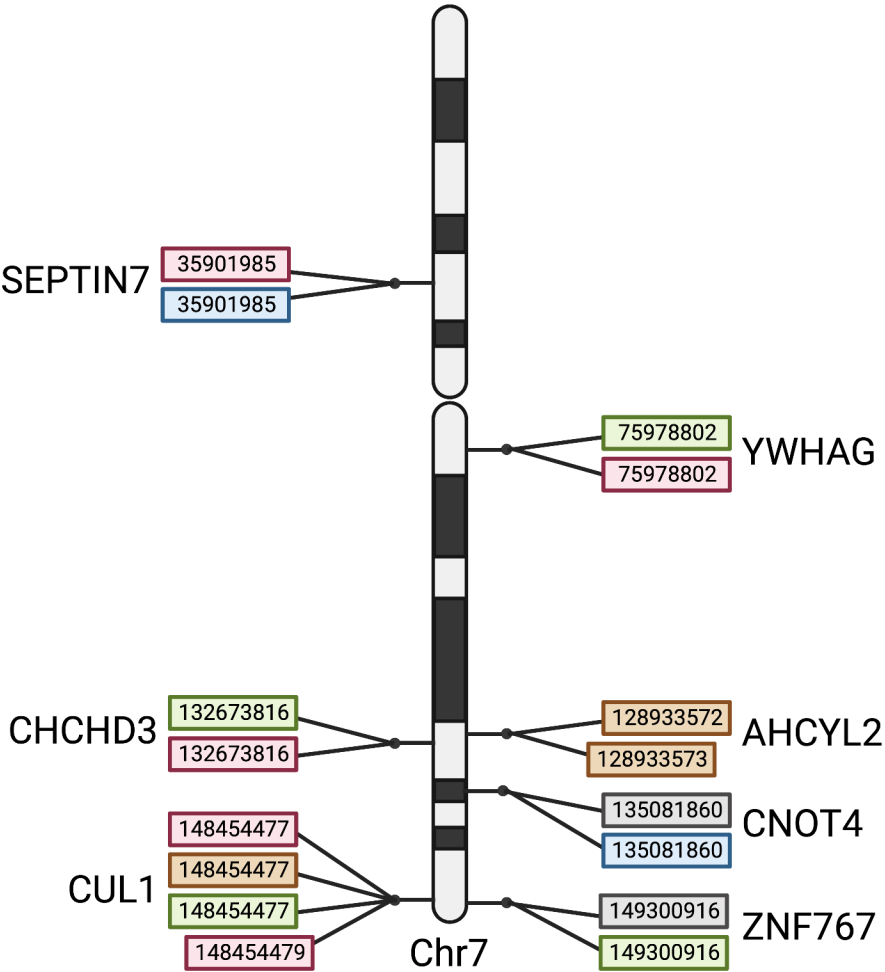


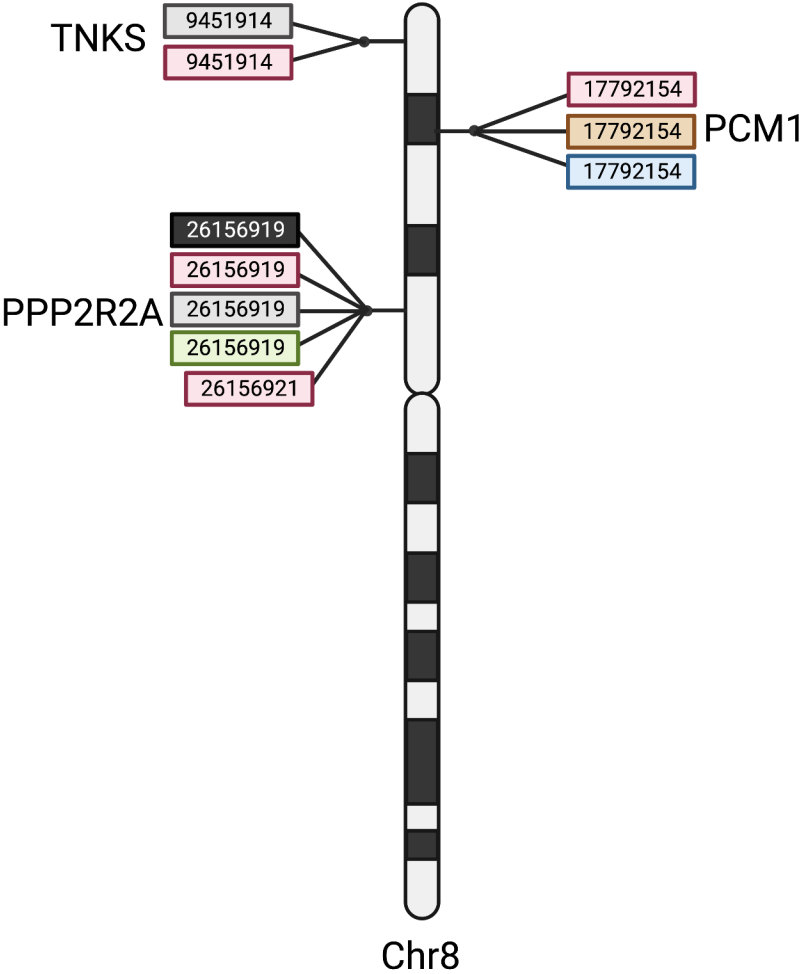


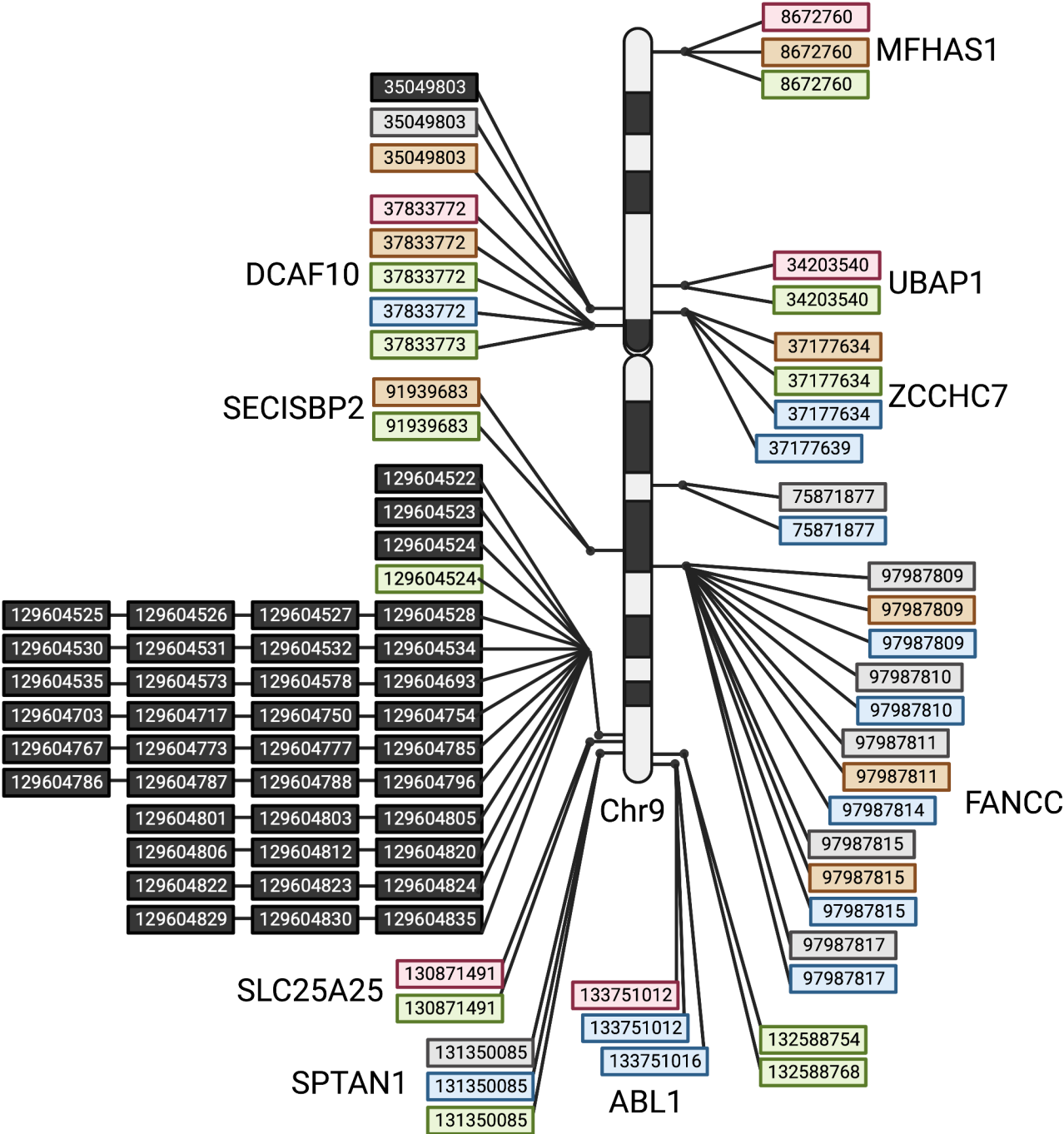


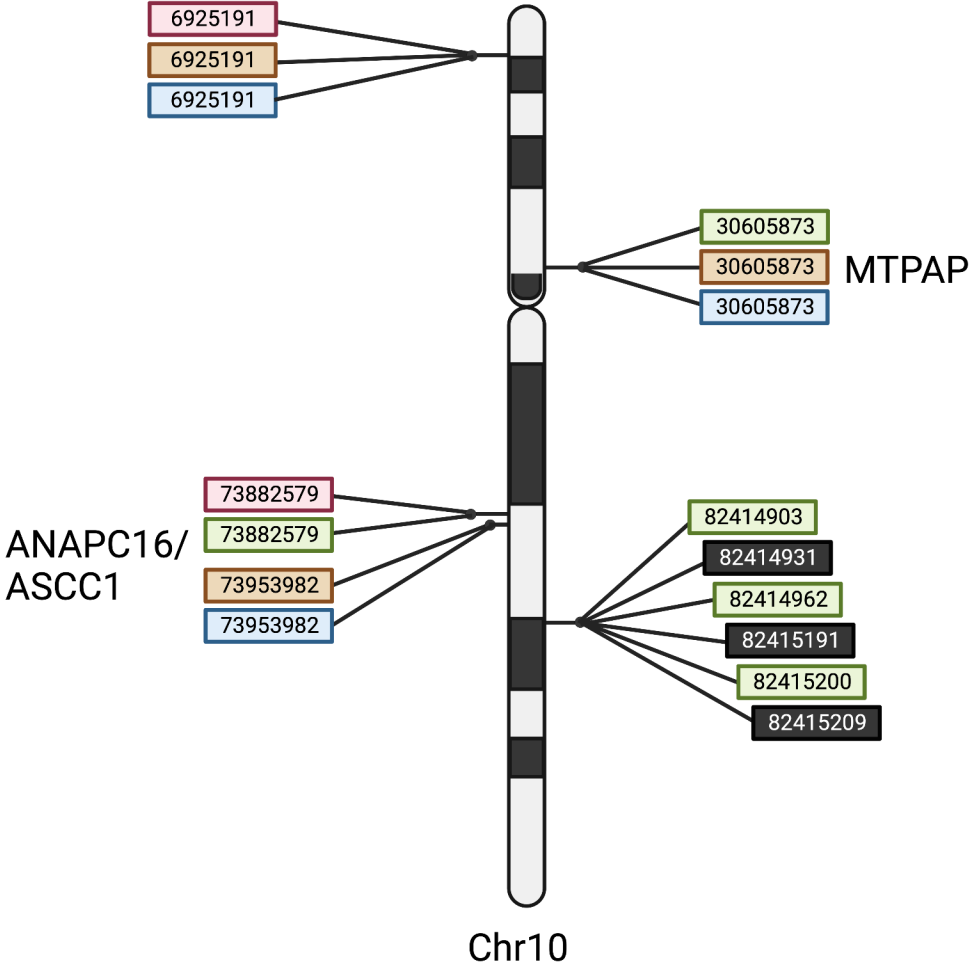


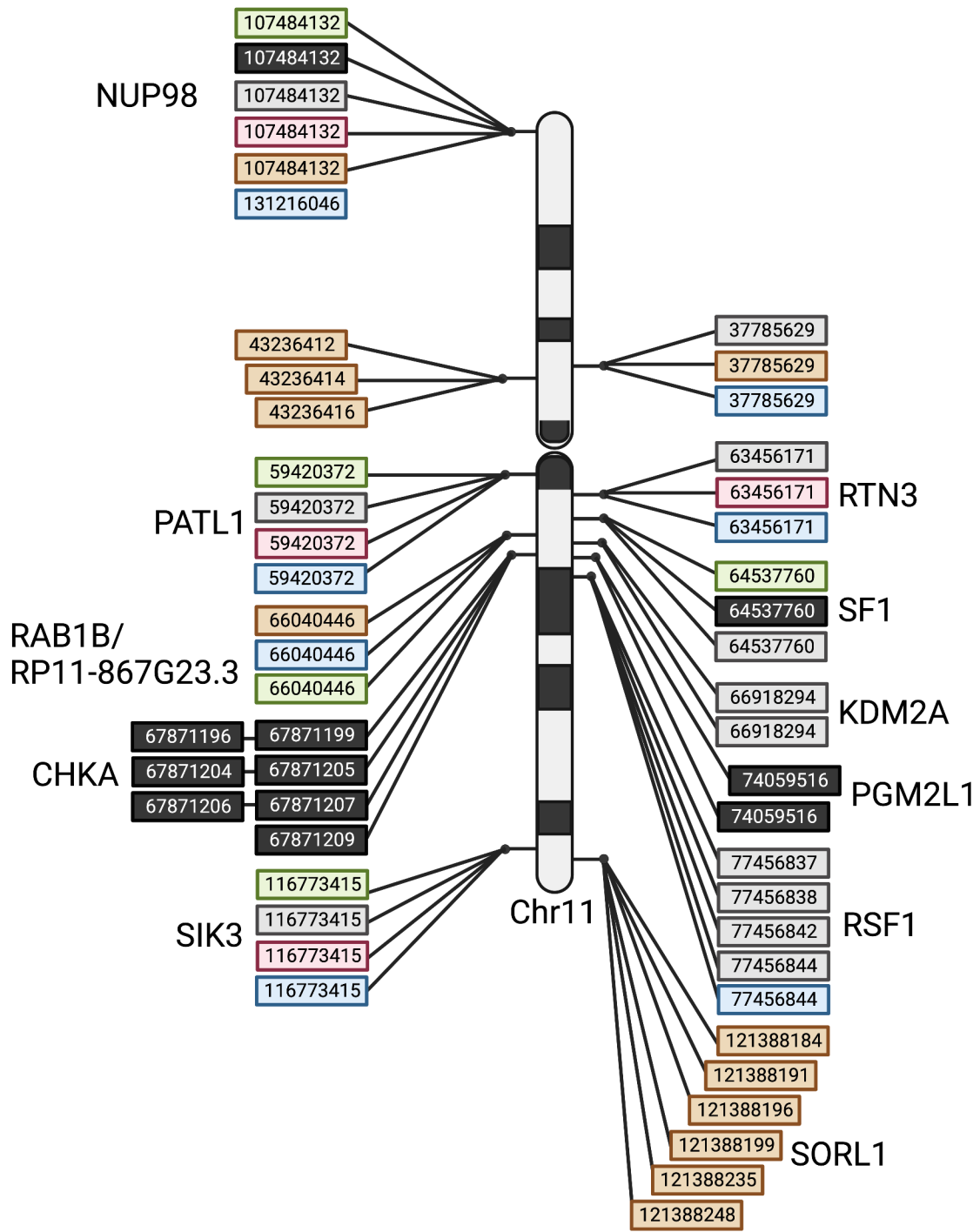
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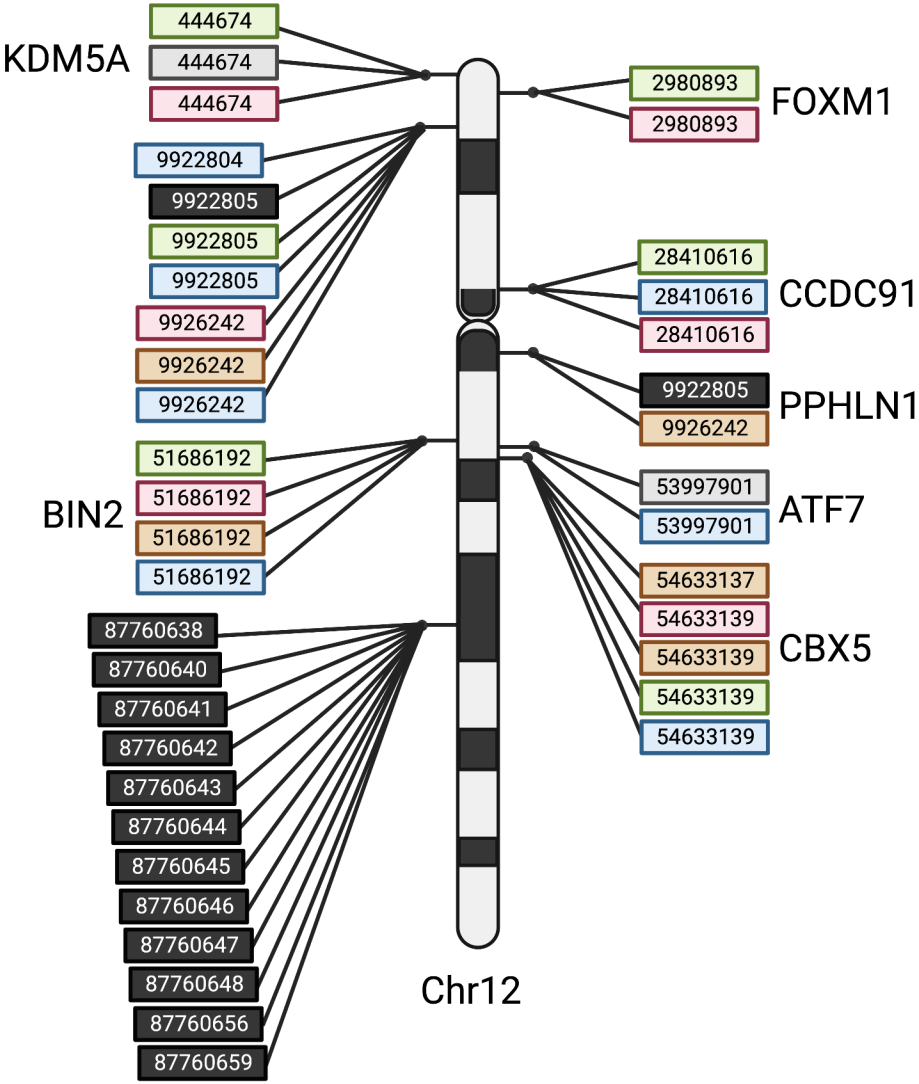


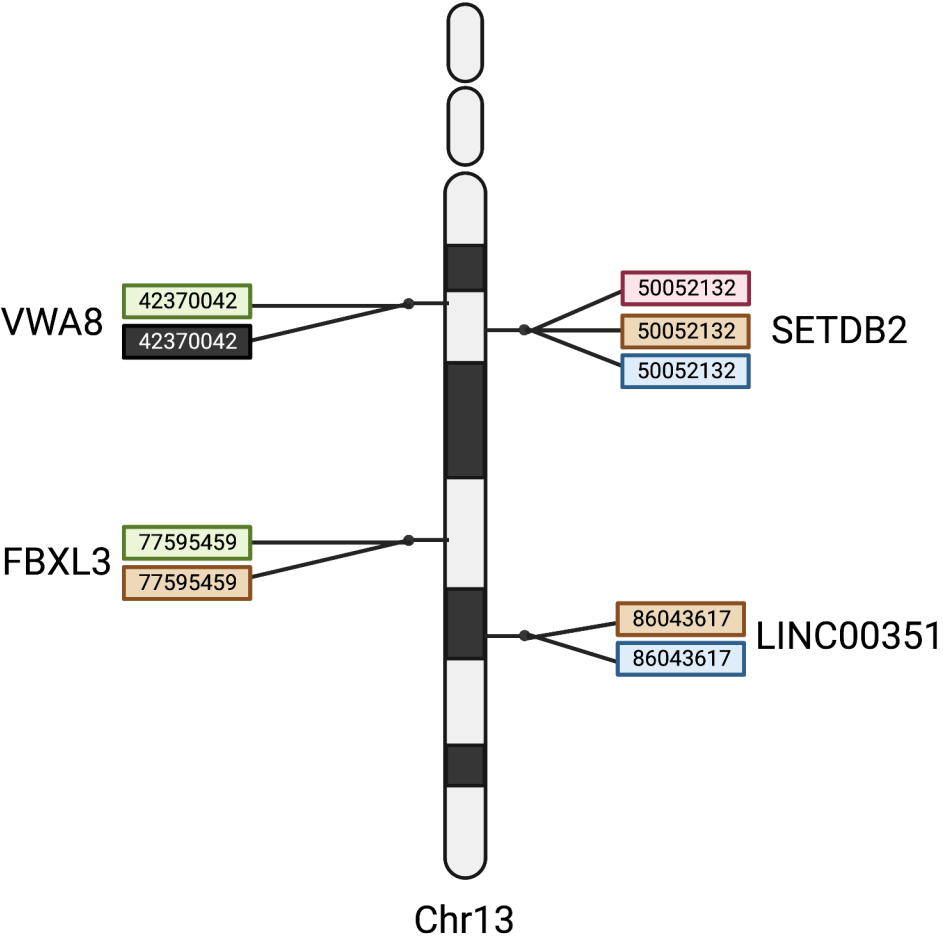






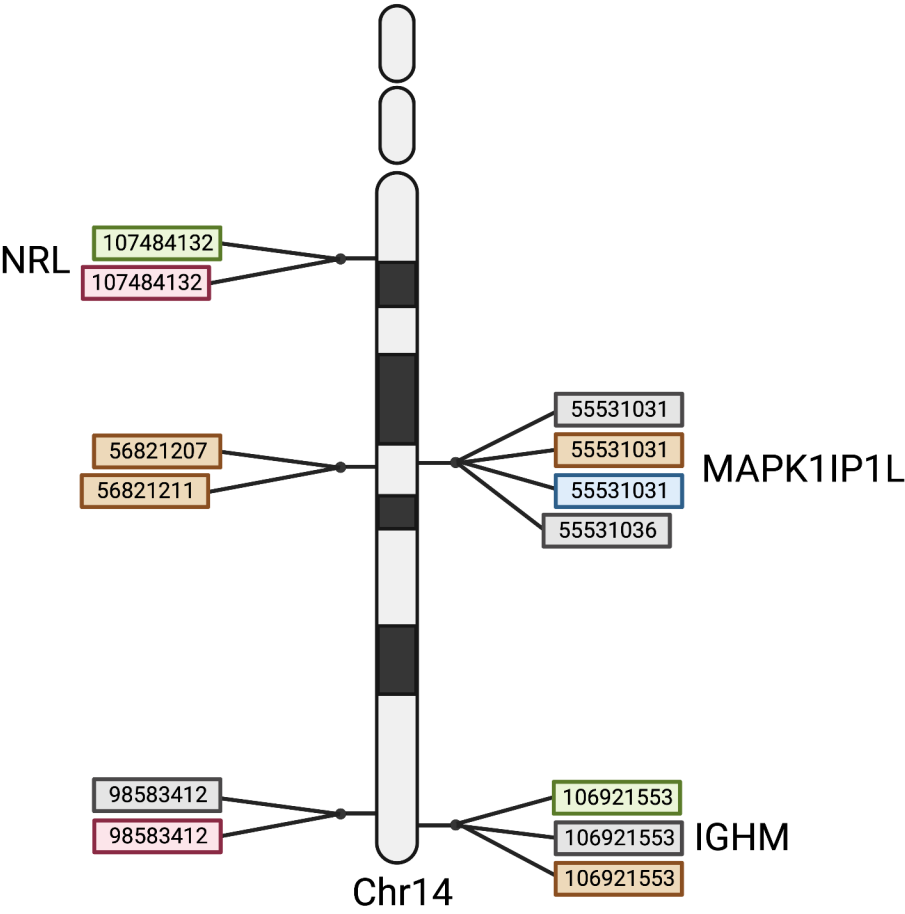


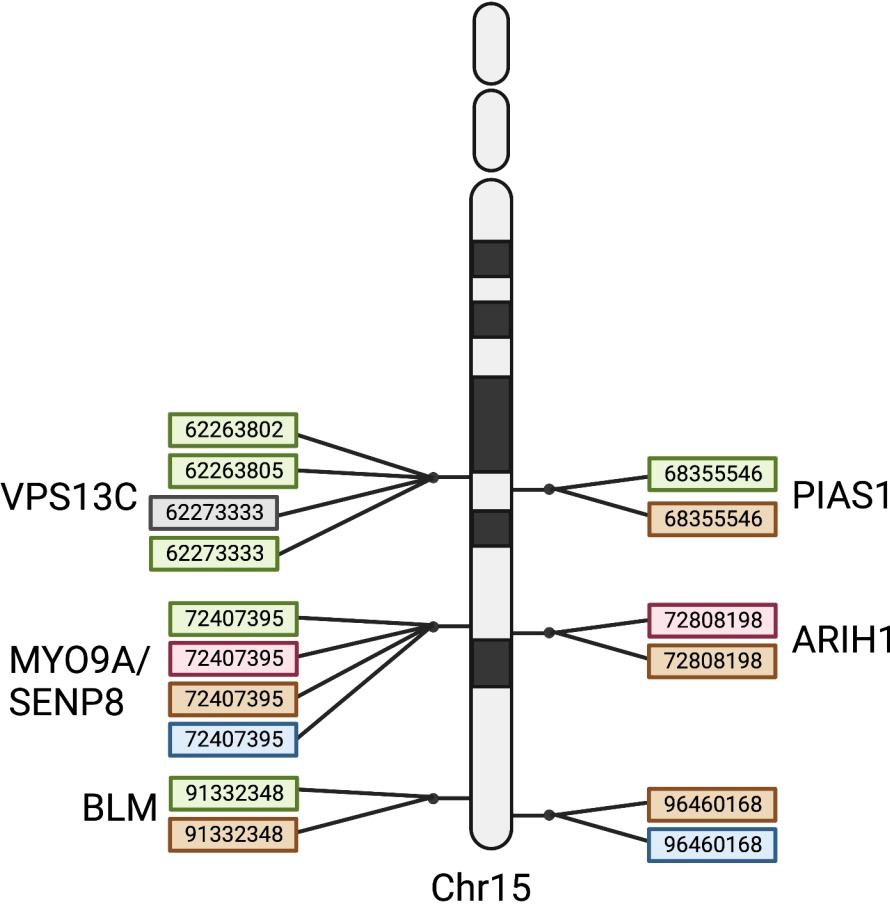


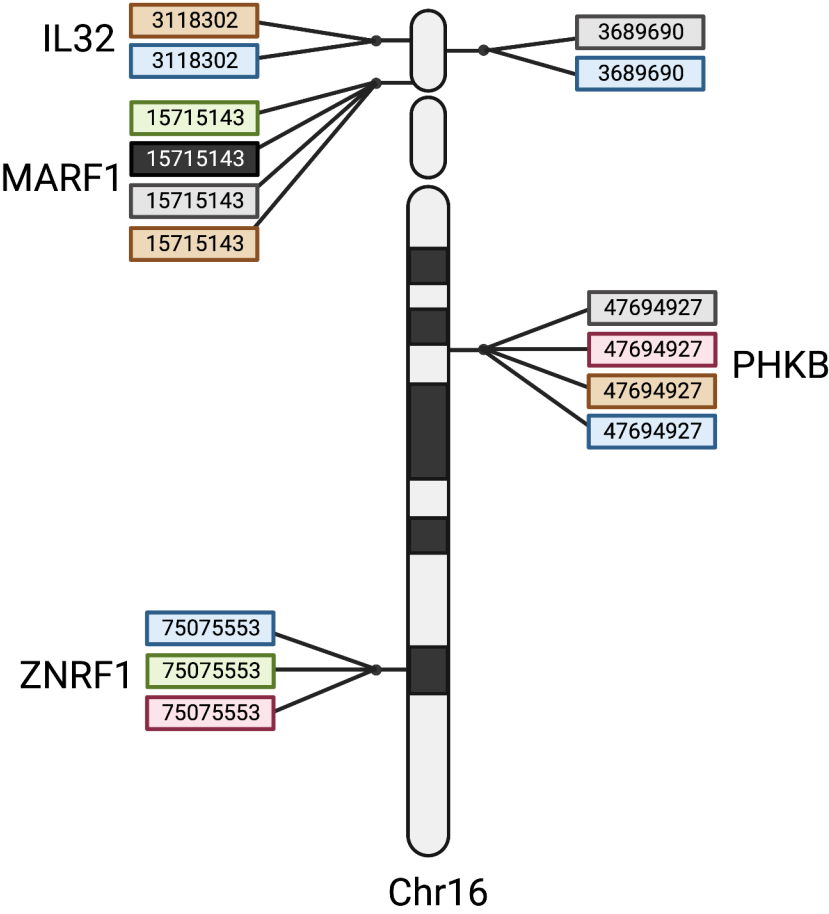


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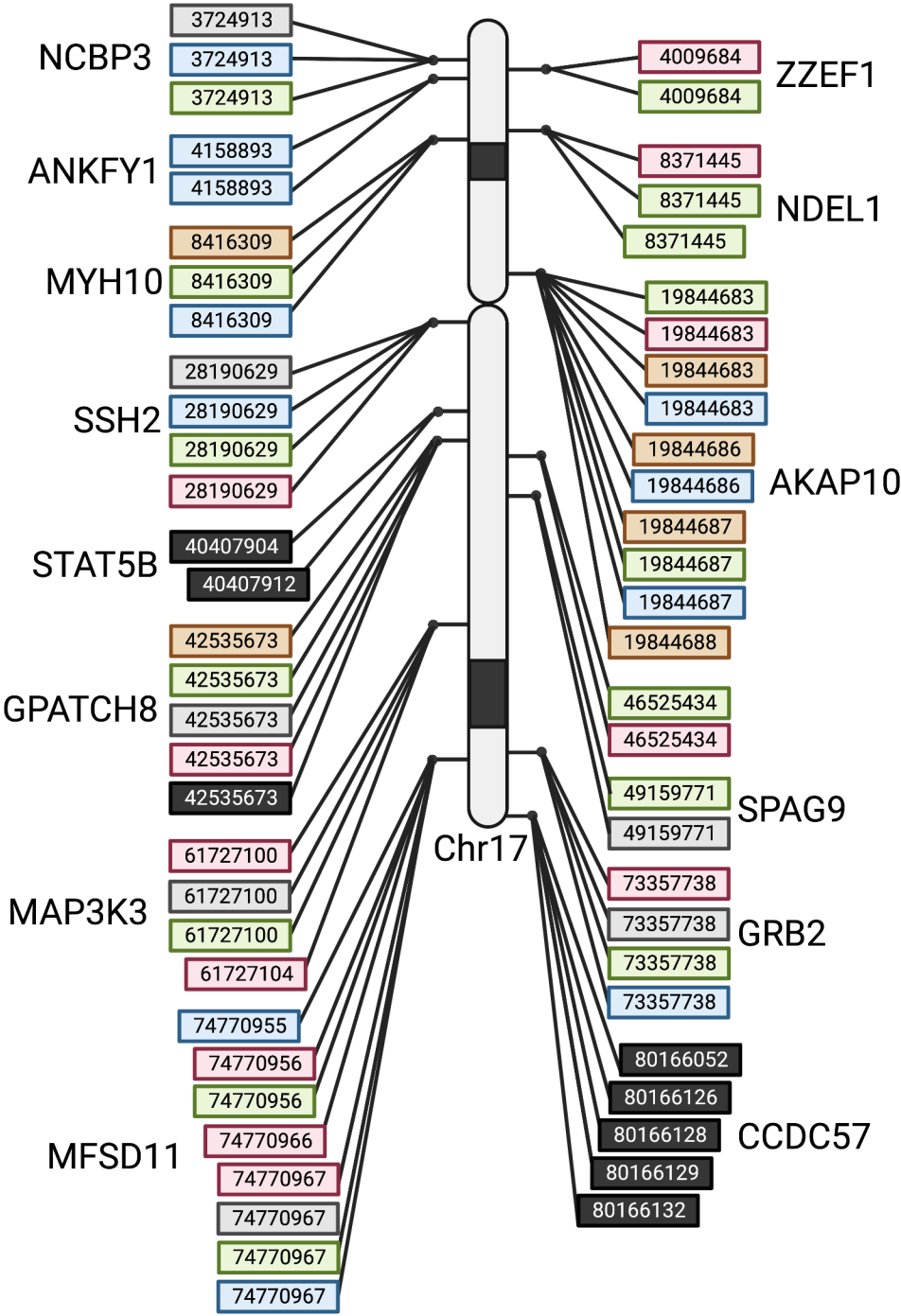


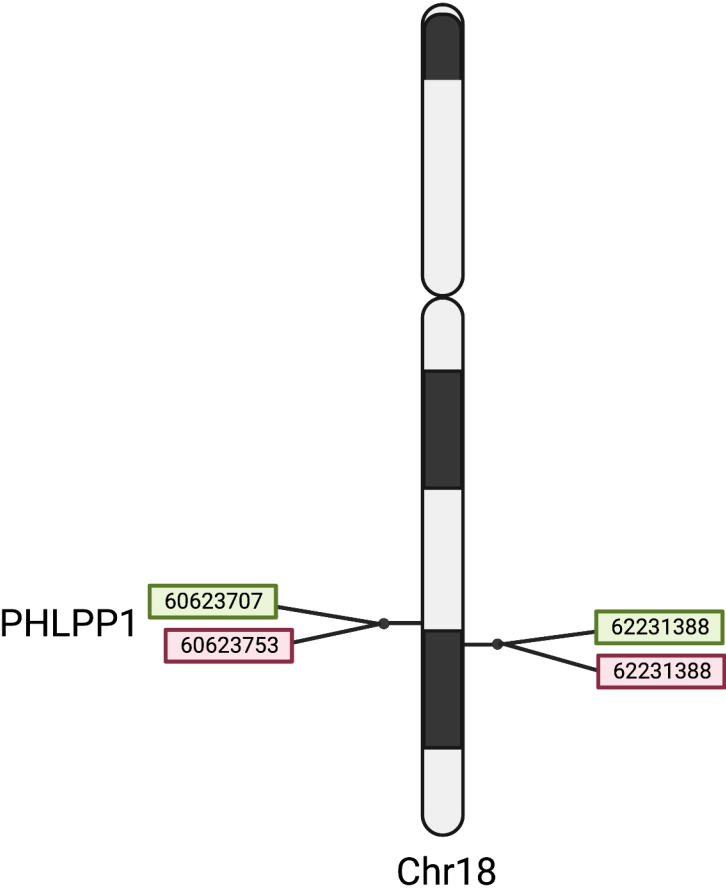




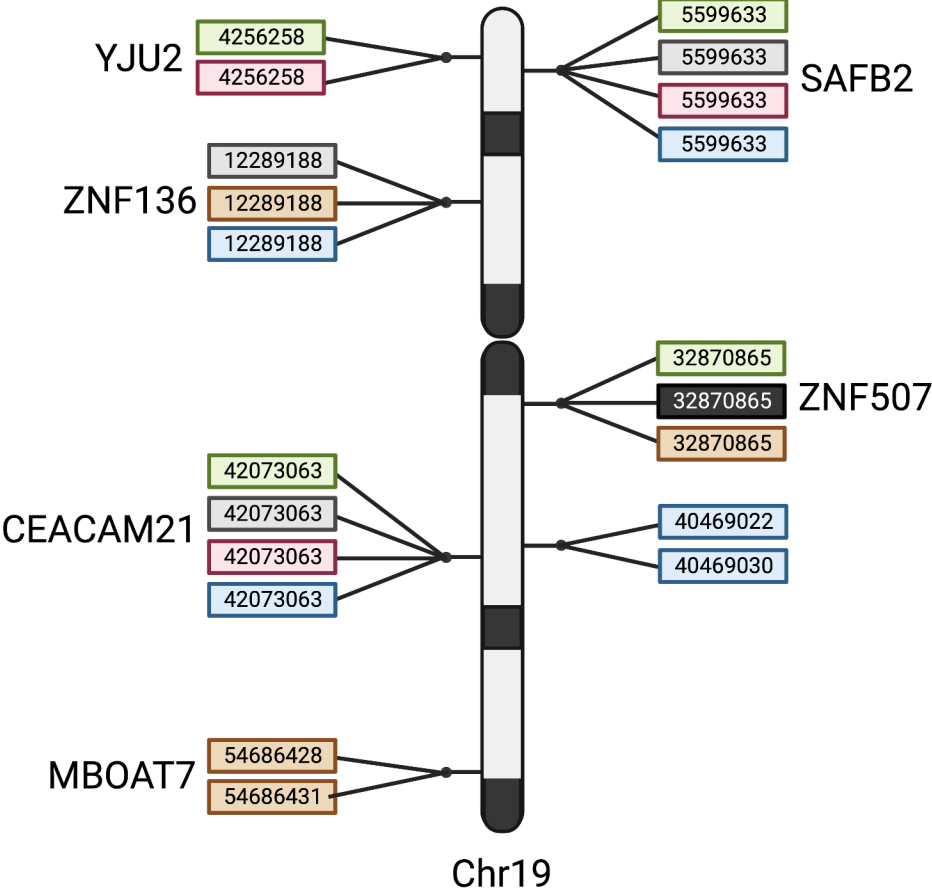
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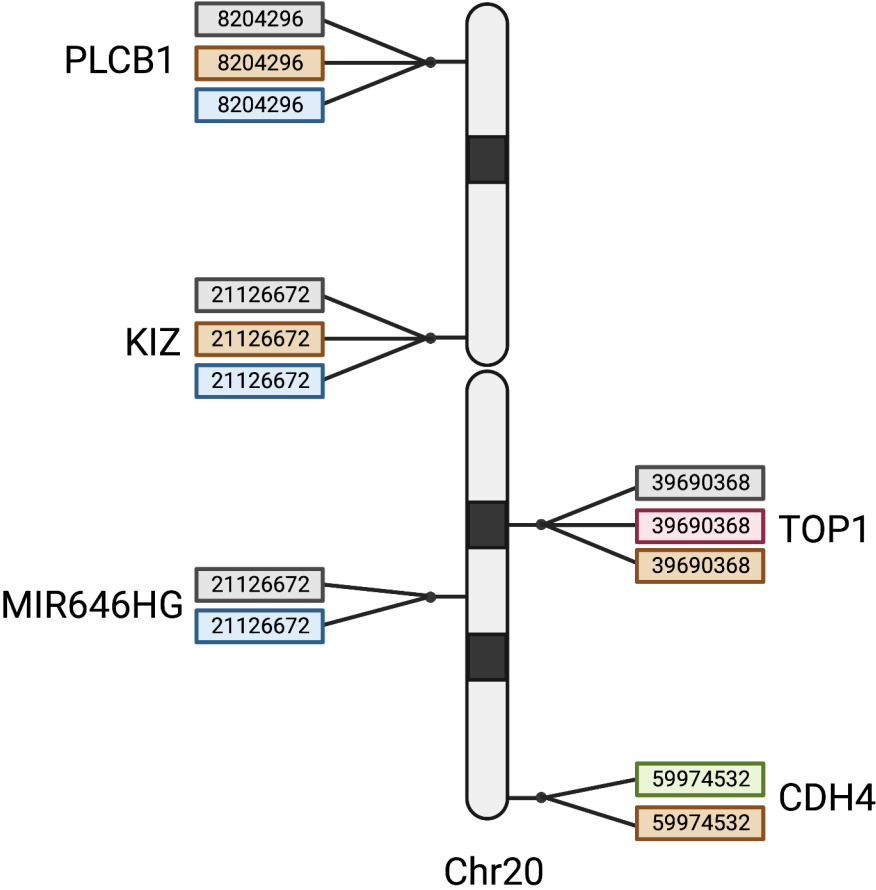


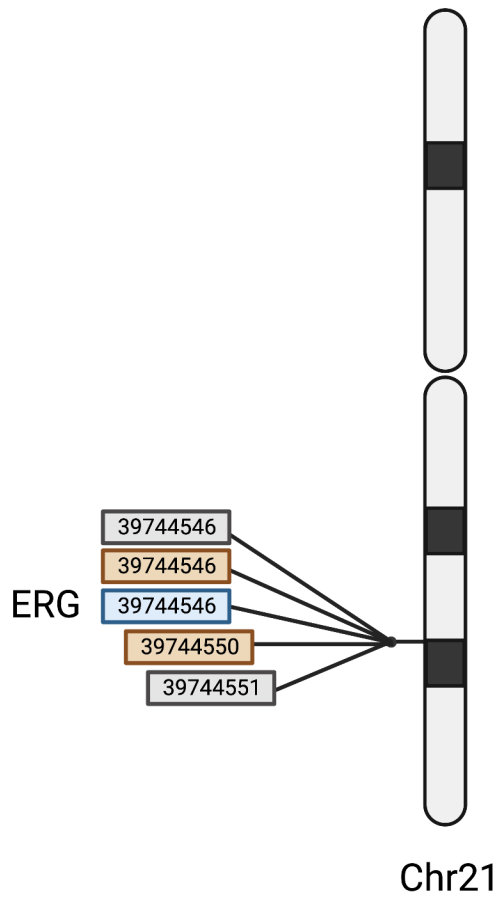


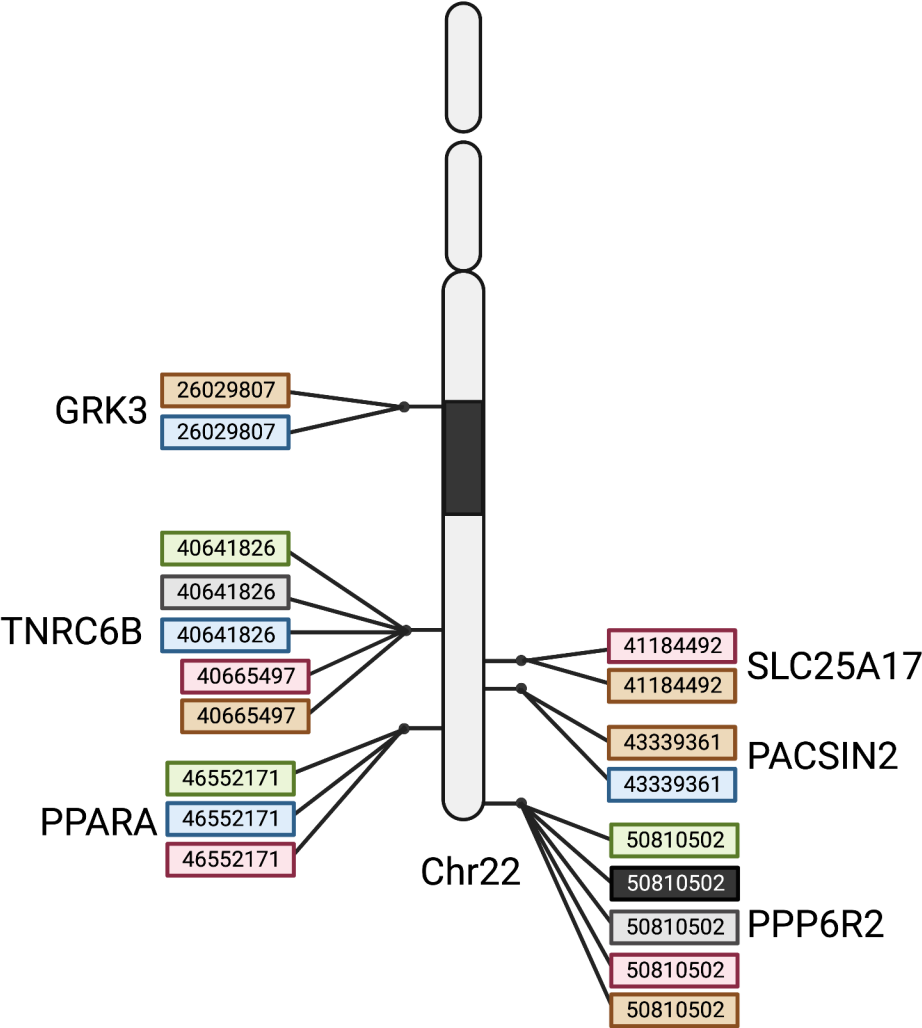
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