**Detection of copy number variants in African goats using whole genome sequence data**

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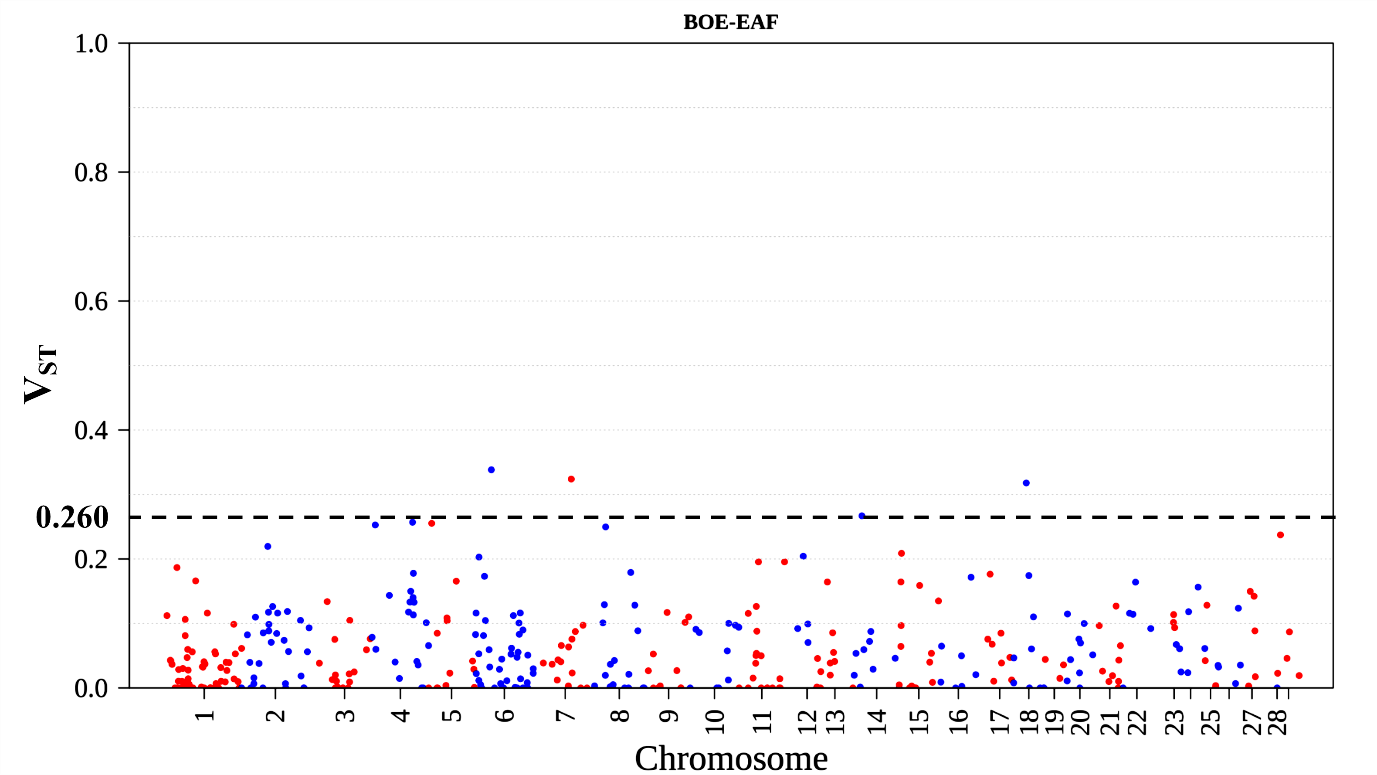
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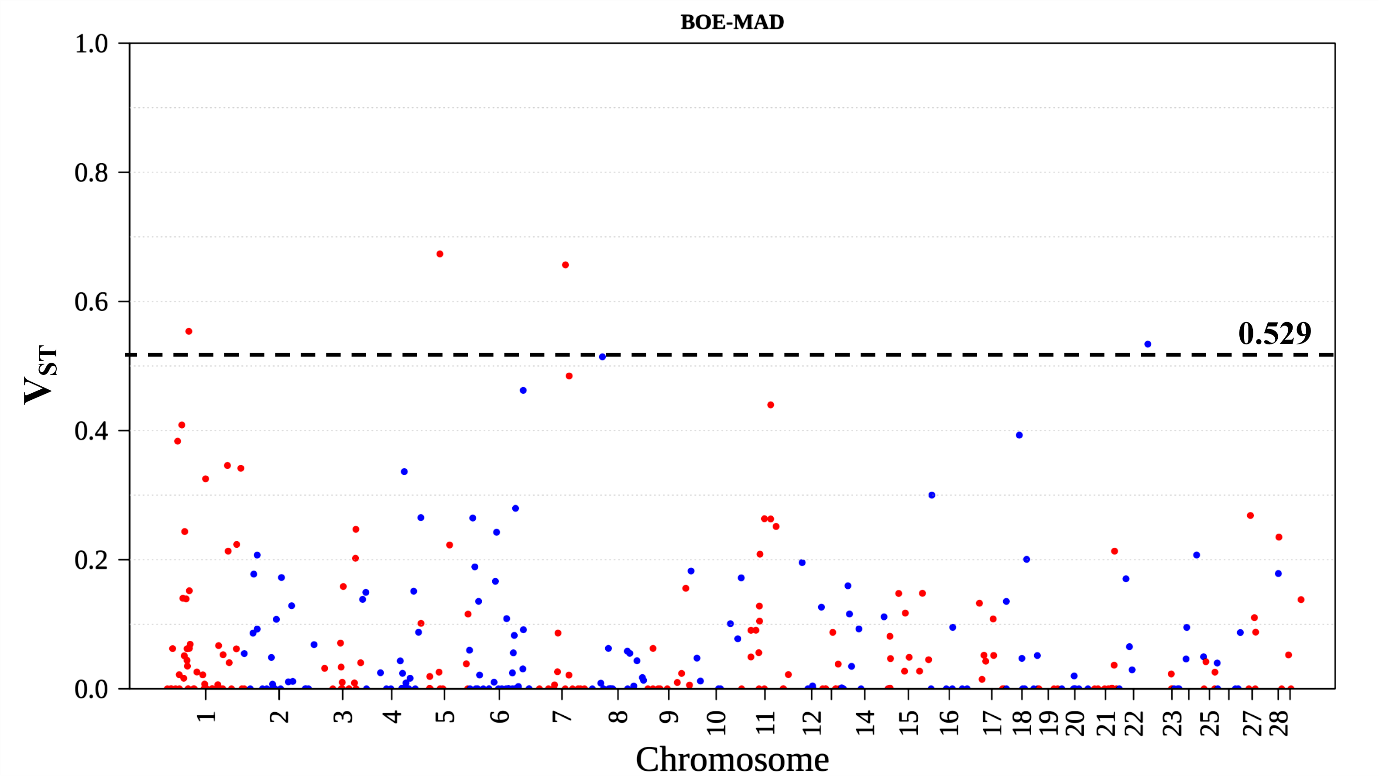
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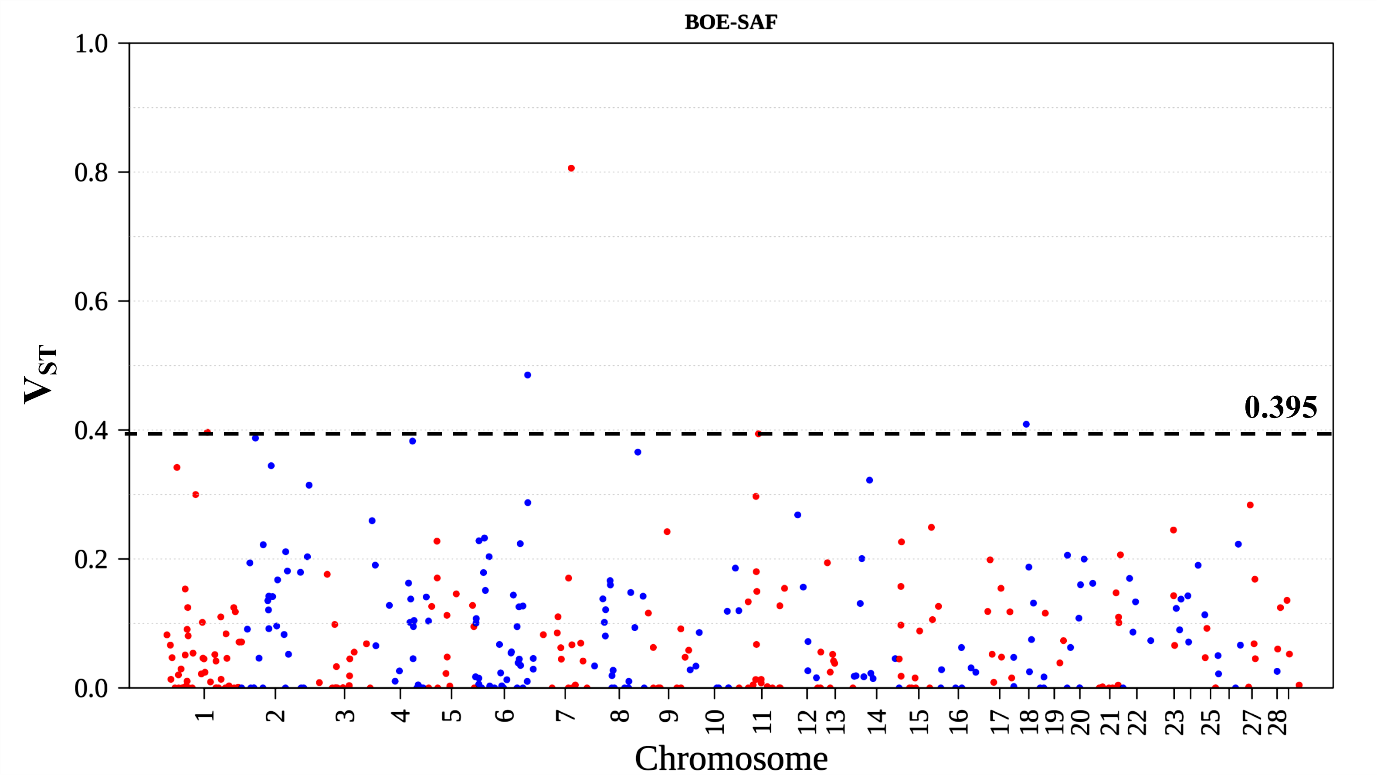
Supplementary Figure 1: CNV differentiation between Boer and East African goat populations.

The dotted line represents the *VST*threshold value for this test.



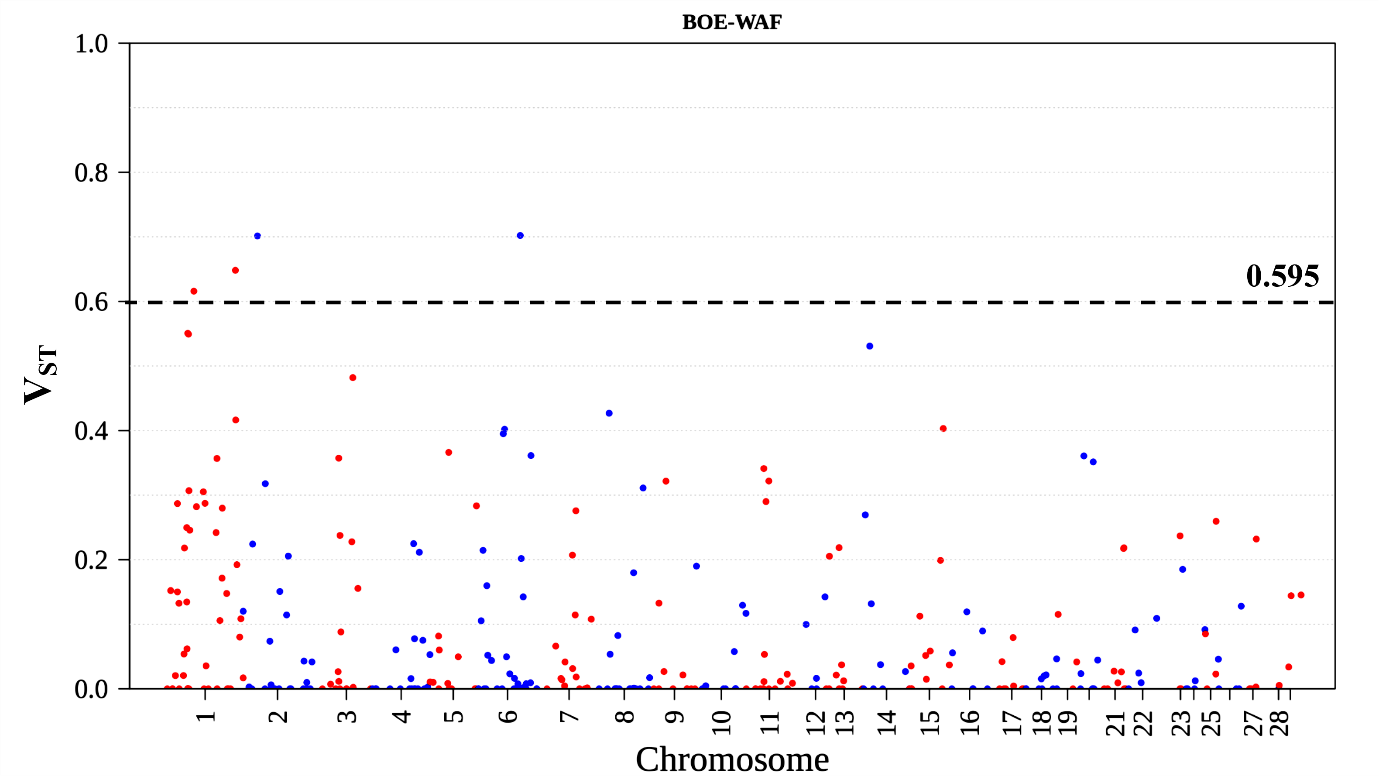
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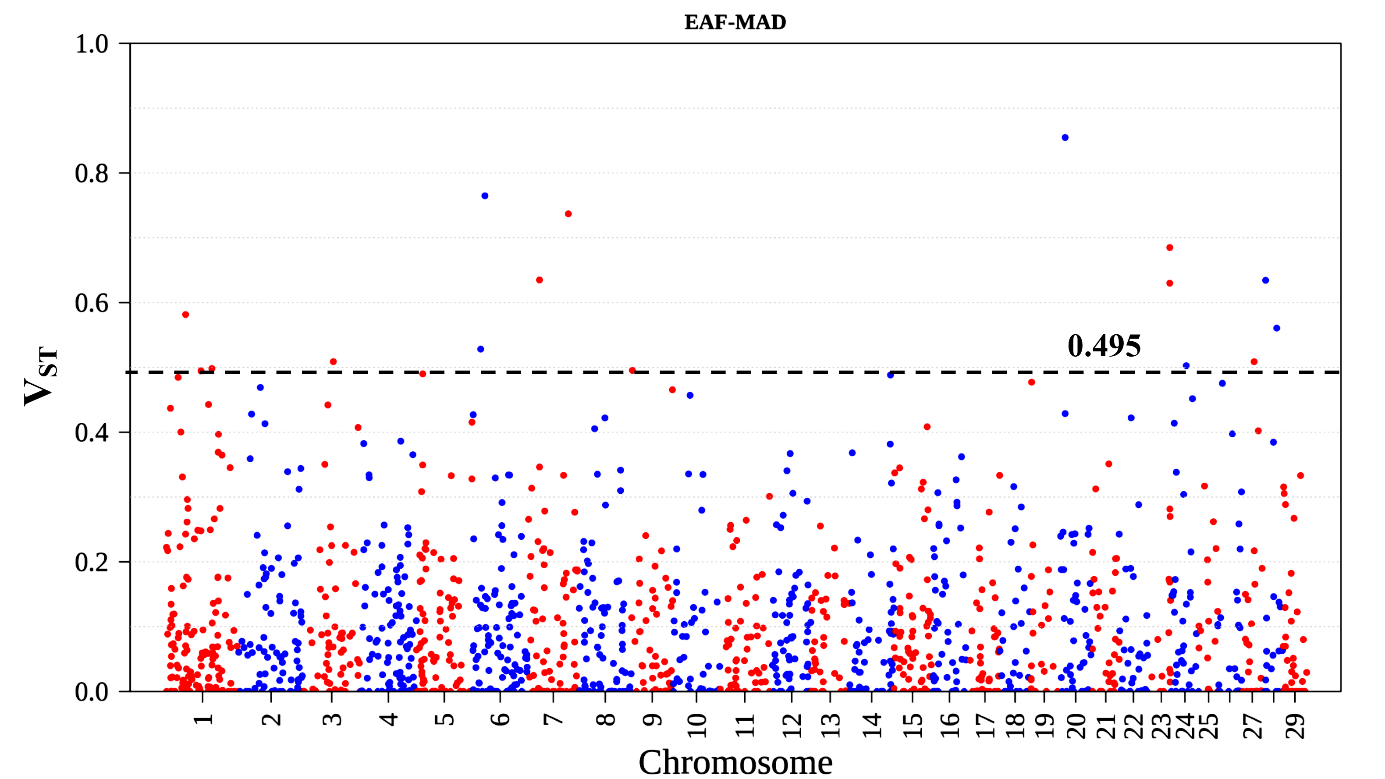
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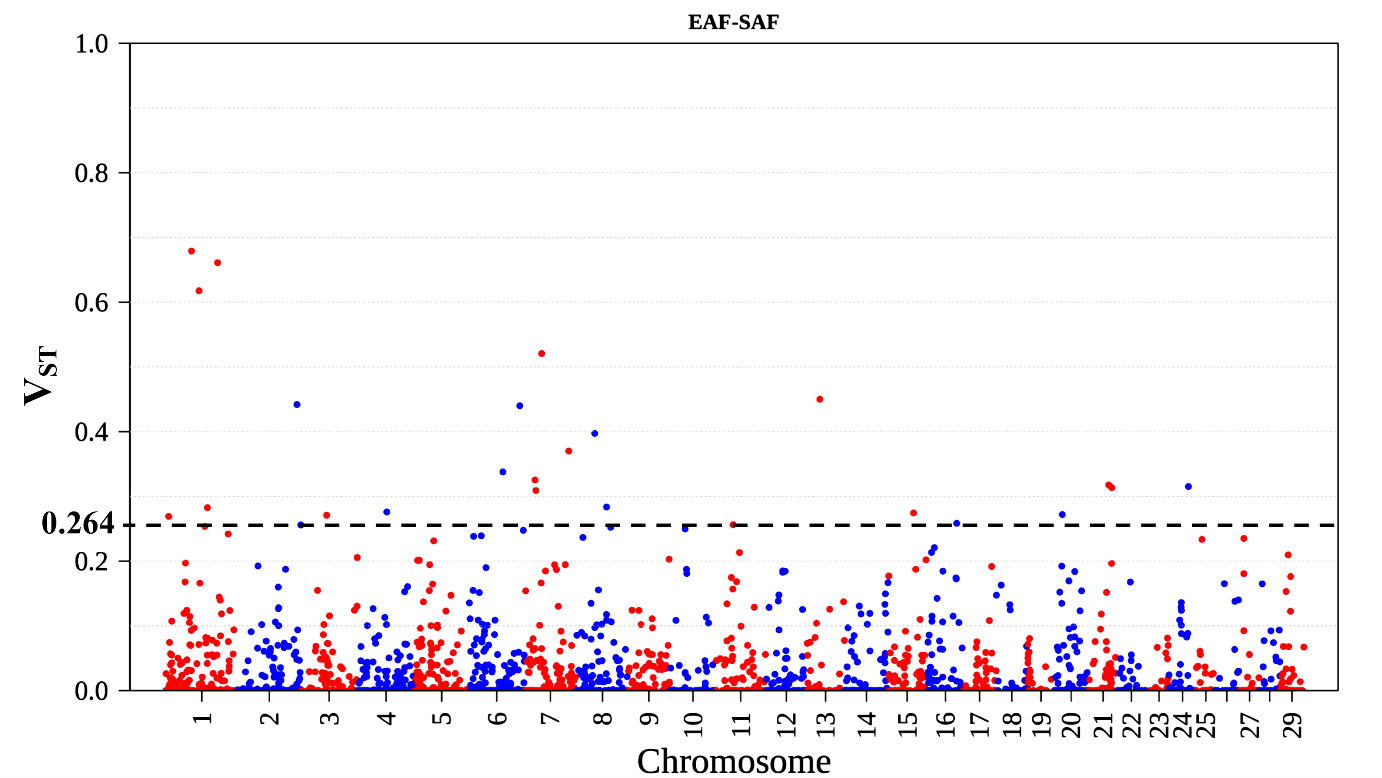
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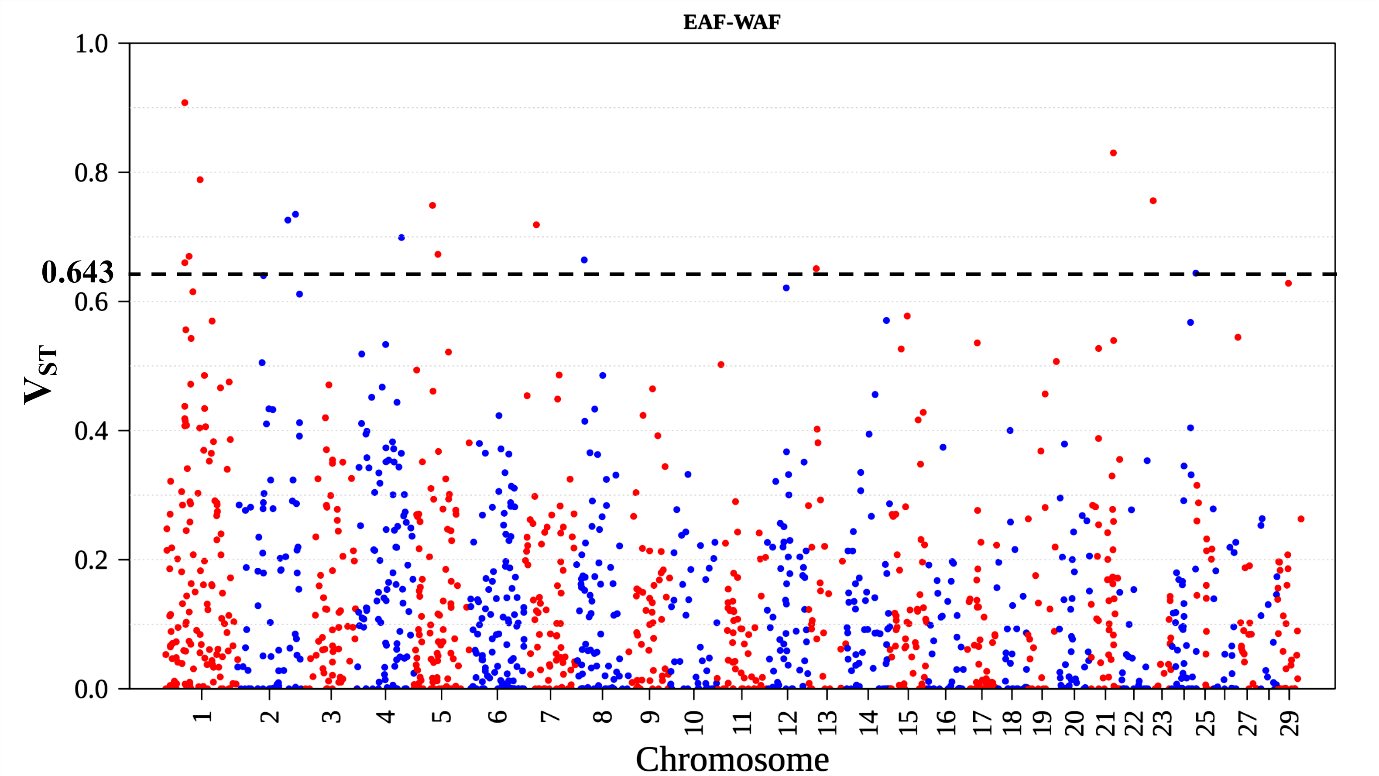
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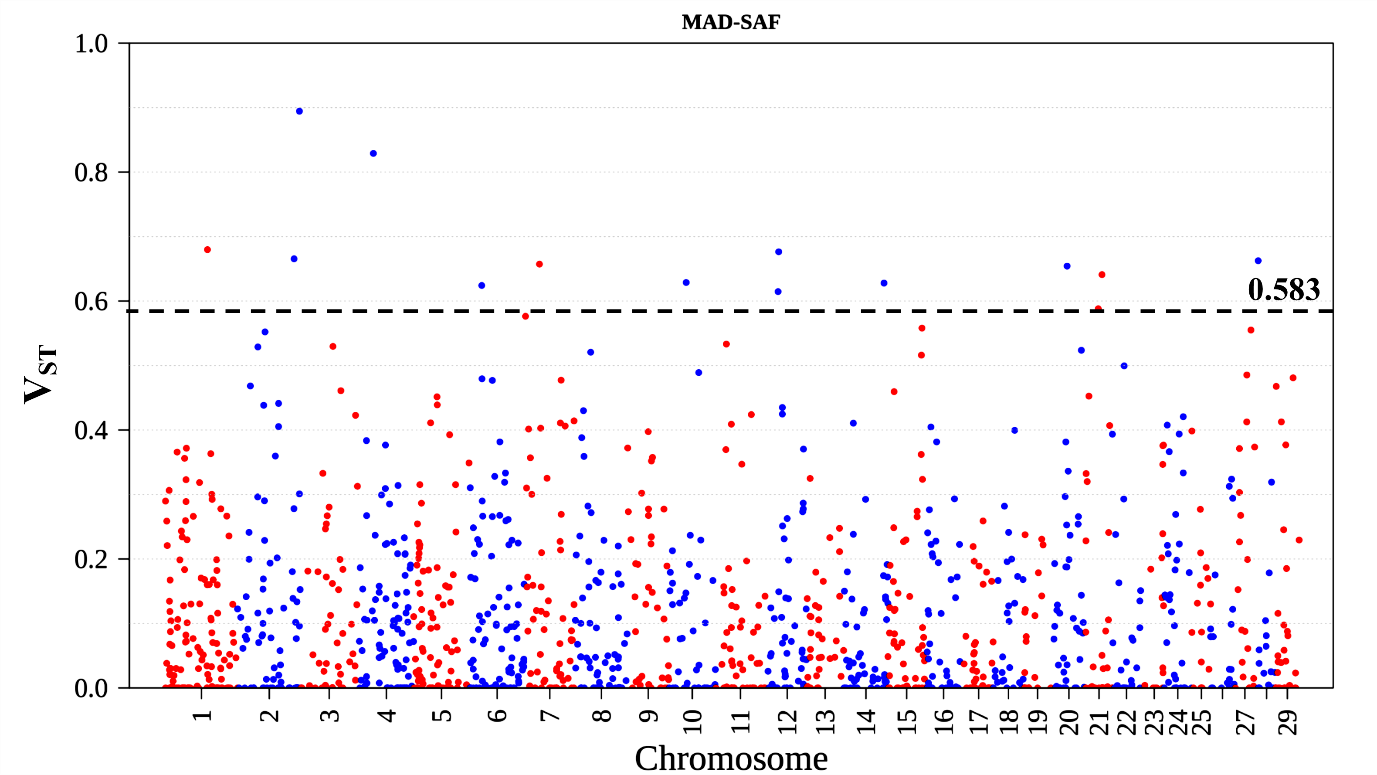
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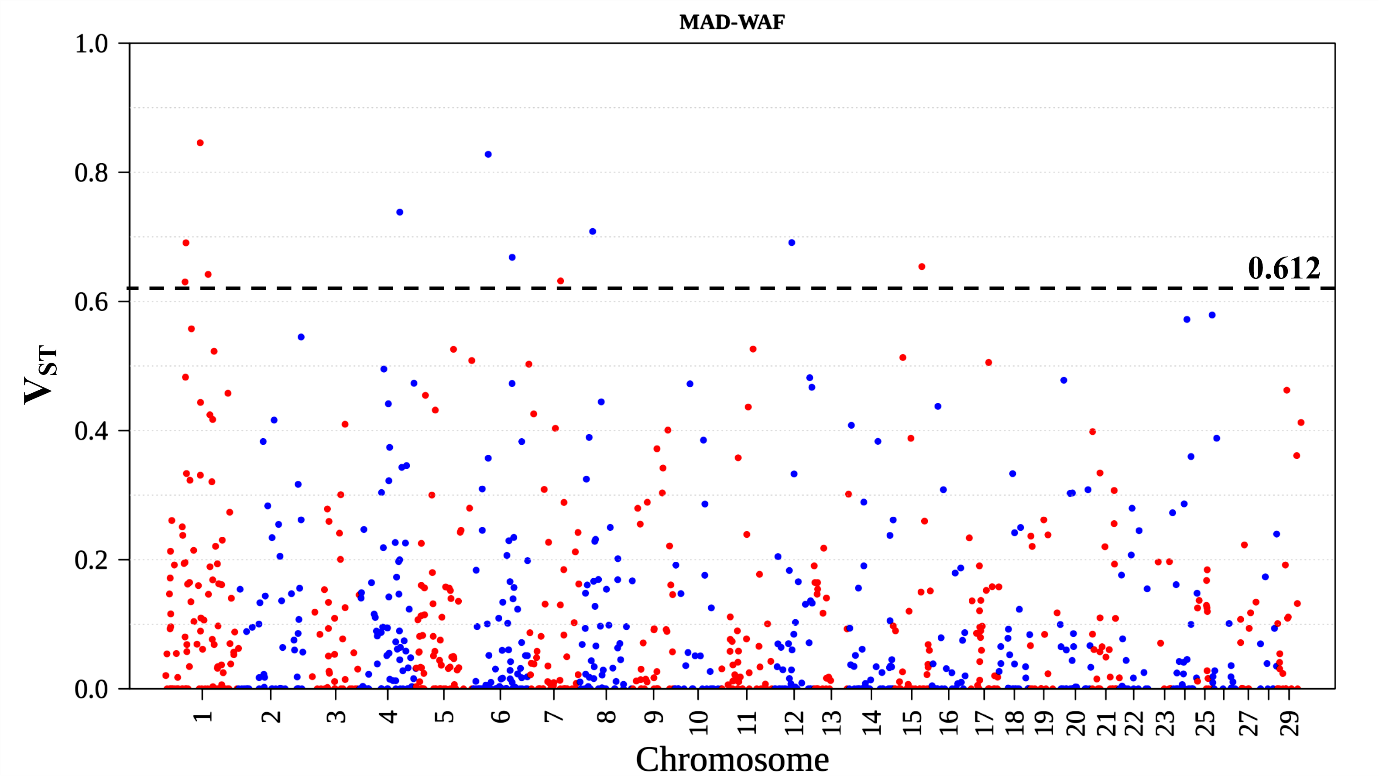
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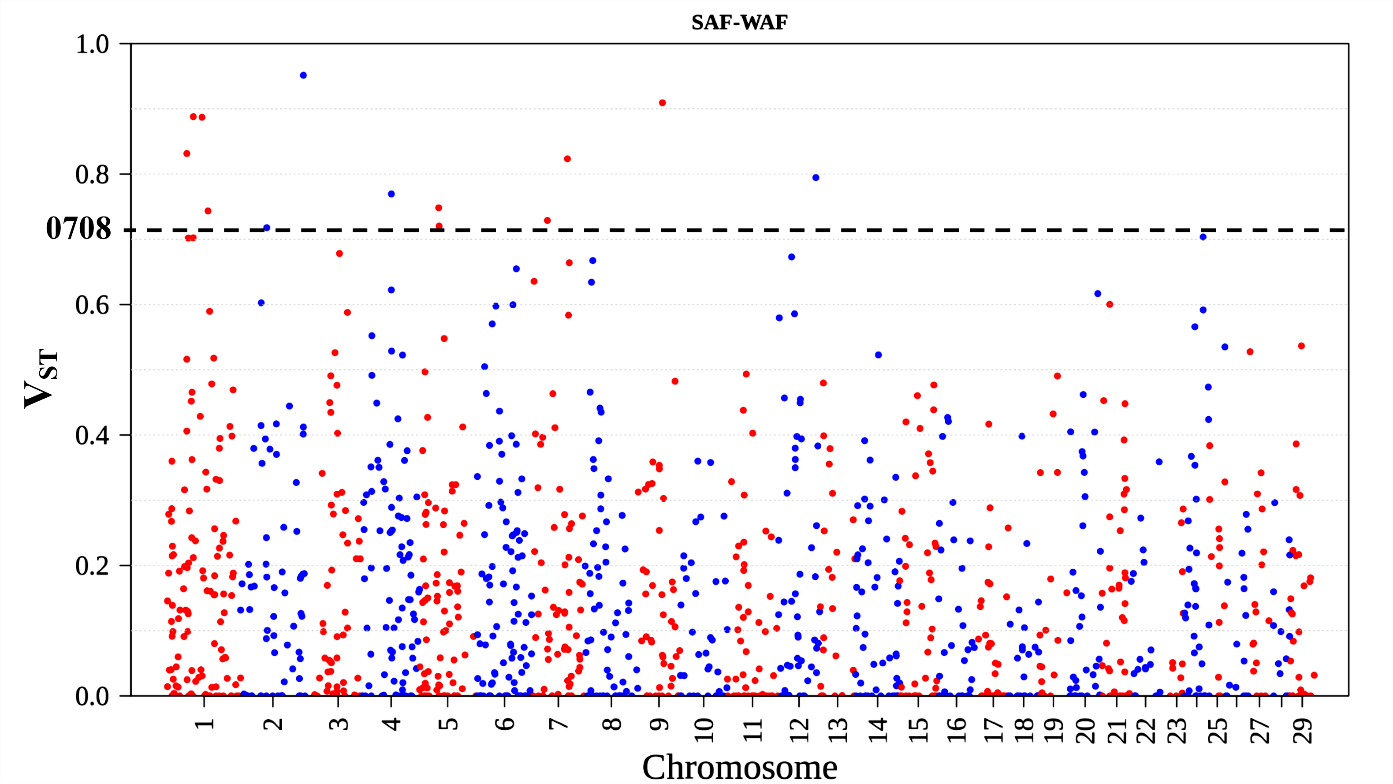
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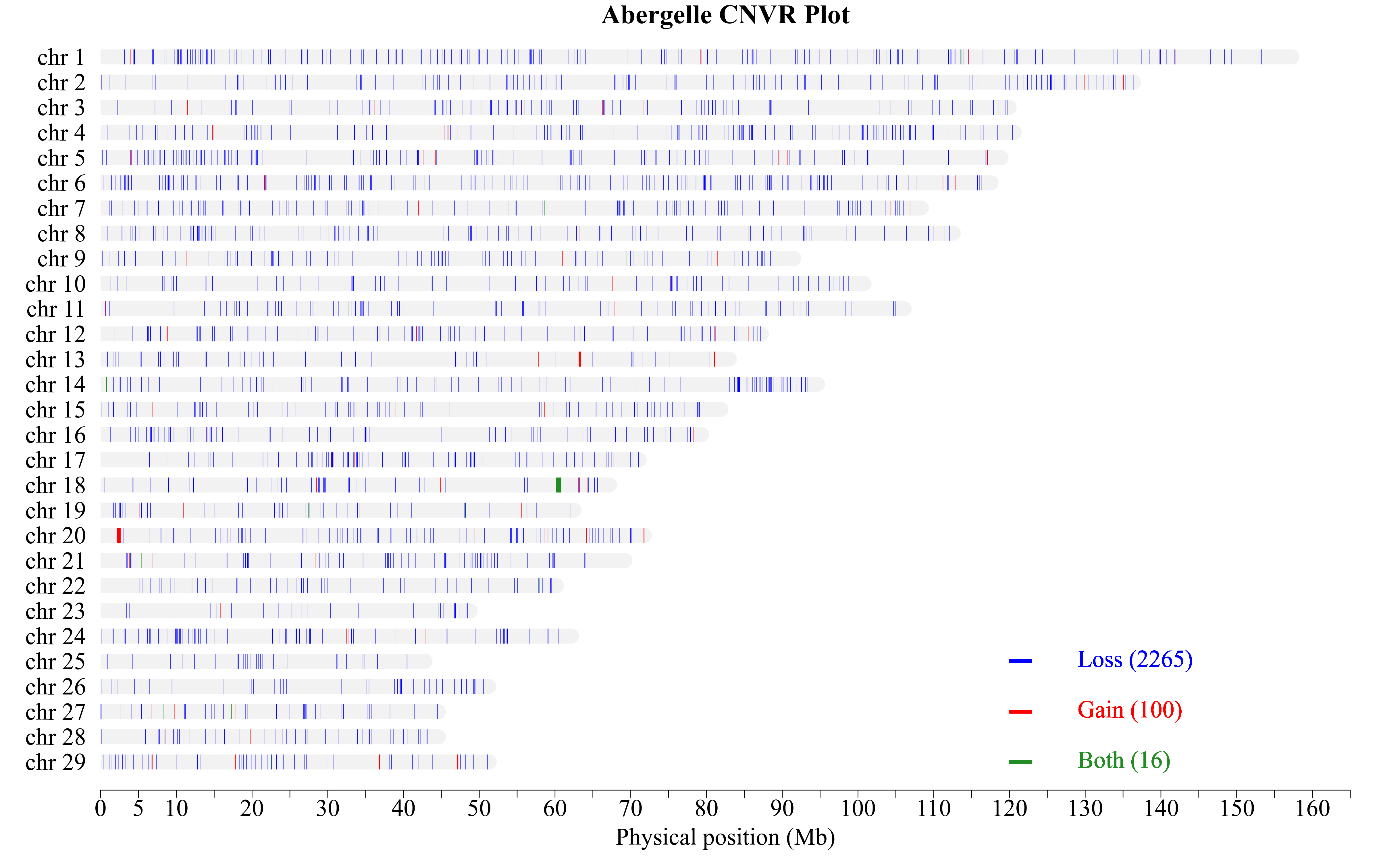
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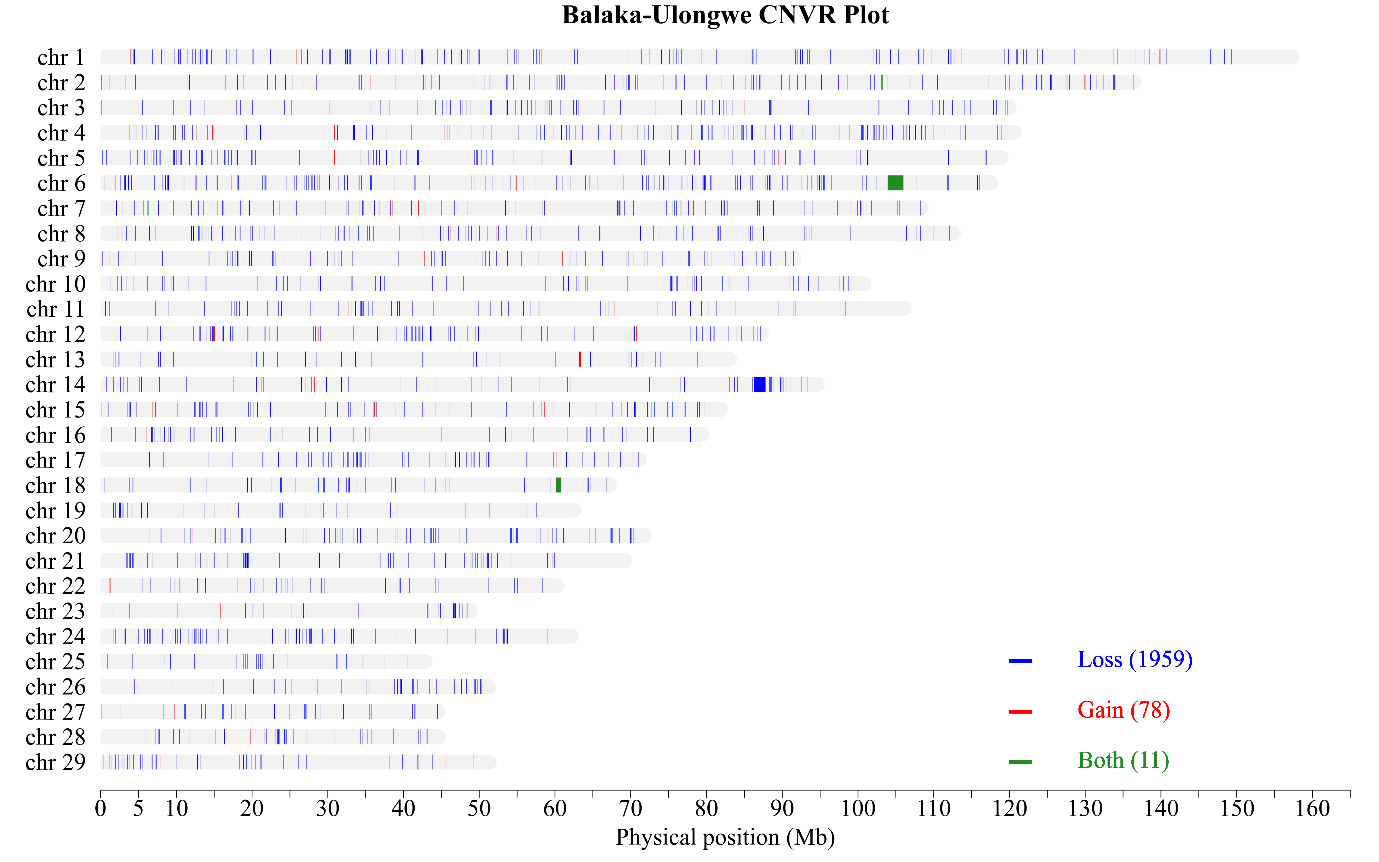
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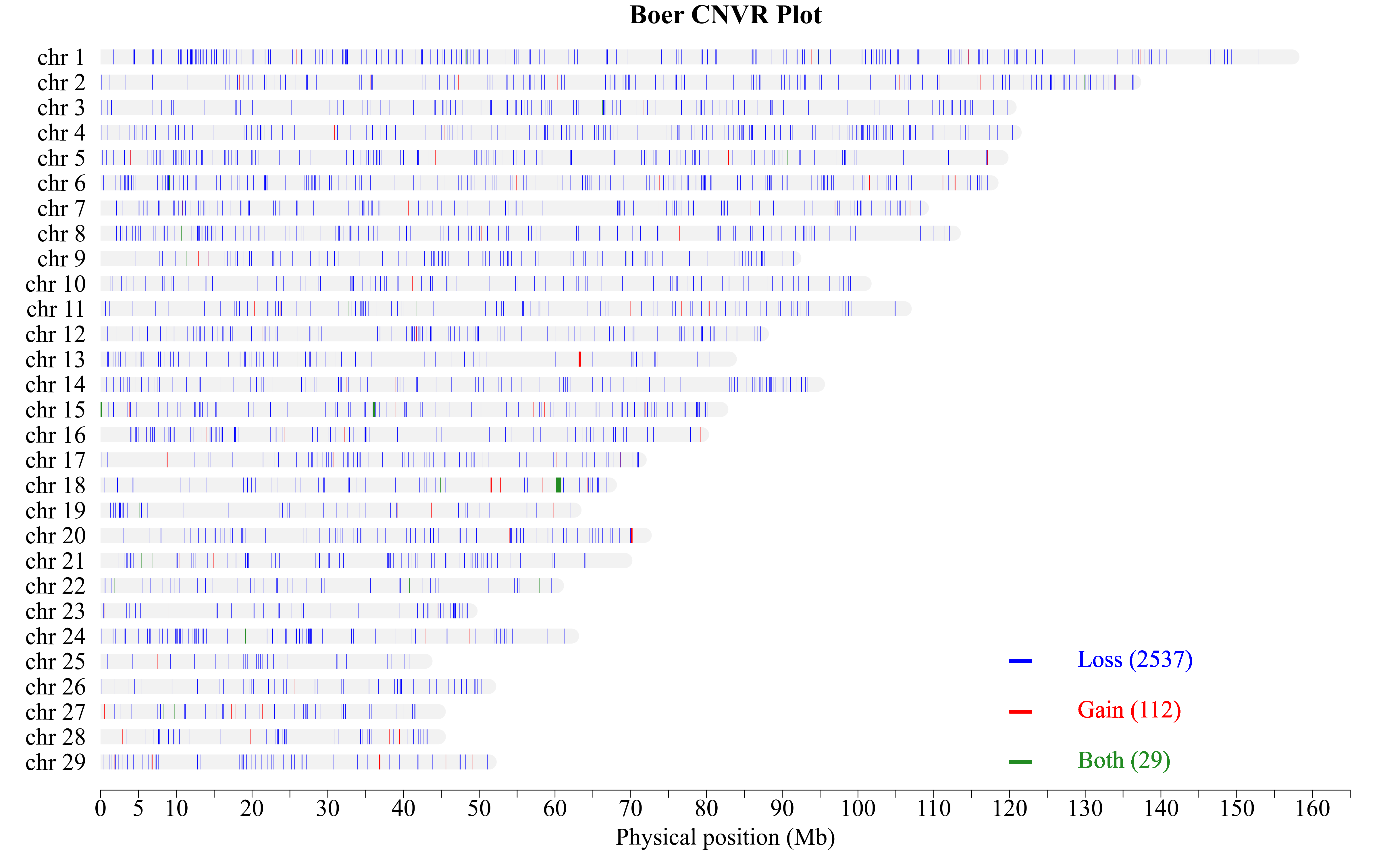
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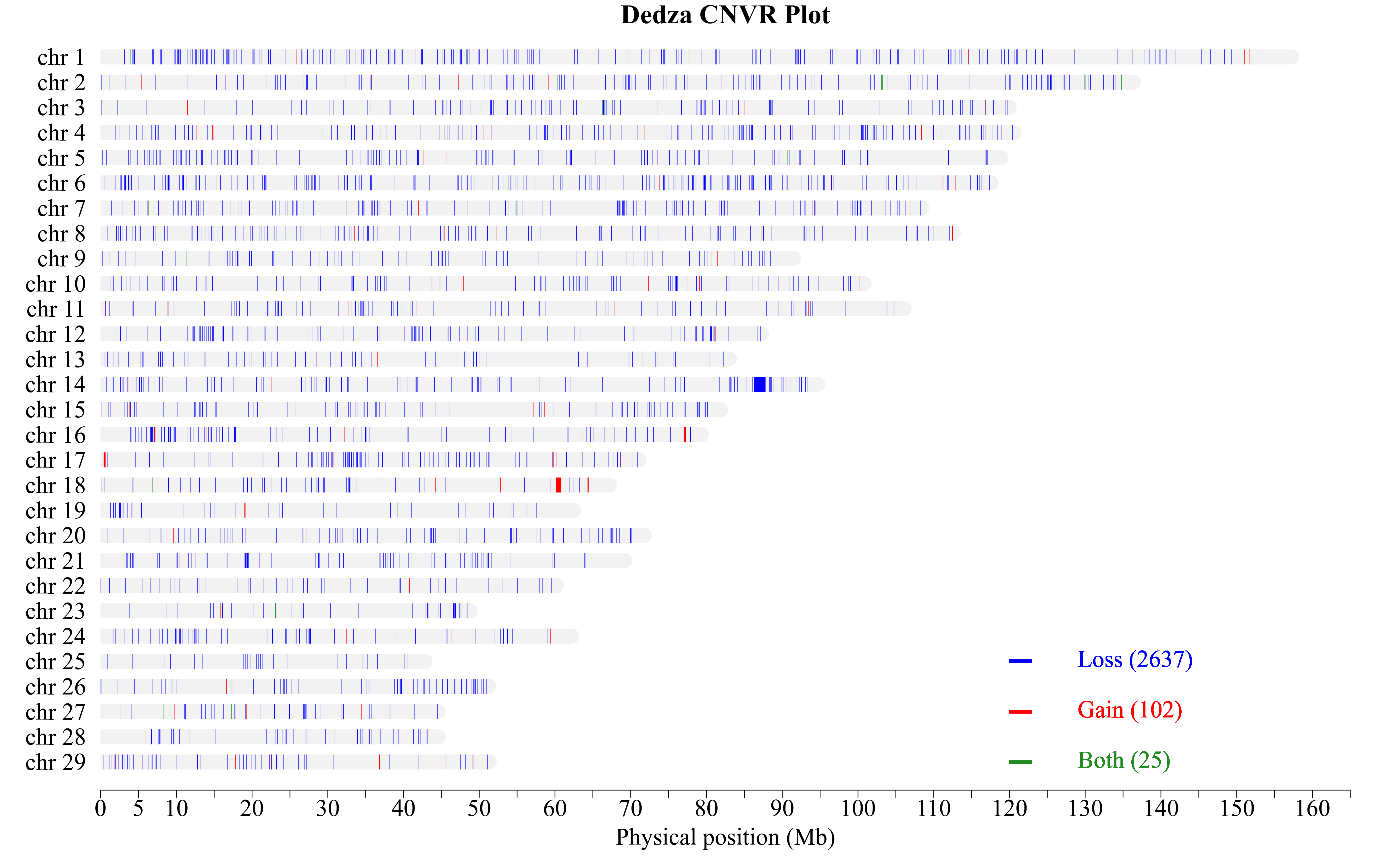
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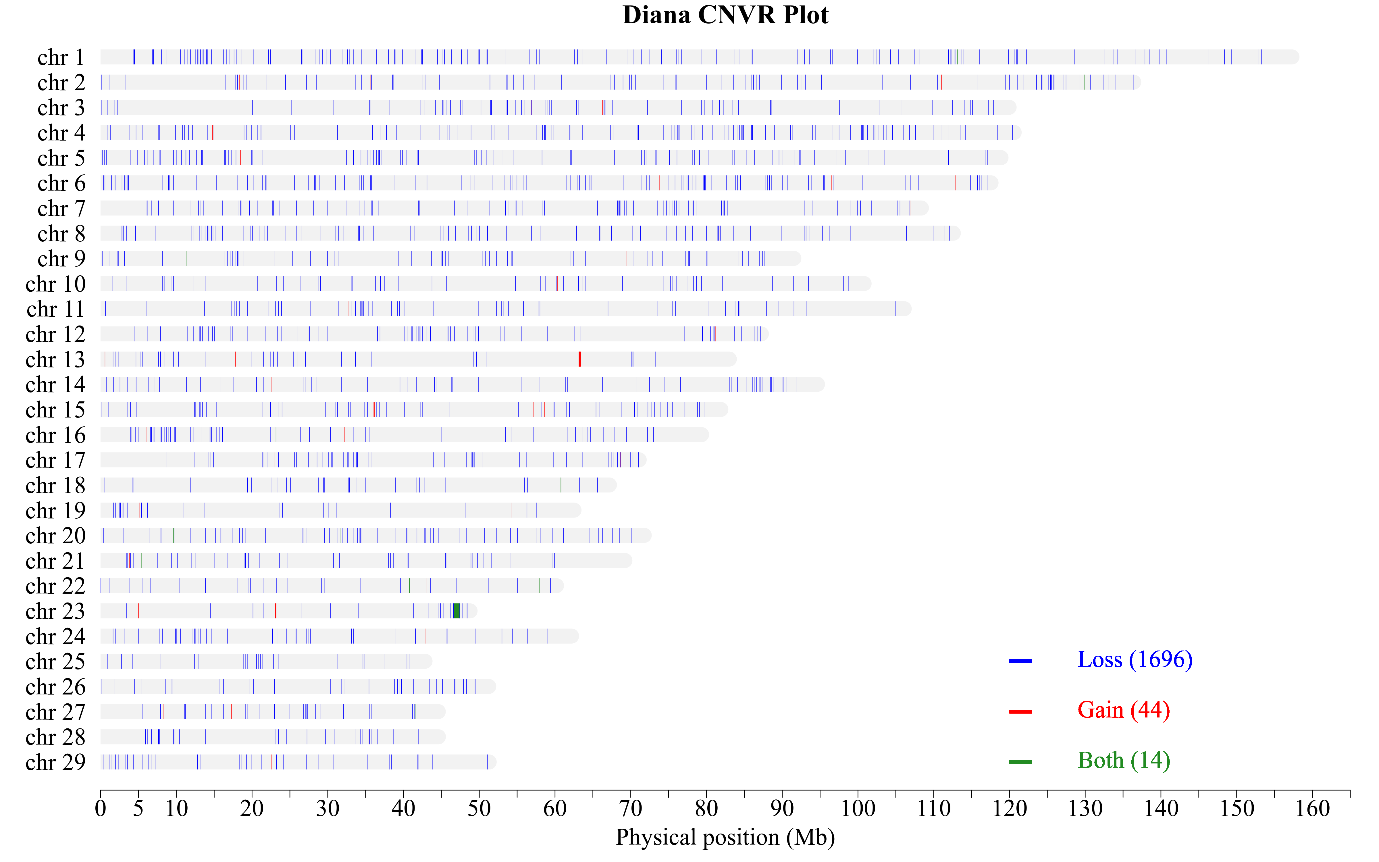
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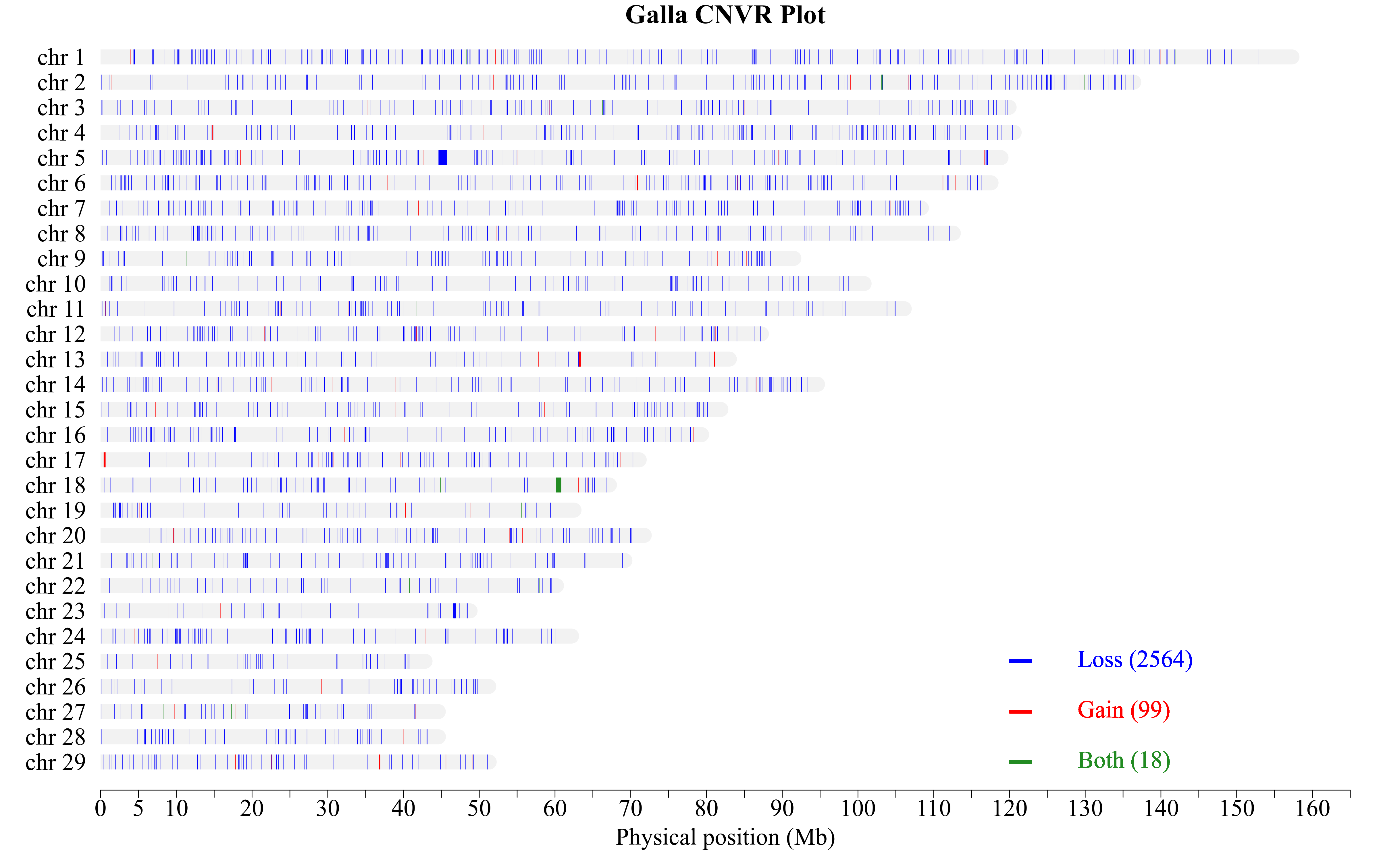
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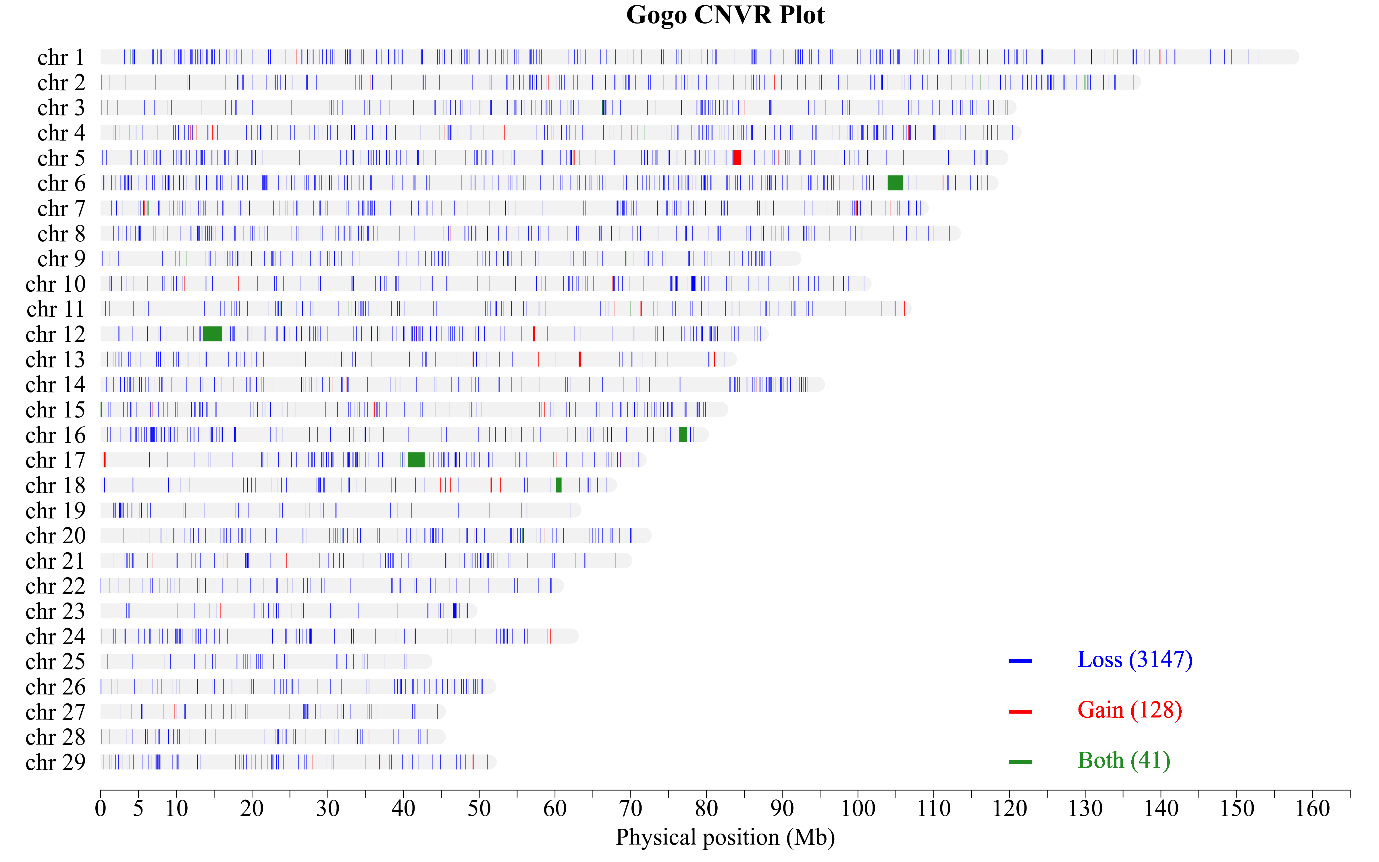
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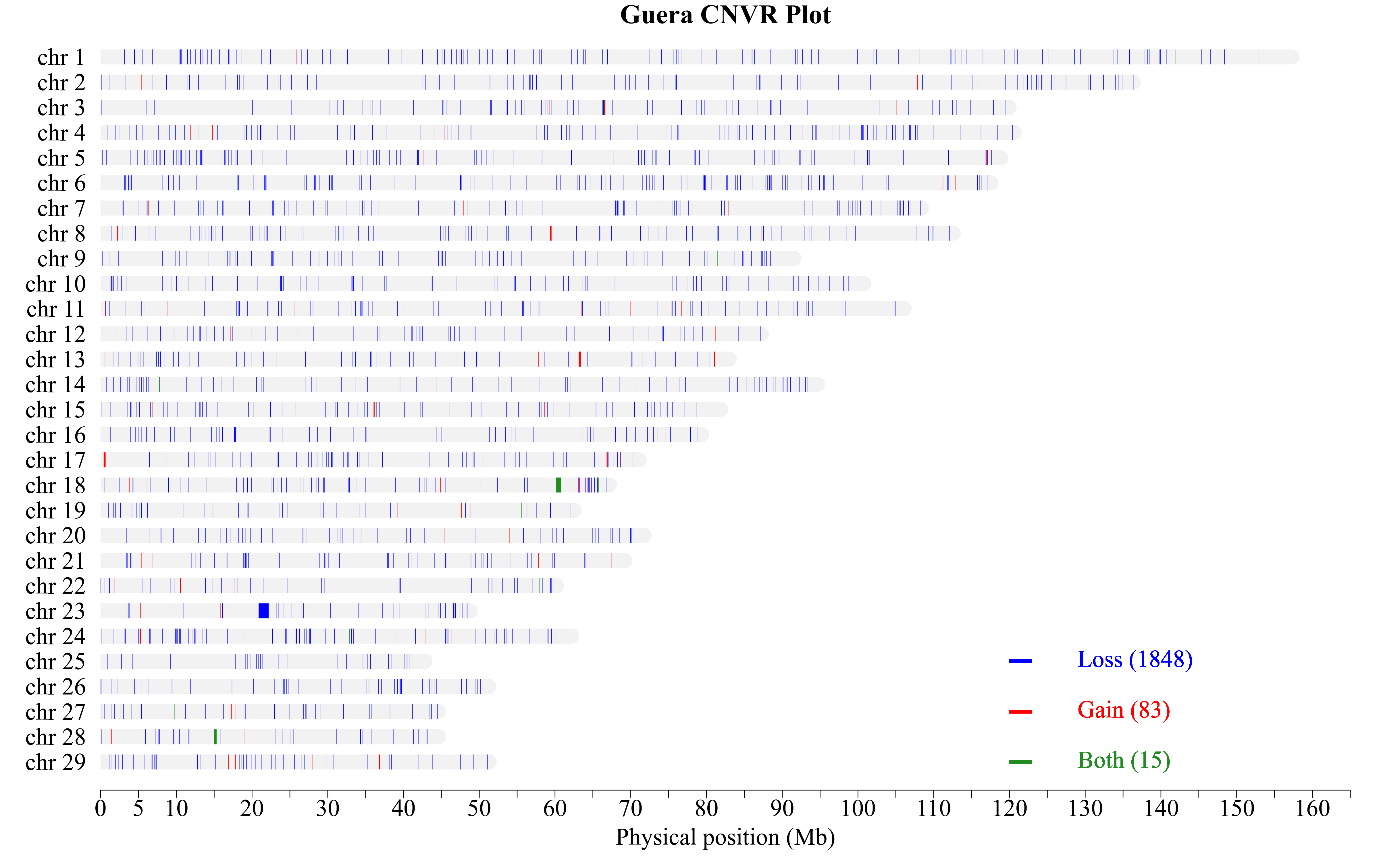
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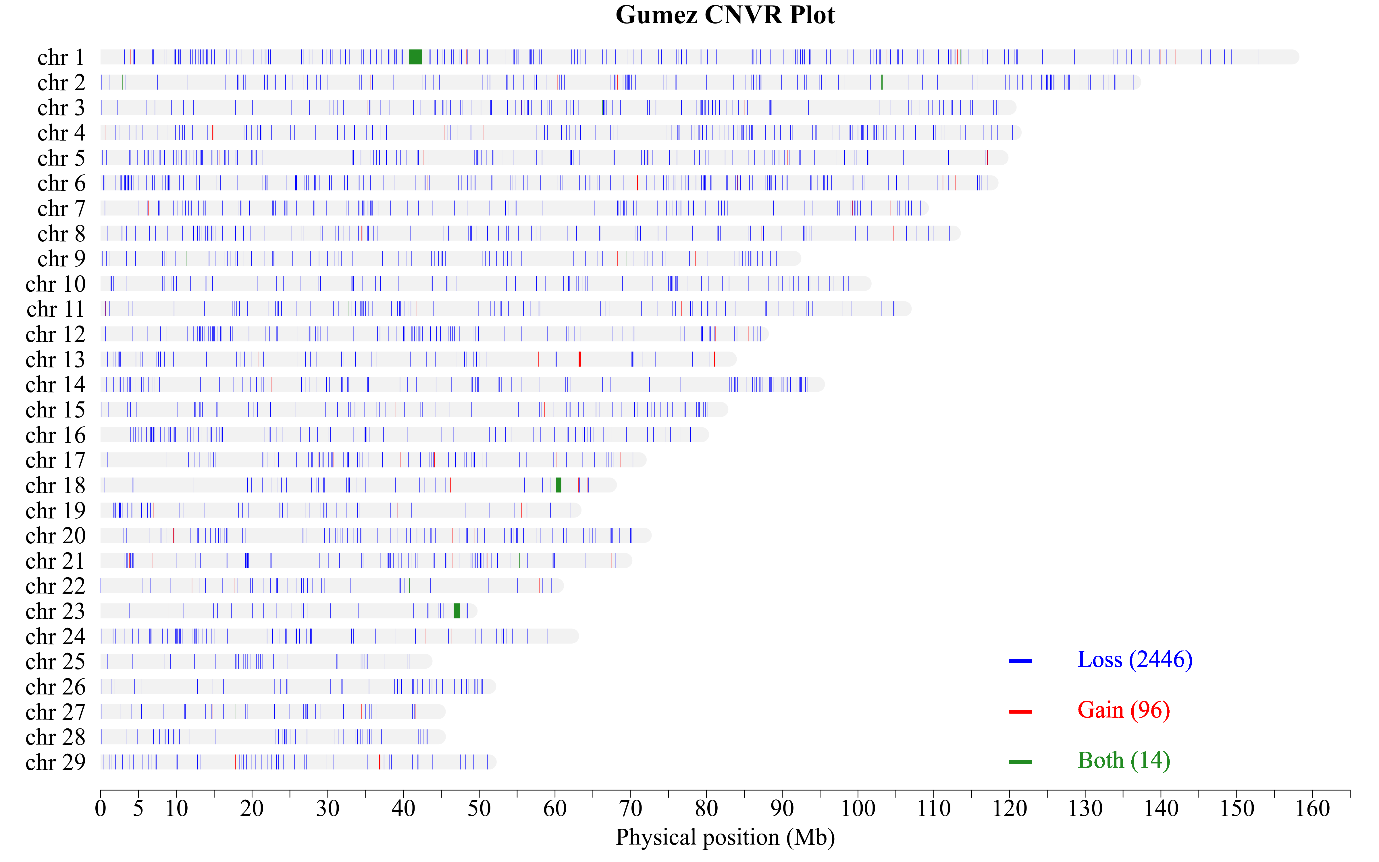
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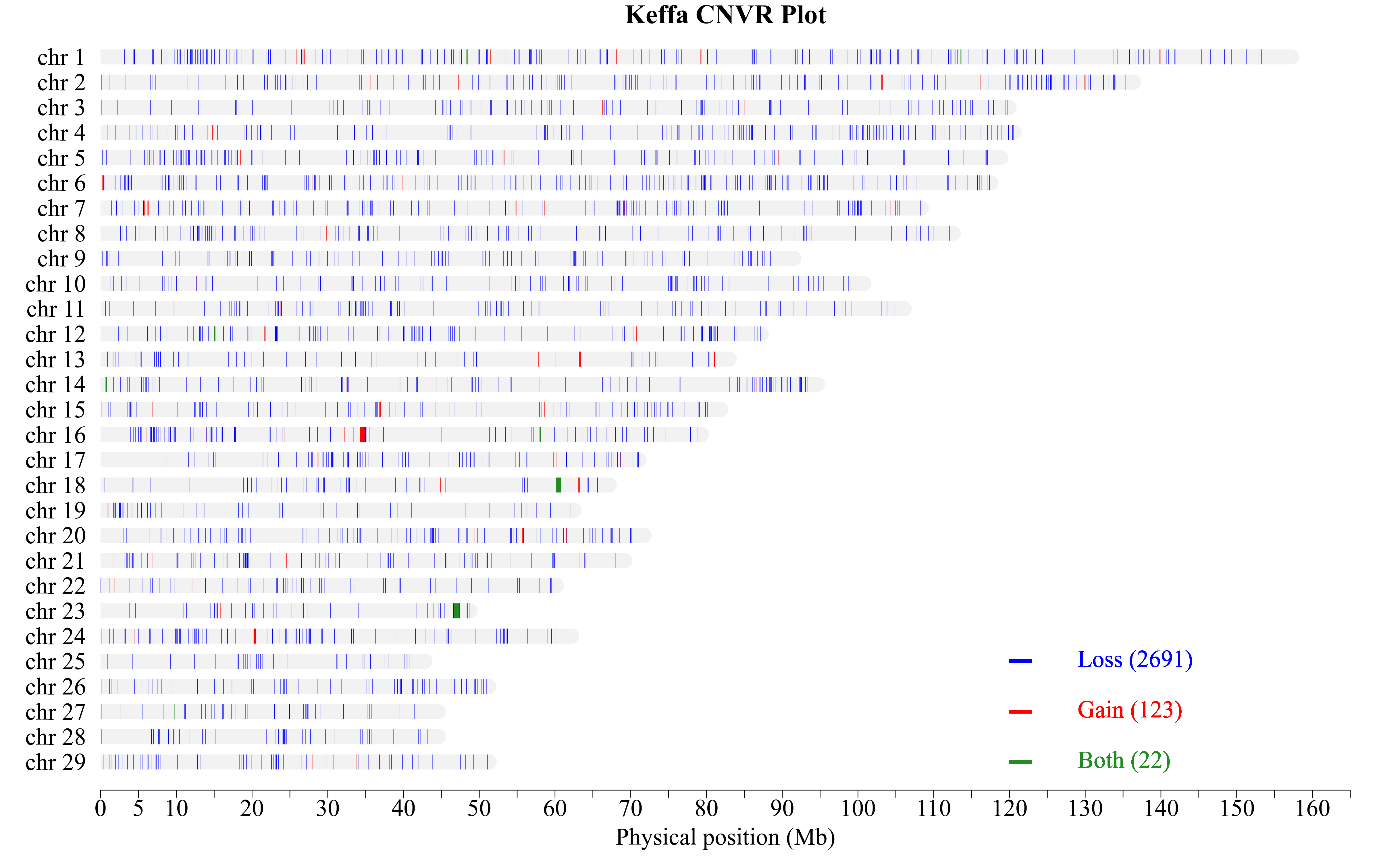
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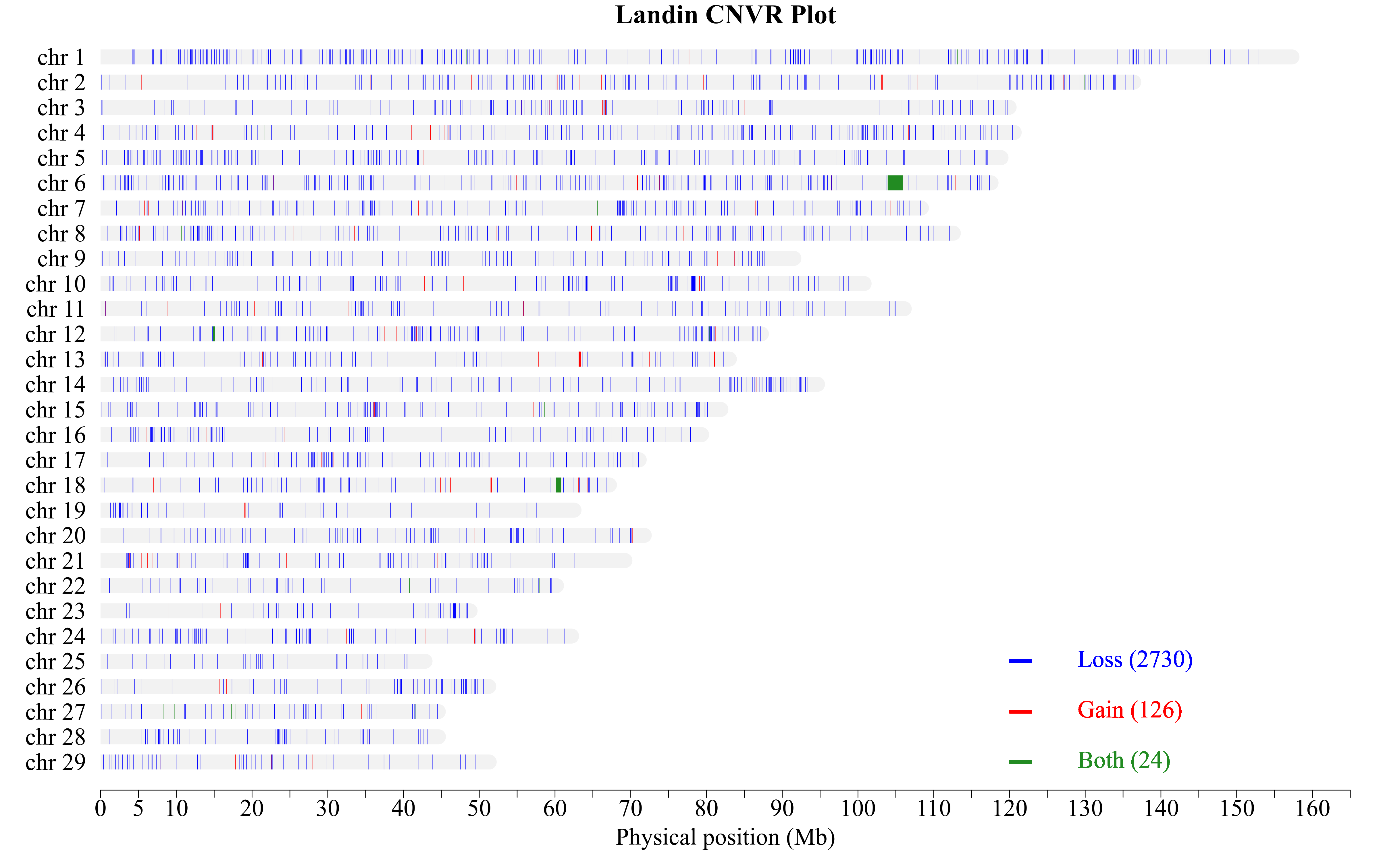
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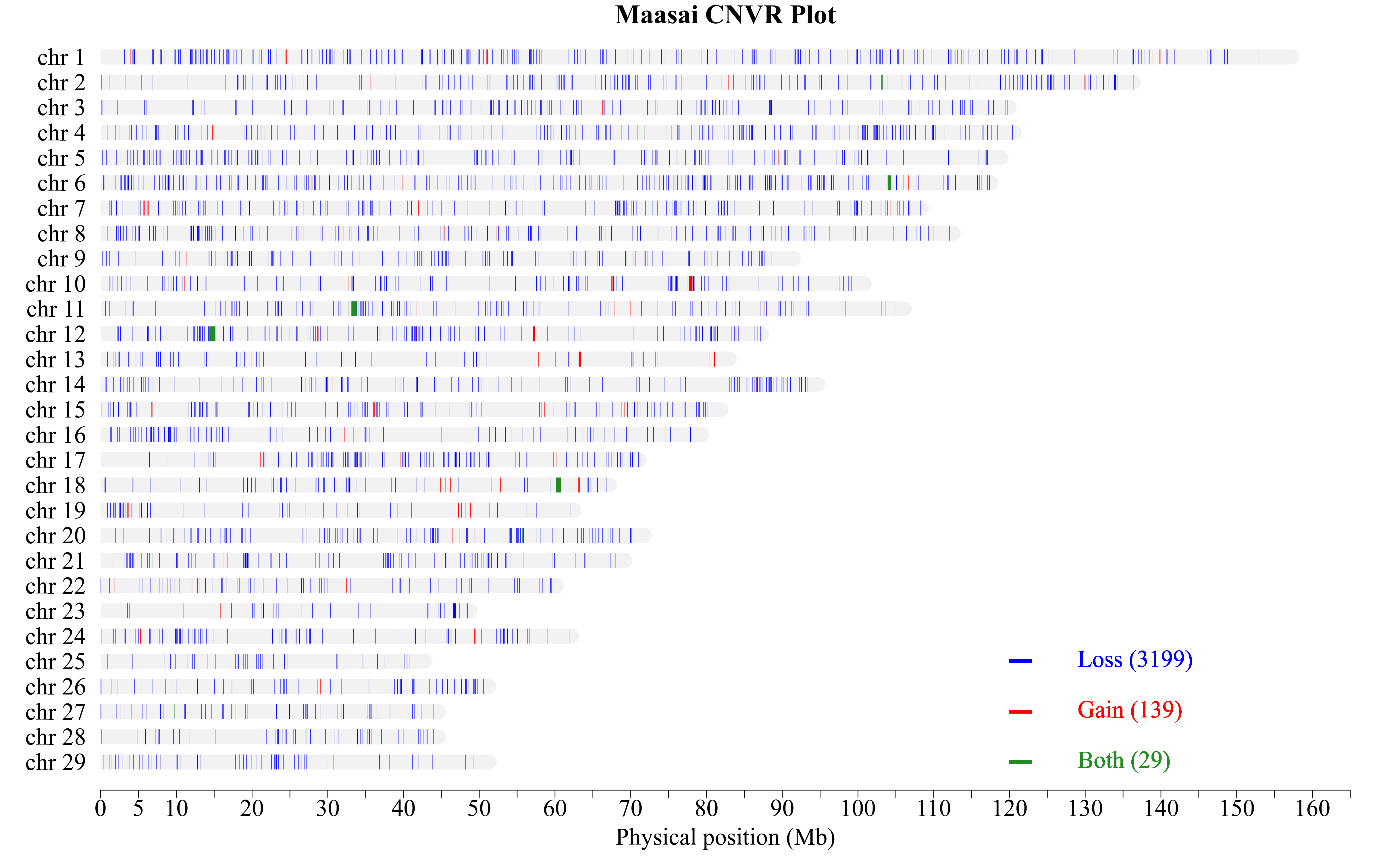
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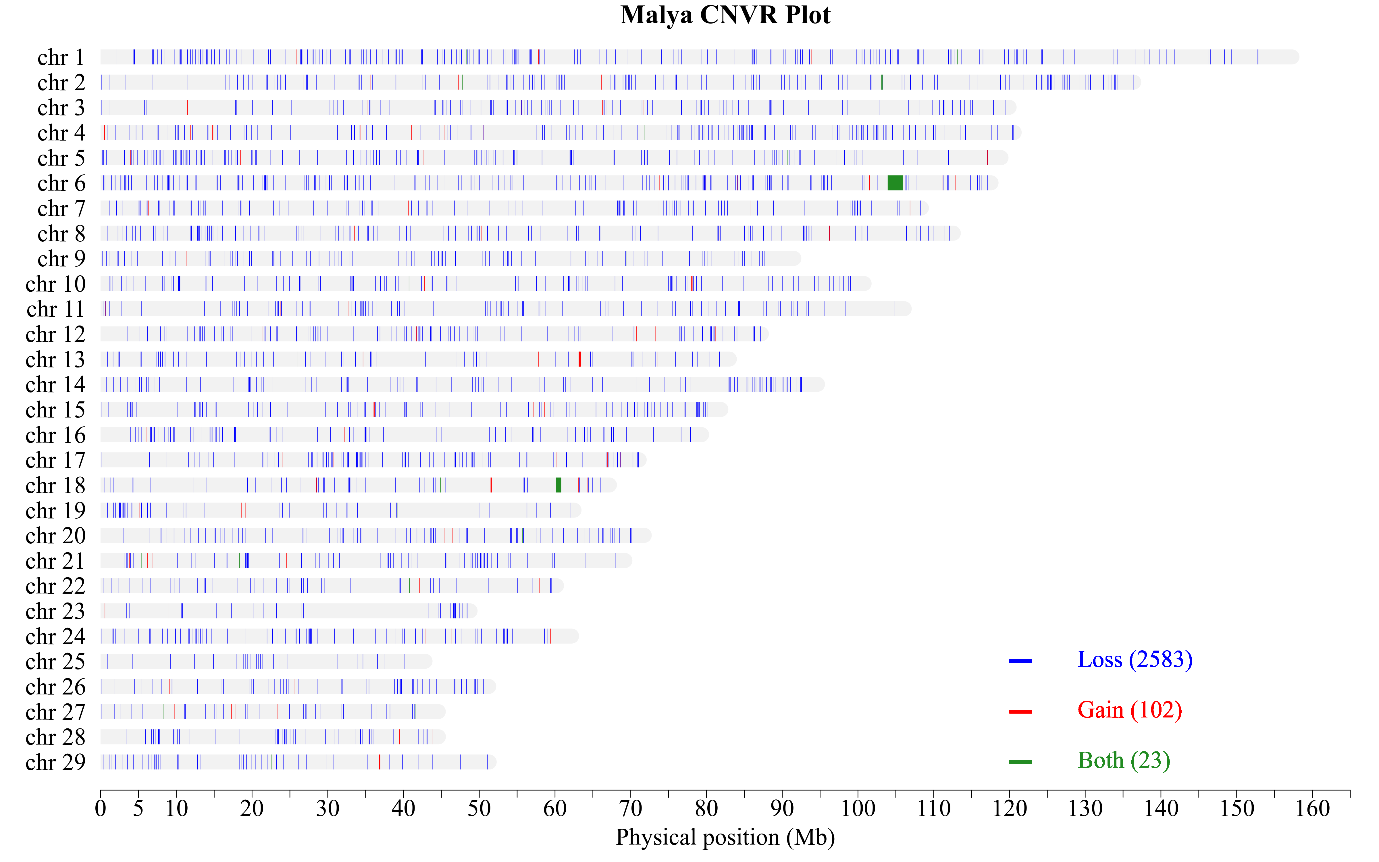
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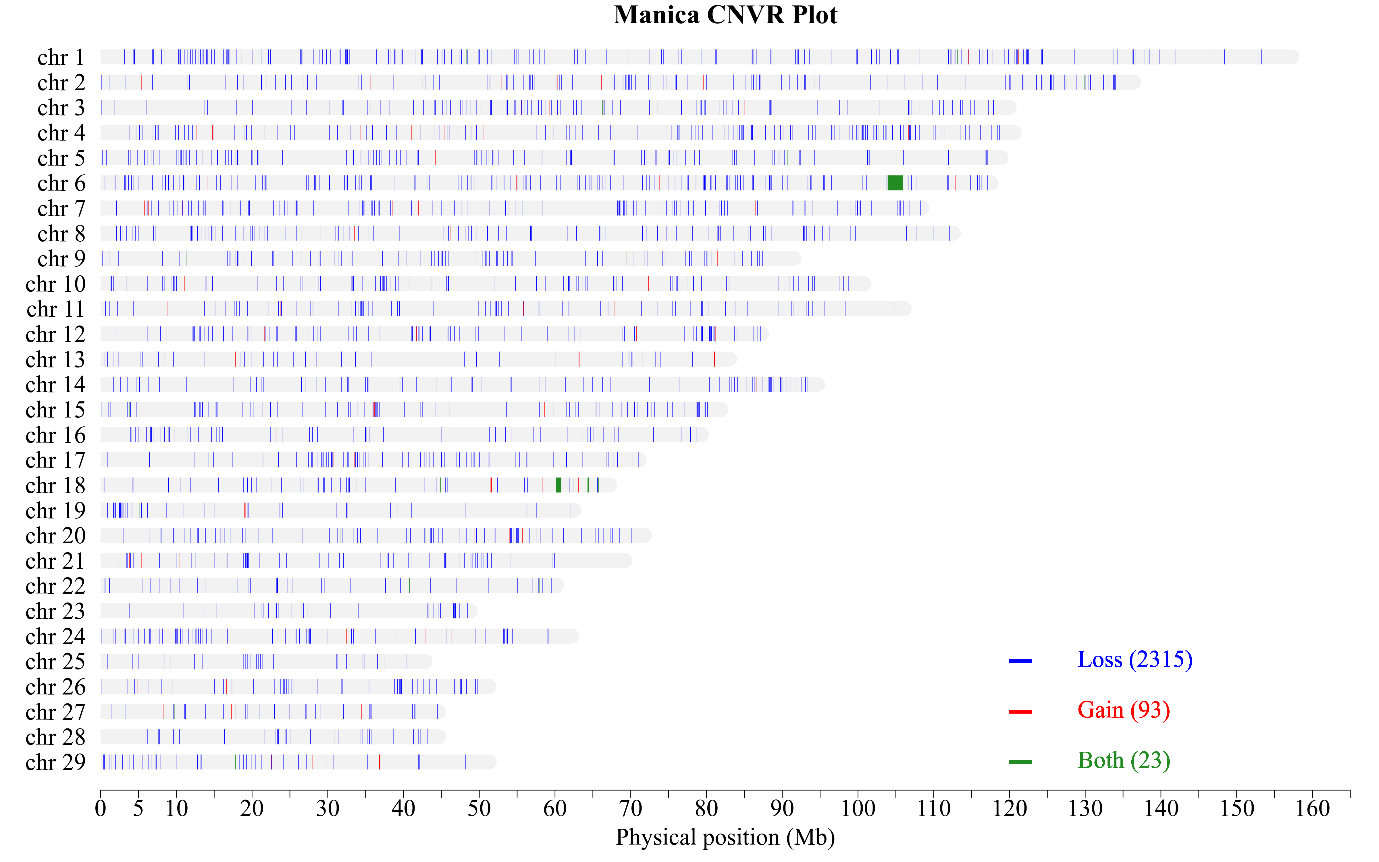
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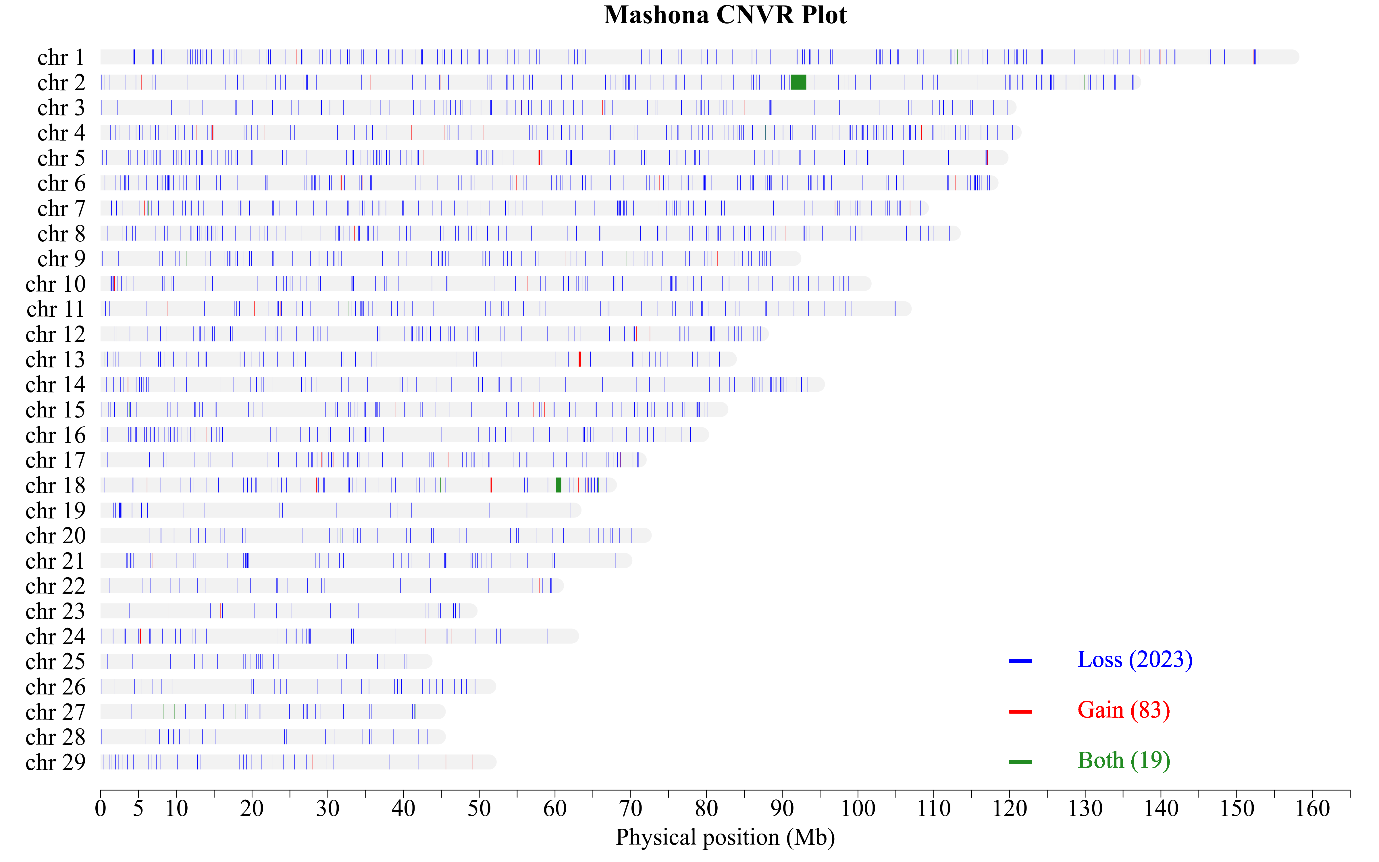
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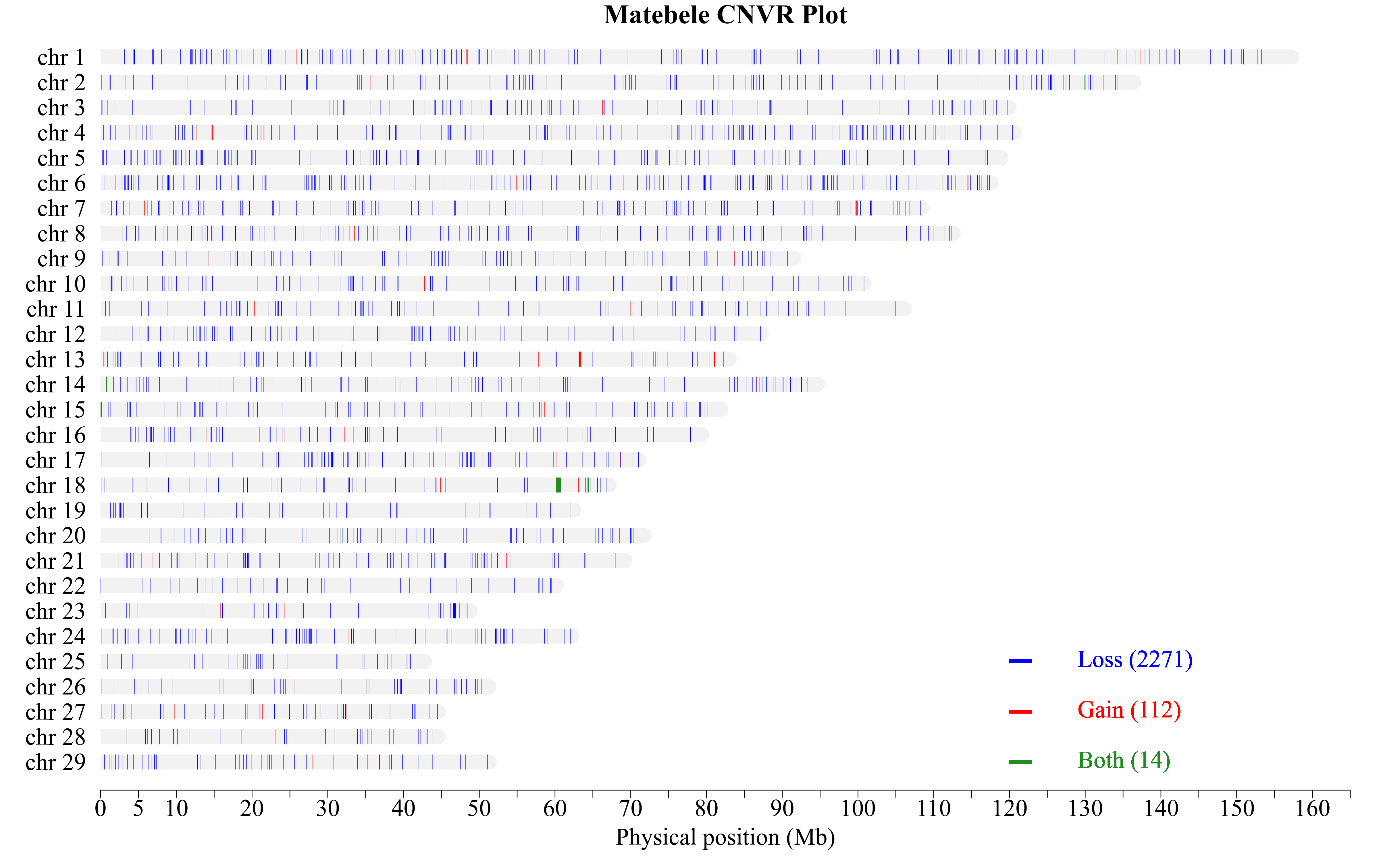
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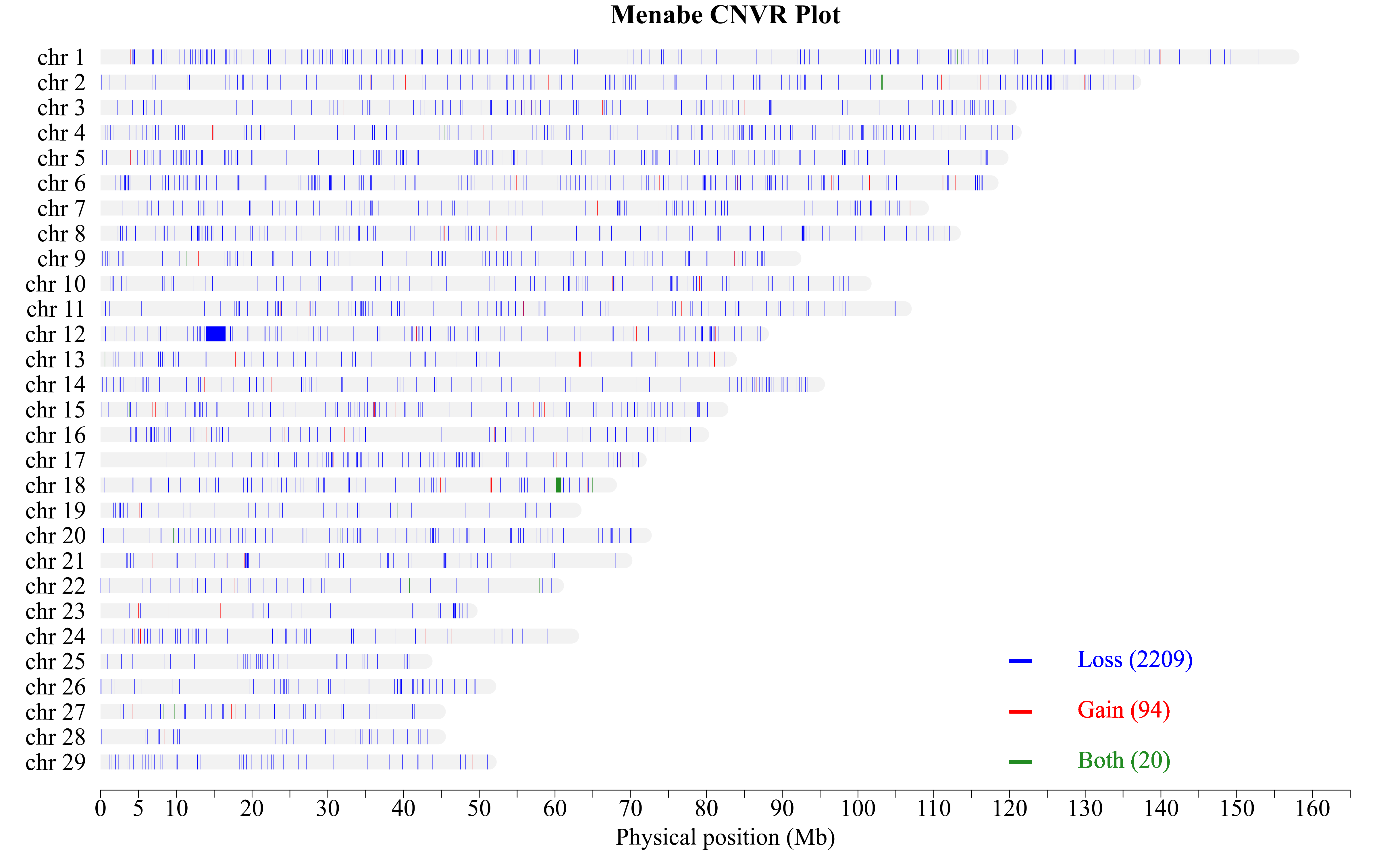
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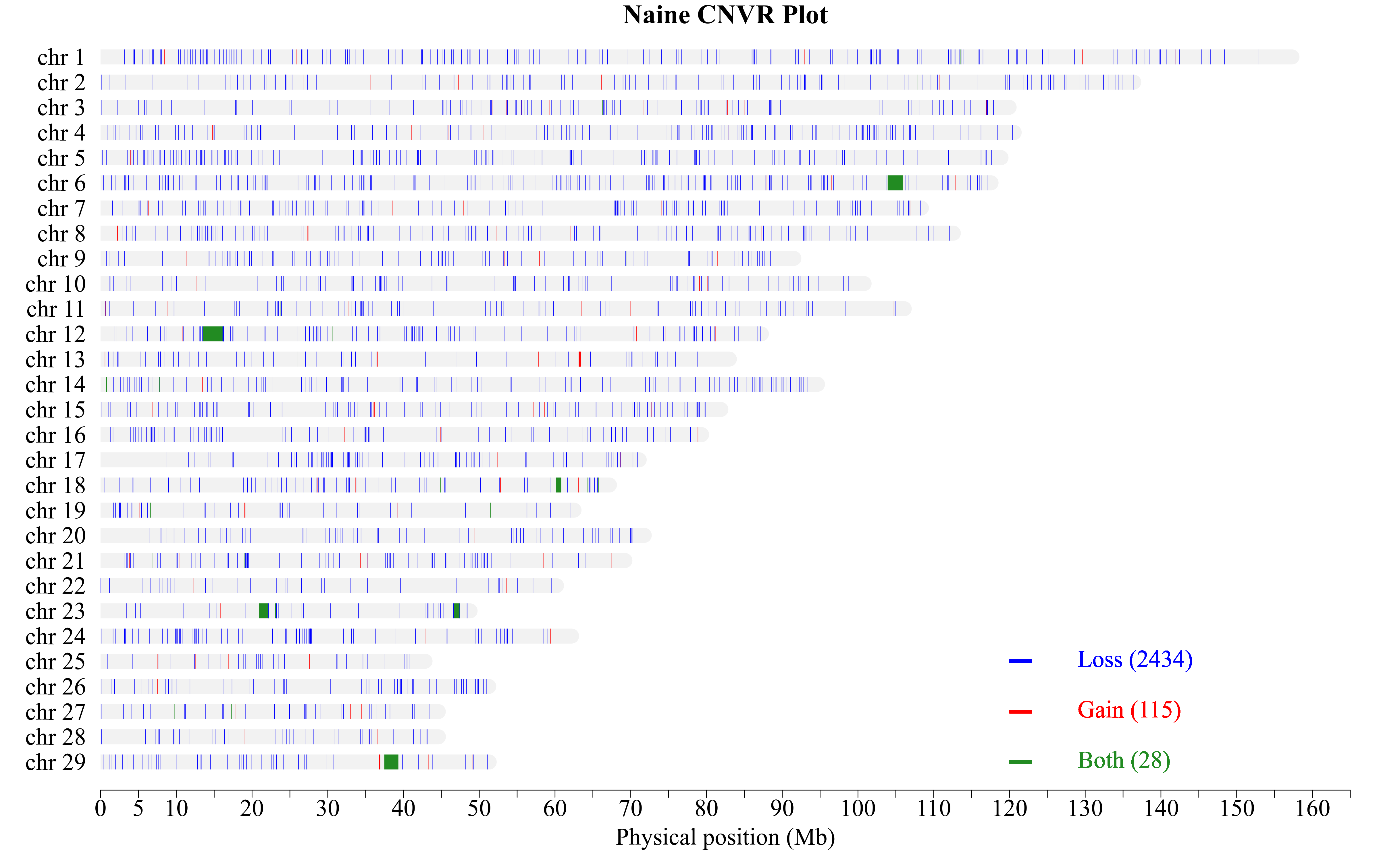
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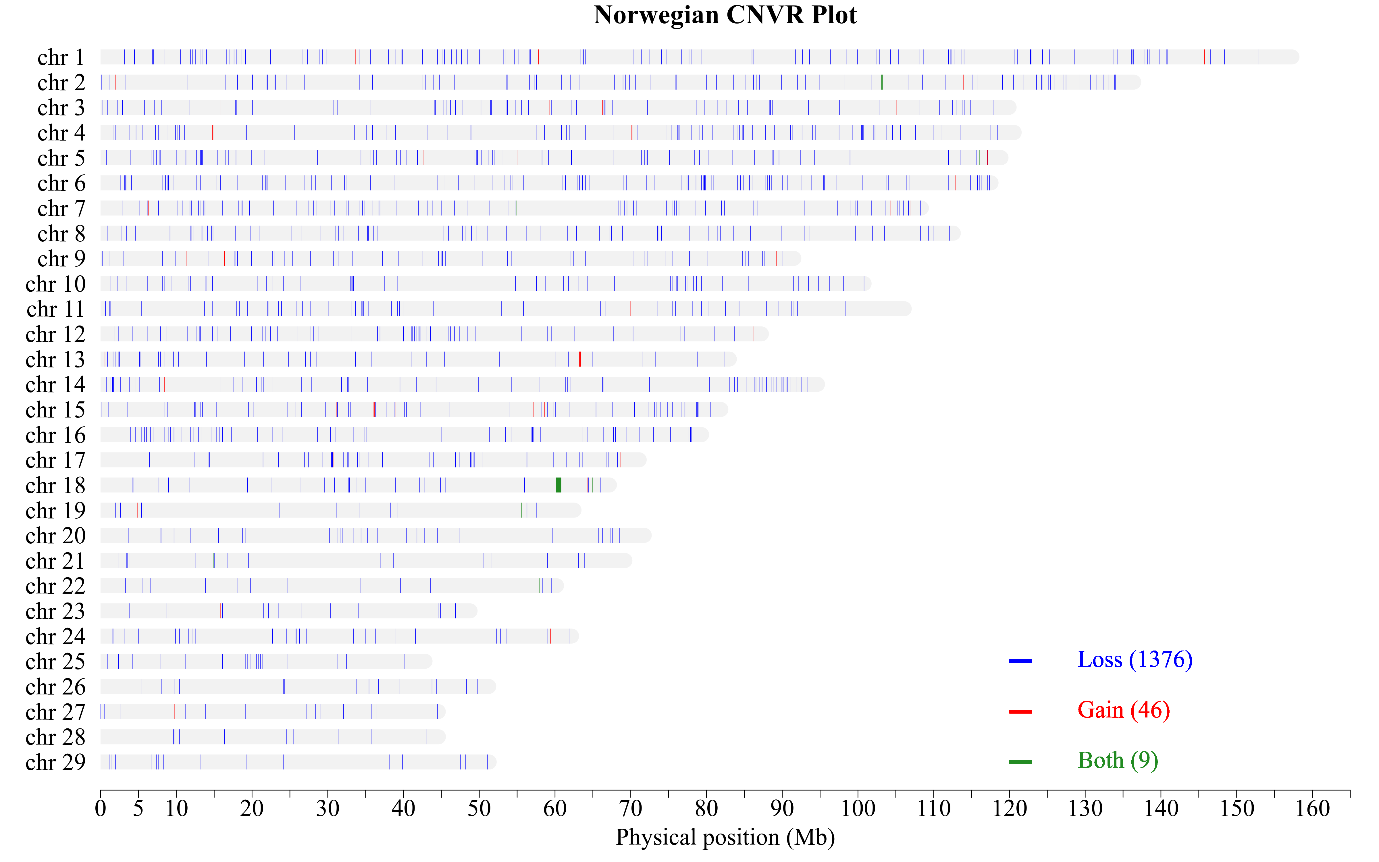
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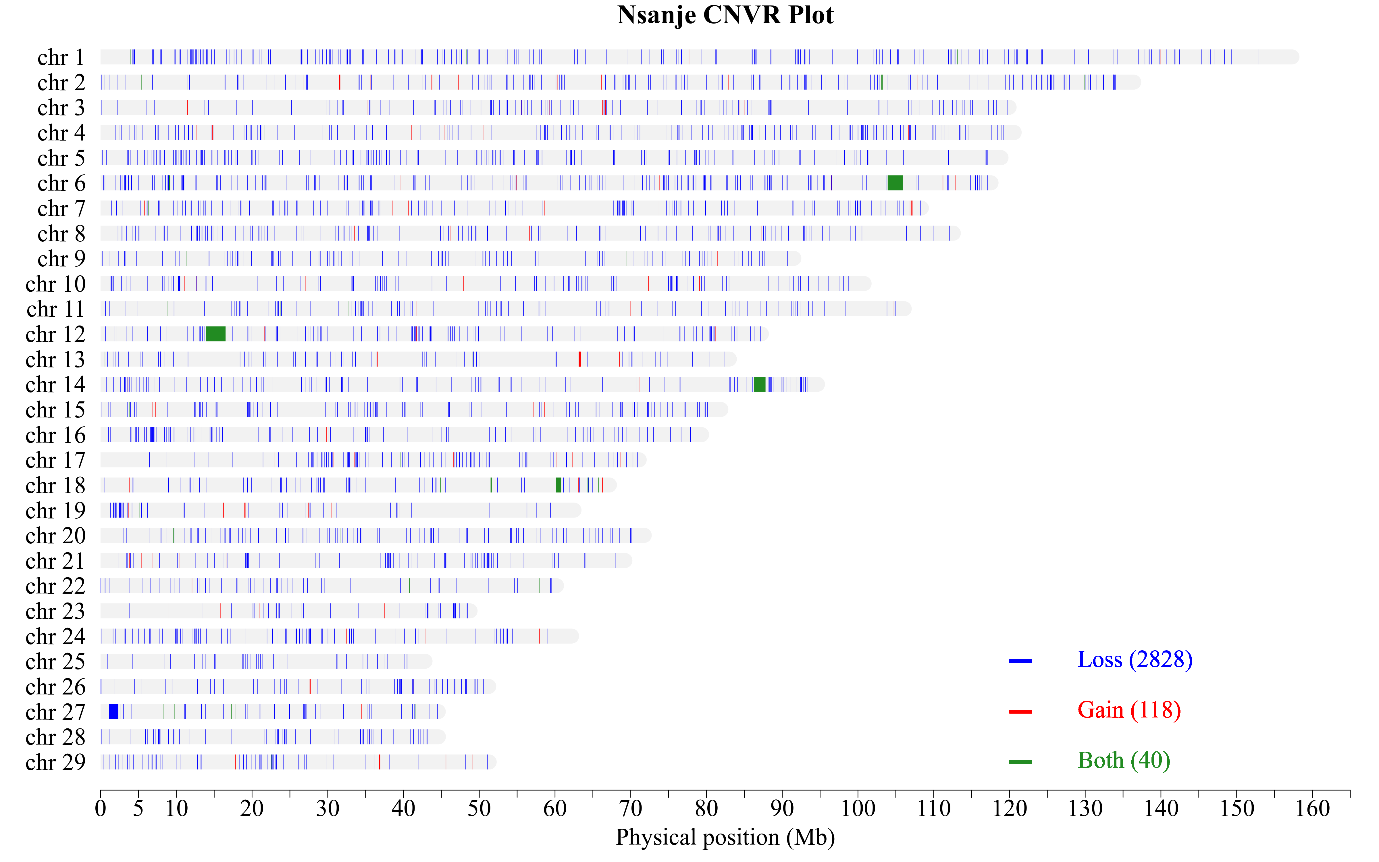
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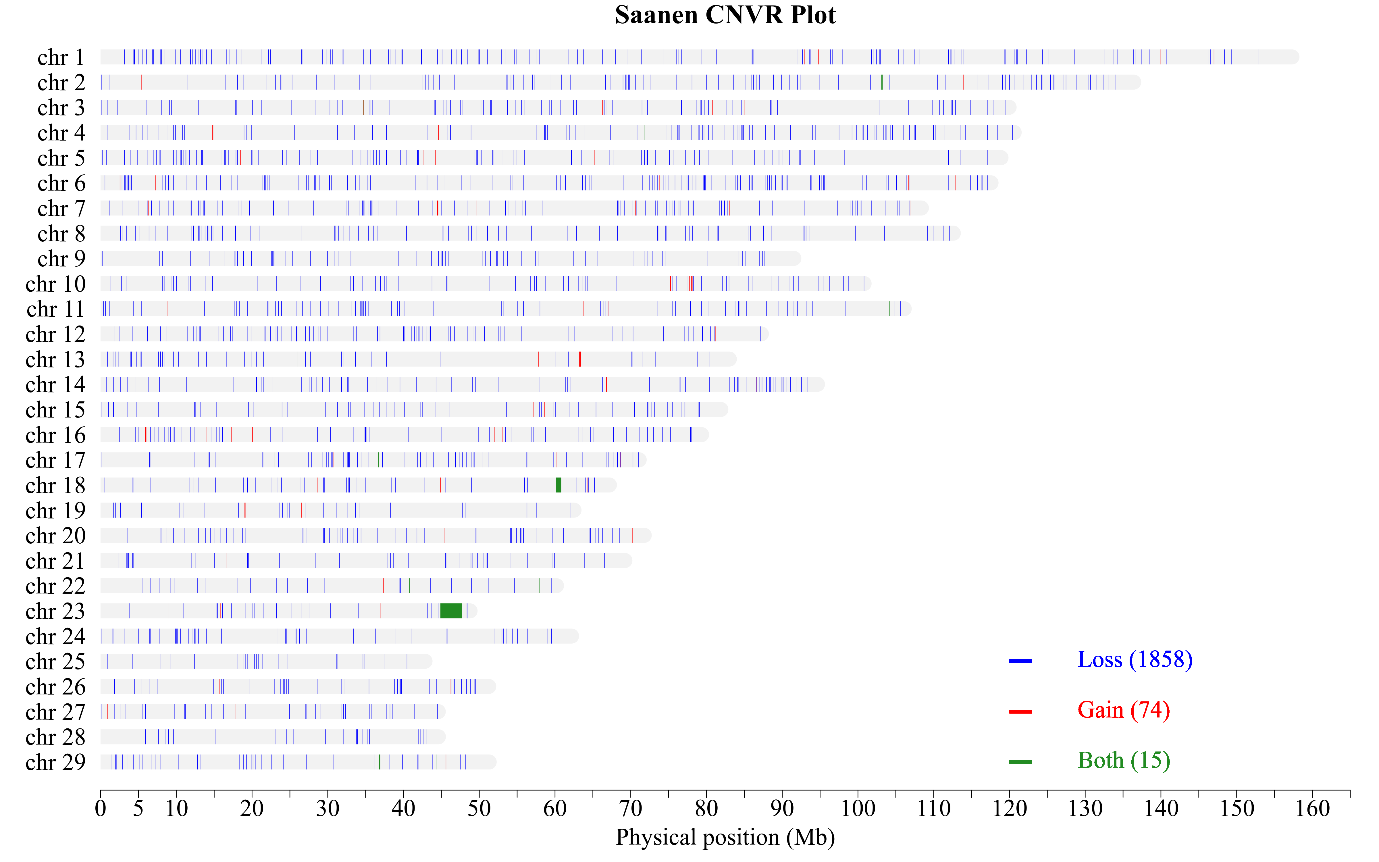
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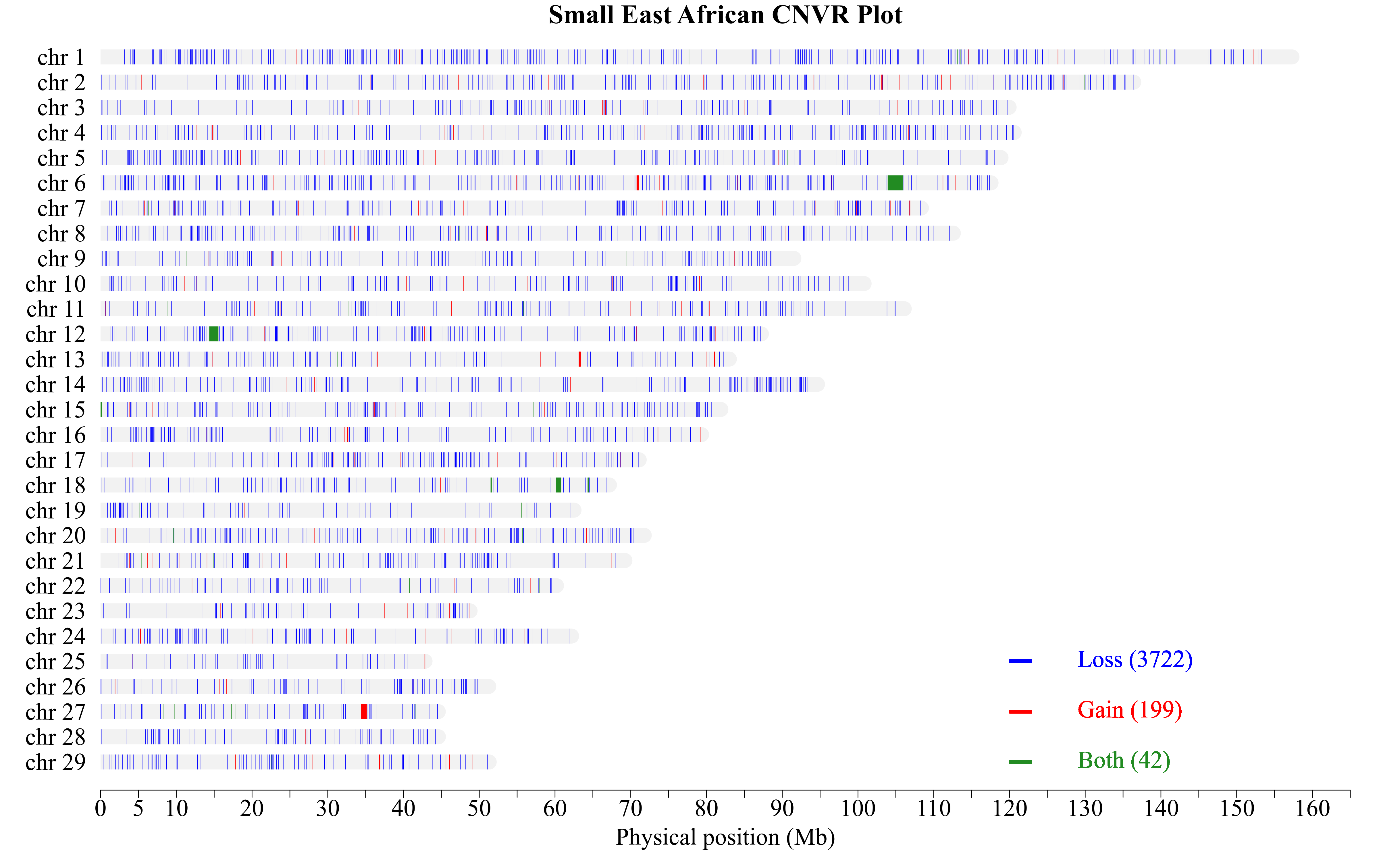
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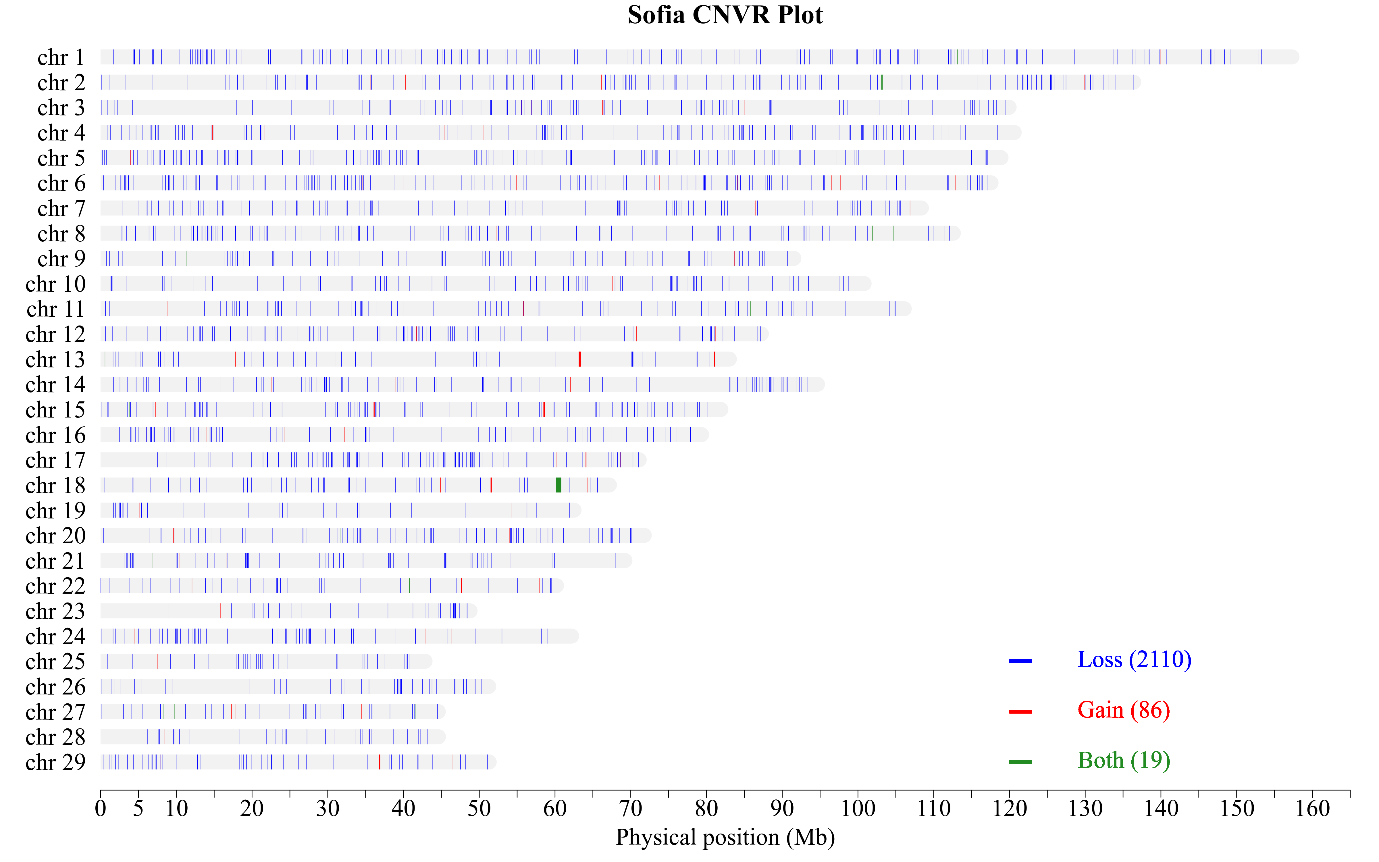
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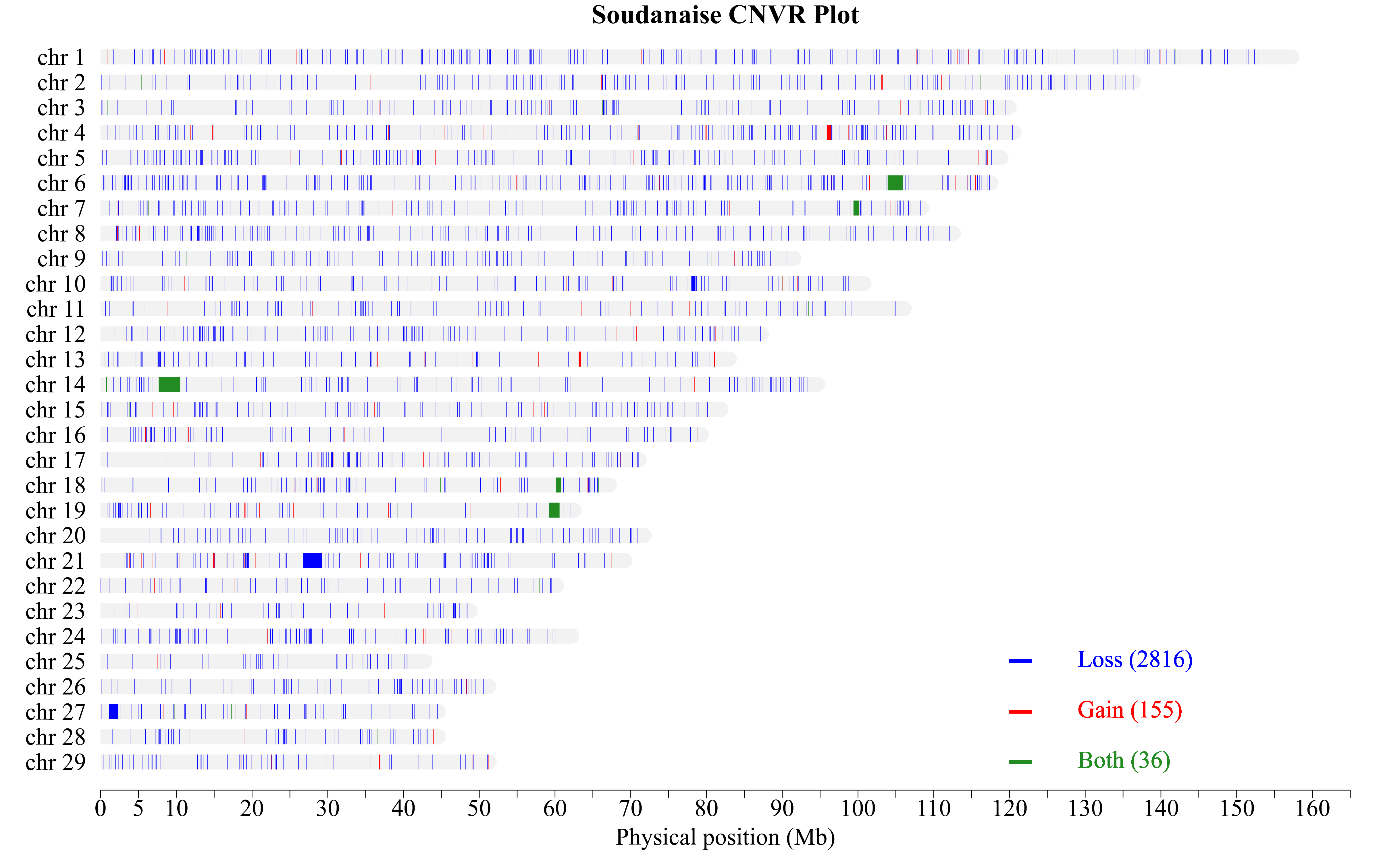
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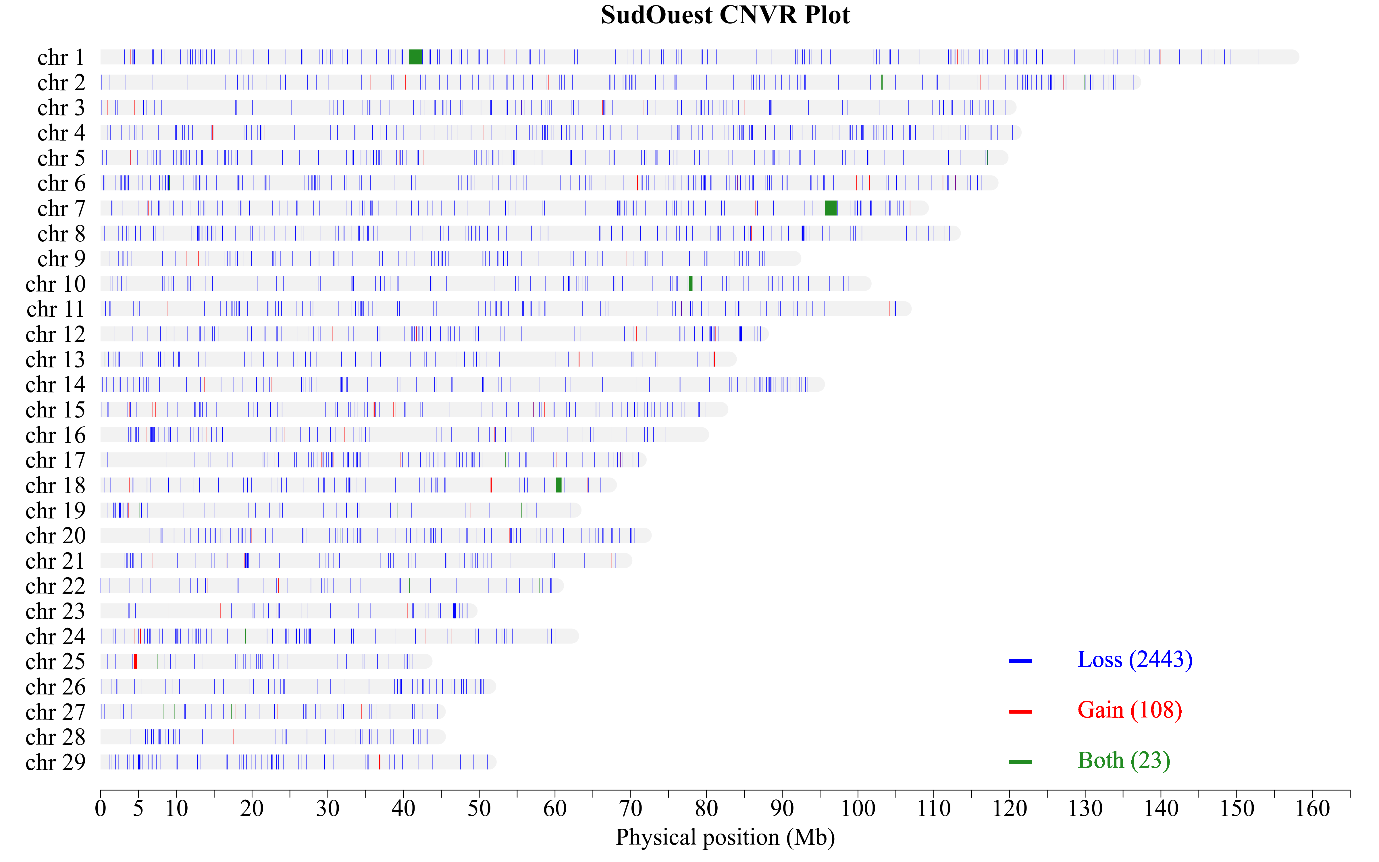
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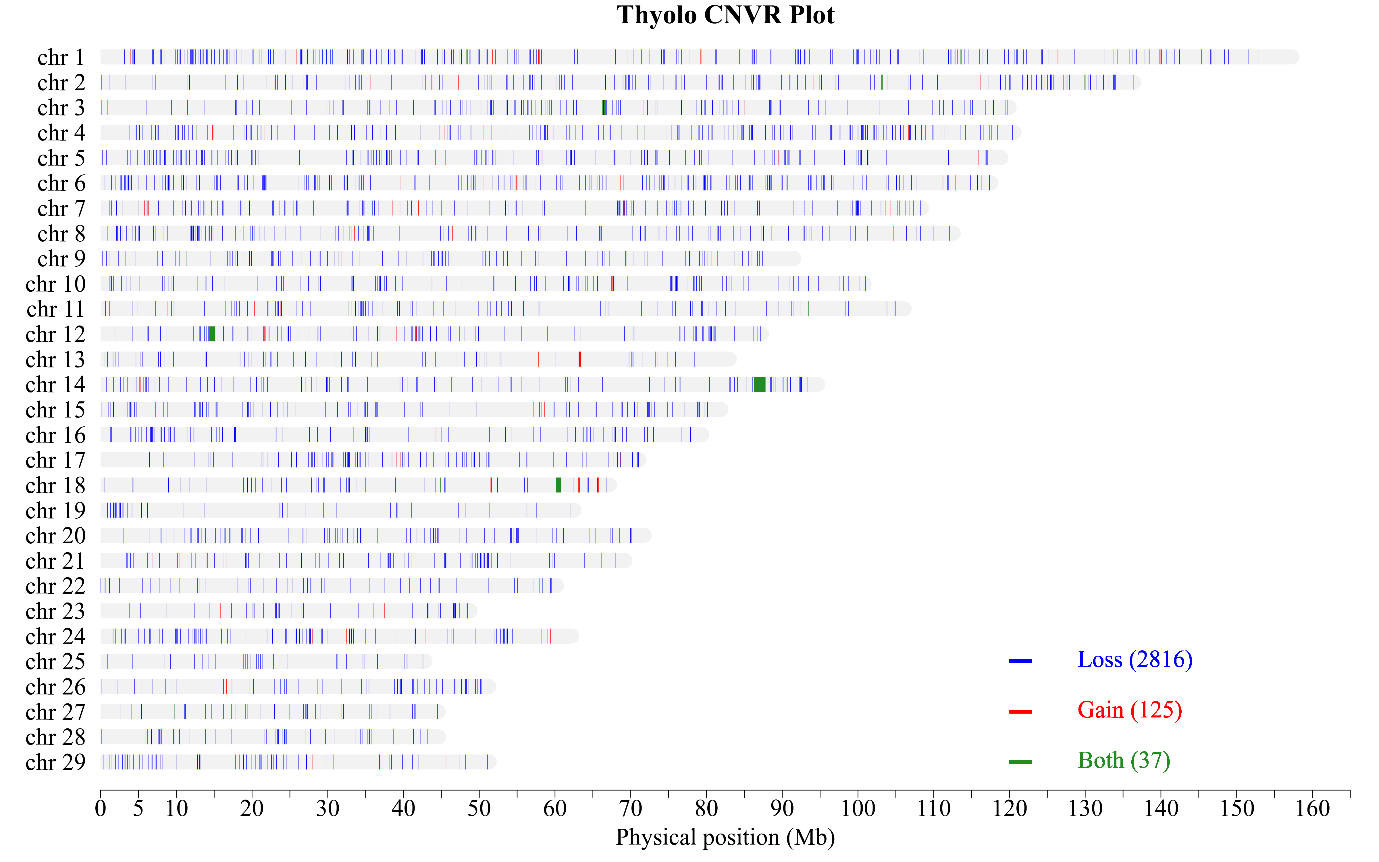
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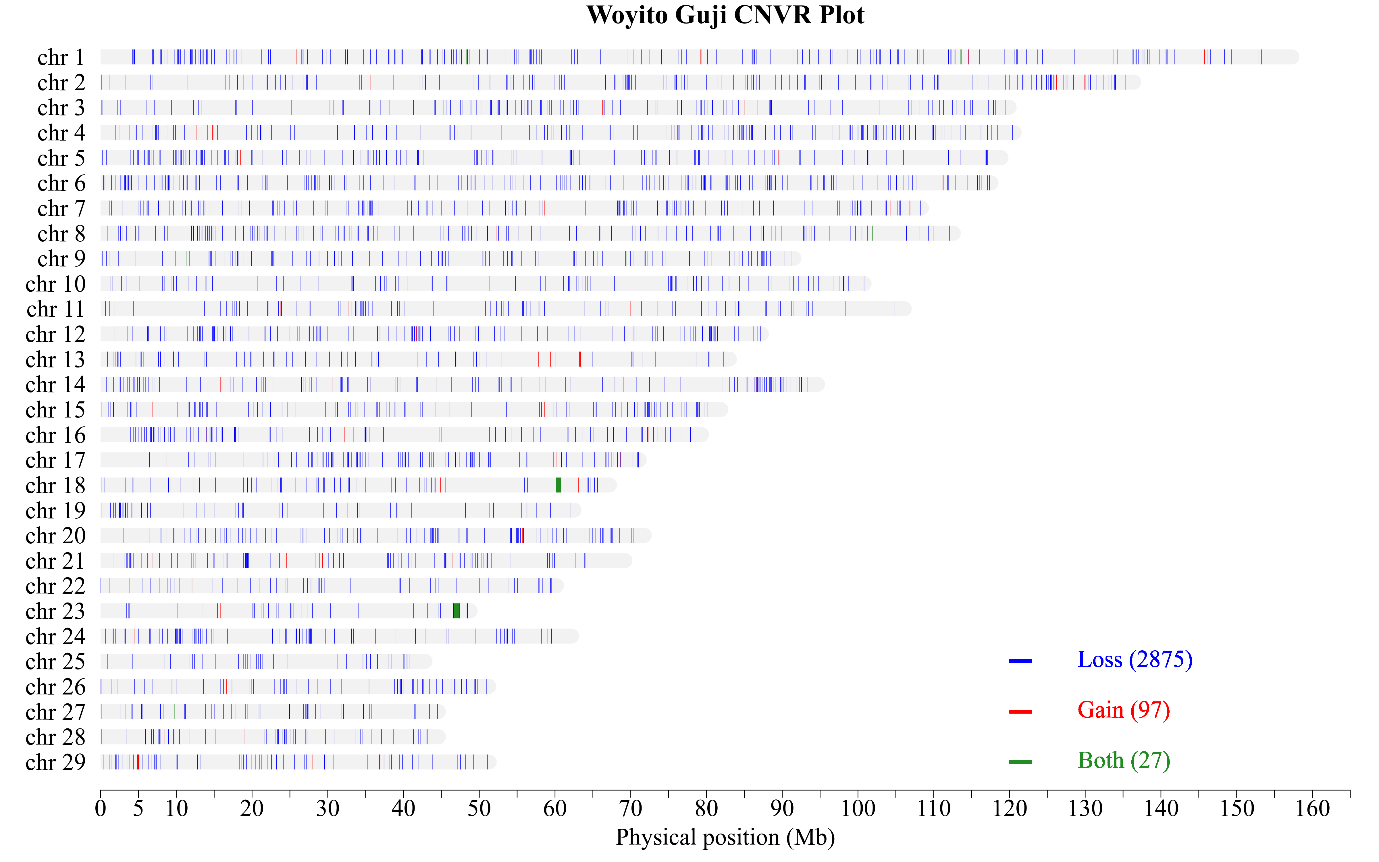
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