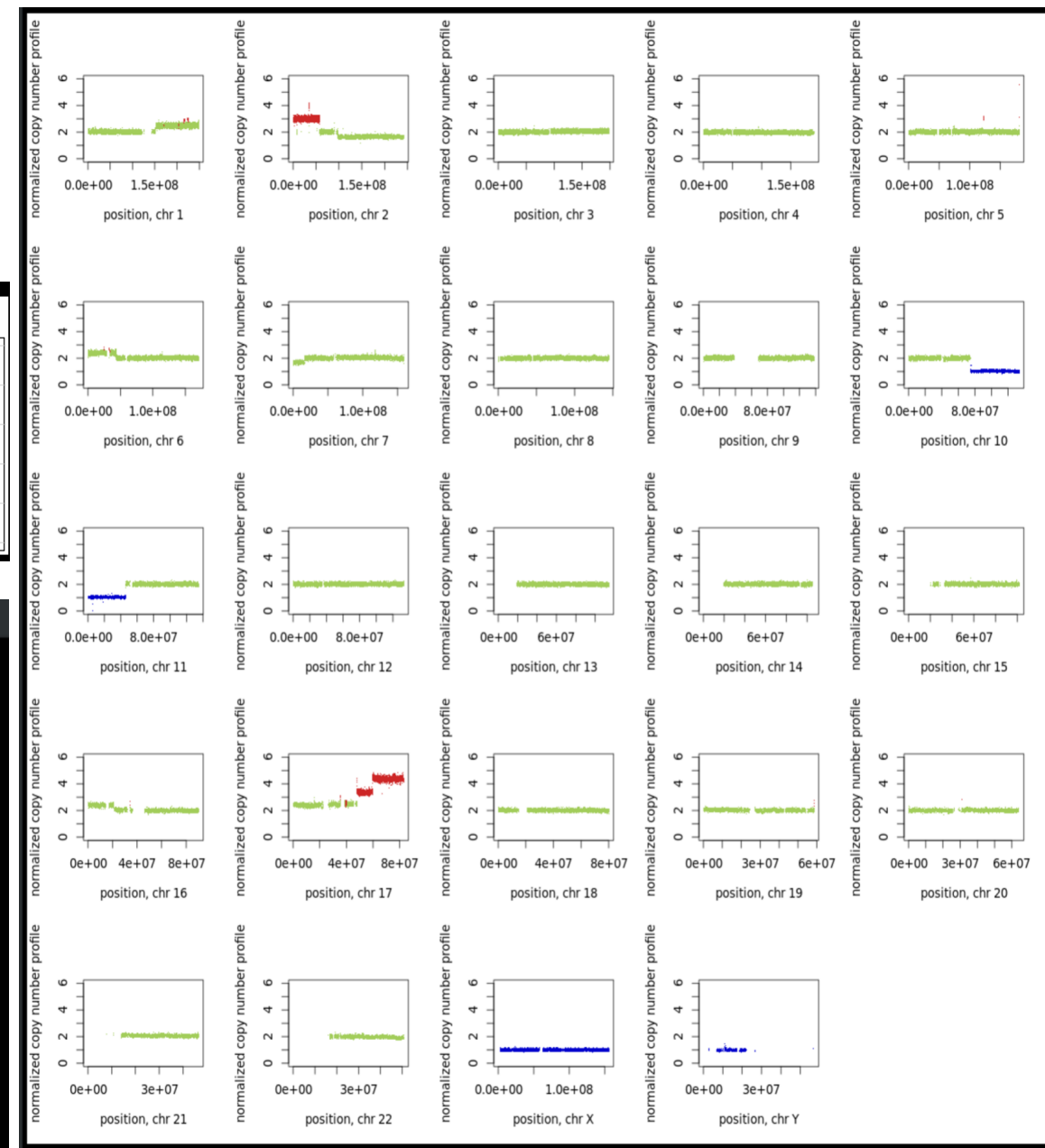
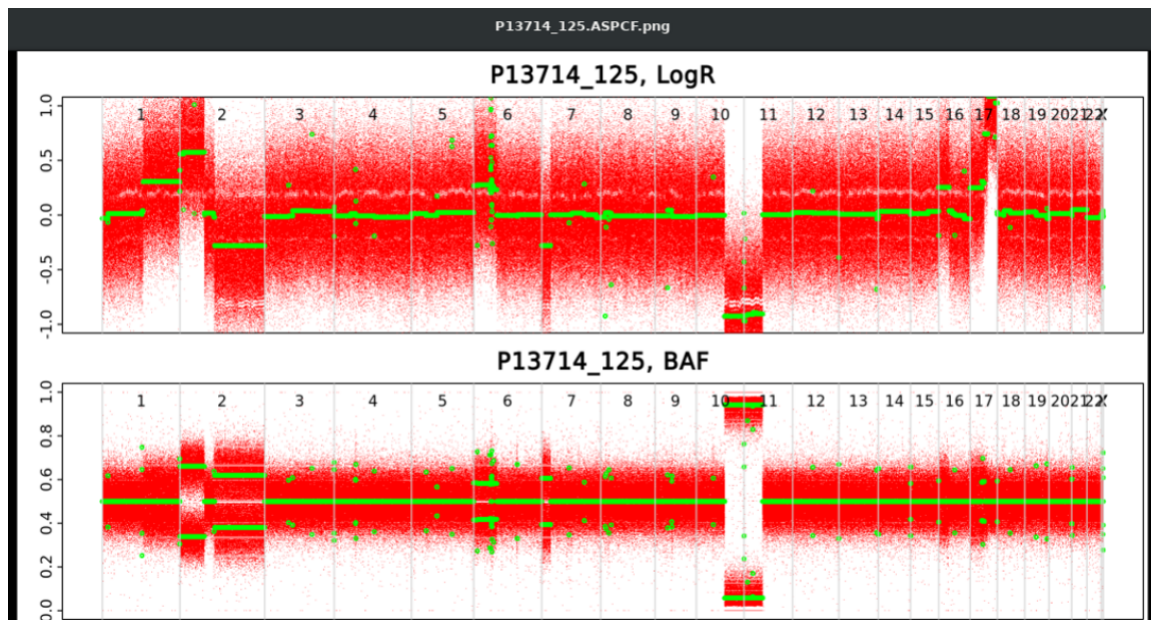
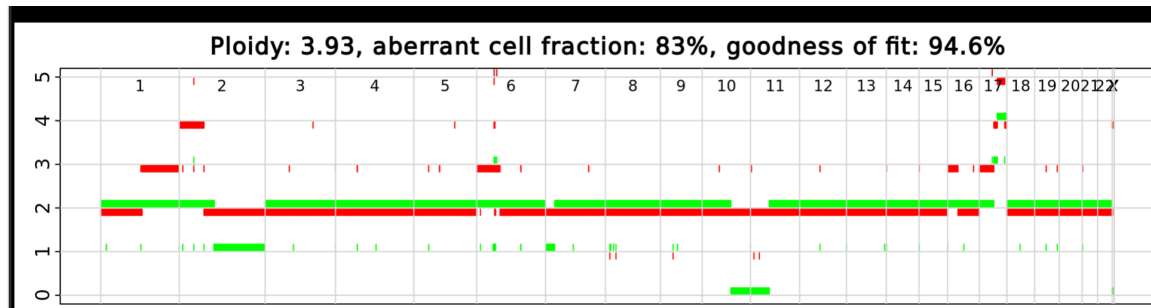
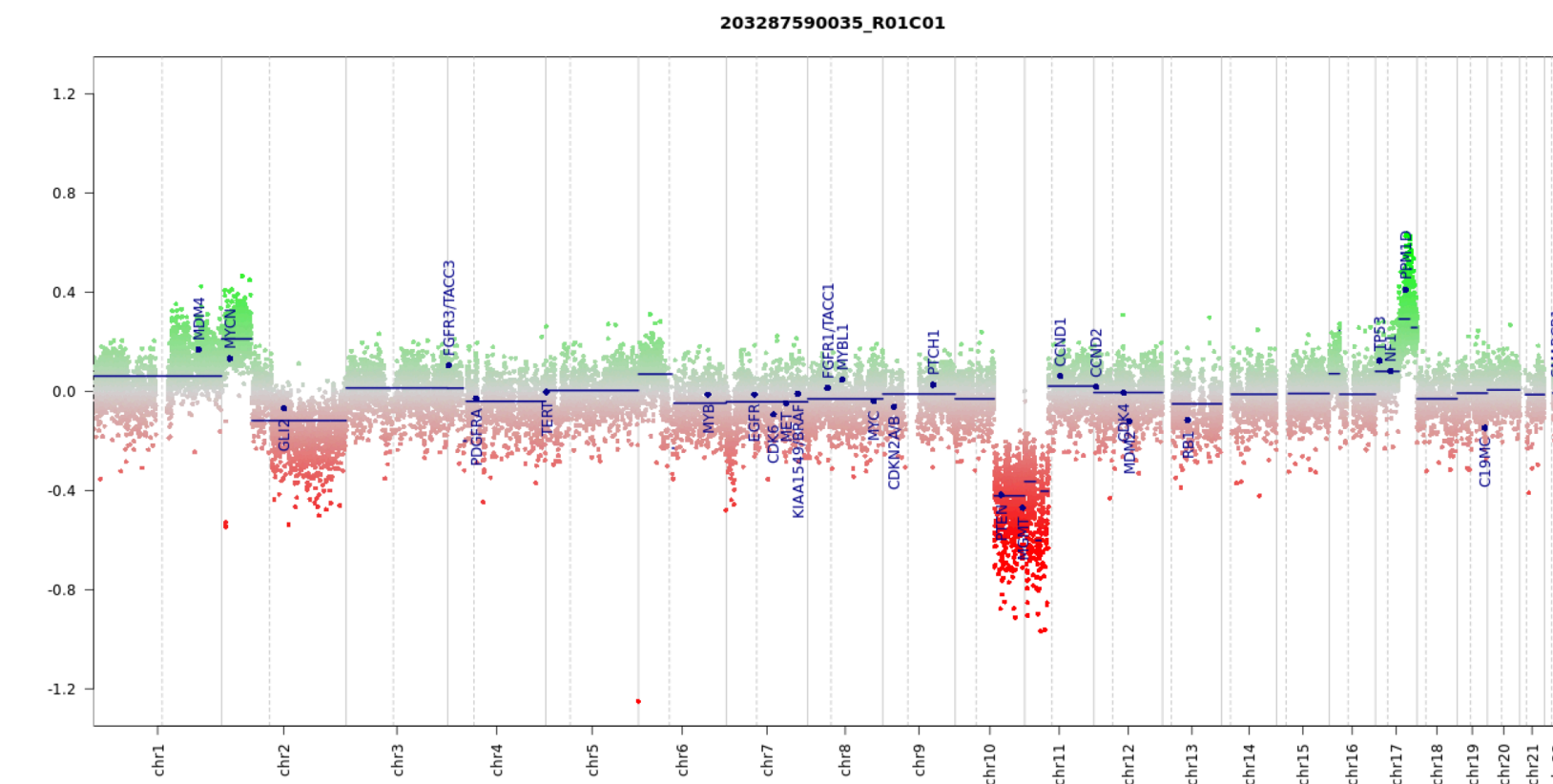


MB1

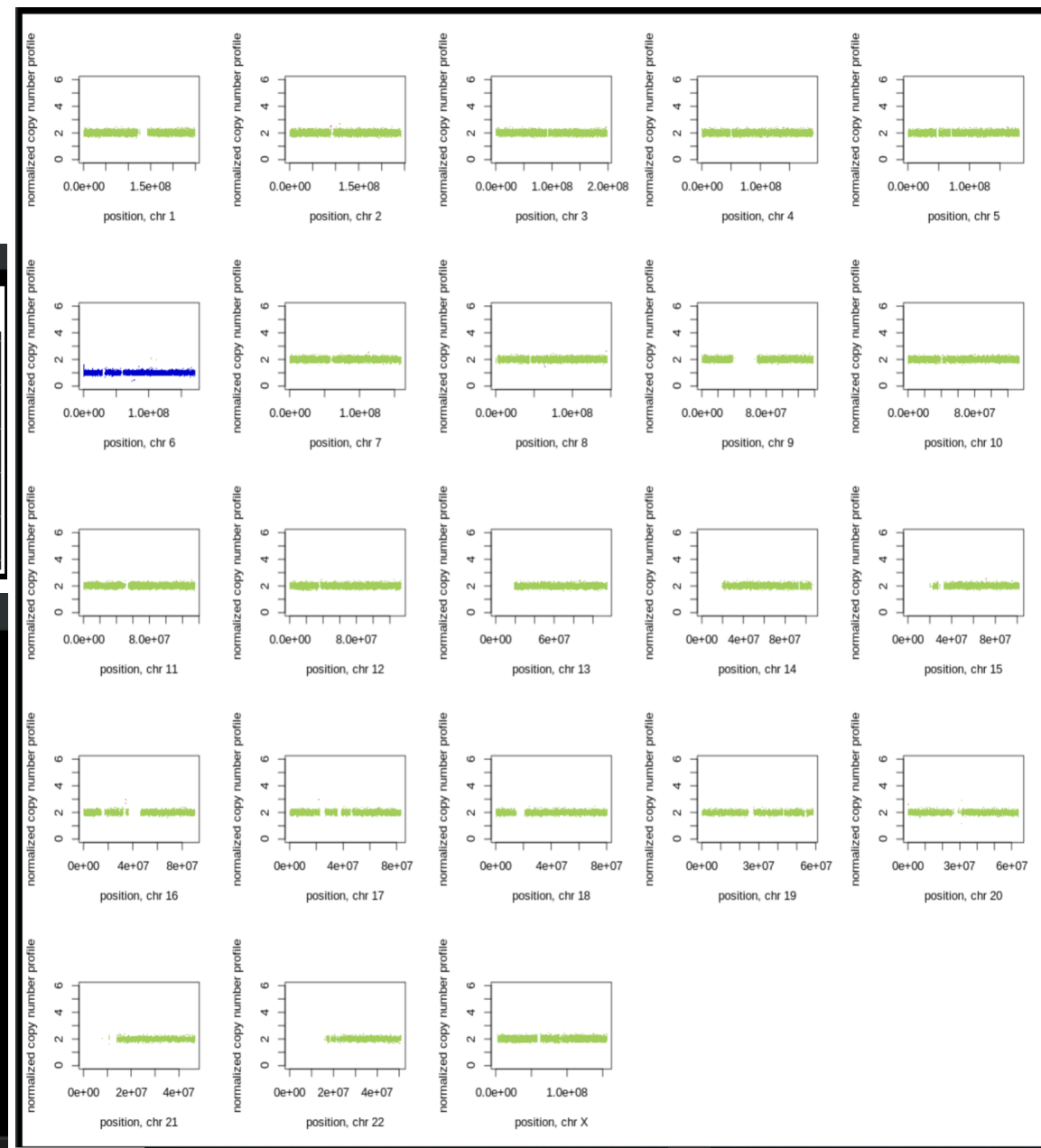
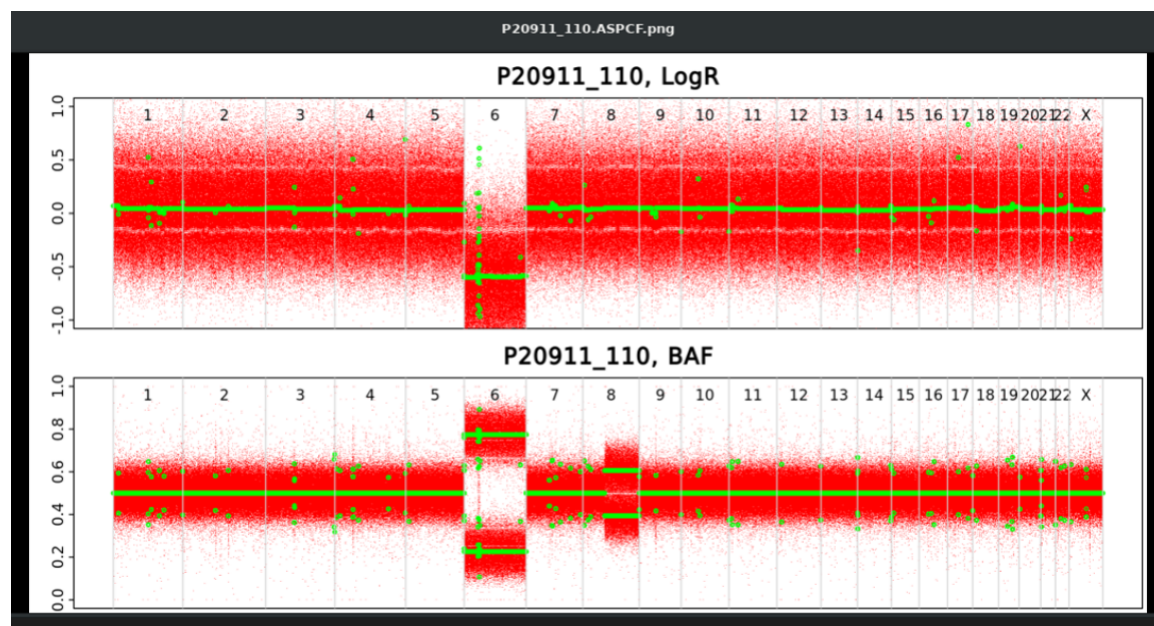
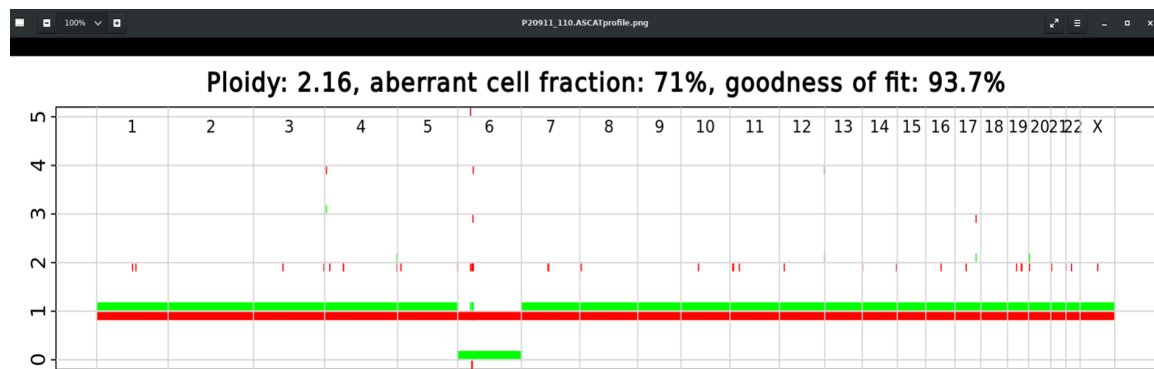


MB1

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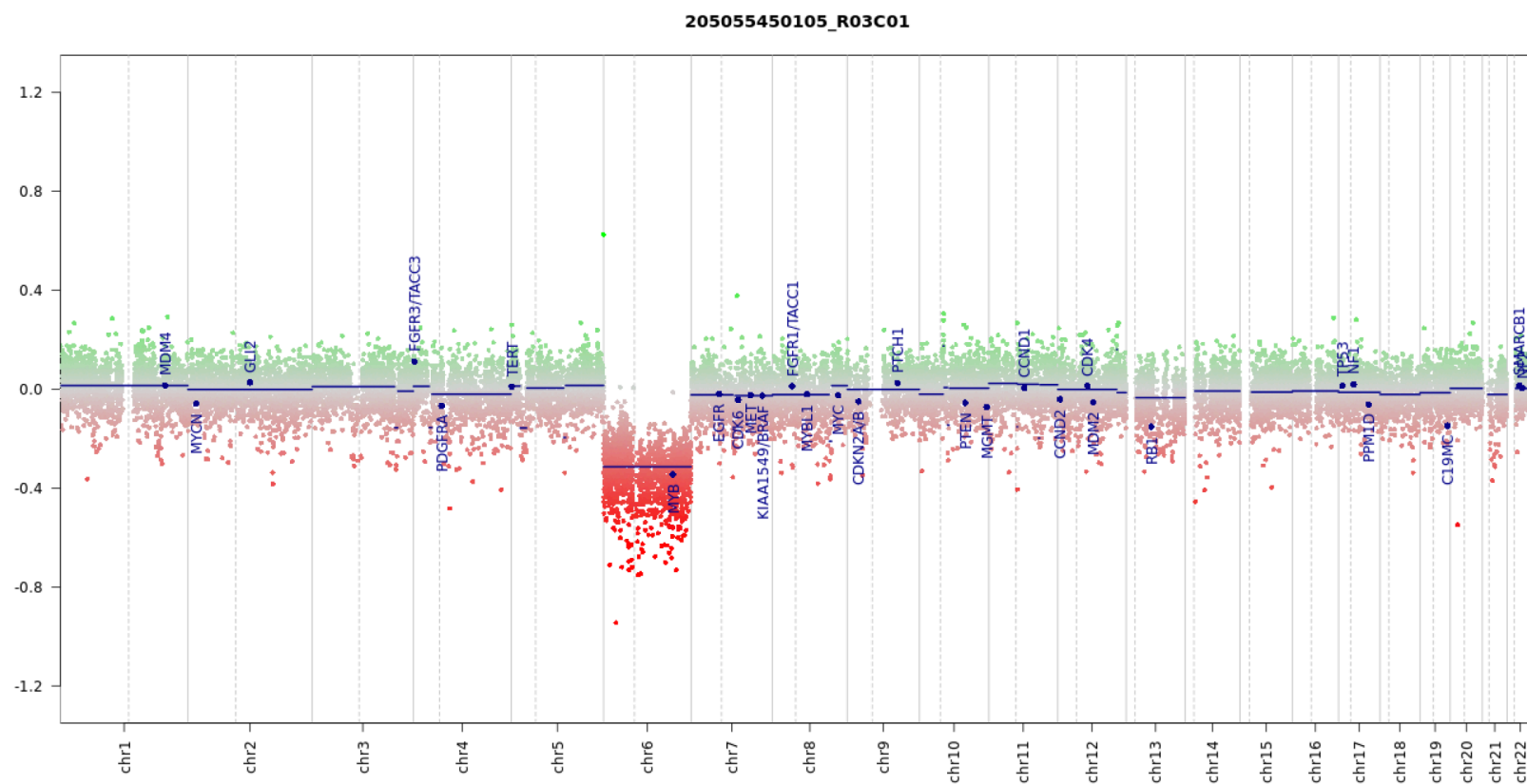


MB9

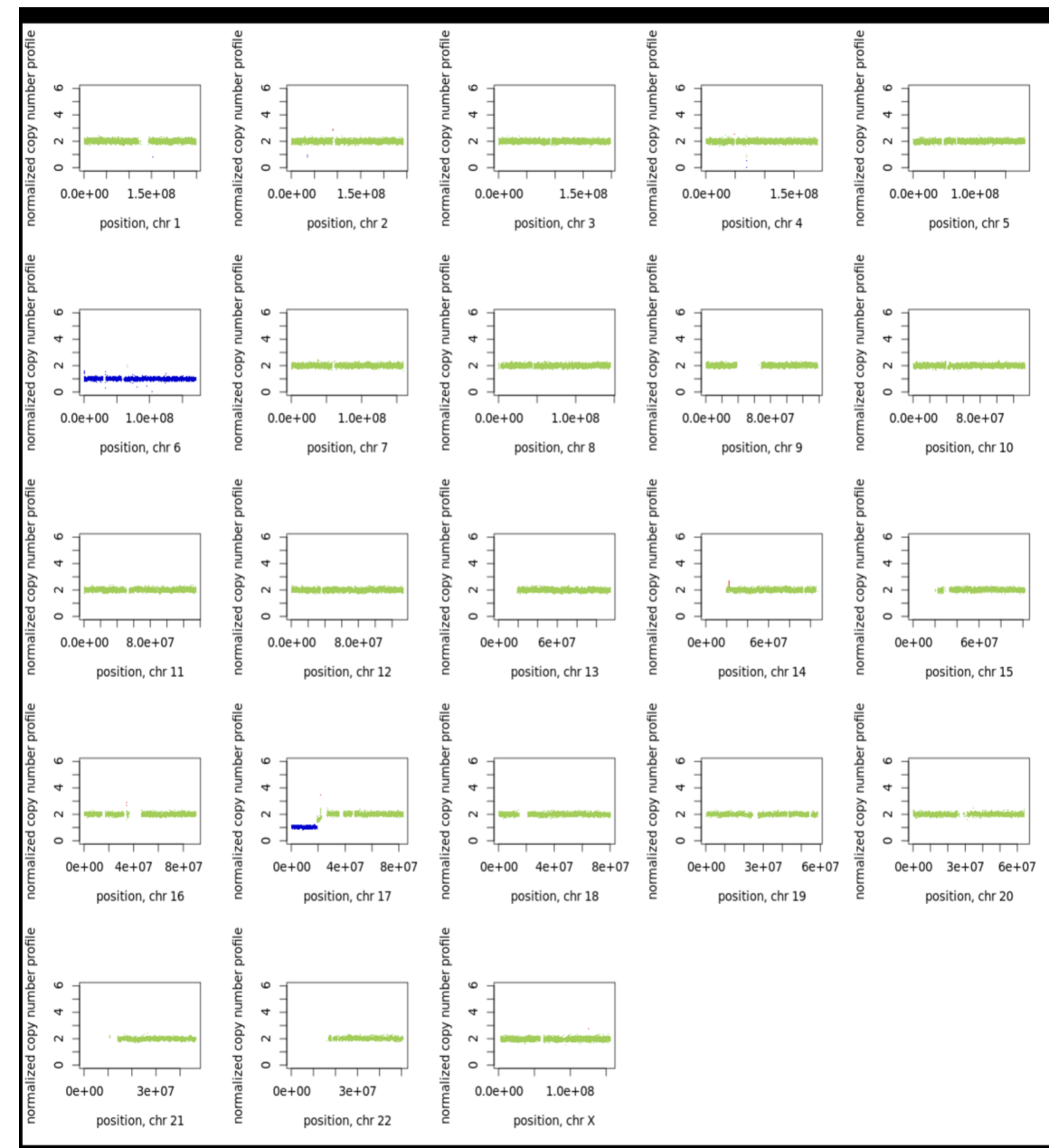
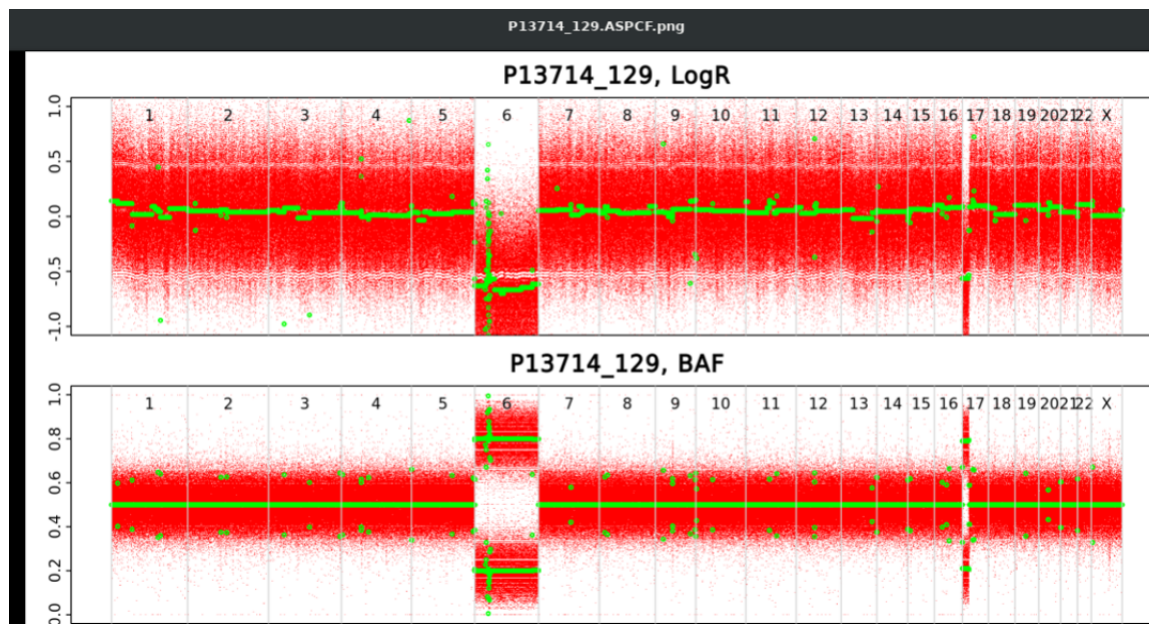
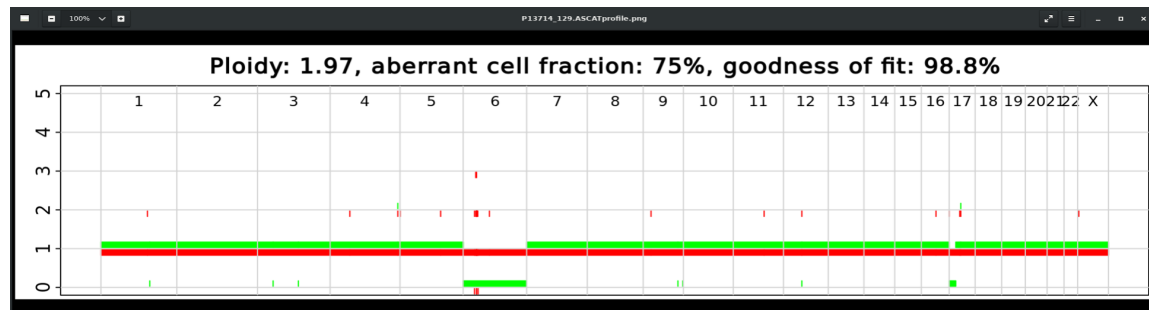


MB9

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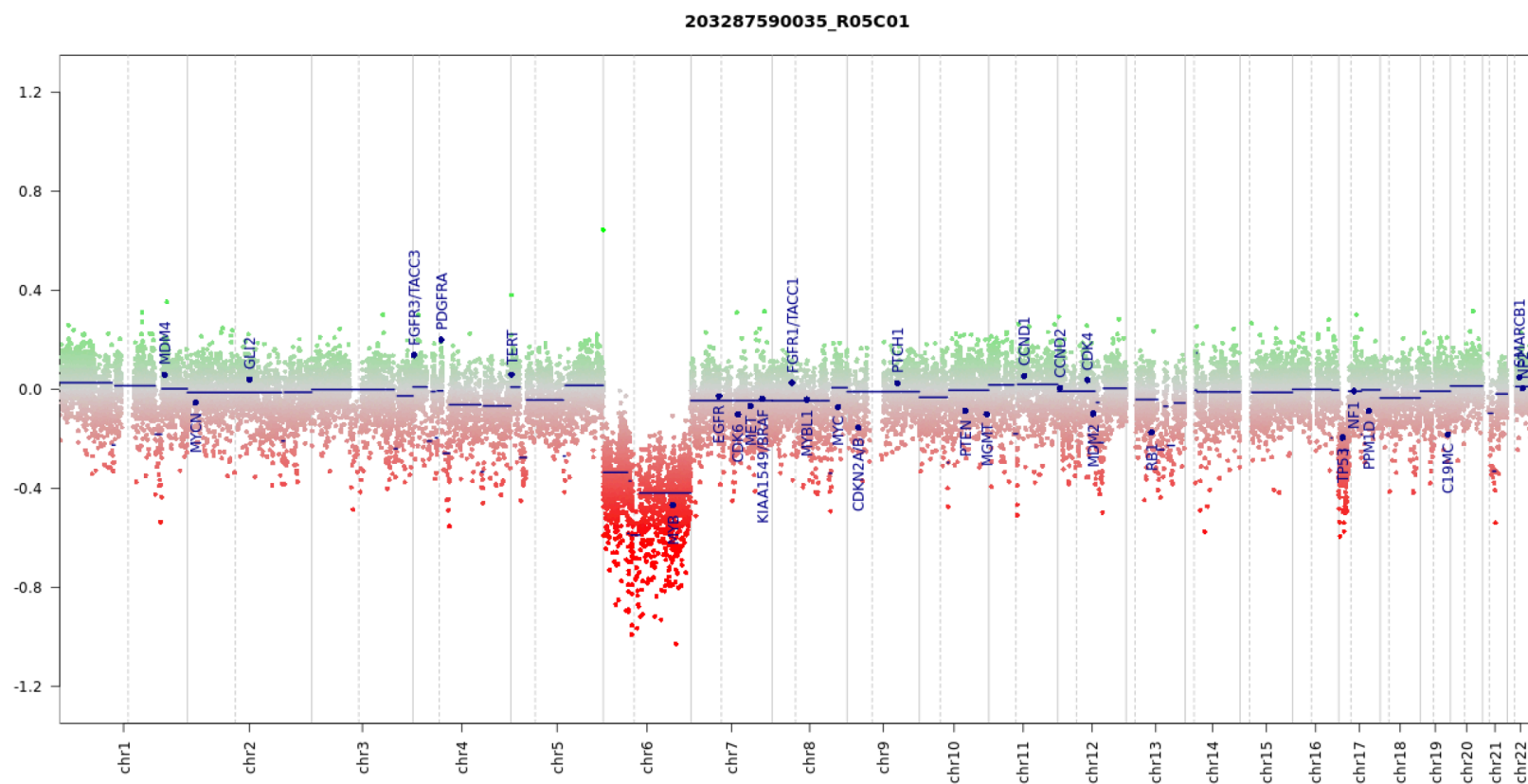


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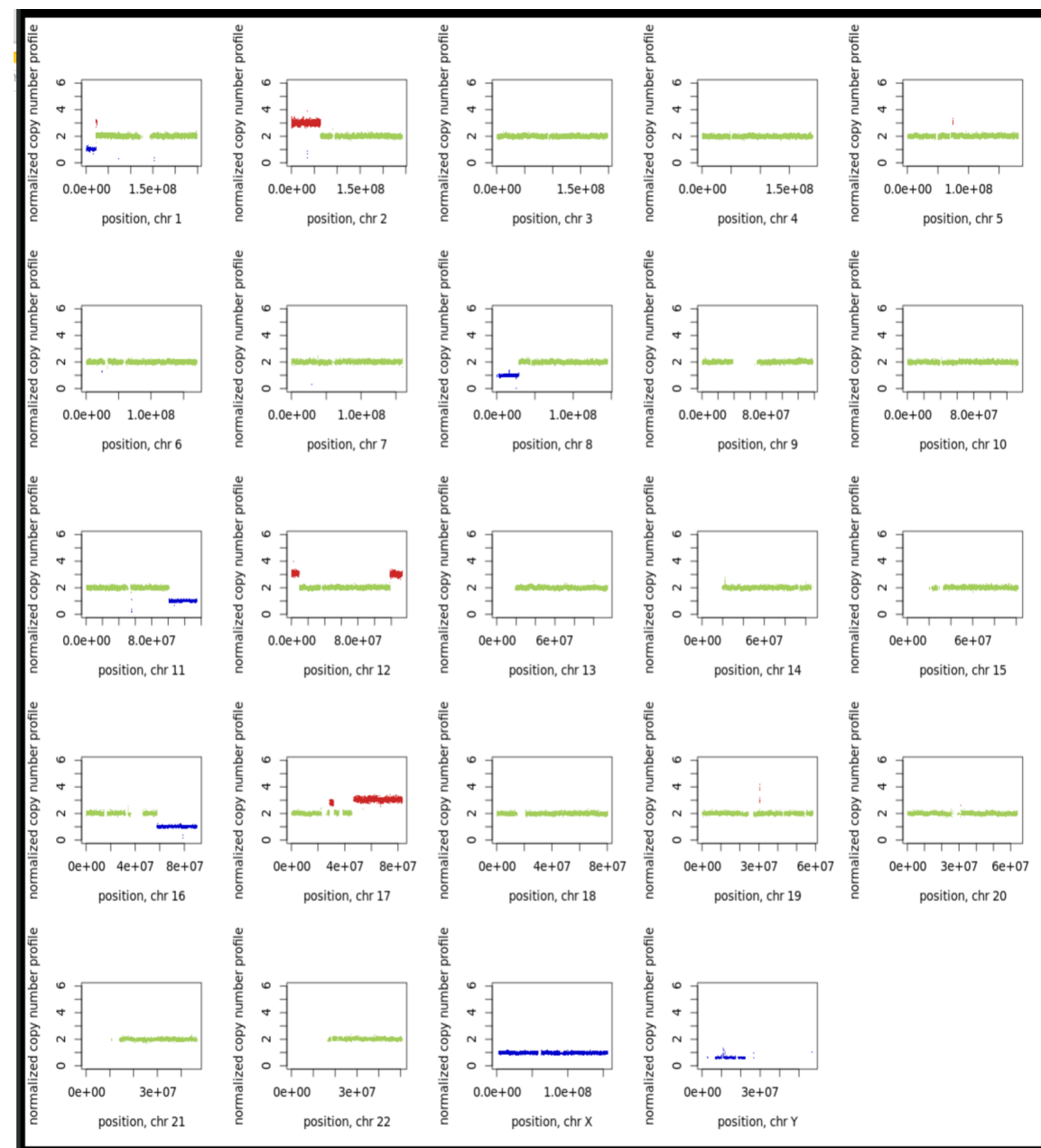
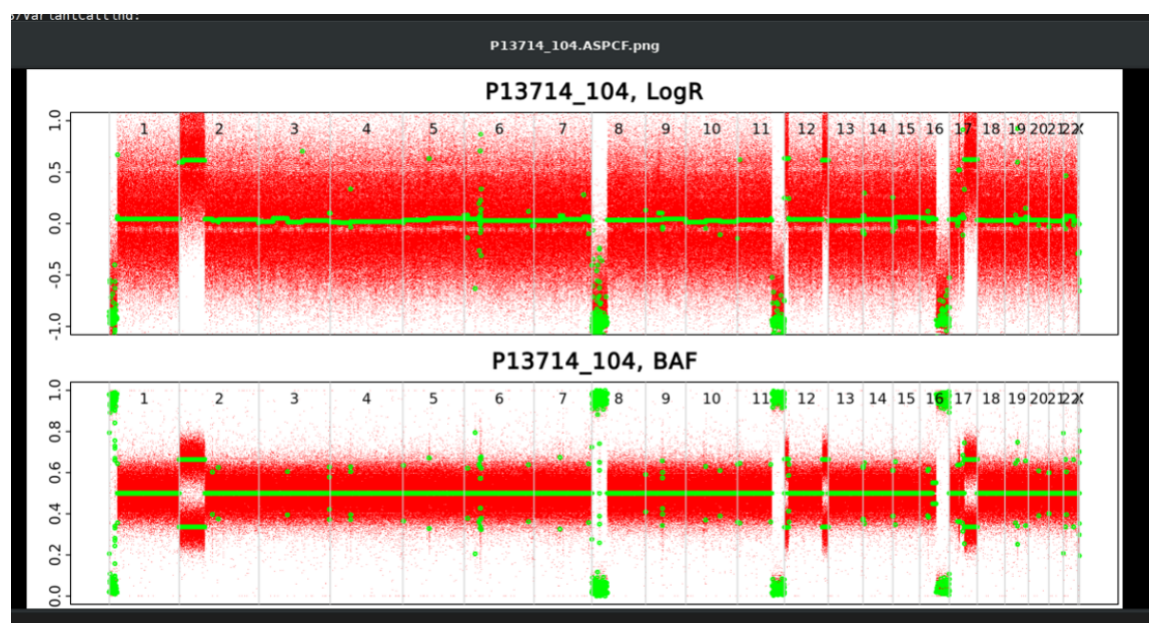
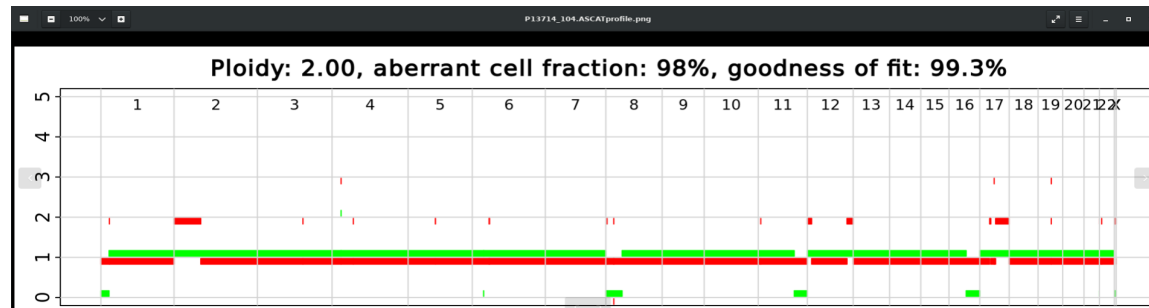


MB13

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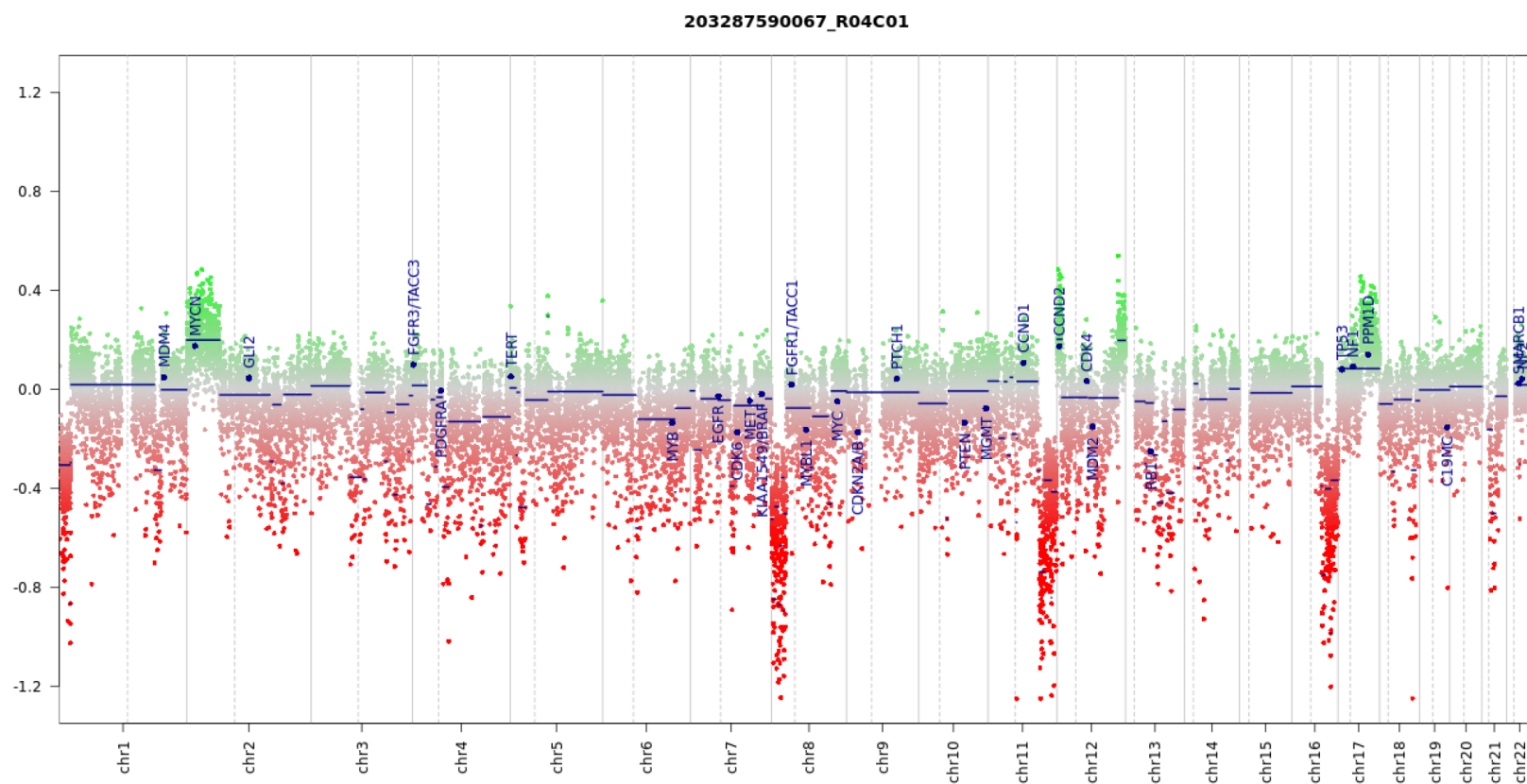


MB15p

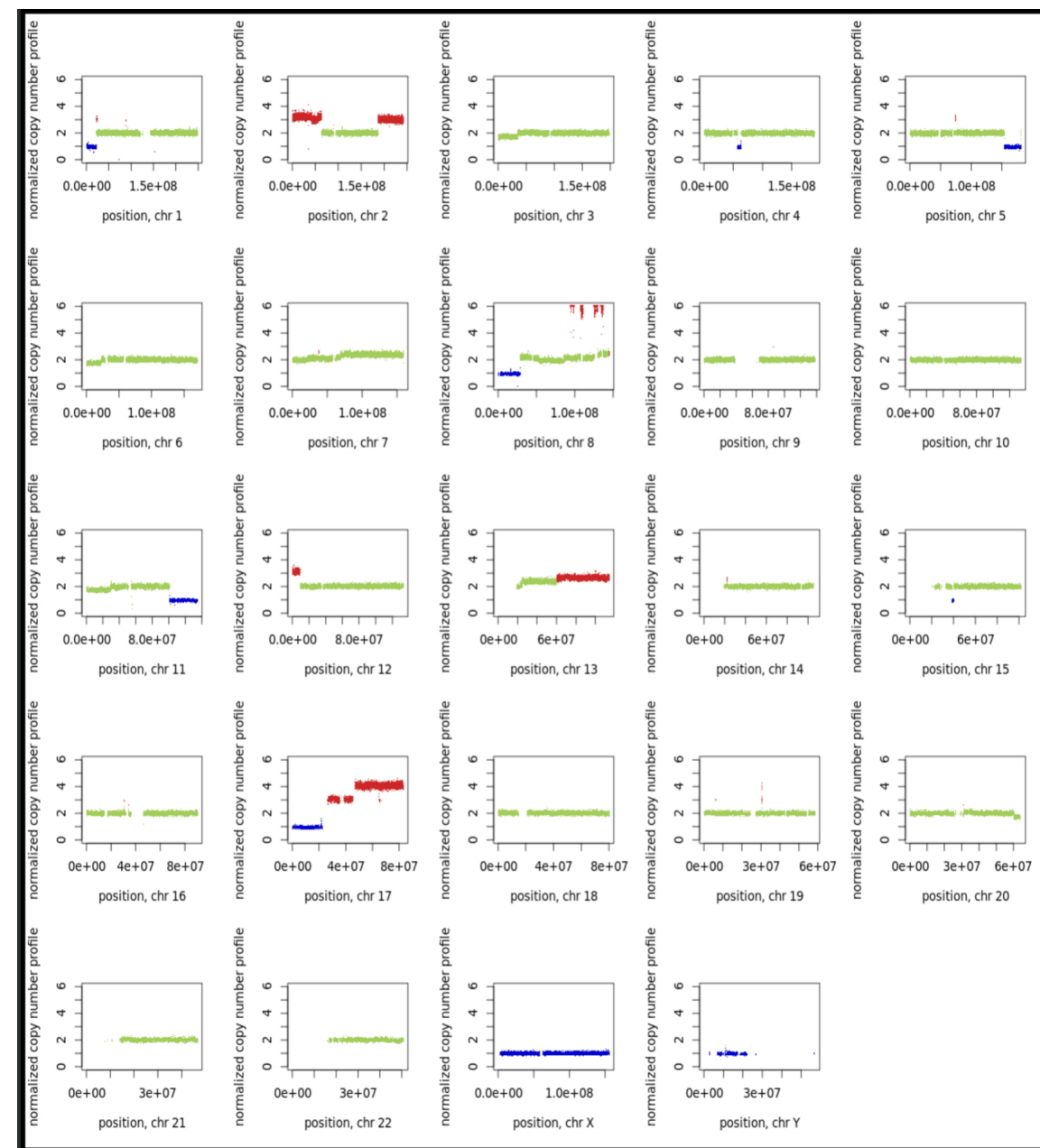
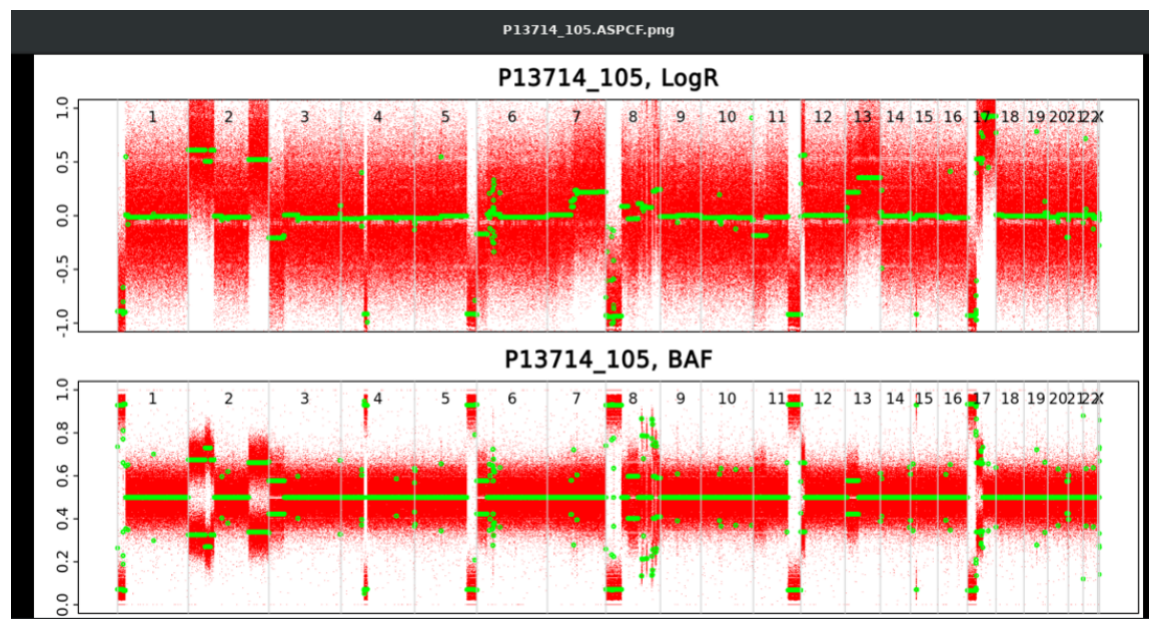
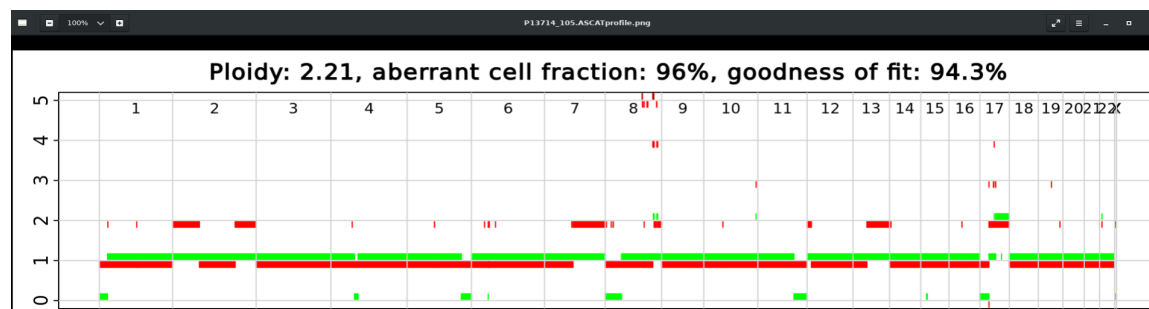


MB15p

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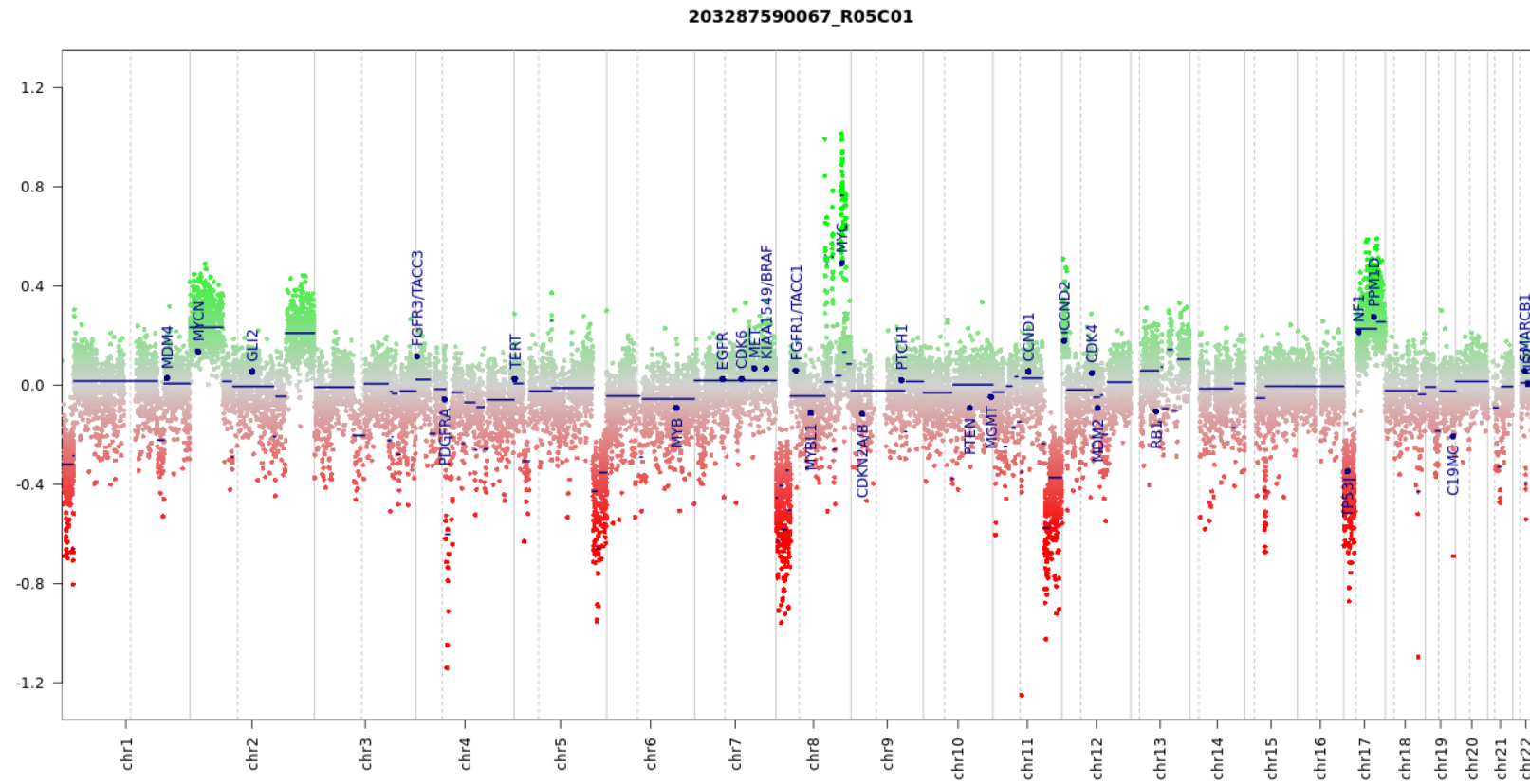


MB15r/m

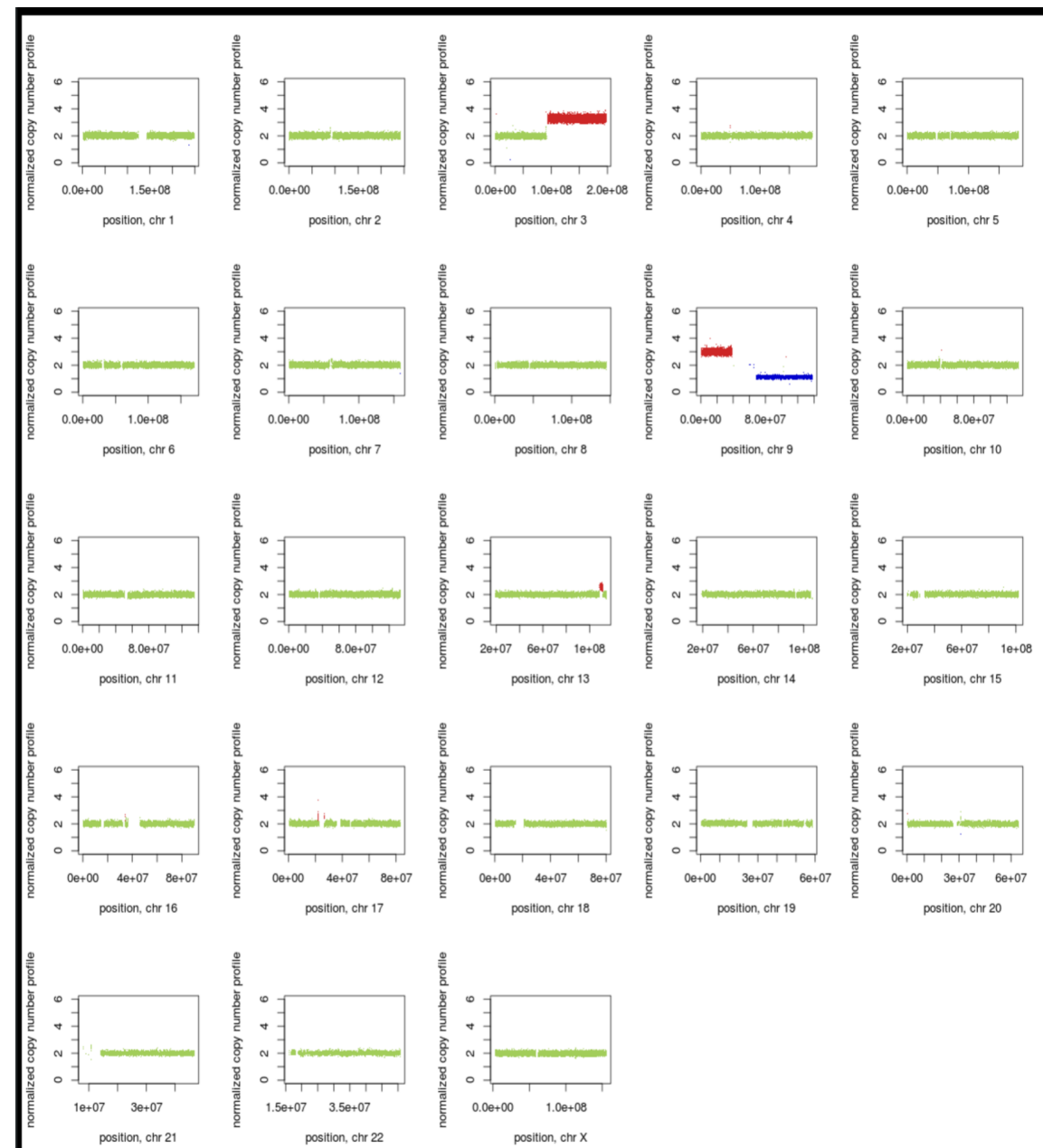
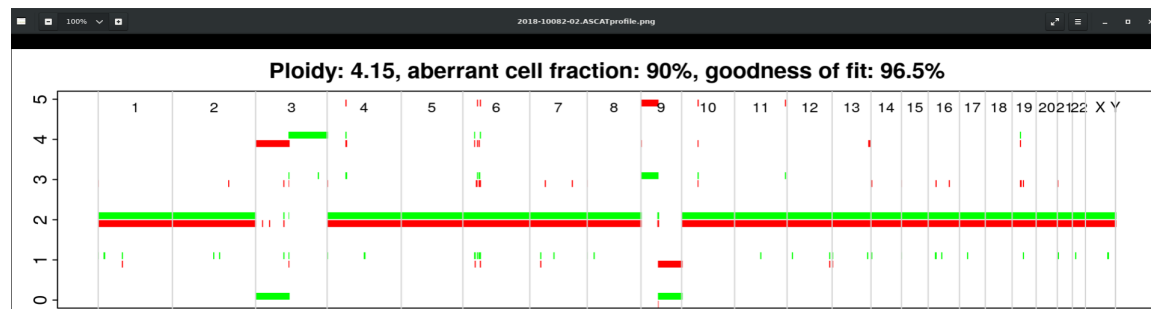


15r/m

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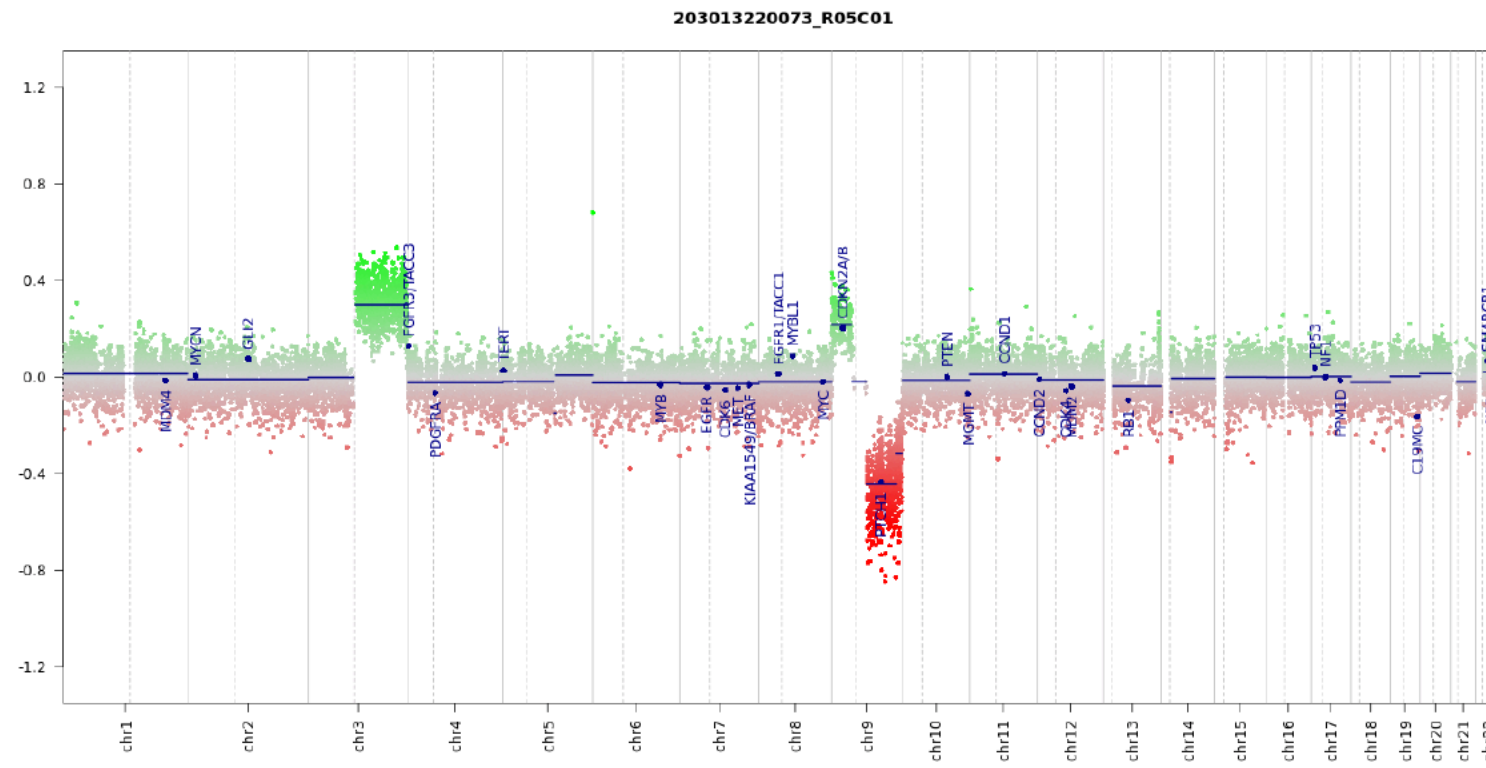


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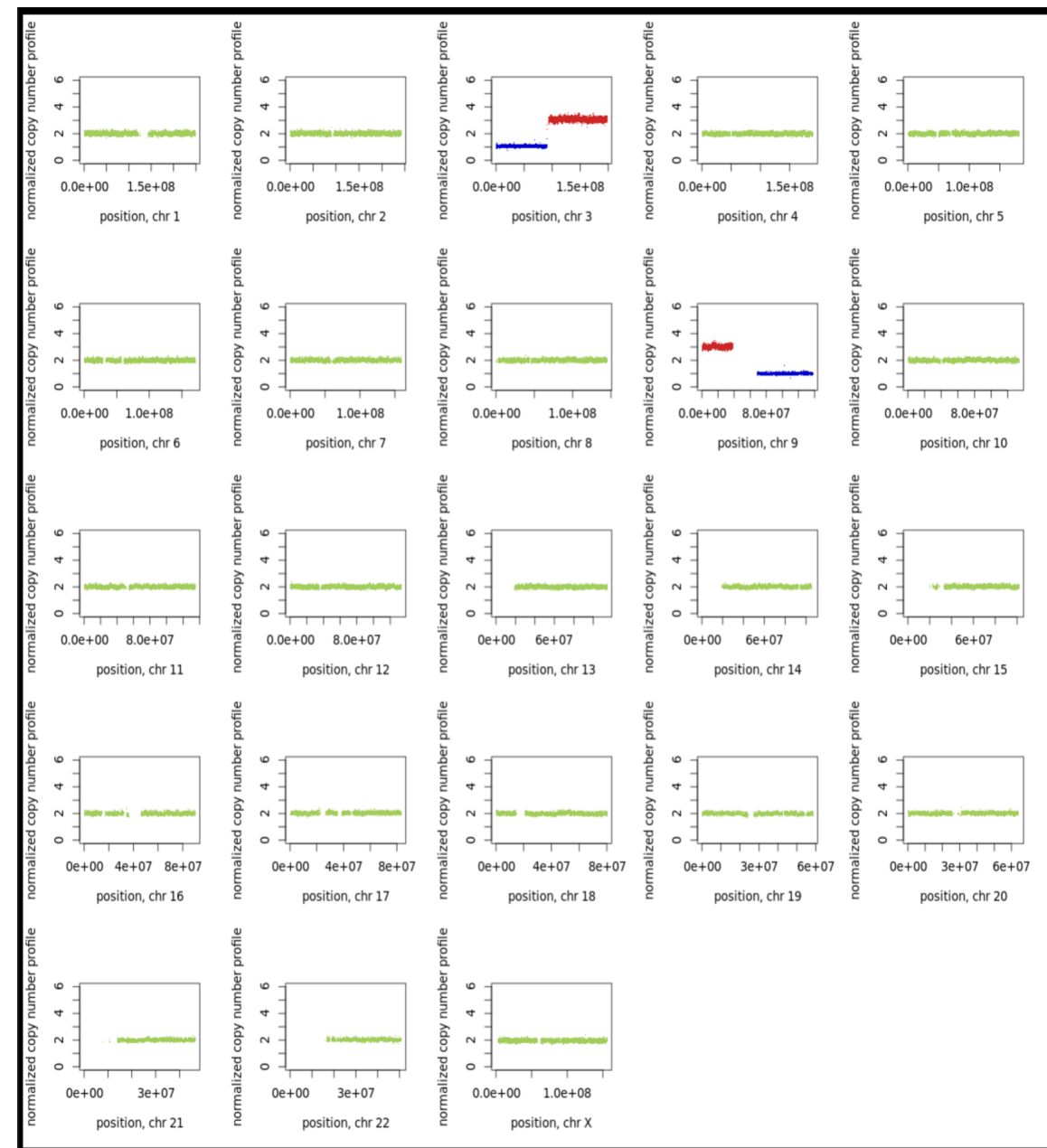


MB19p

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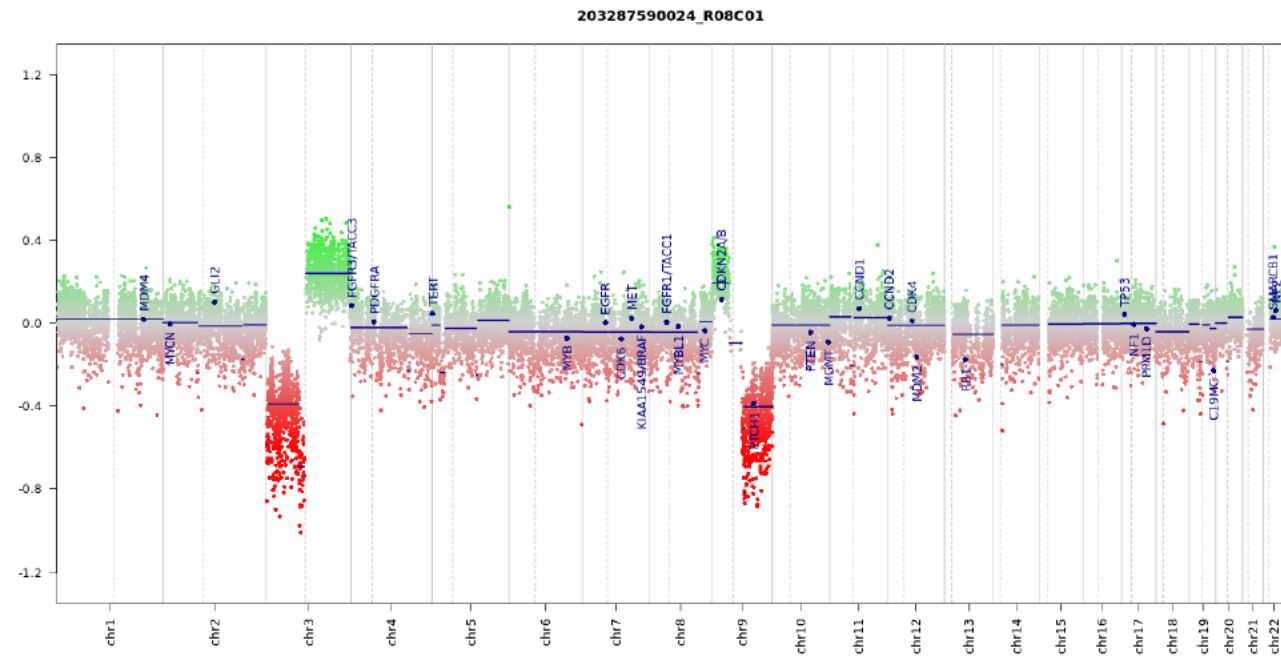


MB19r



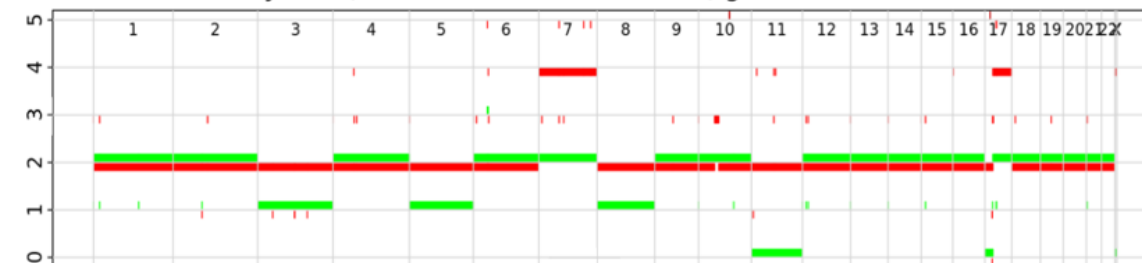
MB19r

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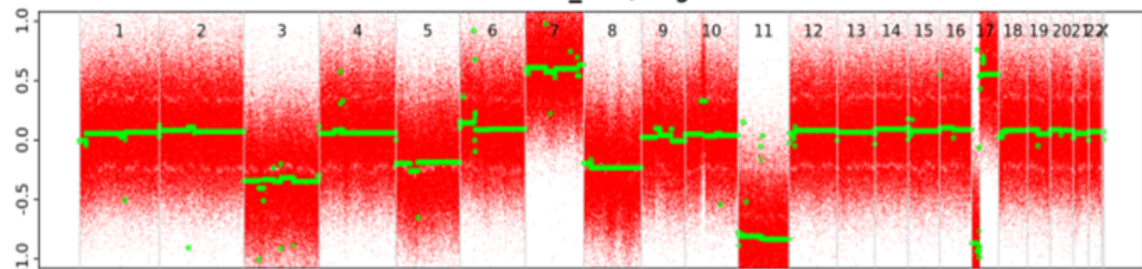


MB21

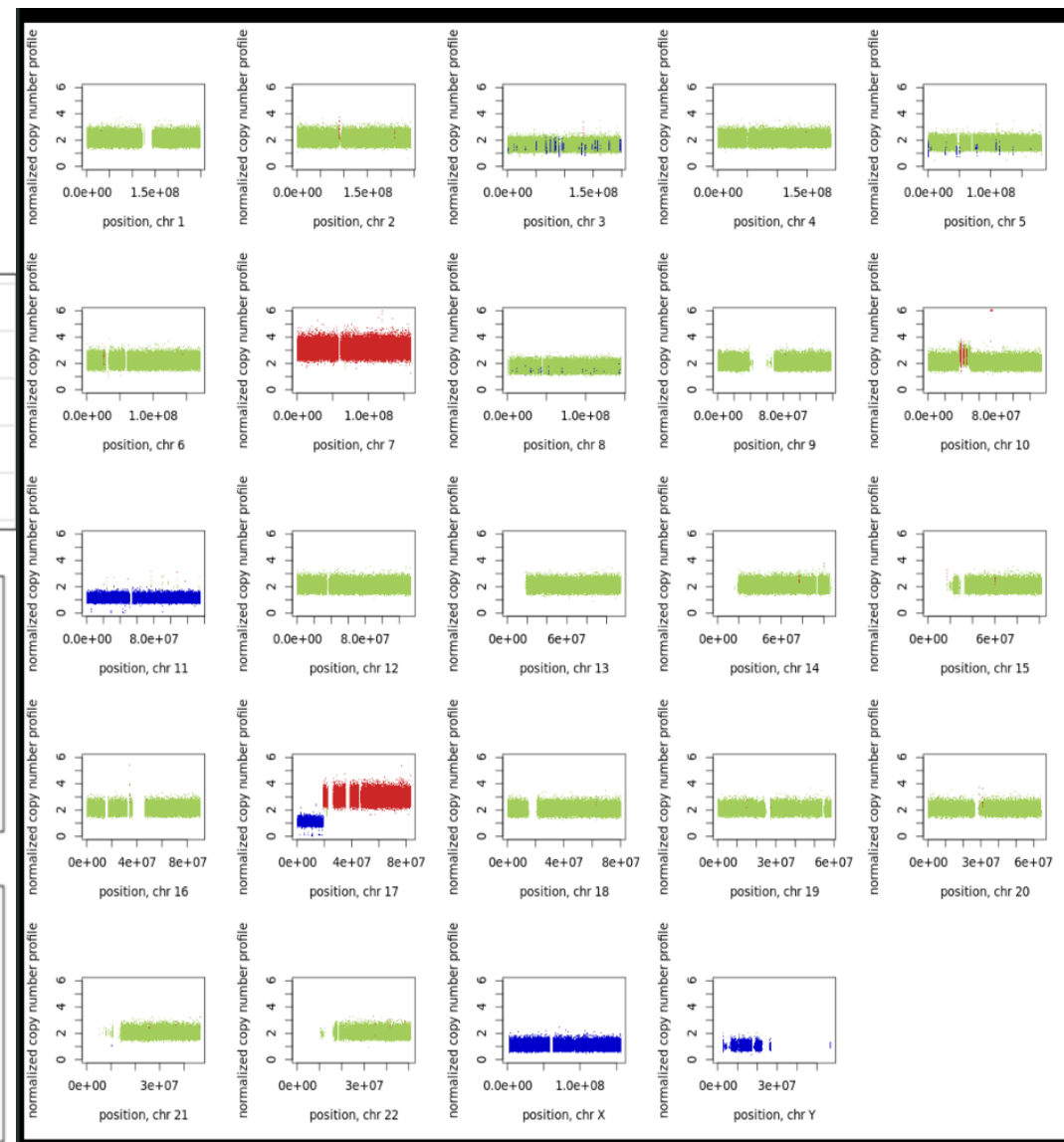
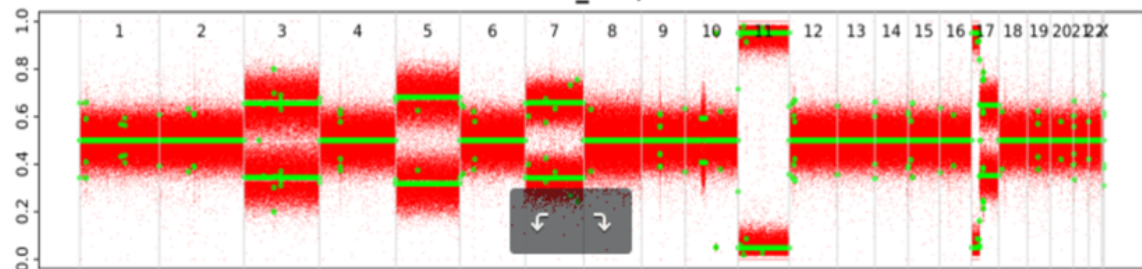
Ploidy: 3.92, aberrant cell fraction: 90%, goodness of fit: 98.6%



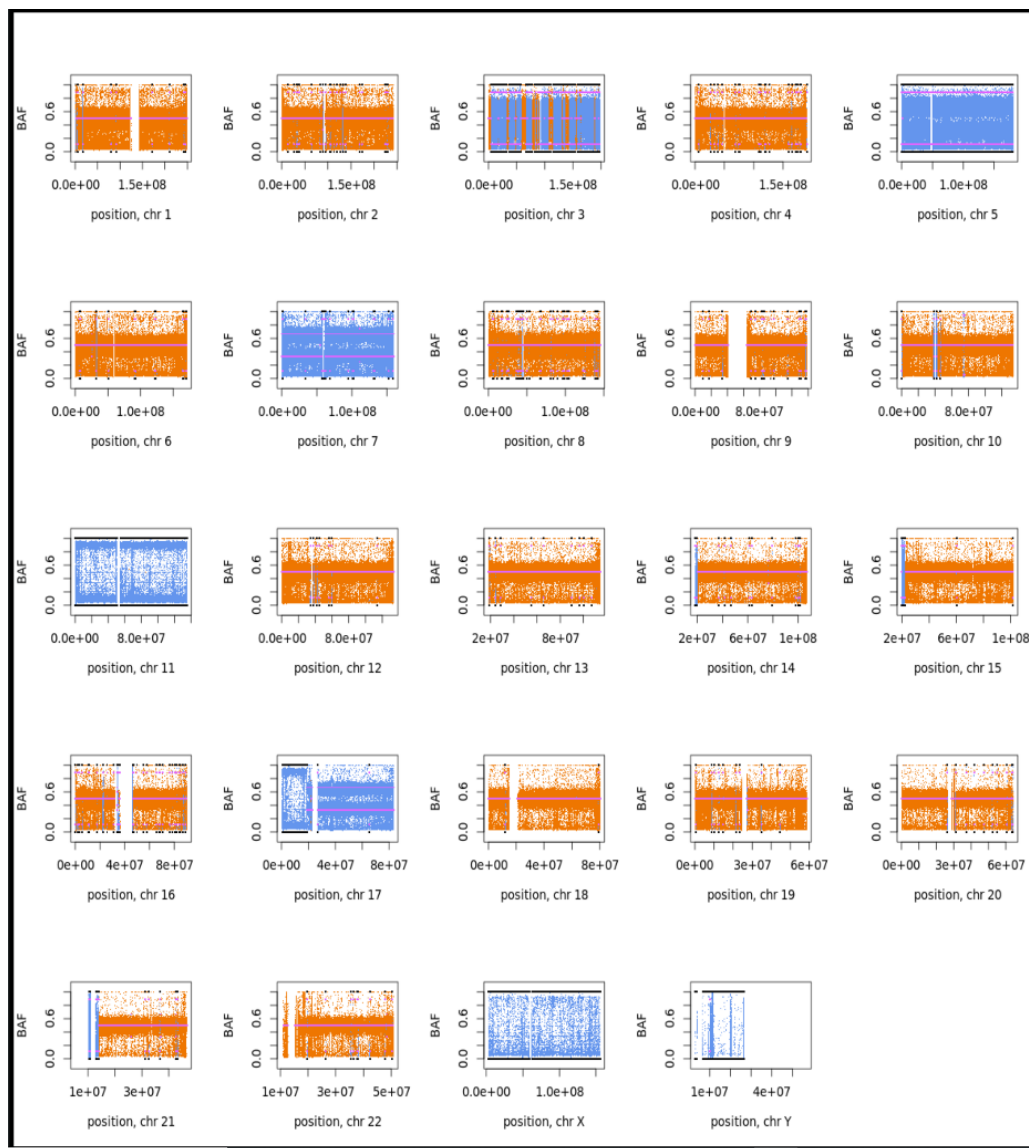
P19060_123, LogR



P19060_123, BAF

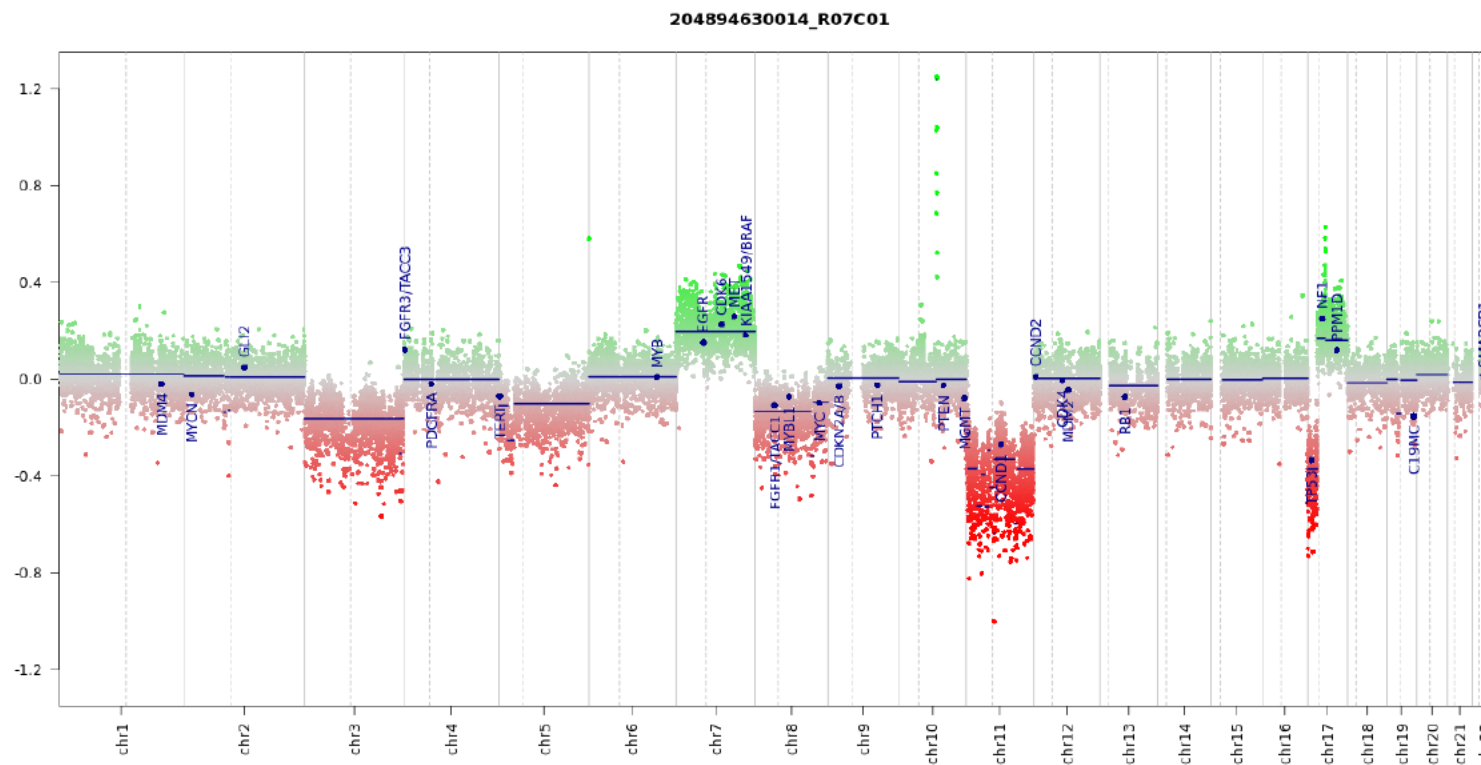


MB21 BAF

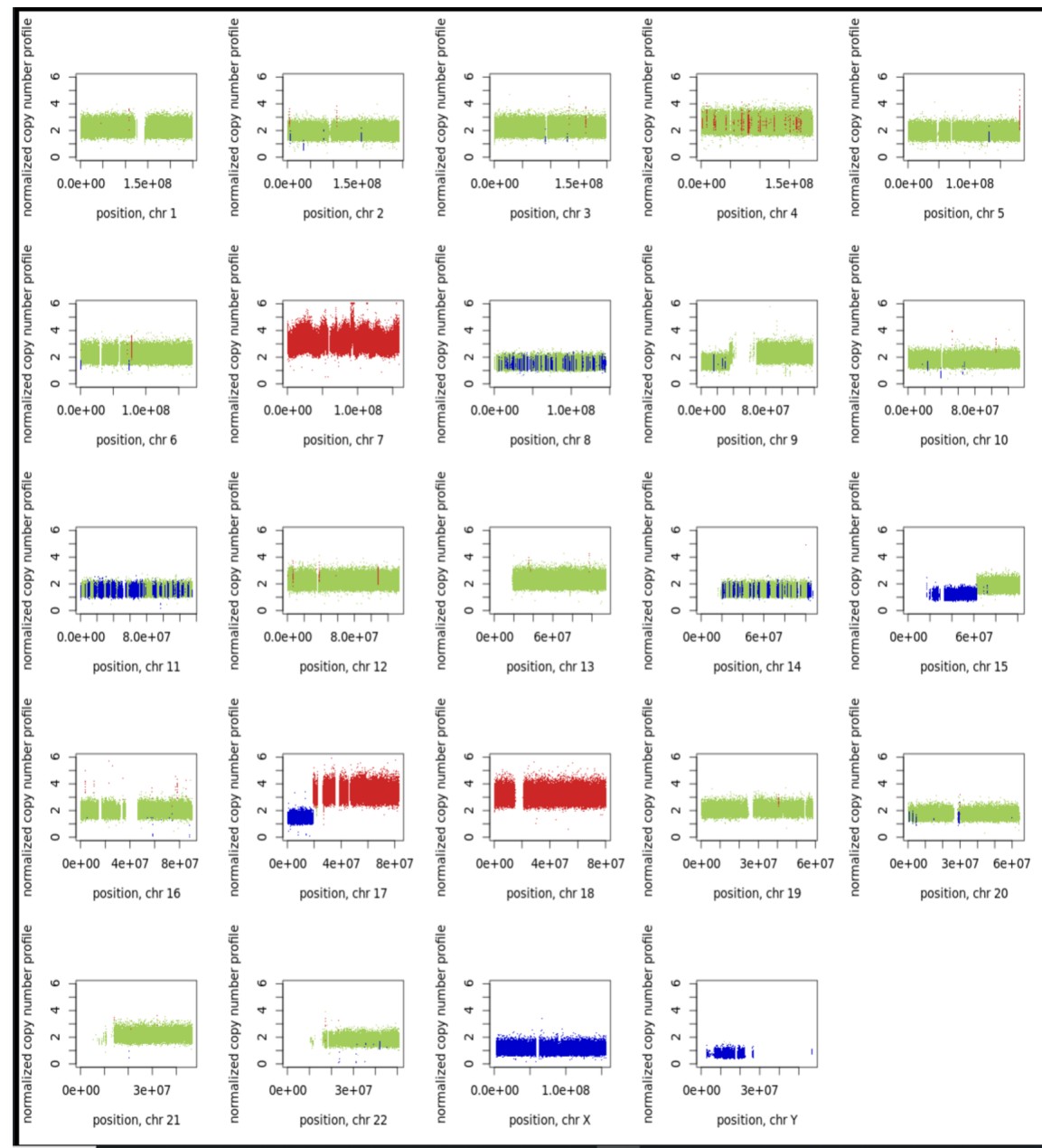
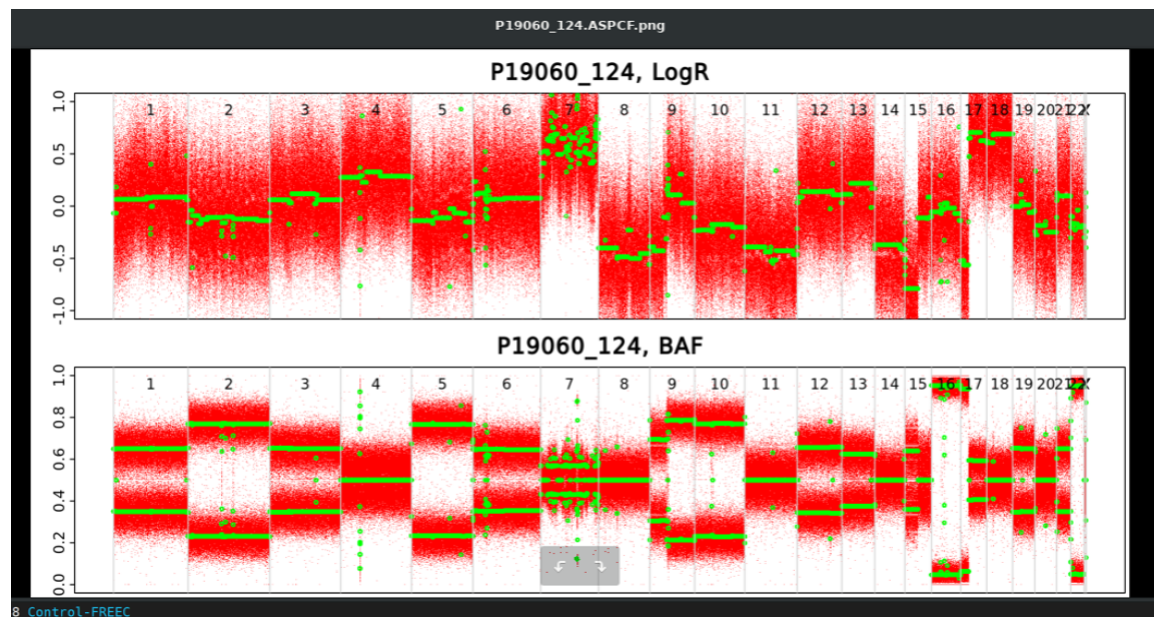
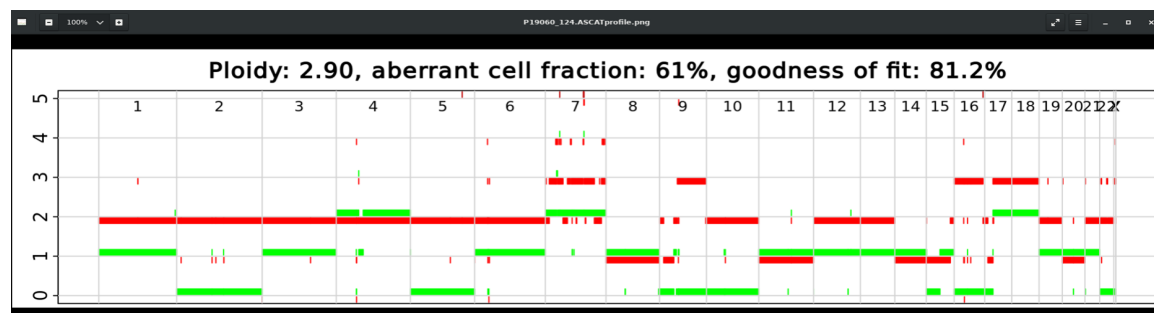


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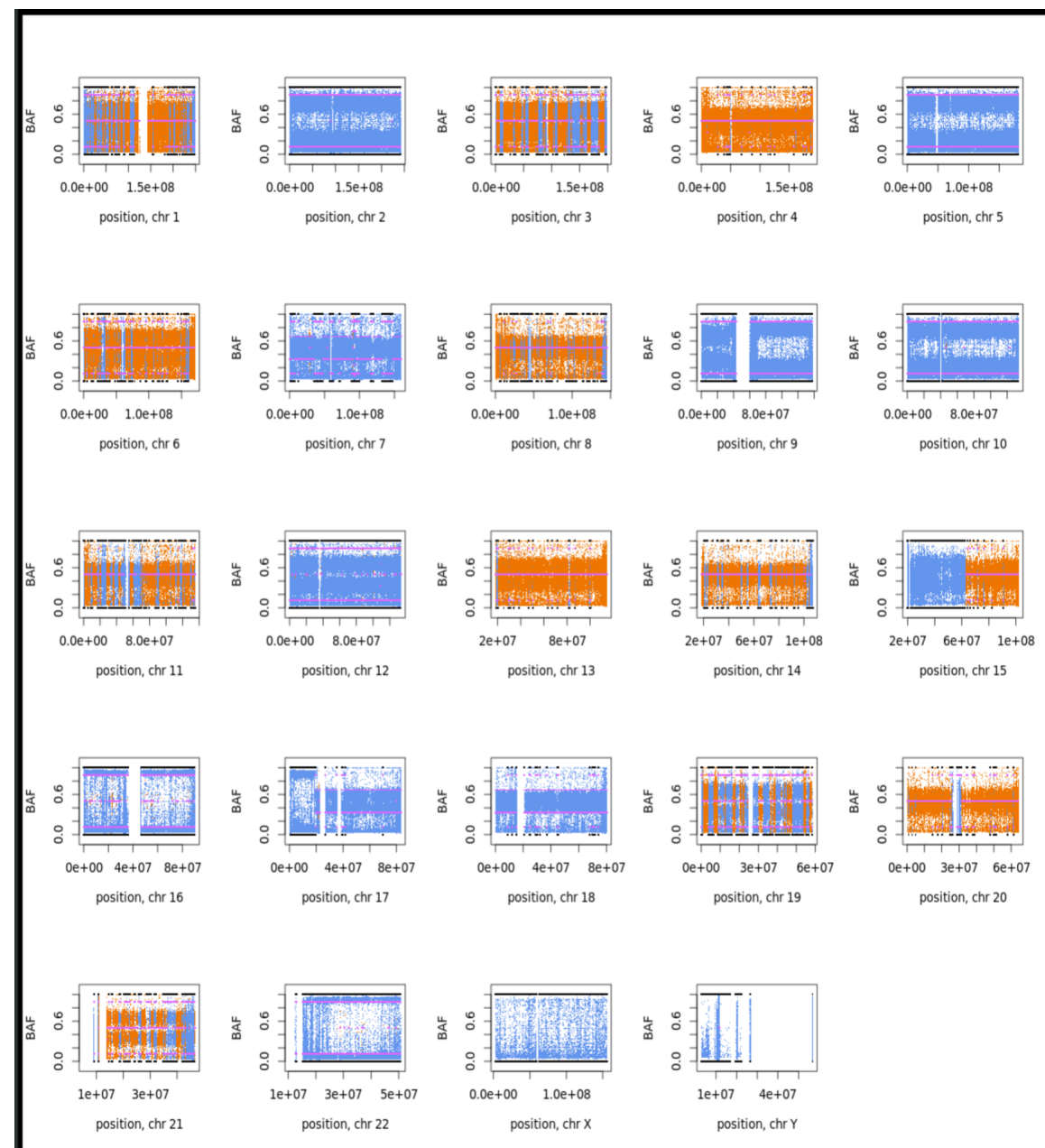
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MB22

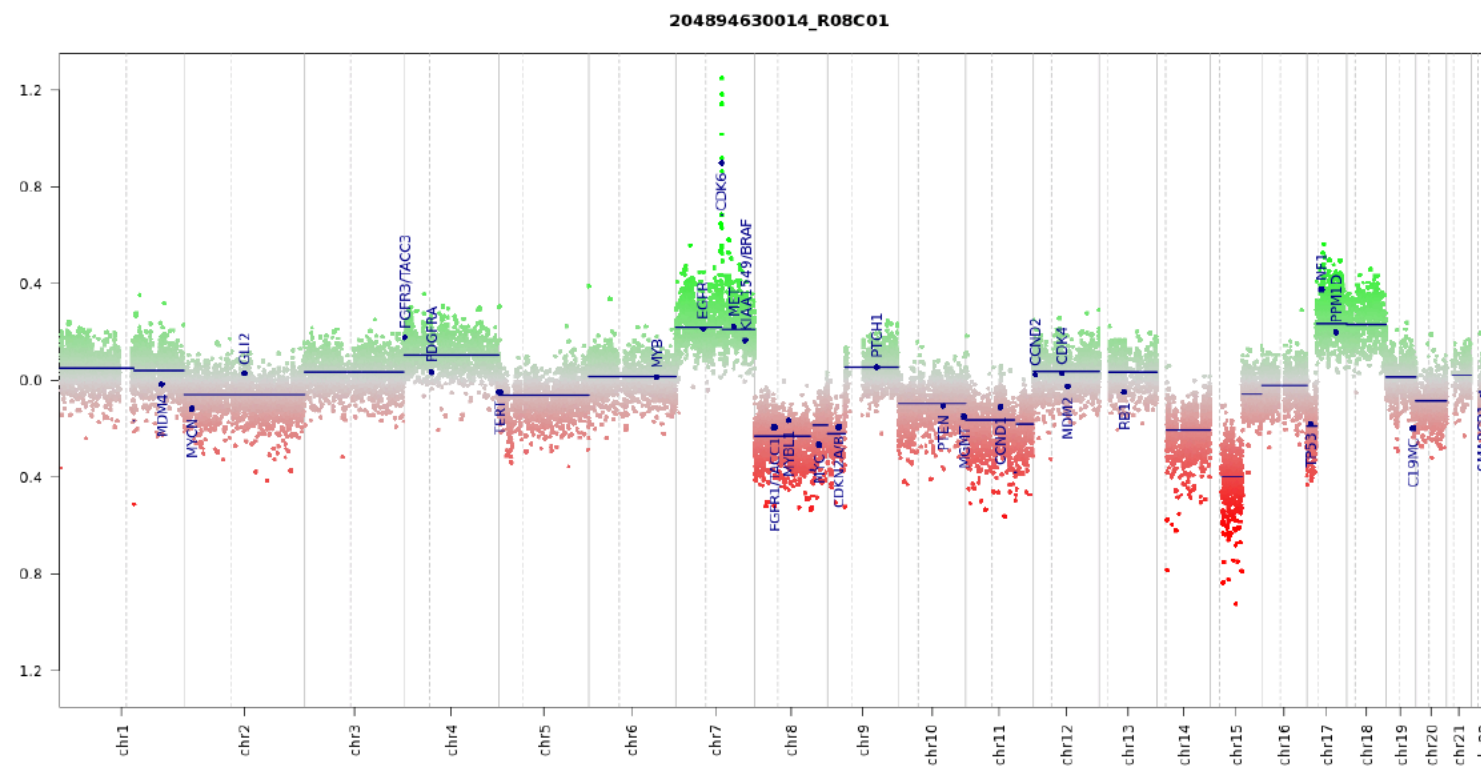


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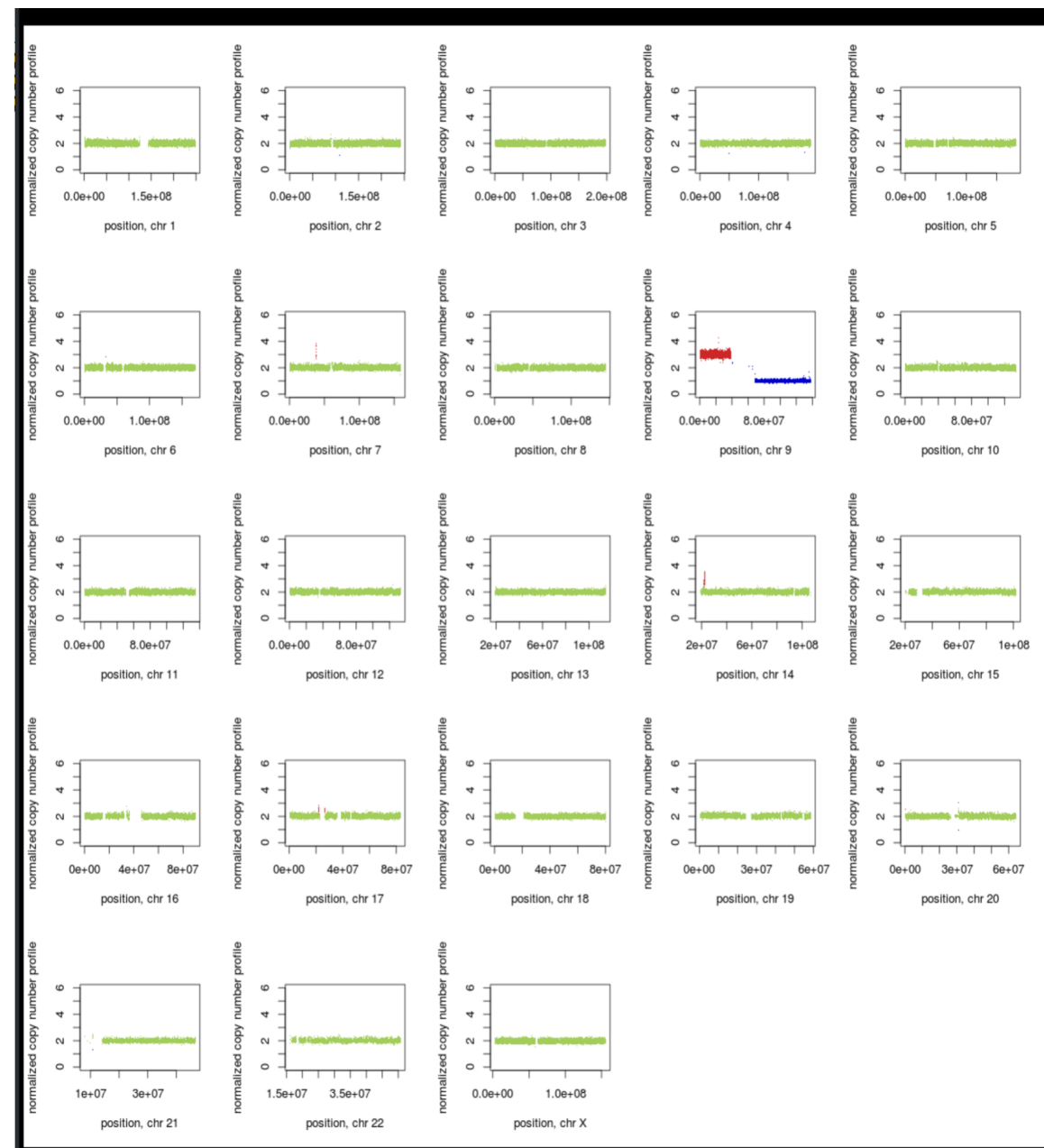


MB22

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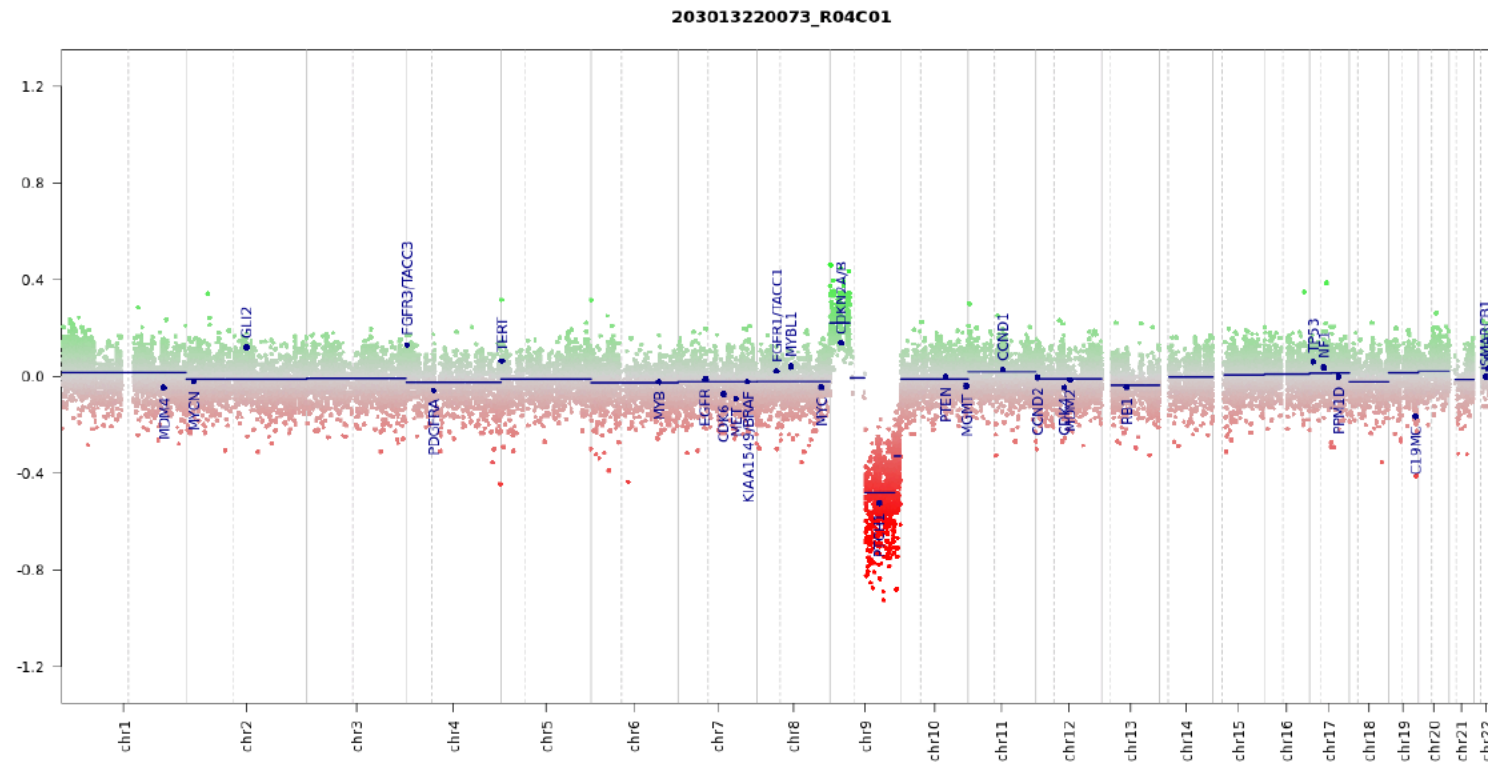


MB24

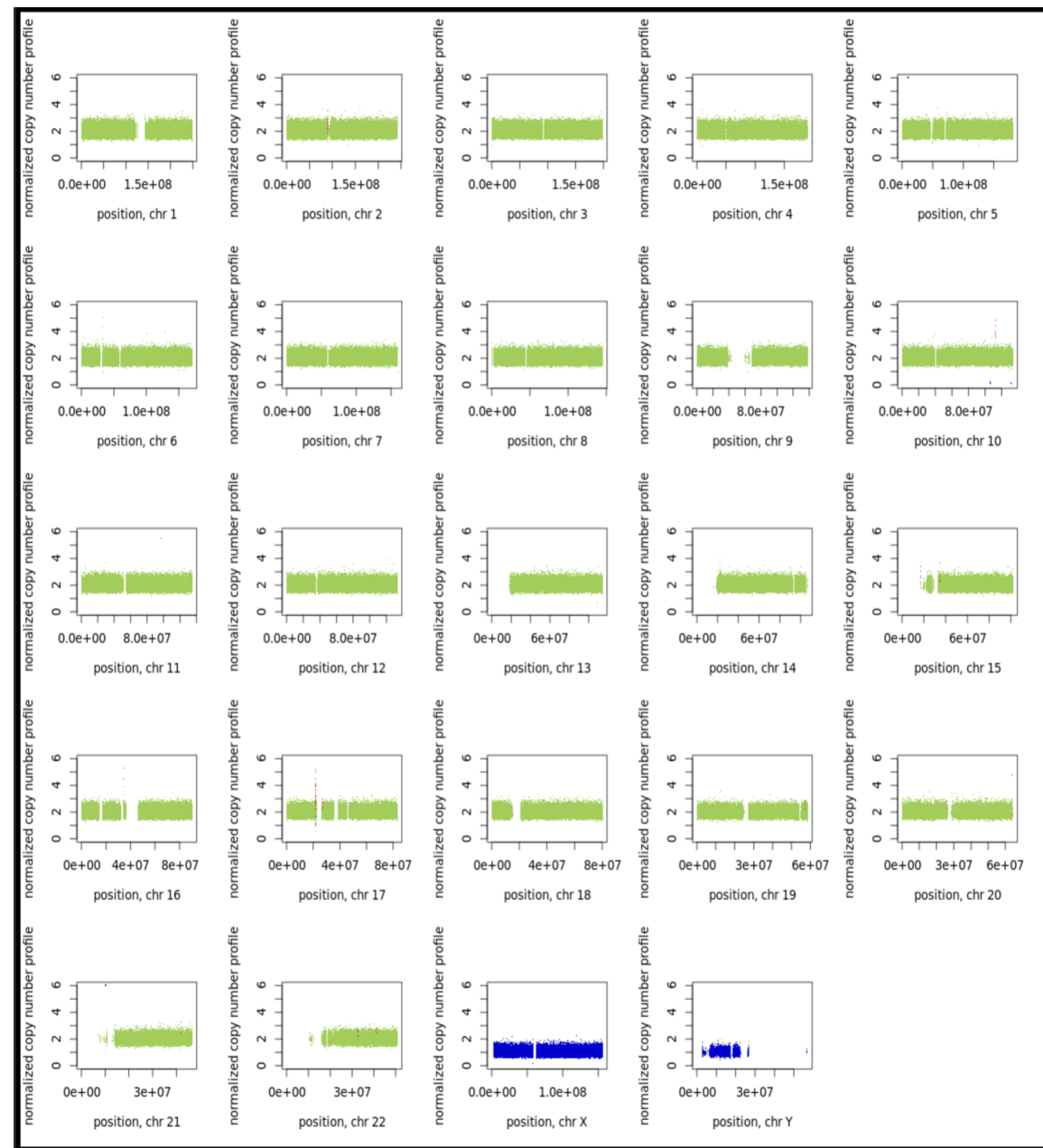
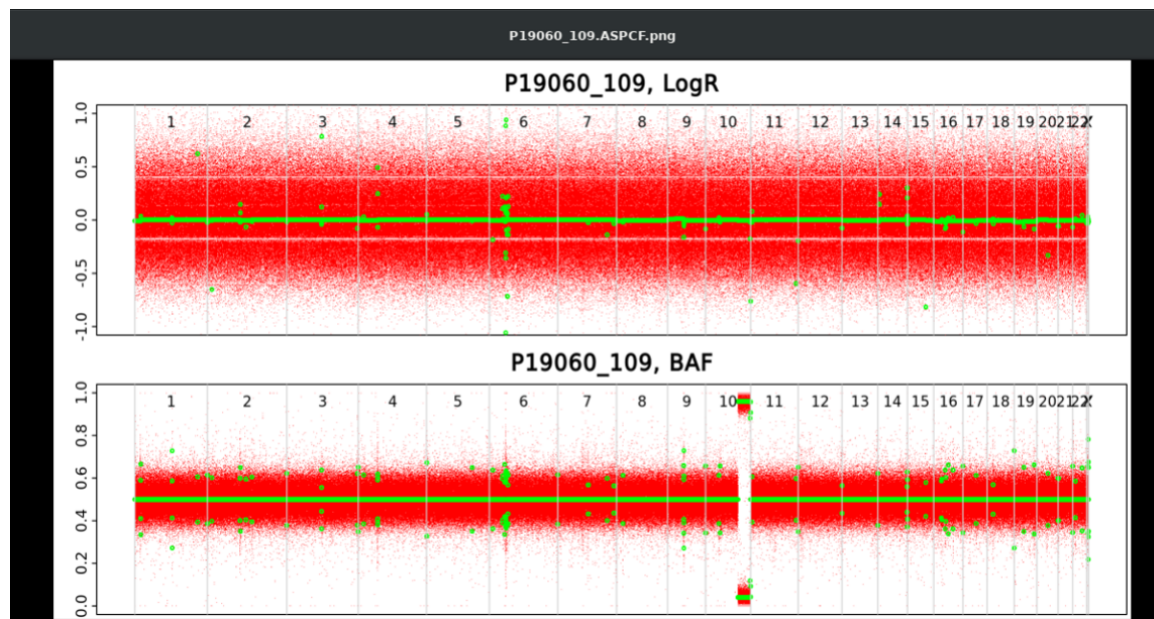
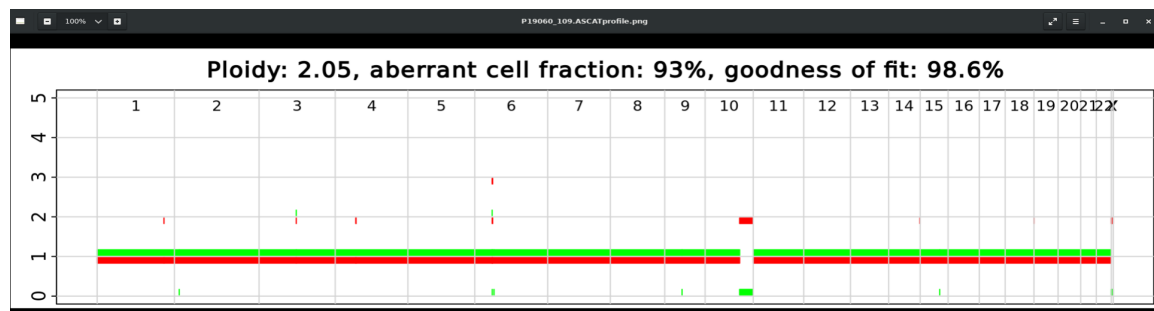


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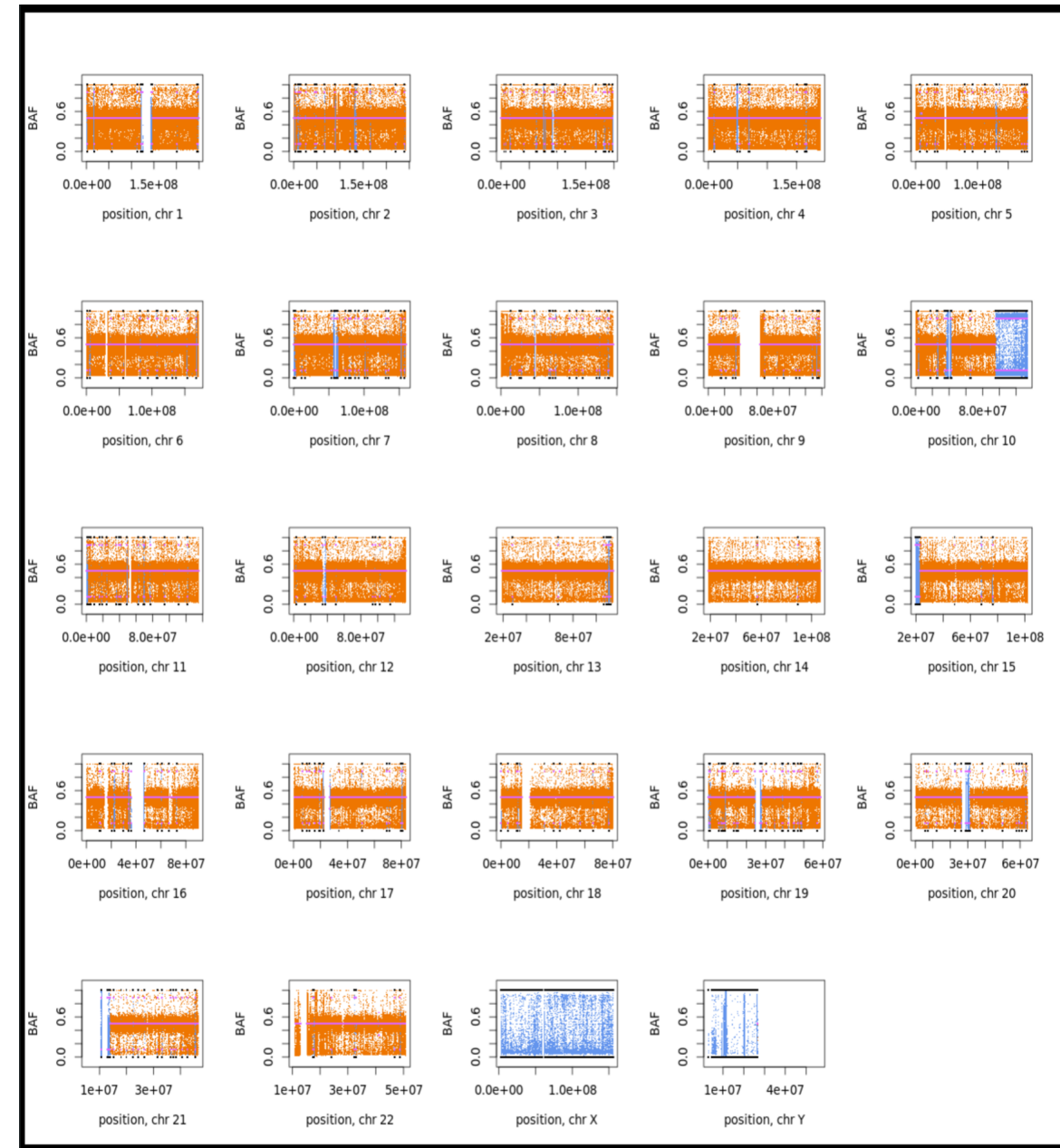
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MB27

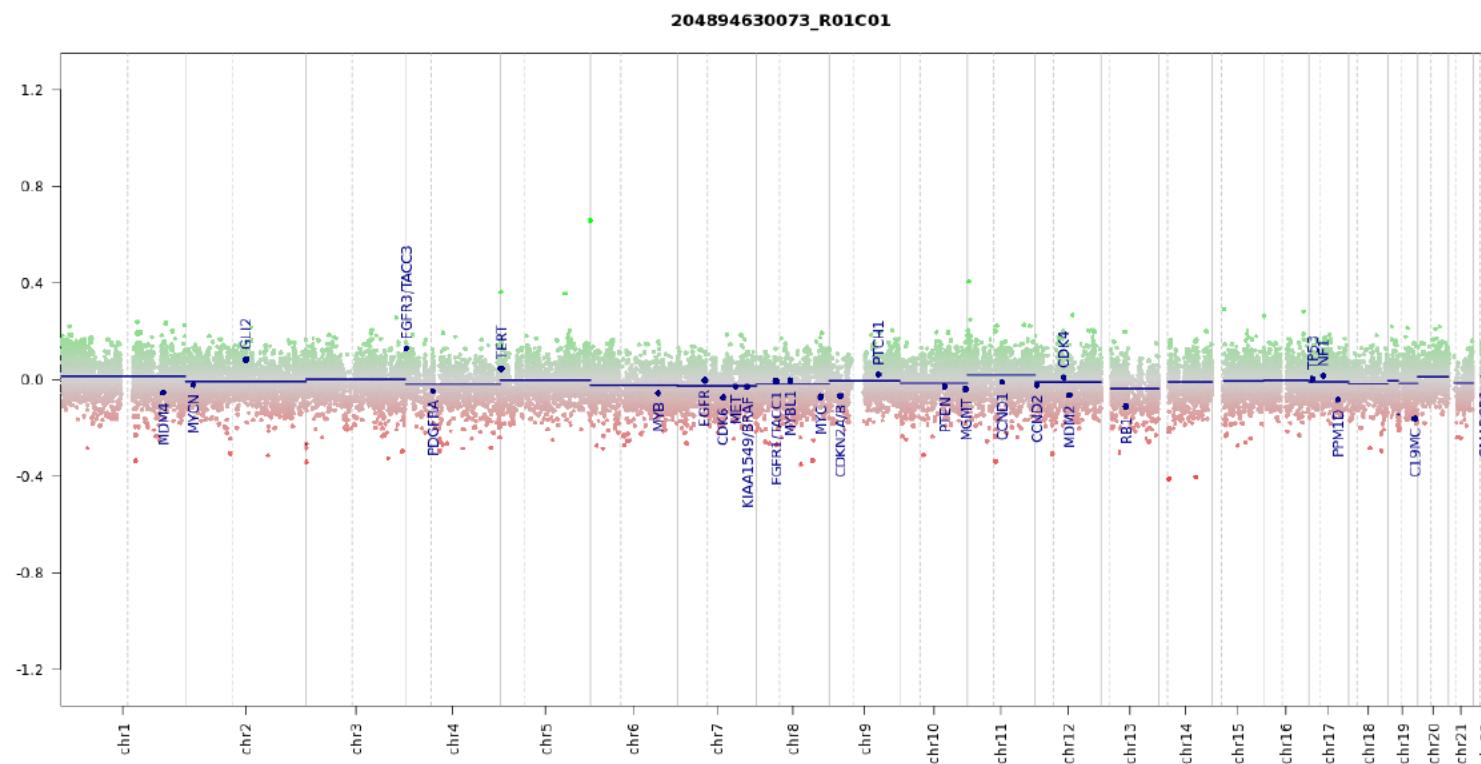


MB27 BAF

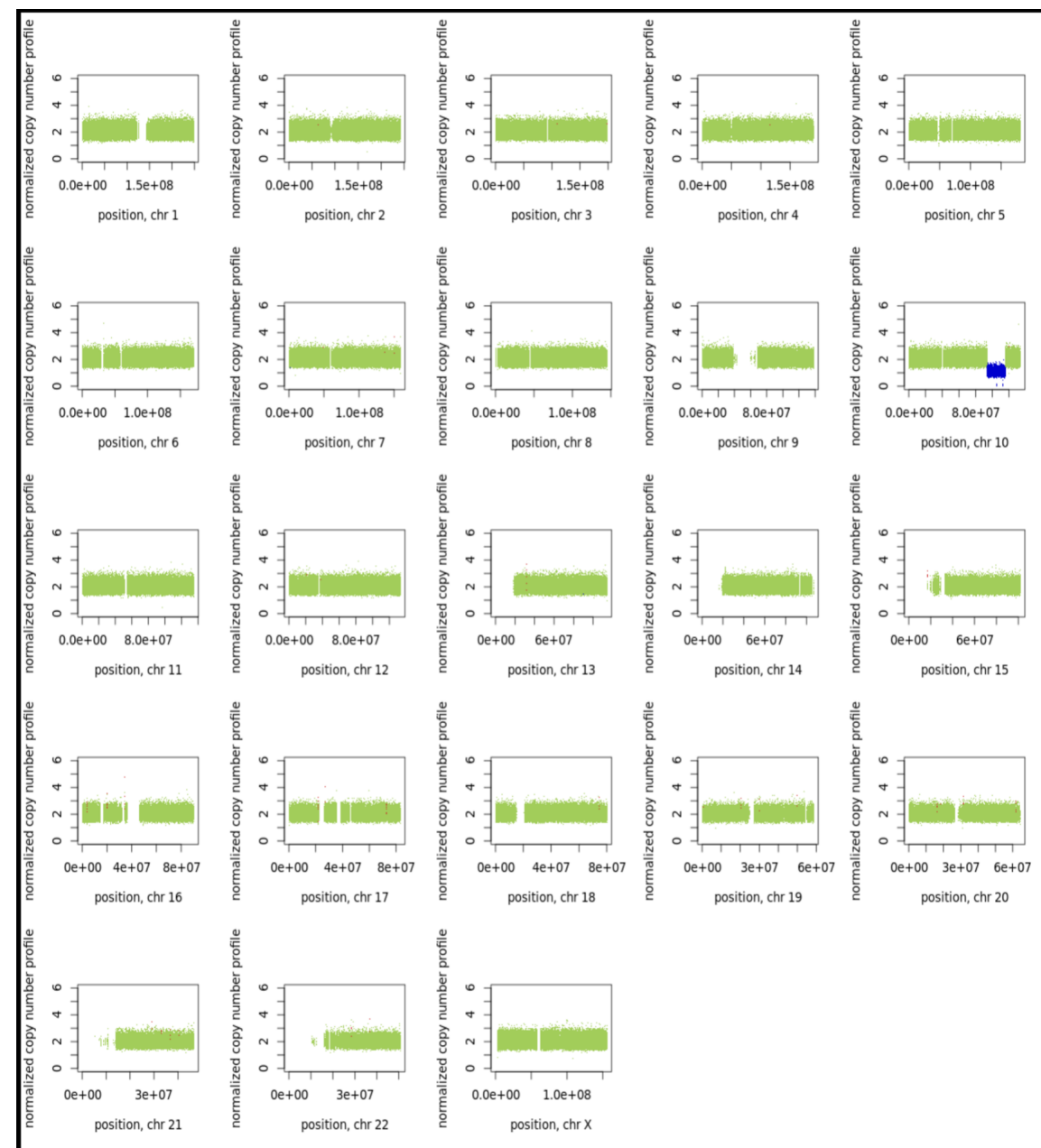
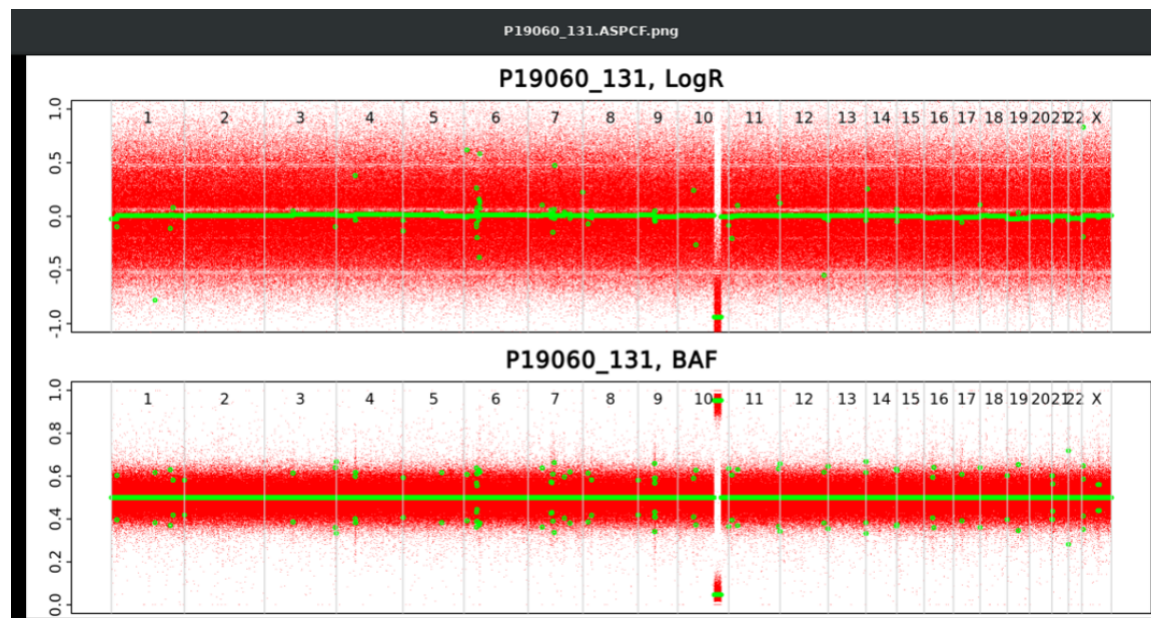
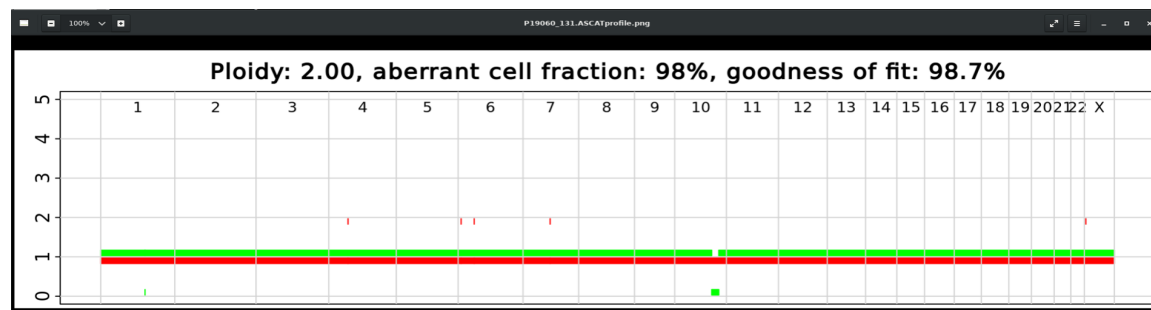


MB27

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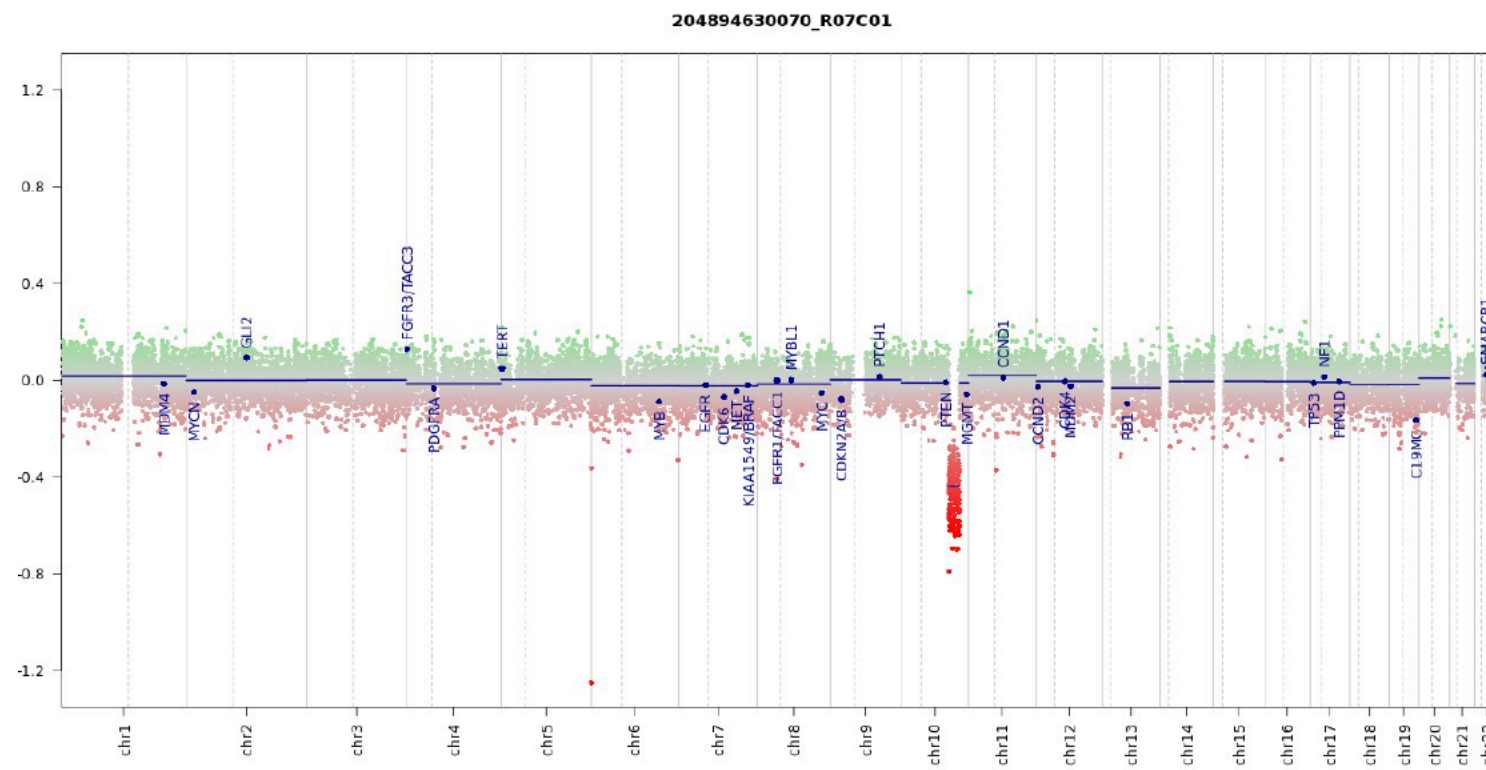


MB31

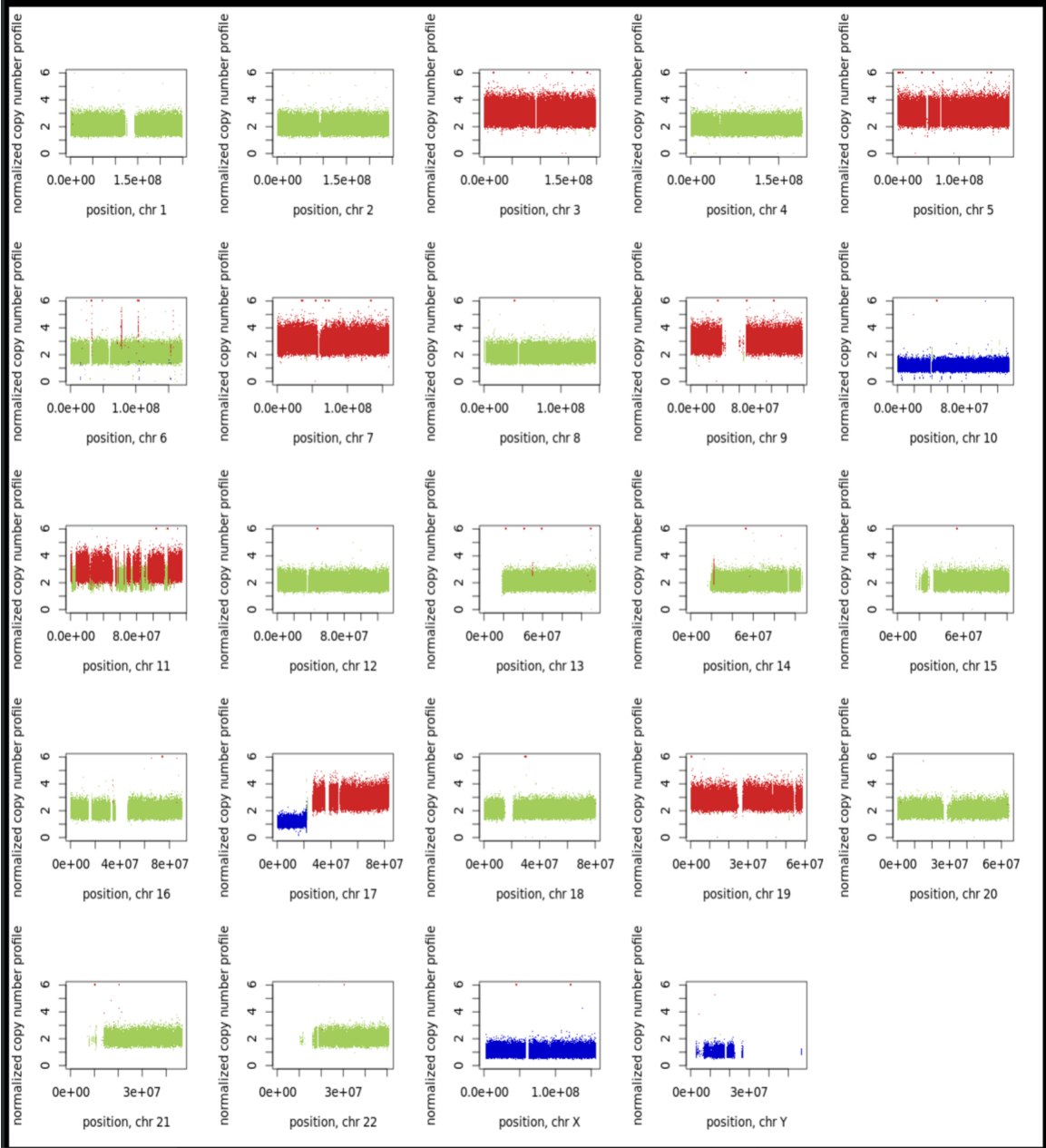
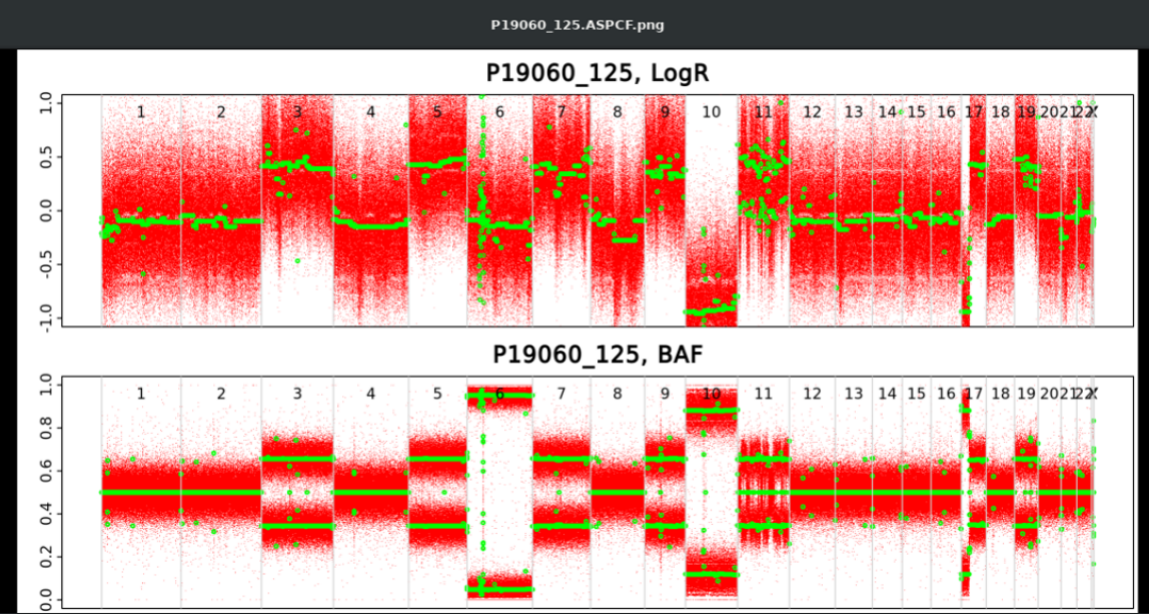
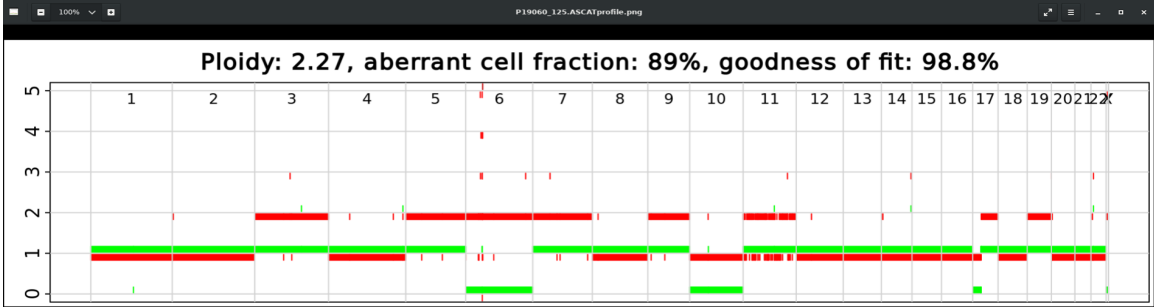


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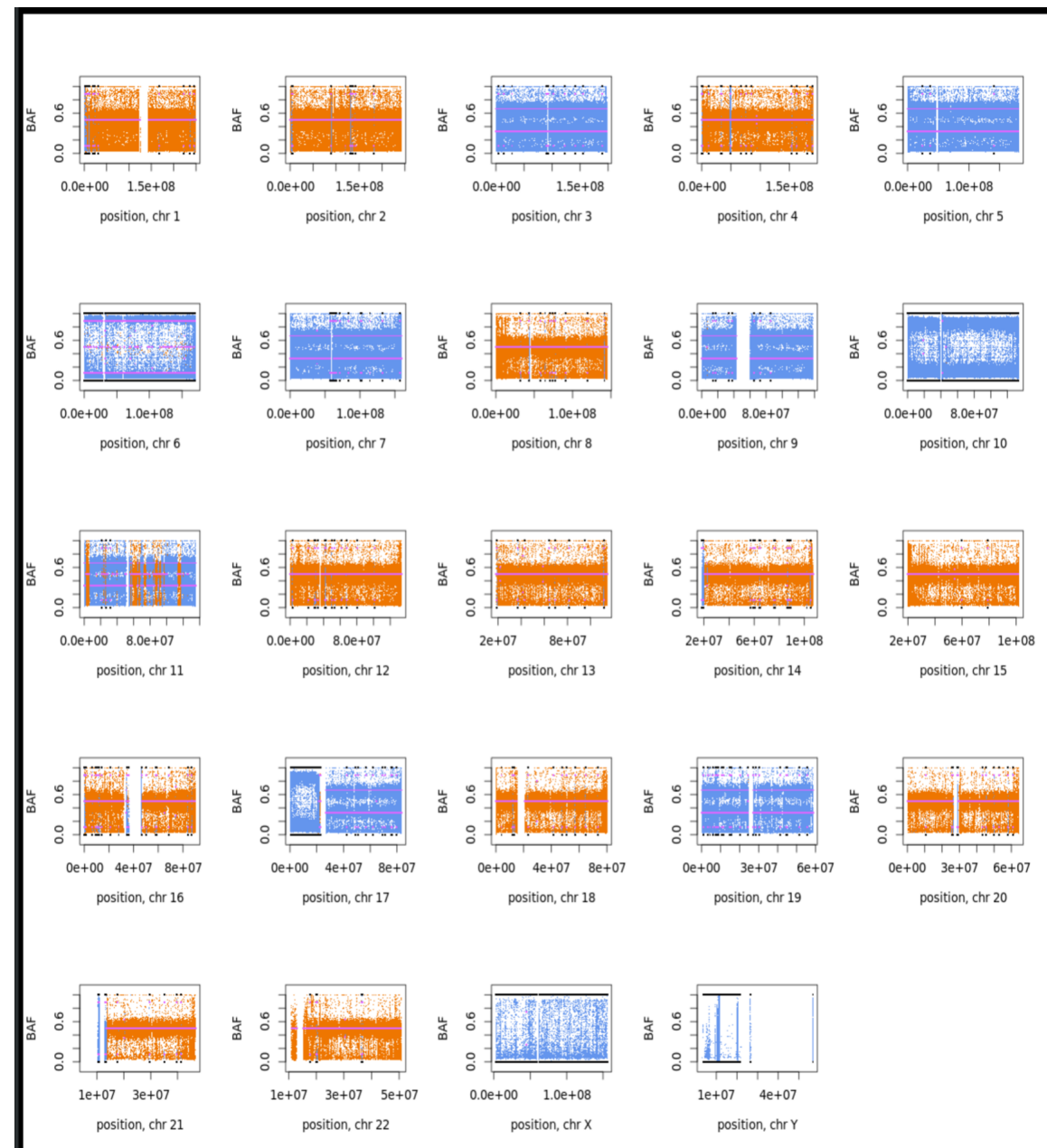
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MB35

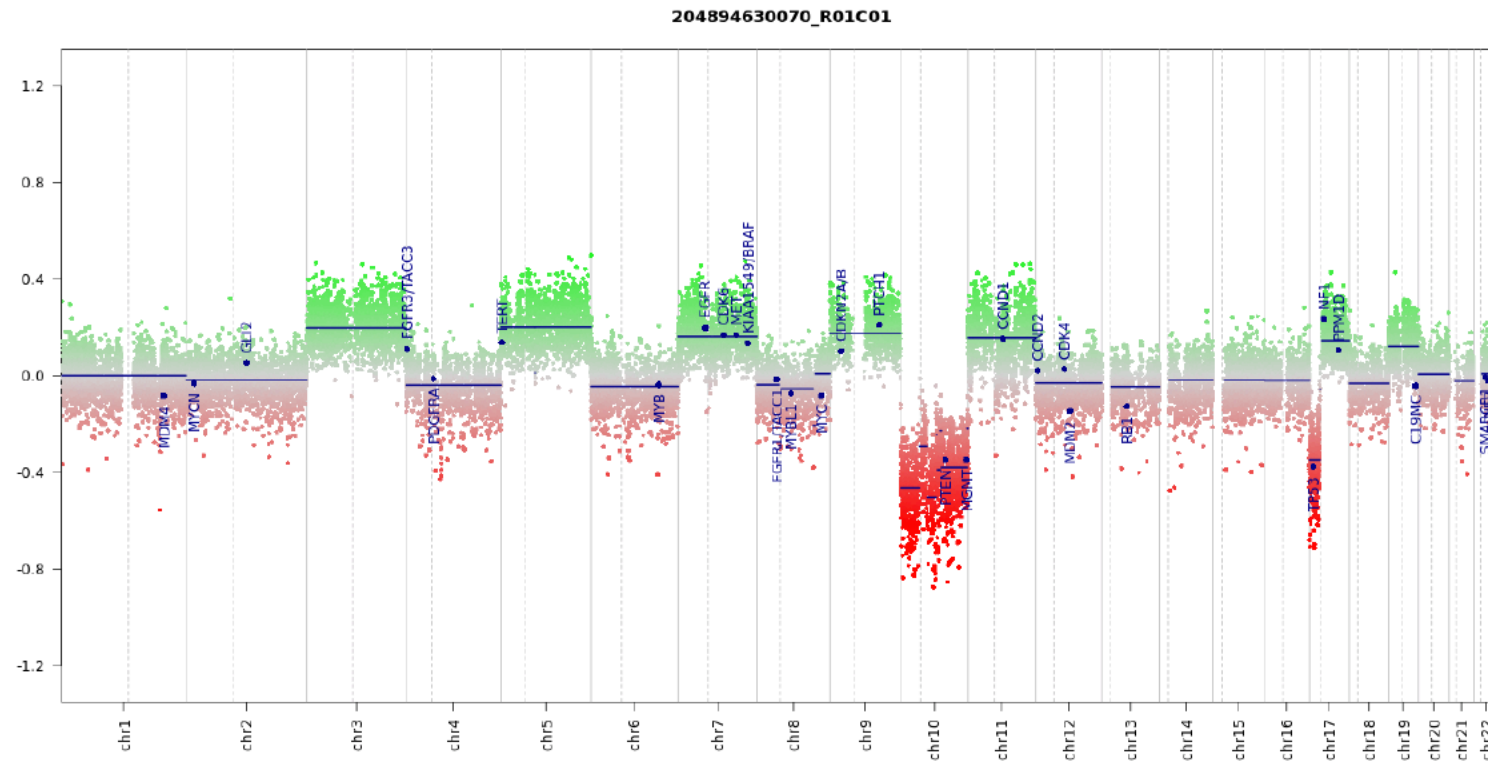


MB35 BAF



MB35

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MB43

