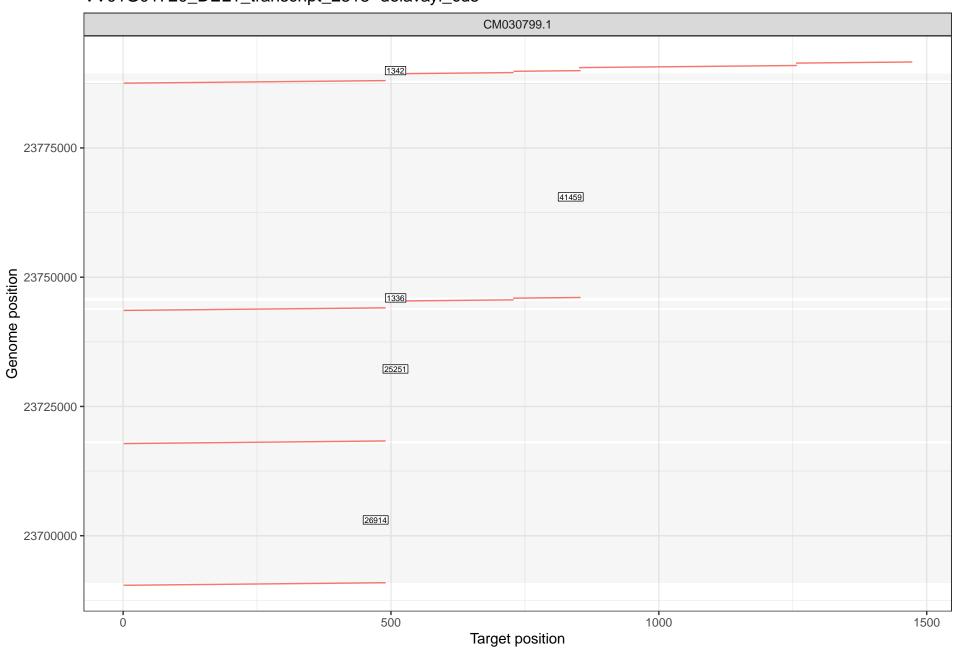
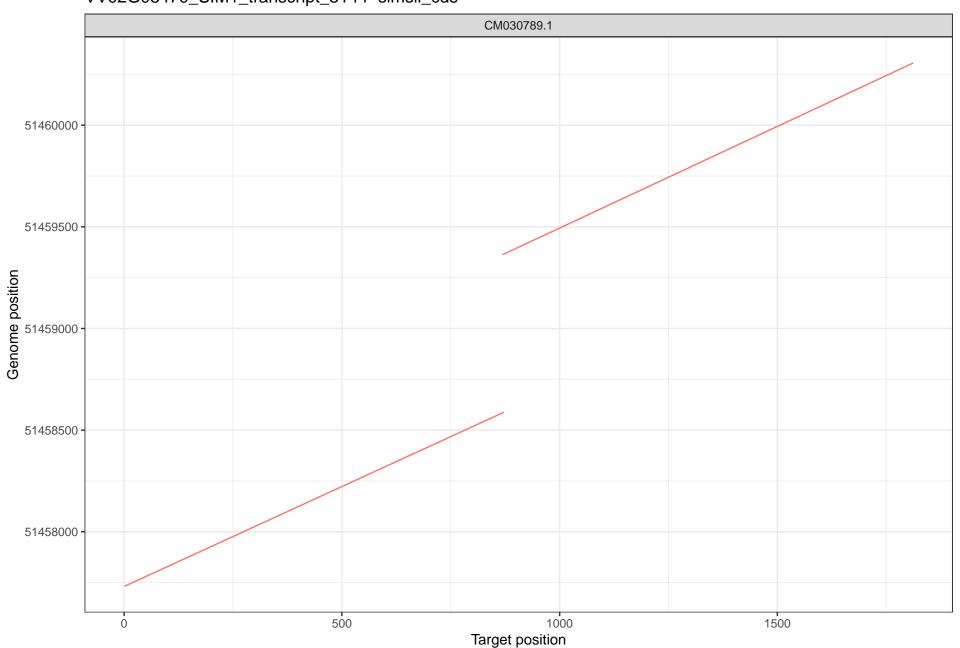
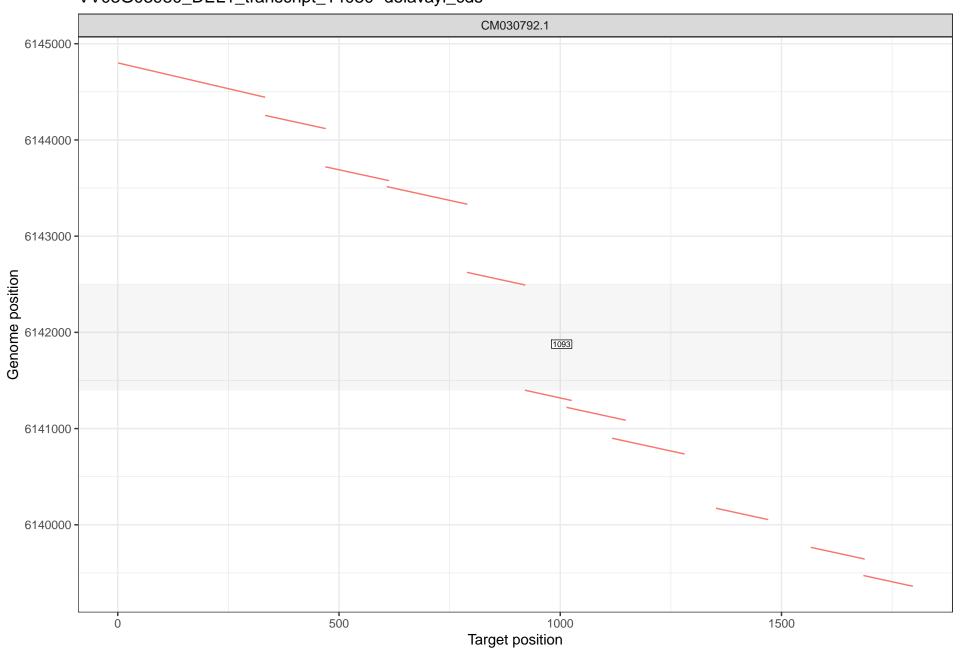
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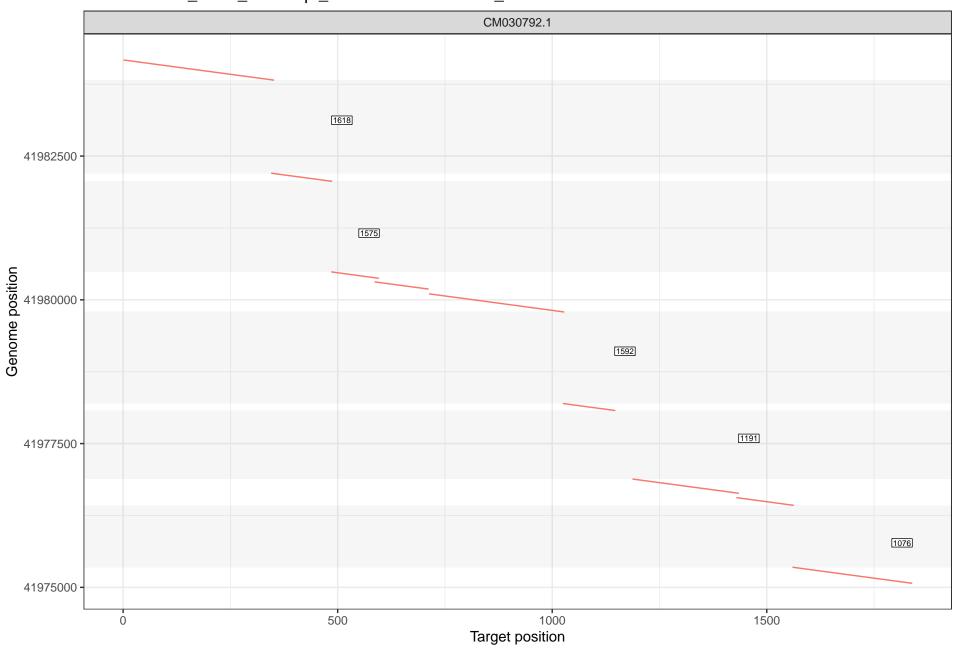


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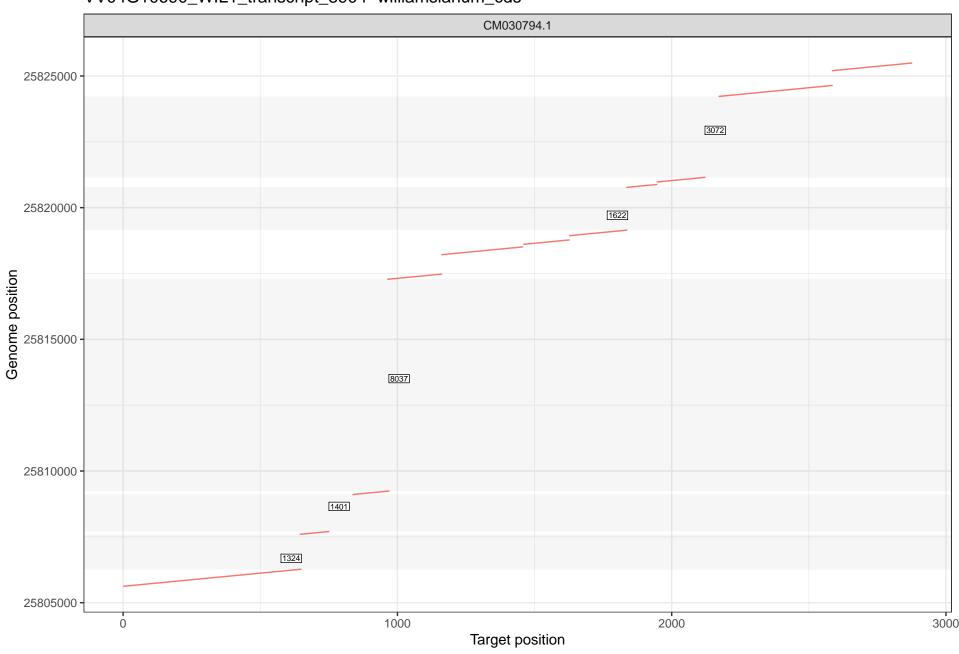


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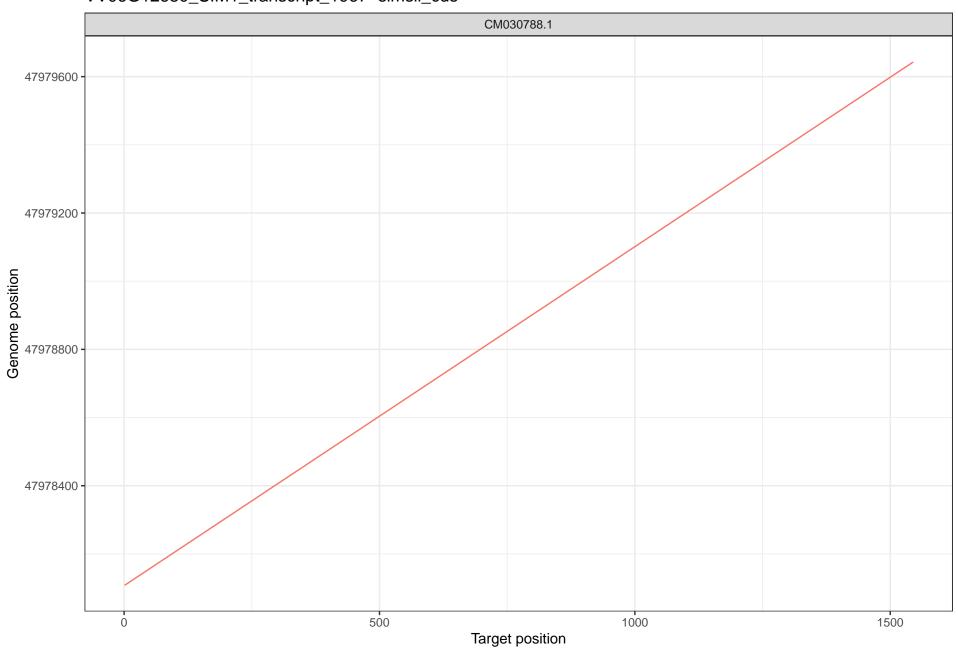
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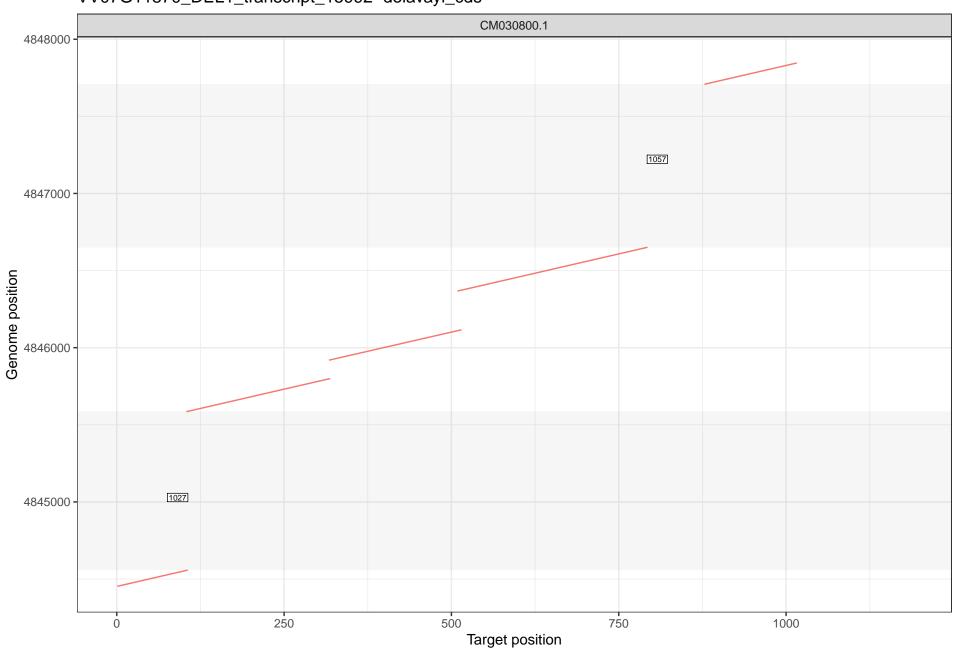
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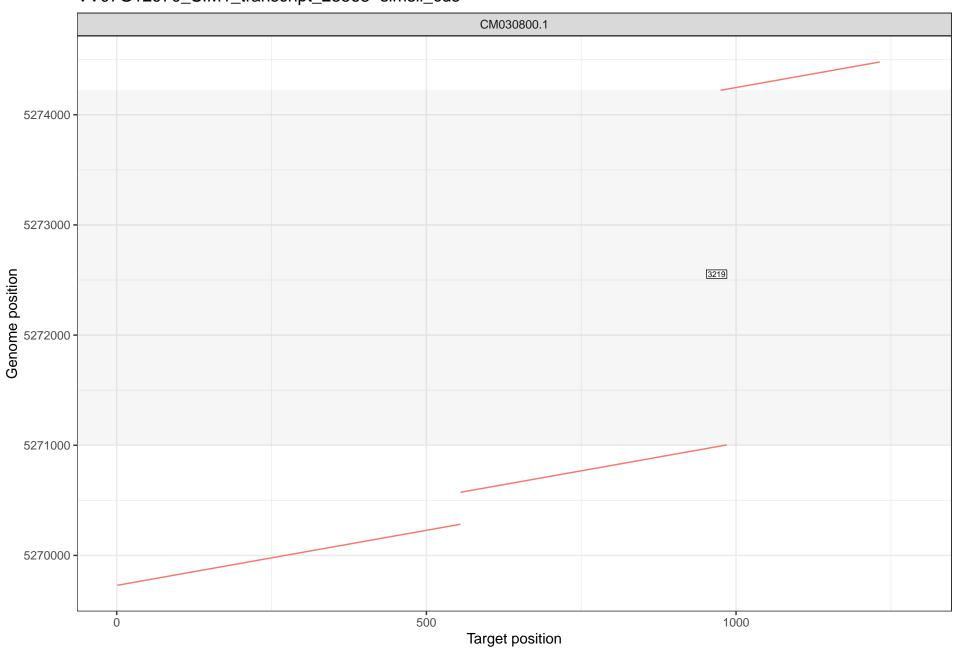
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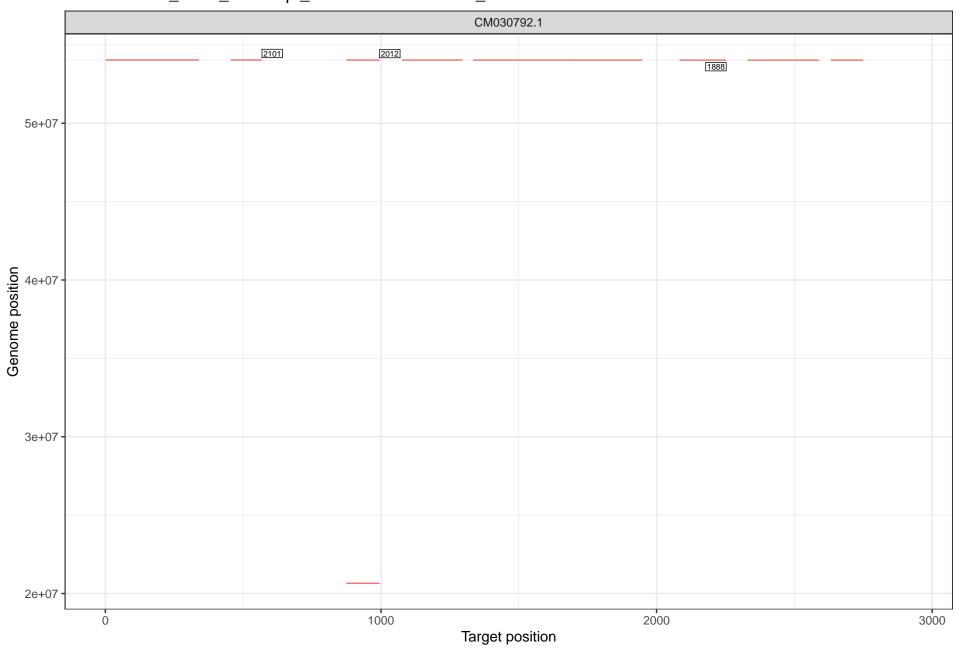
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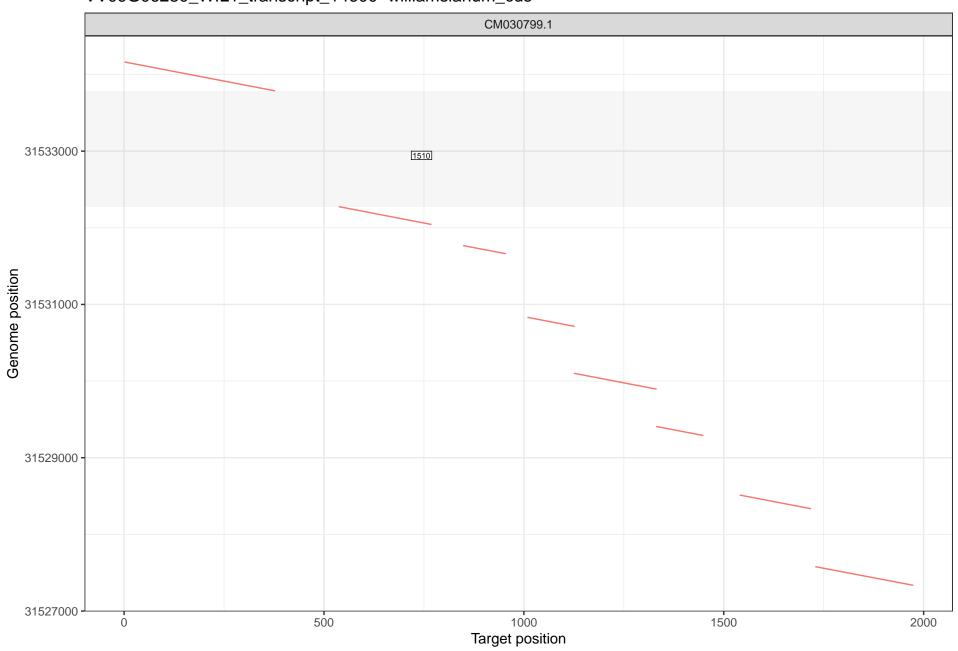
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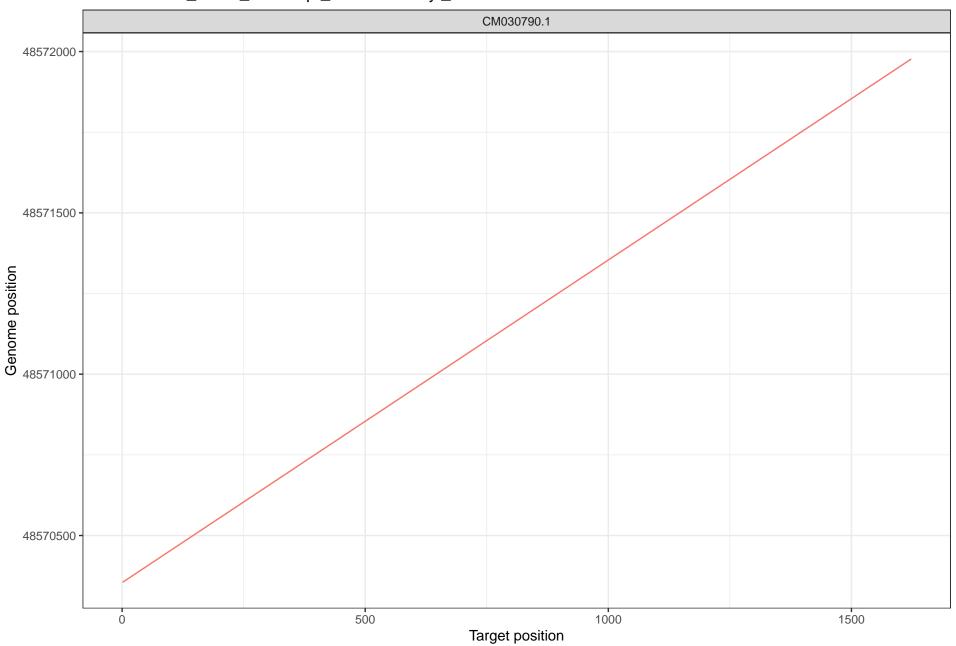
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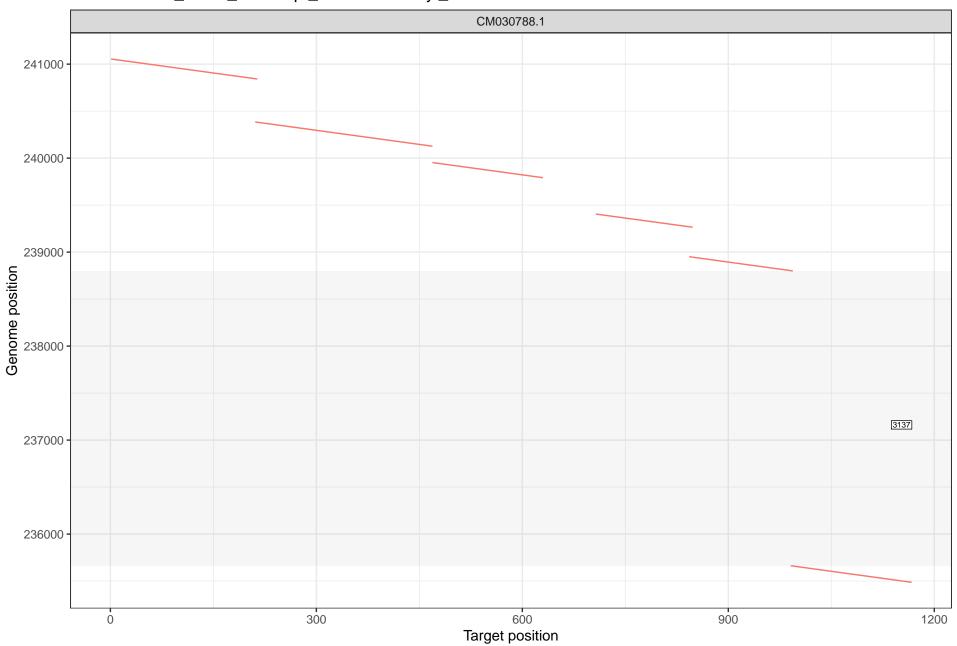
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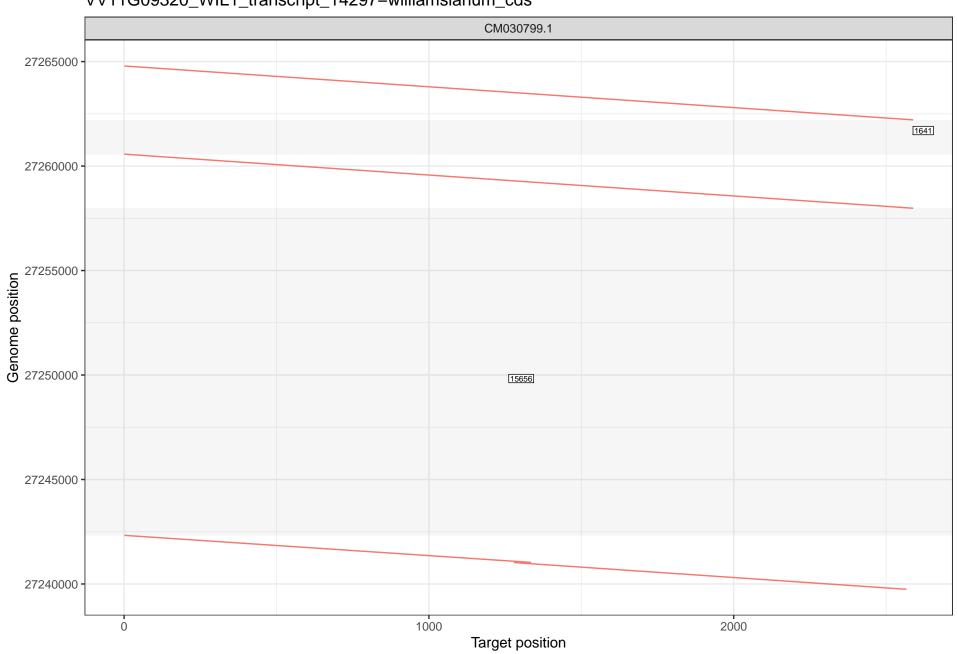
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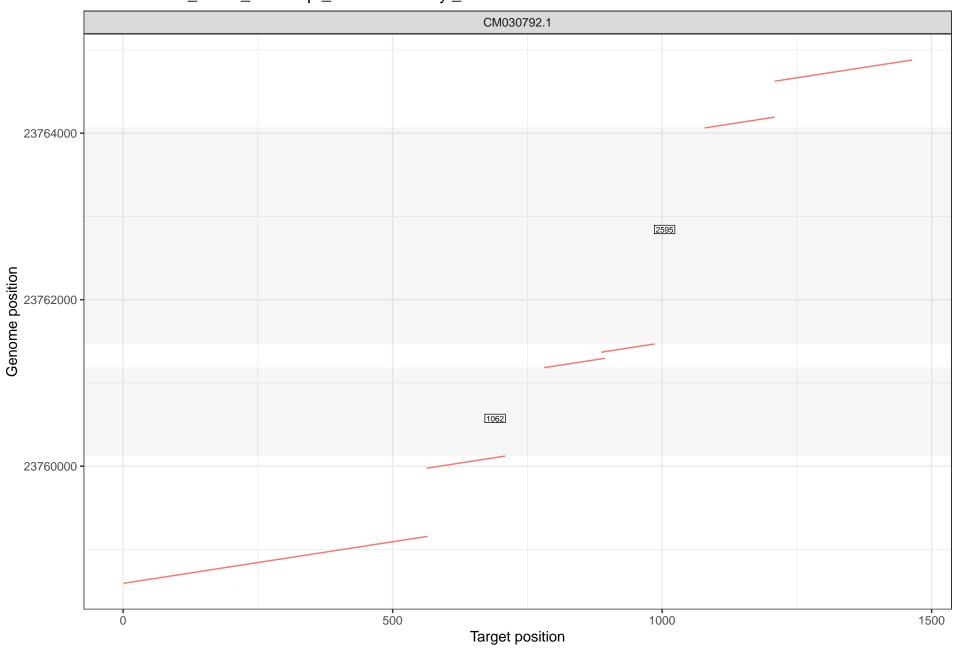
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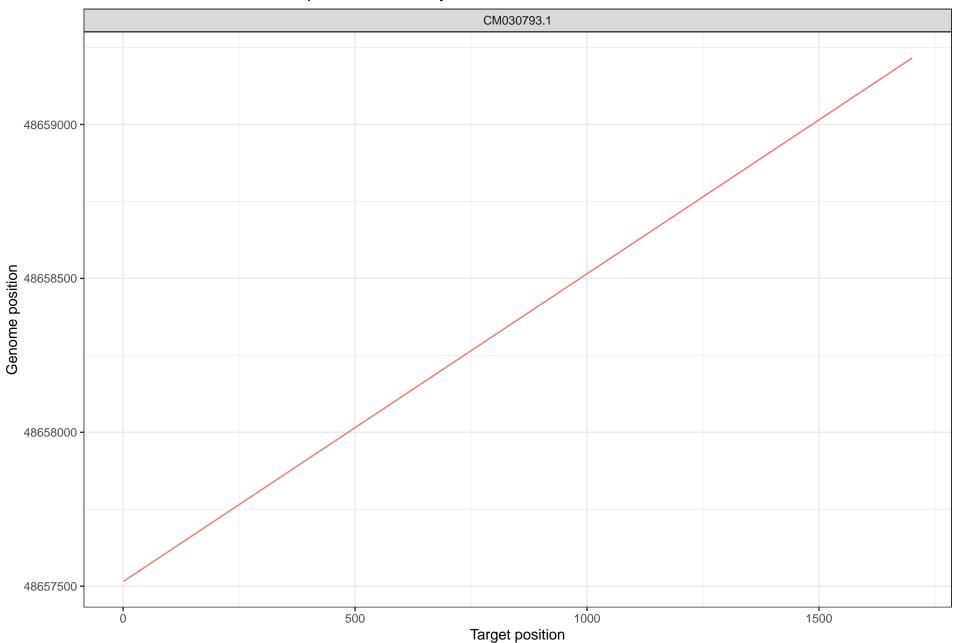
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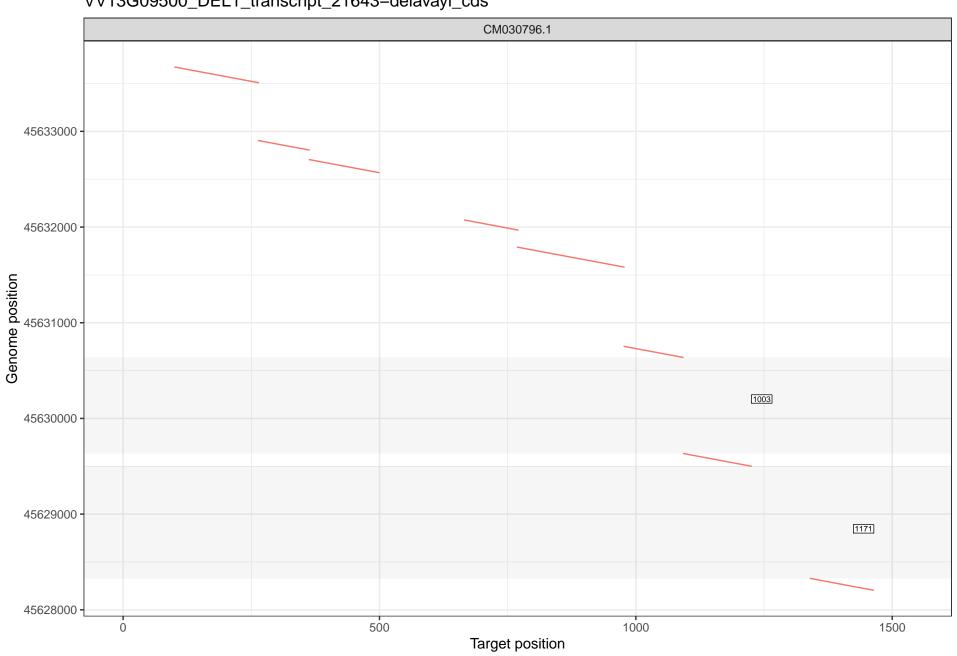
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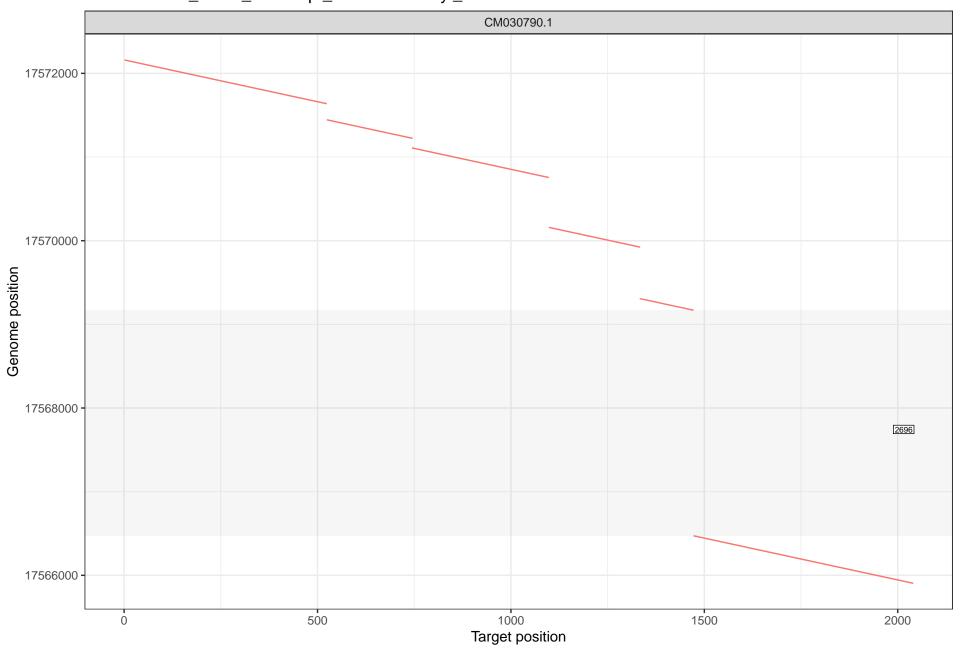
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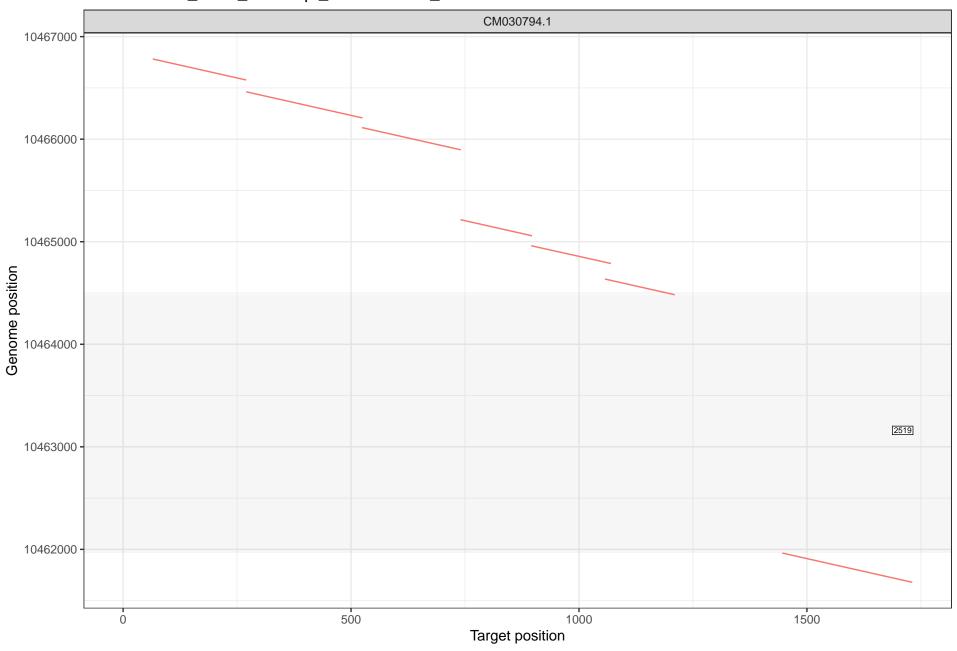
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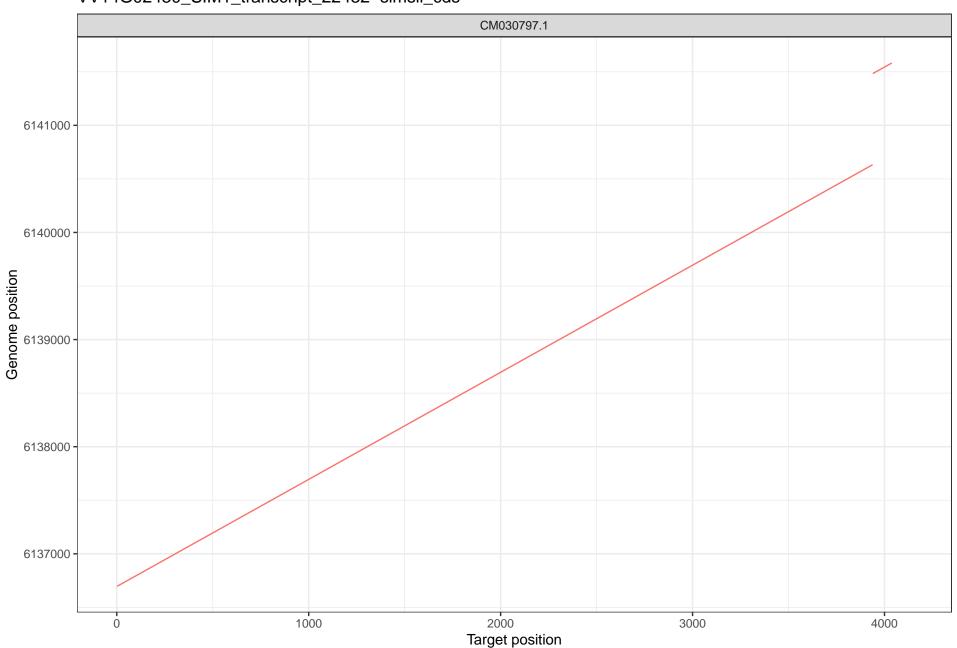
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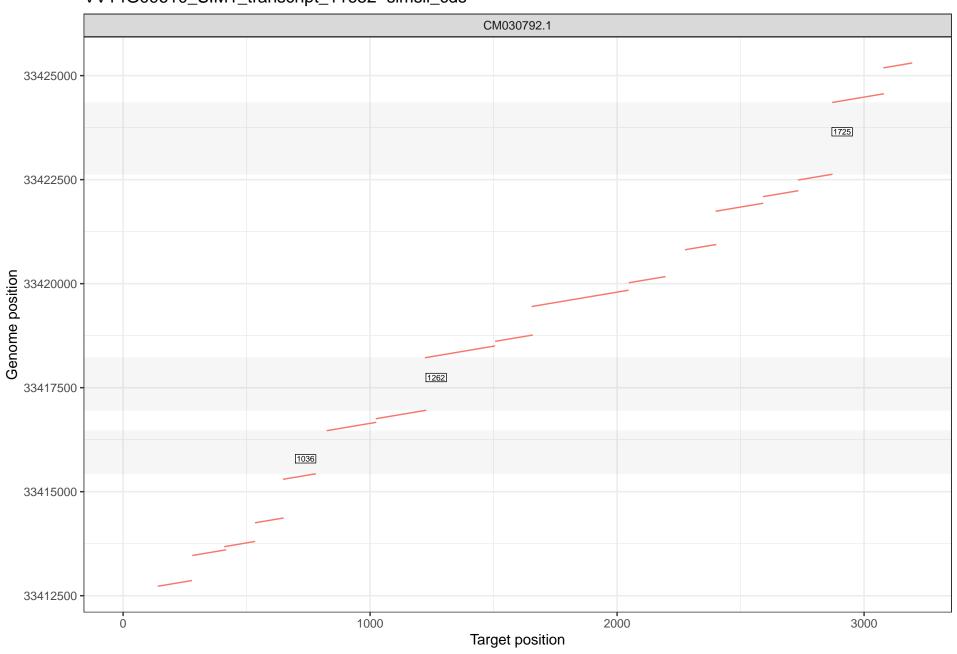
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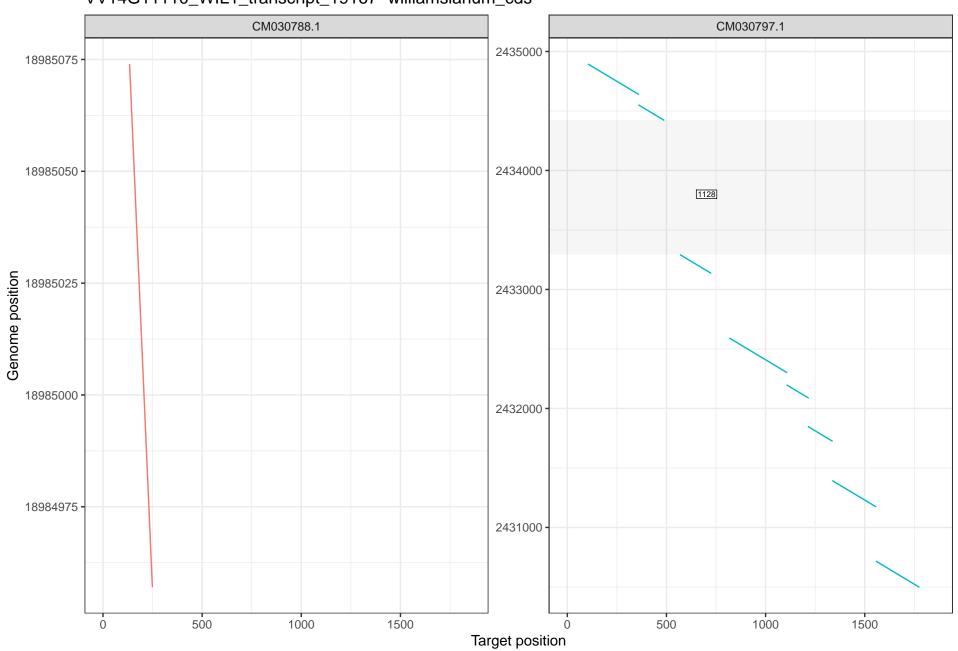
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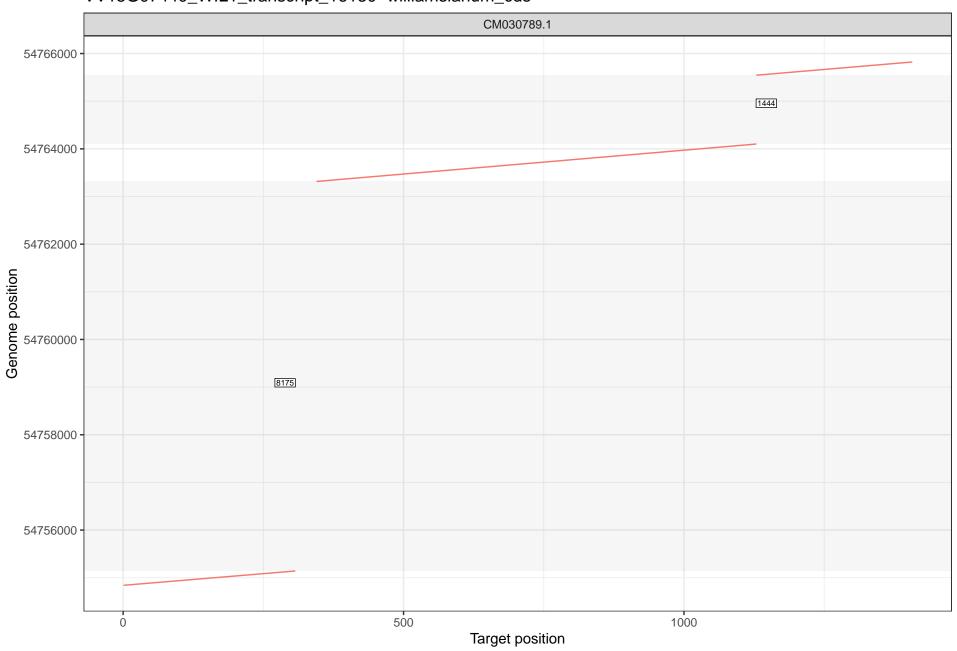
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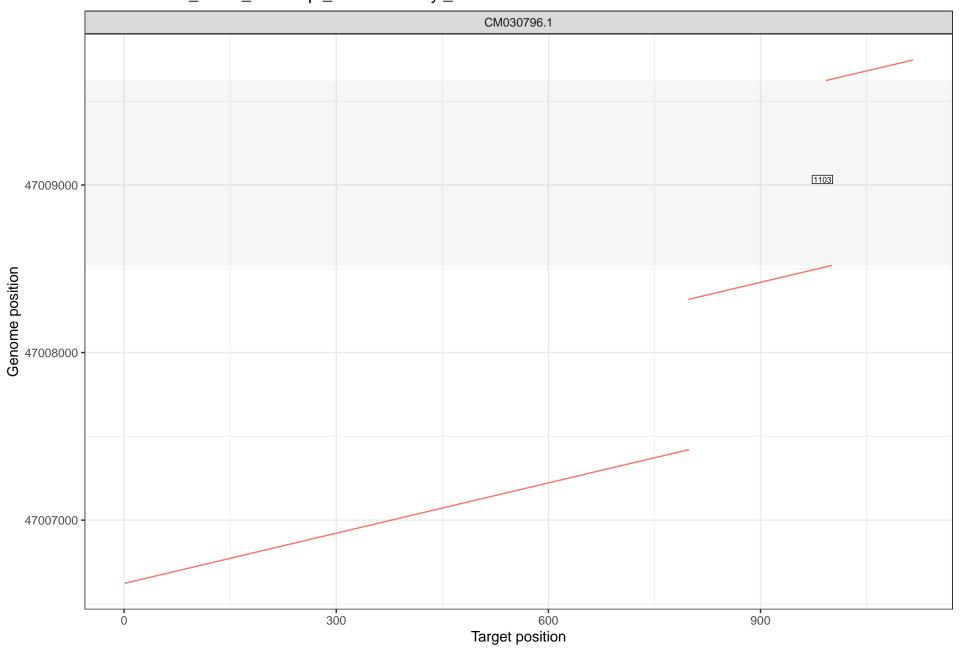
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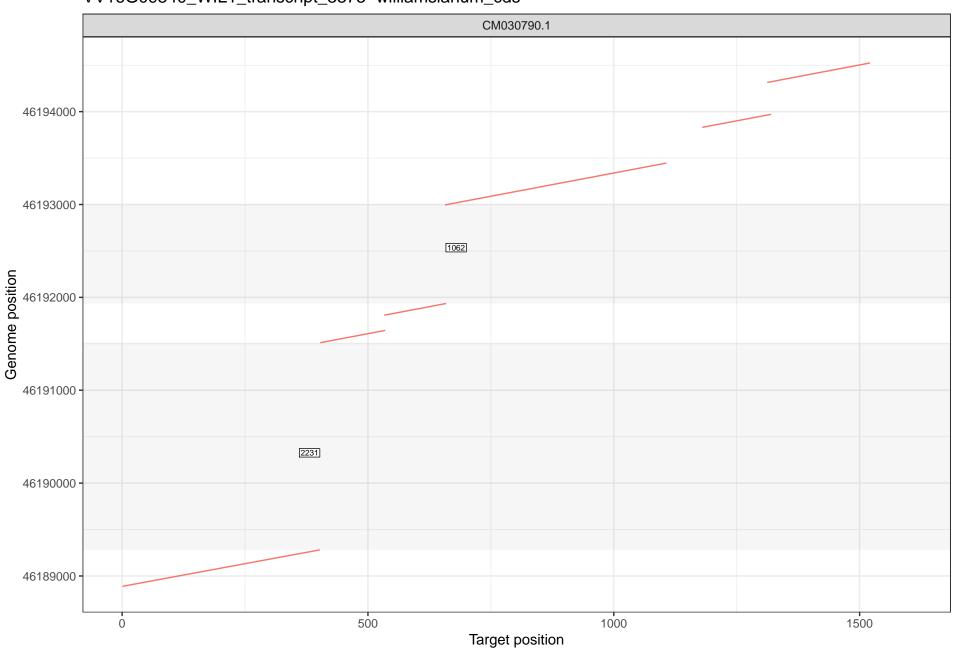
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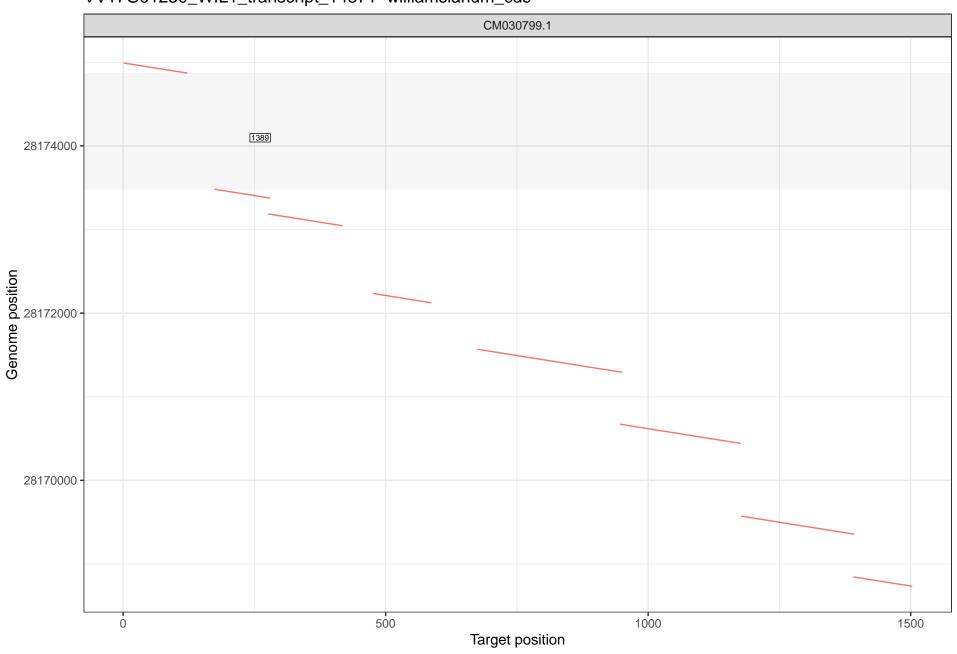
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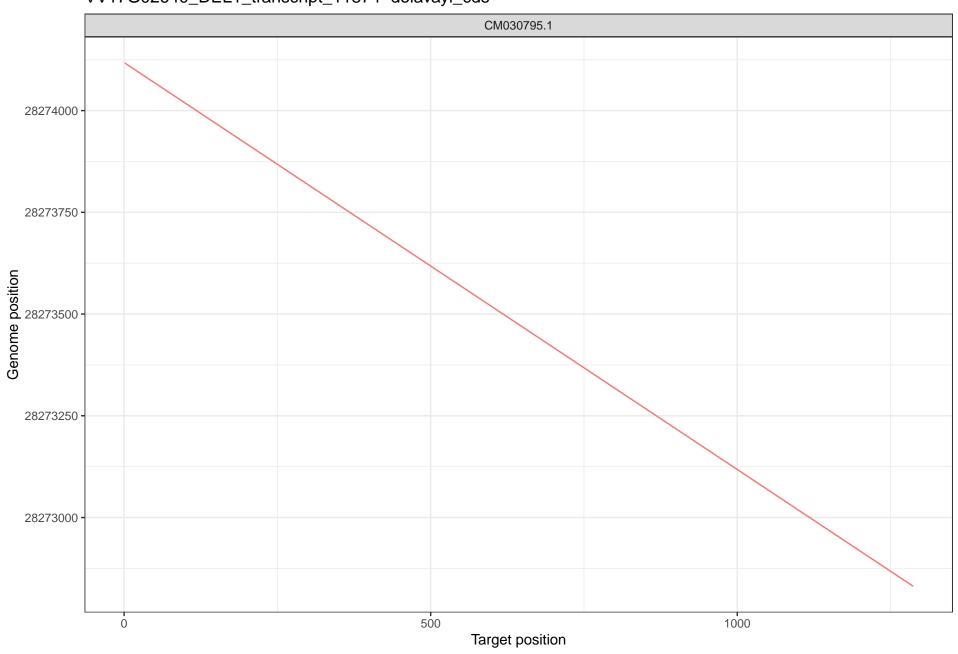
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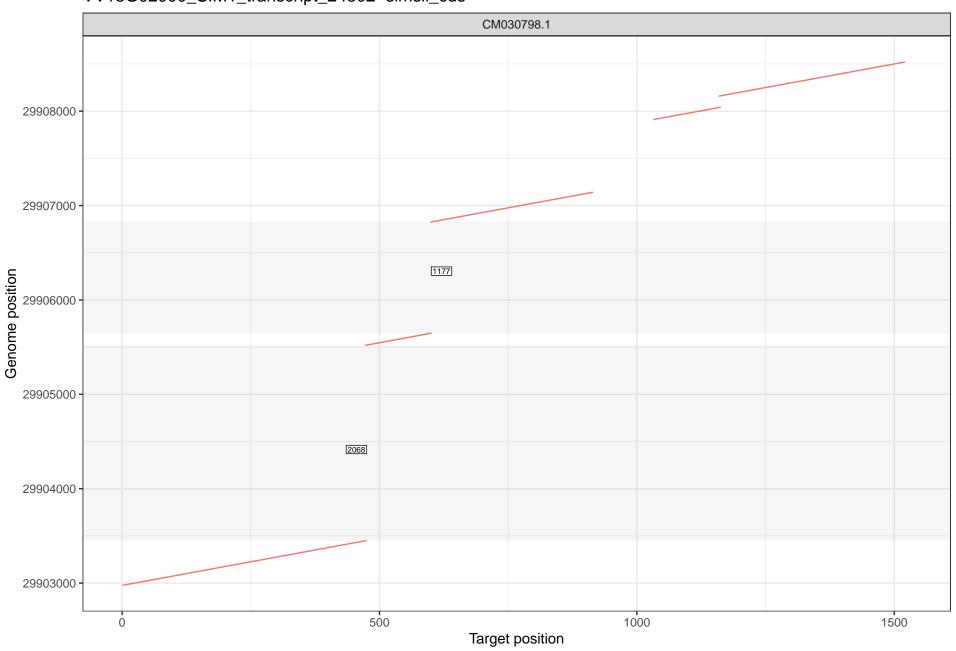
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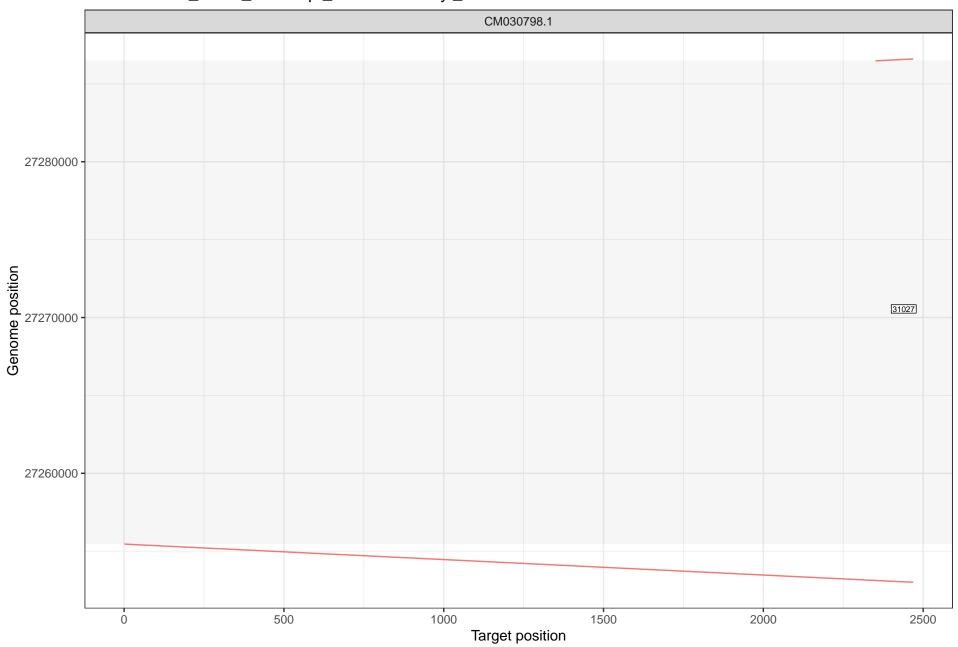


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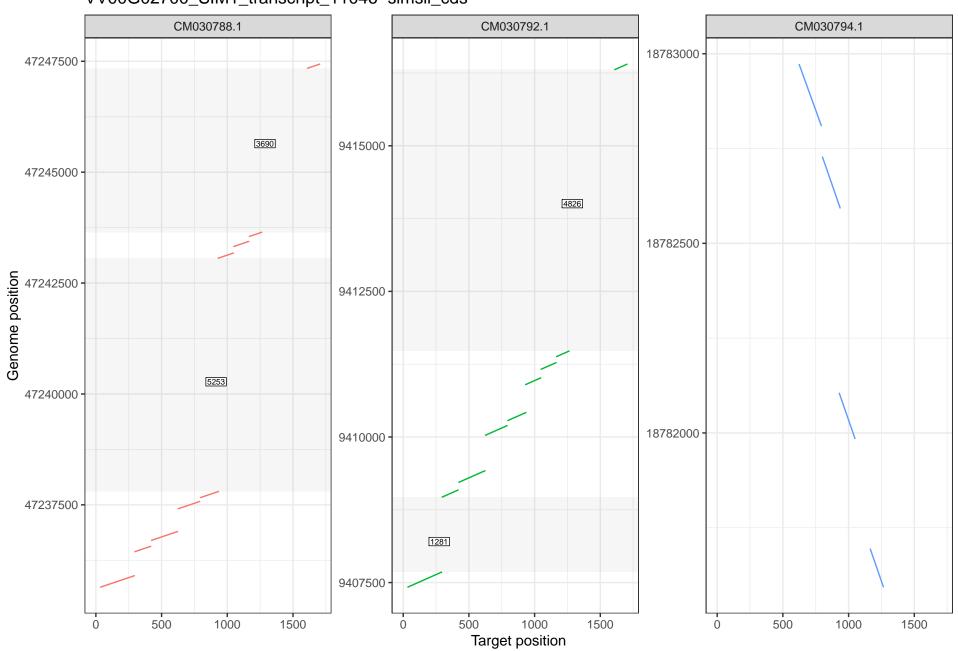


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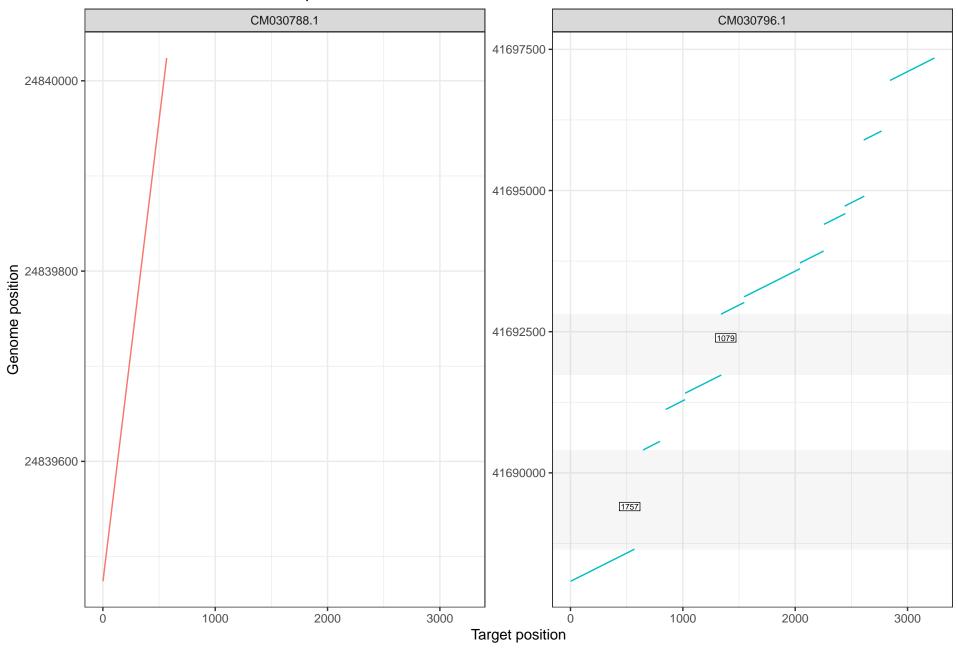
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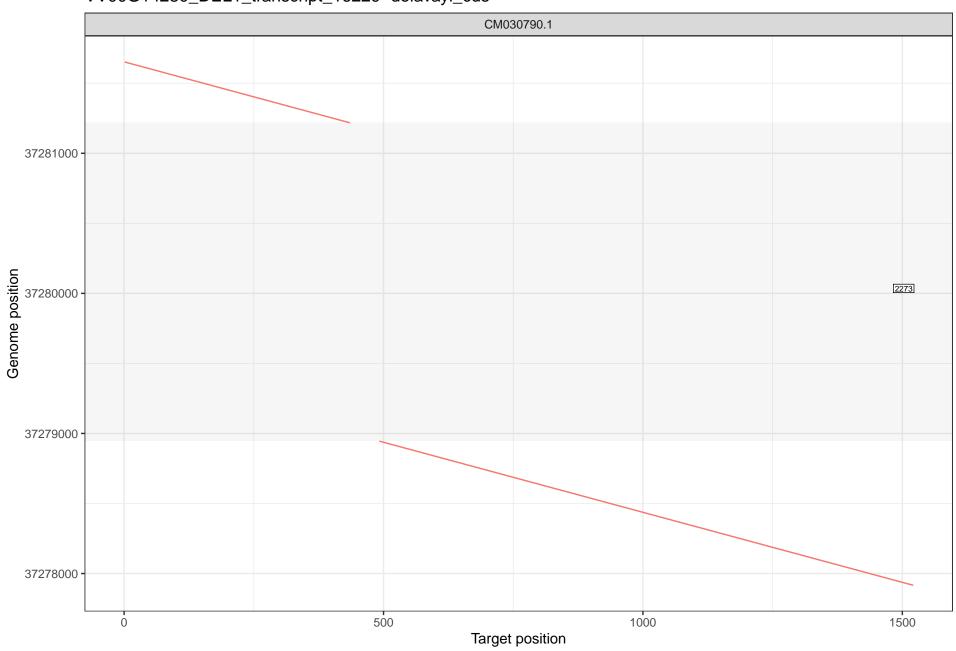
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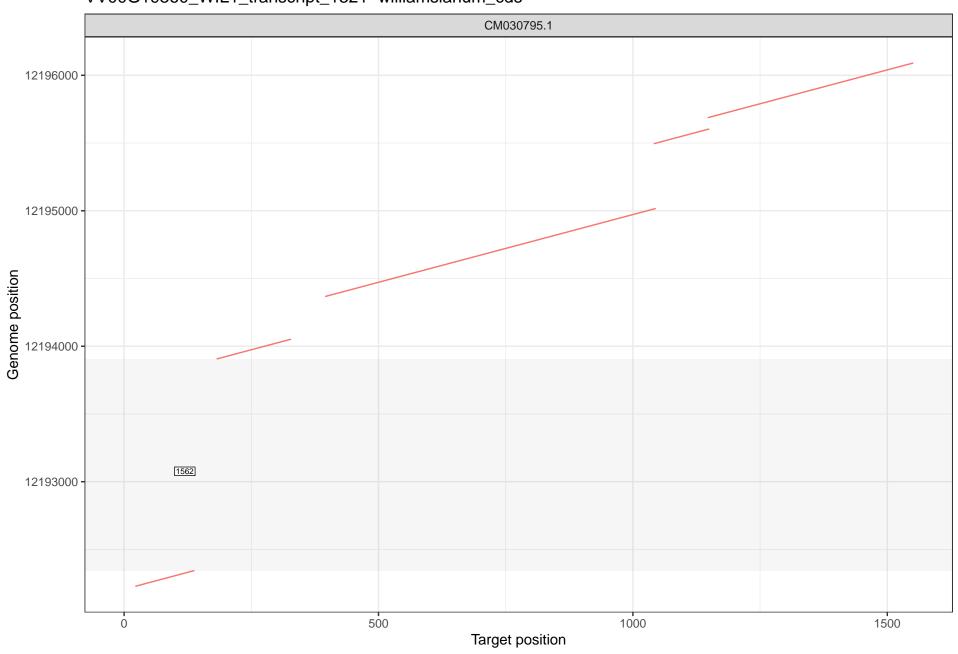
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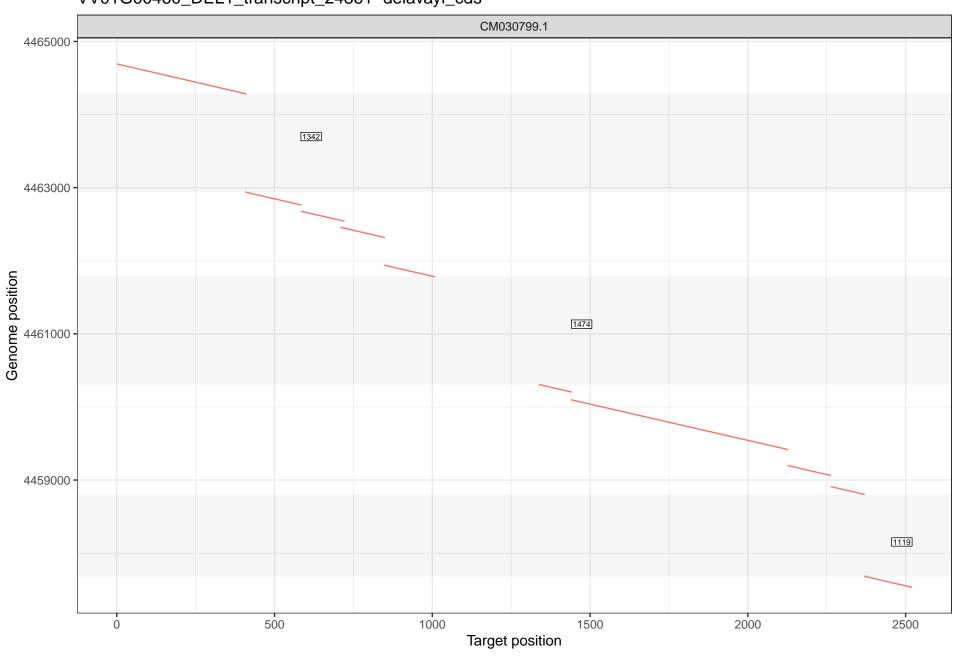


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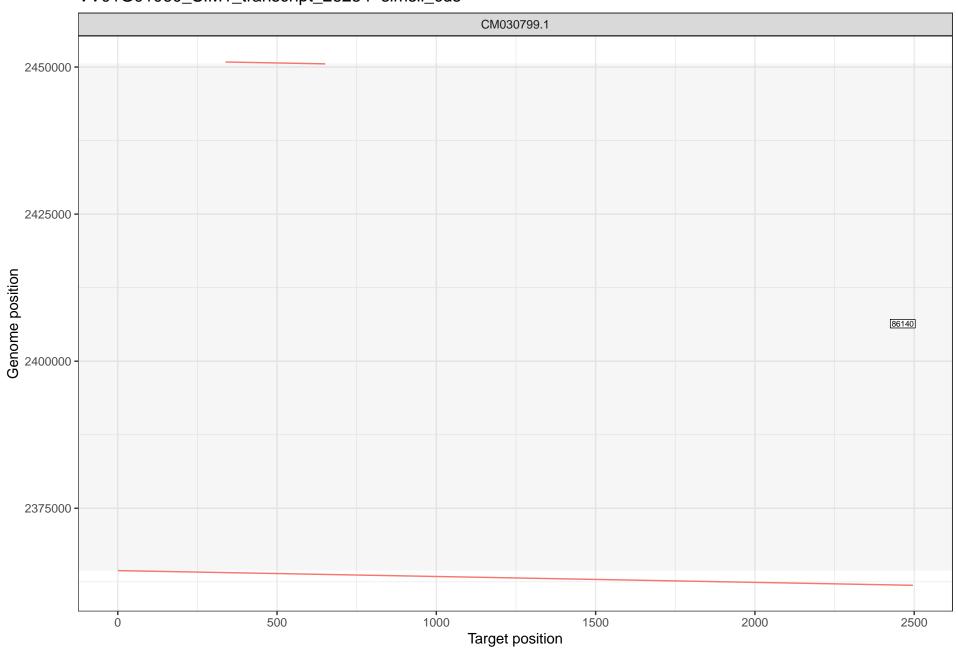


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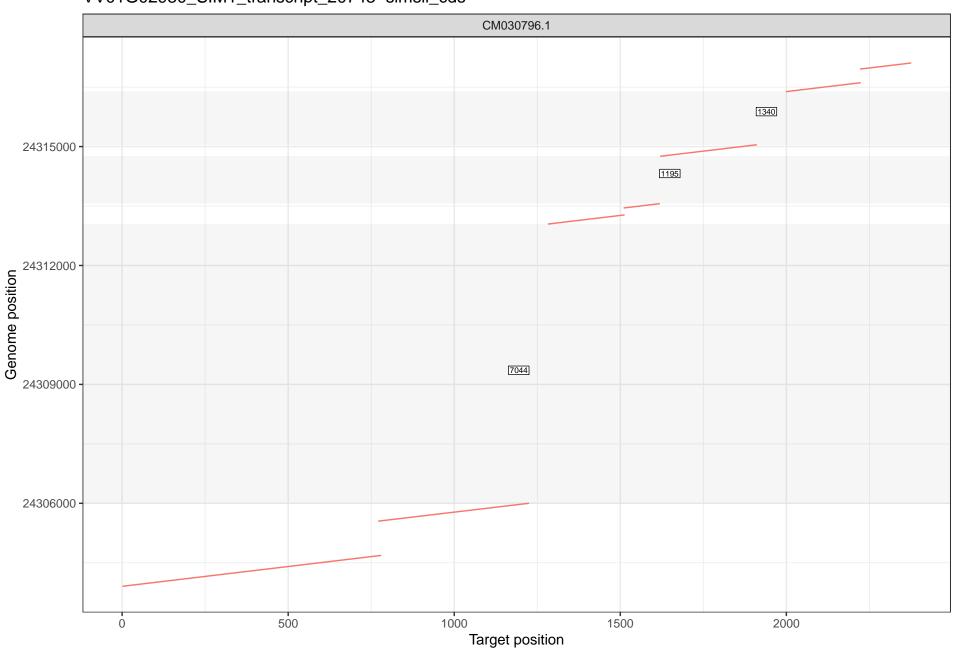
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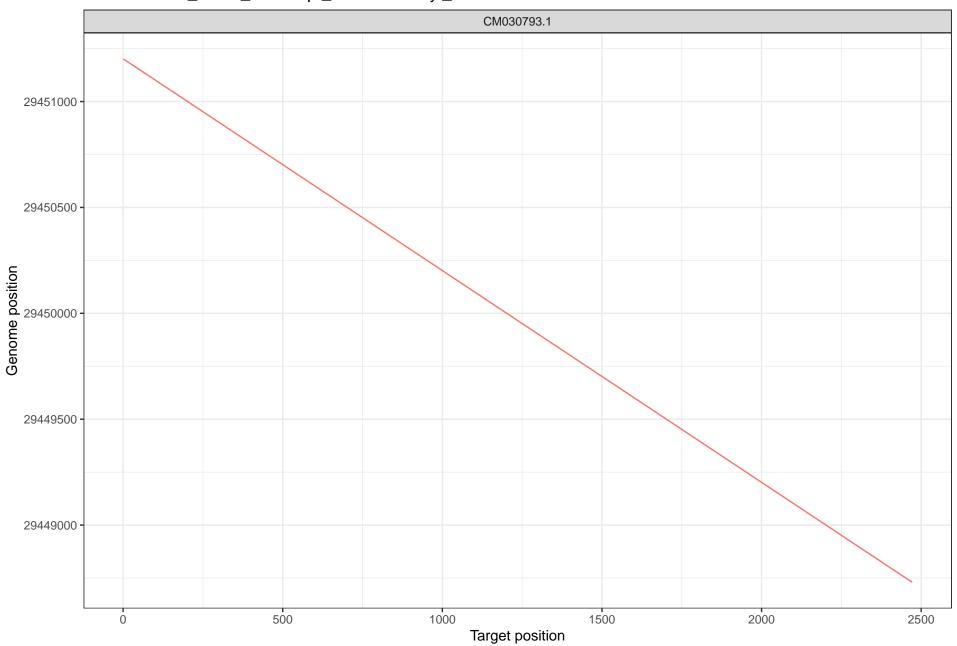
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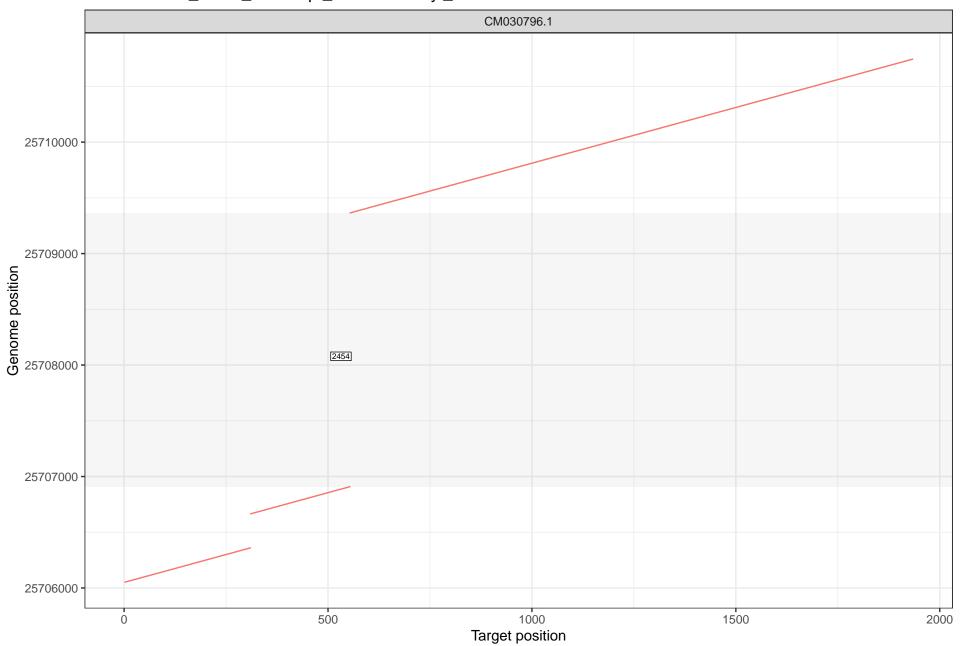
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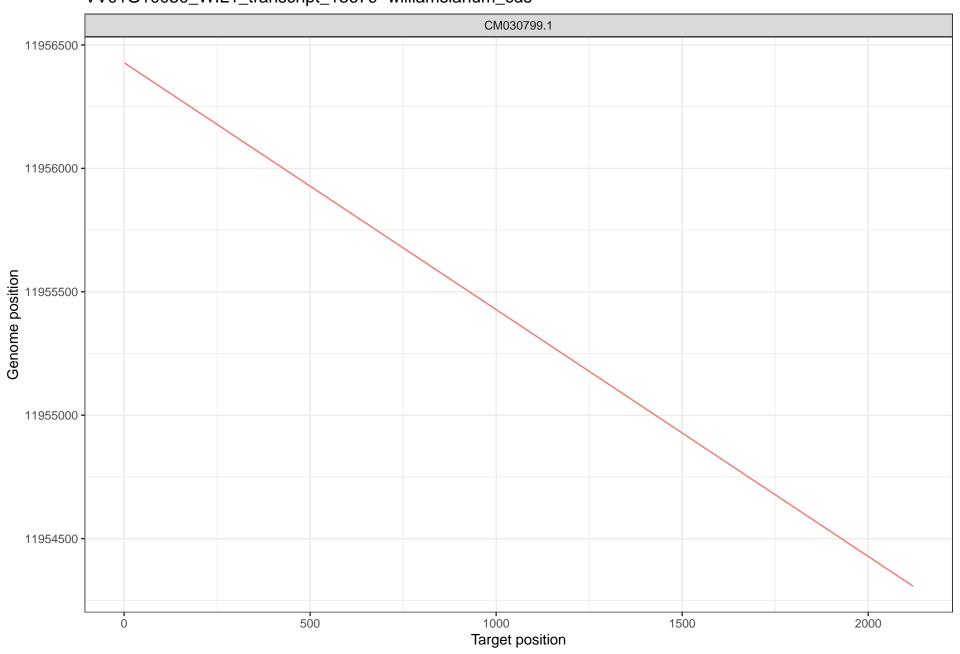
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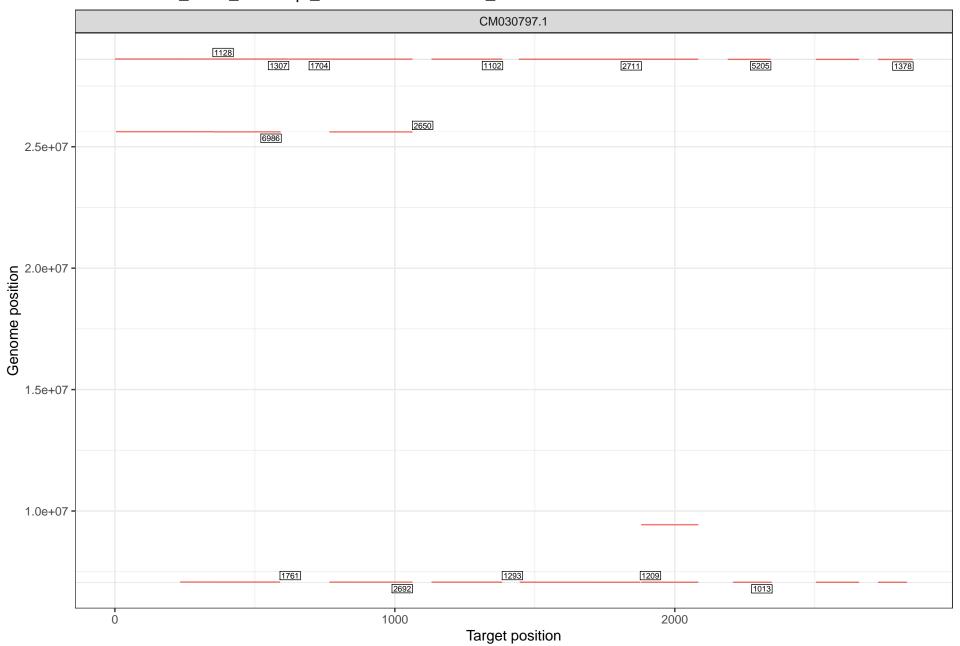
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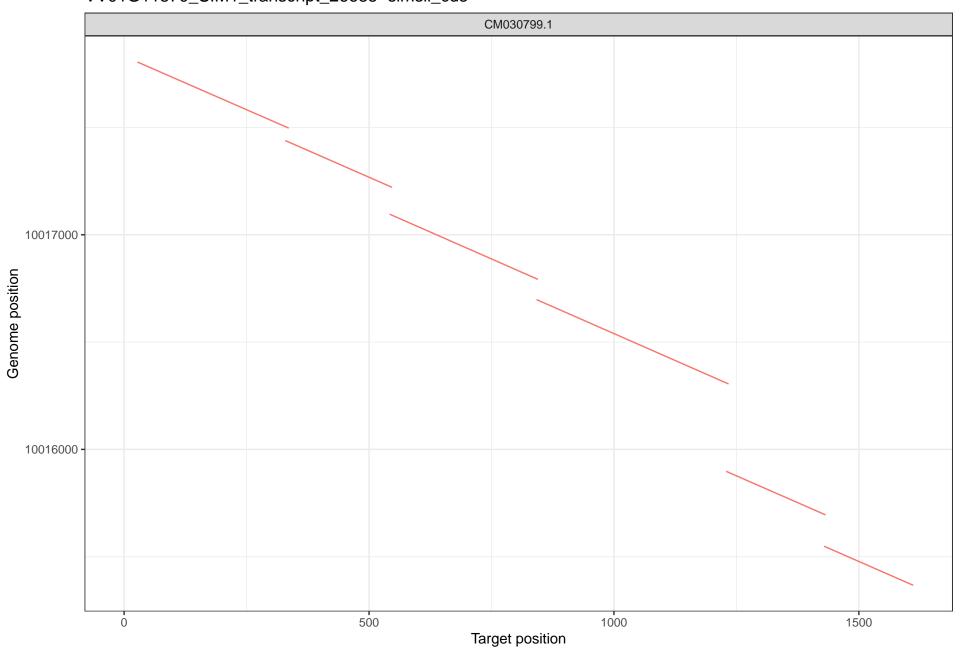
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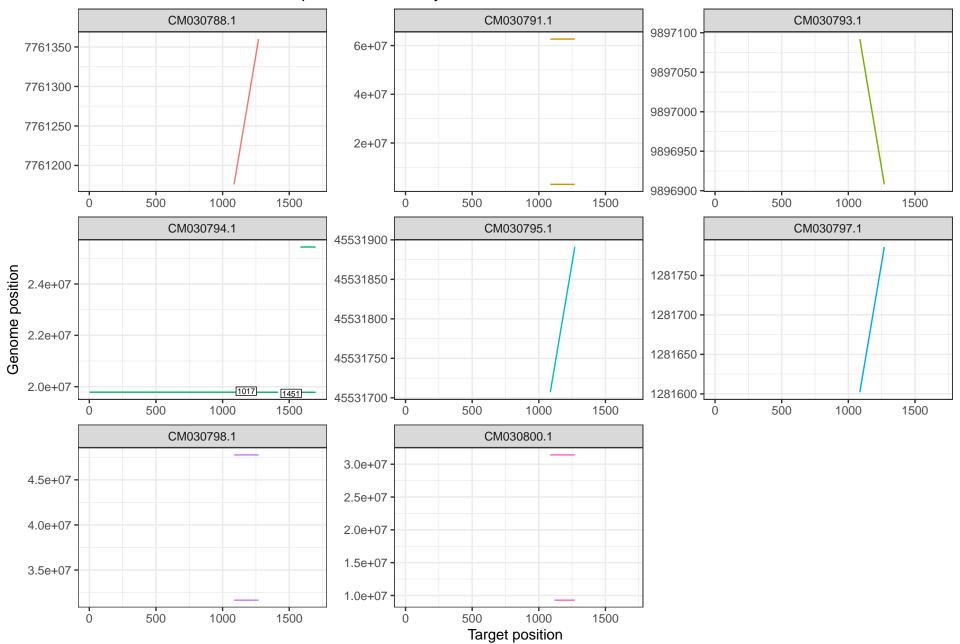
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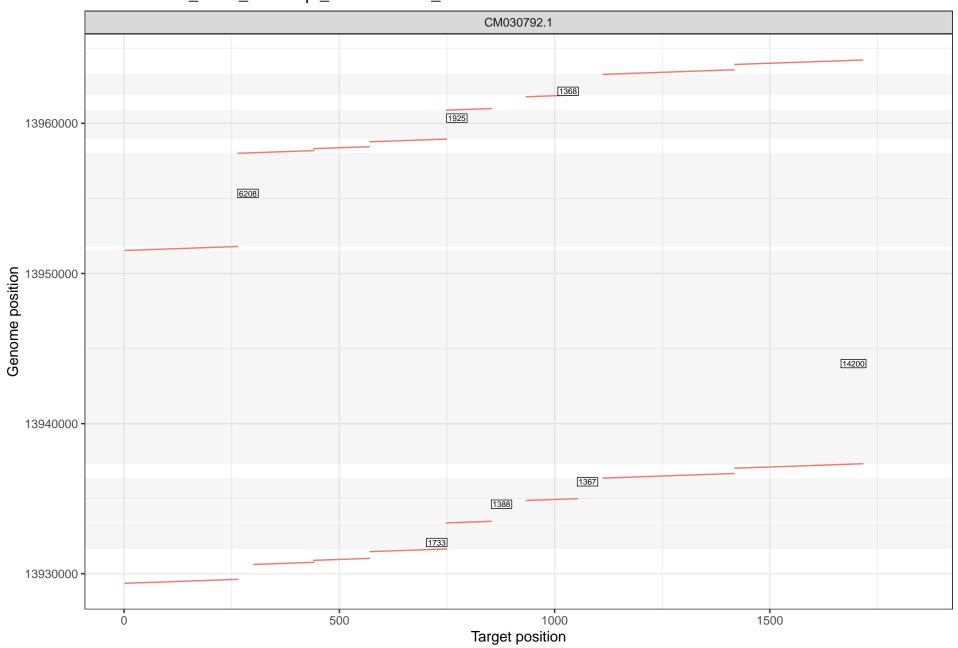
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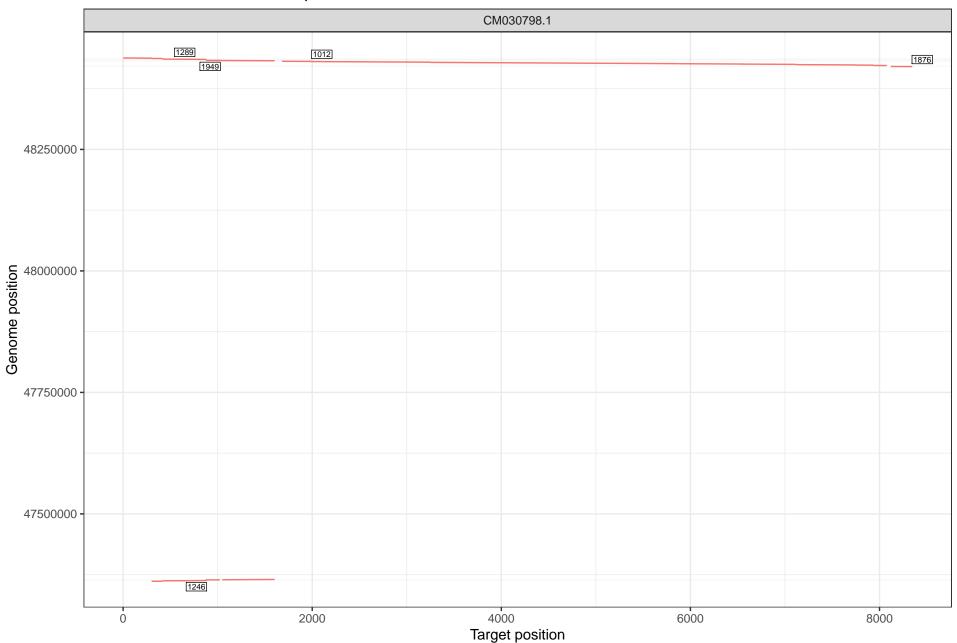
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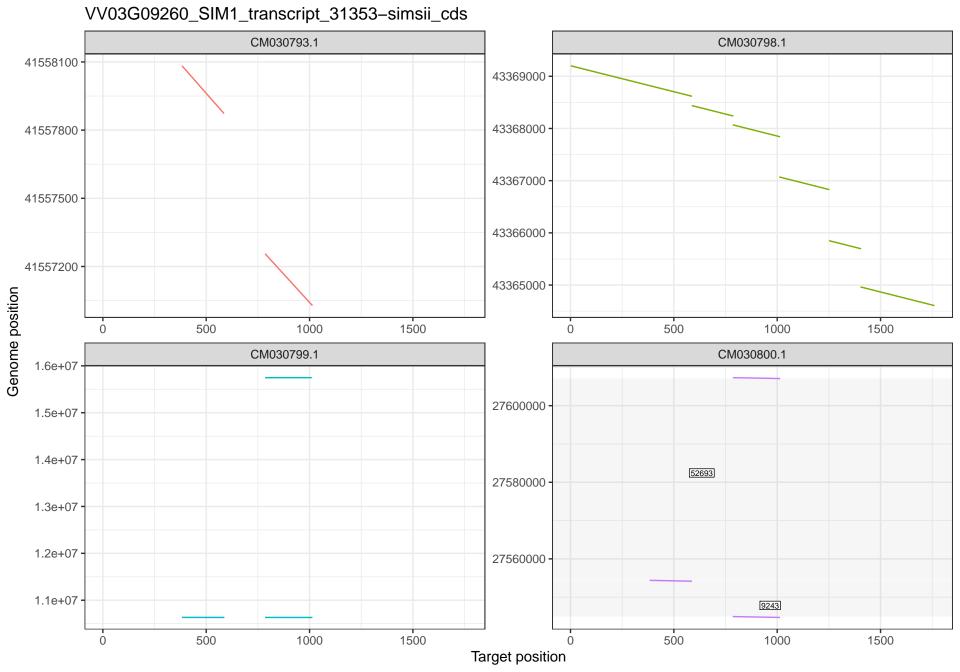


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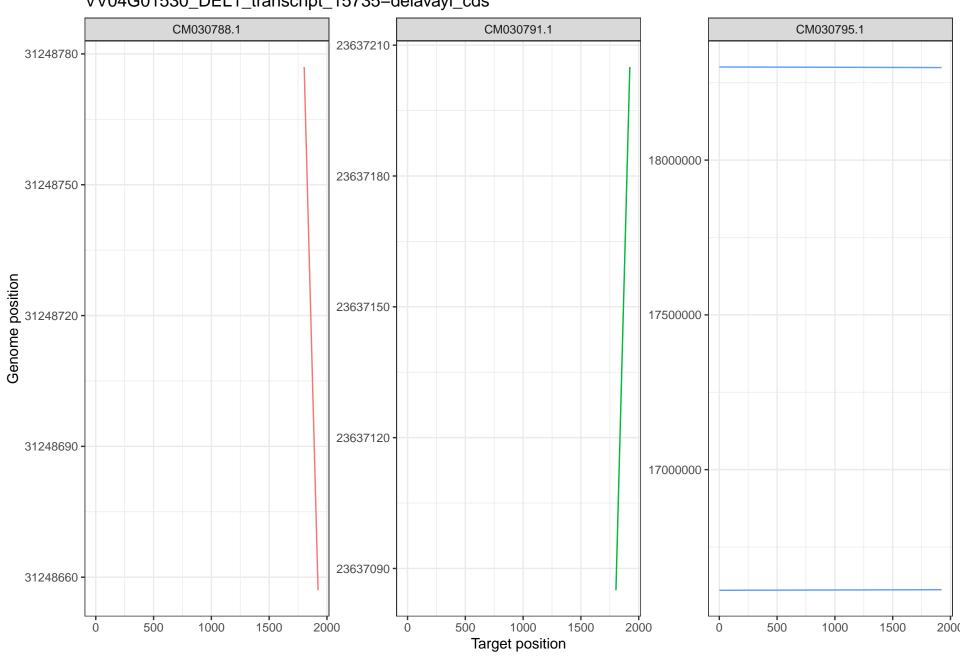
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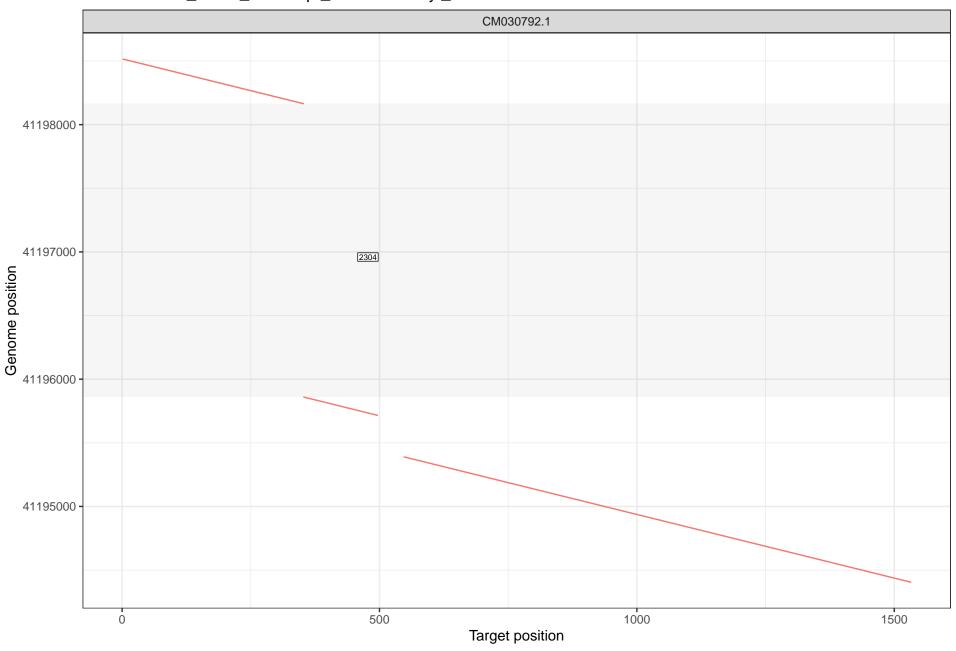


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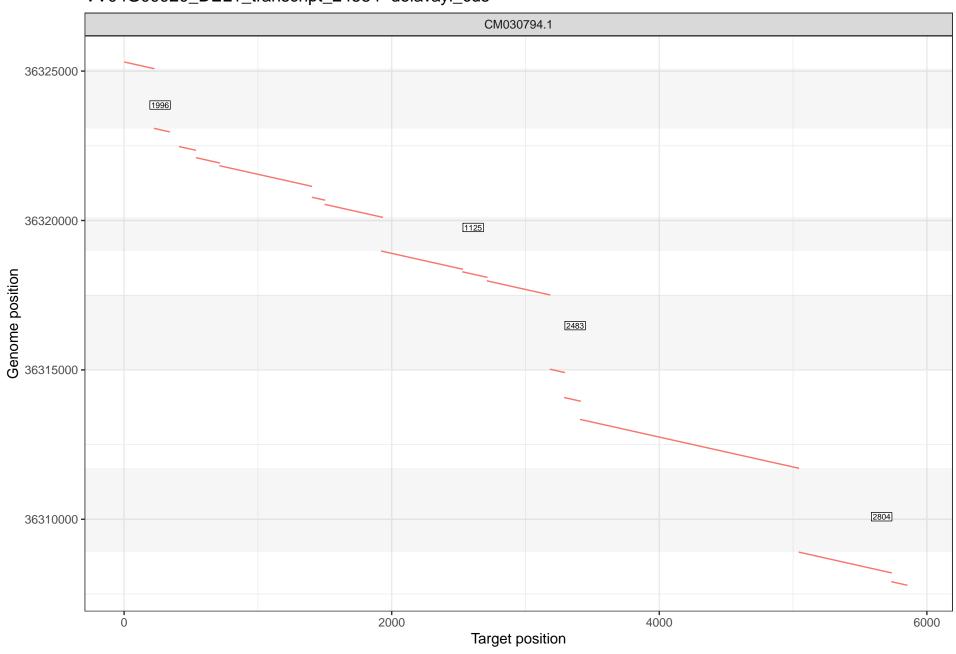
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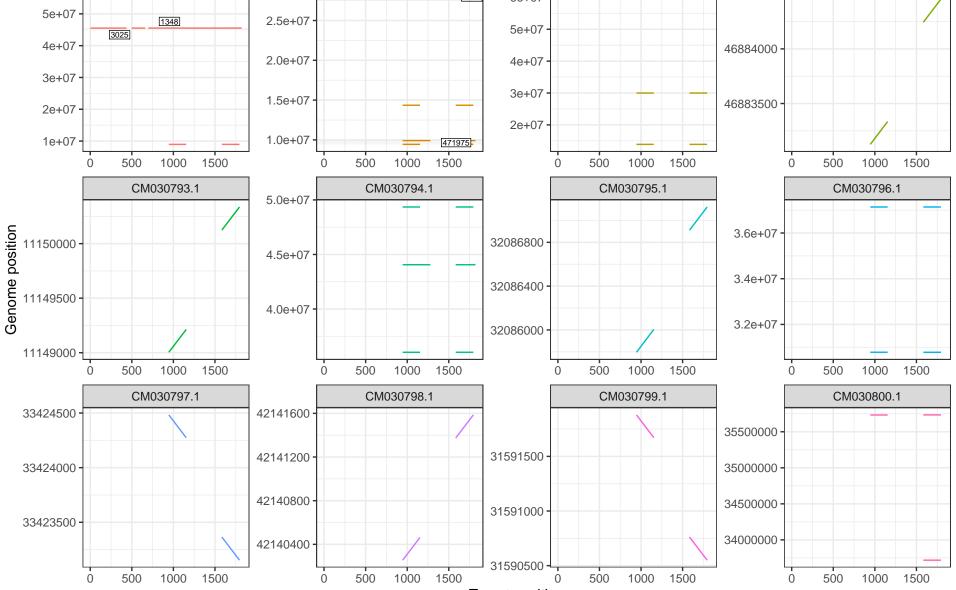
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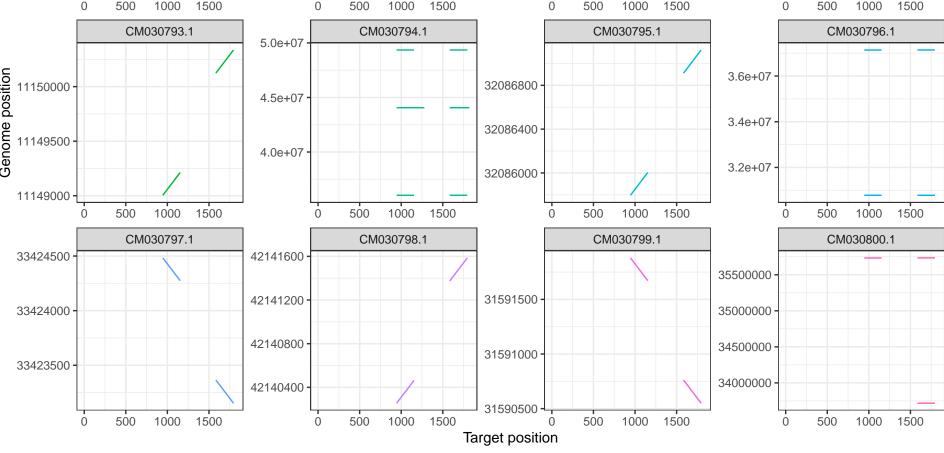


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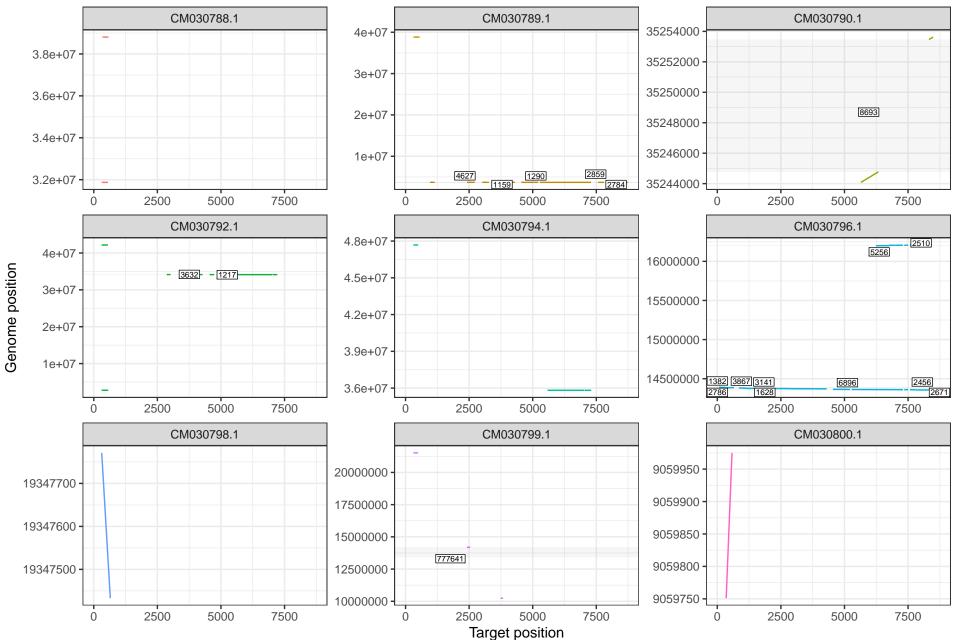


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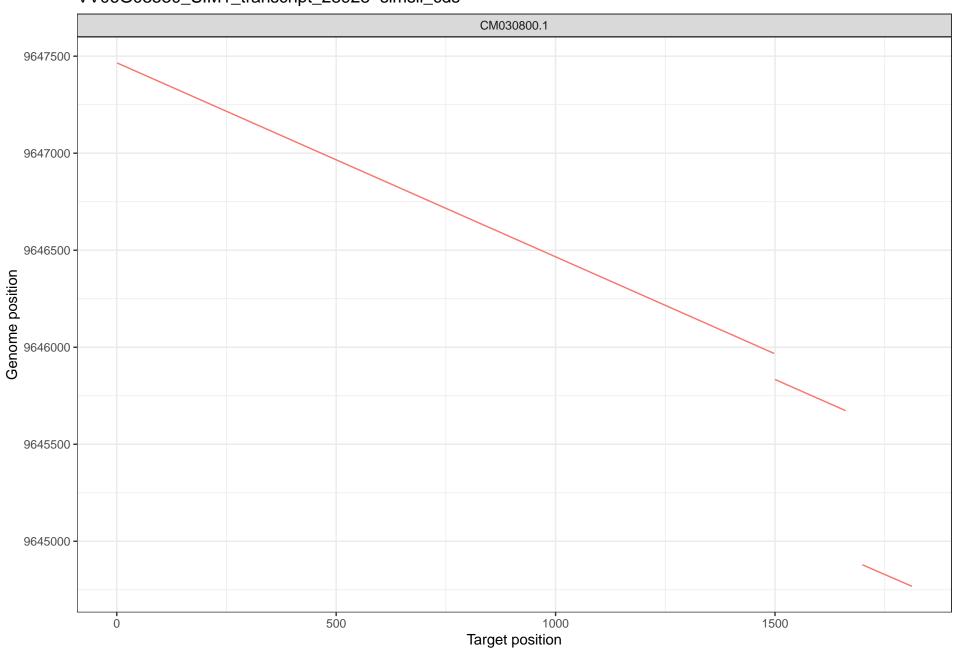
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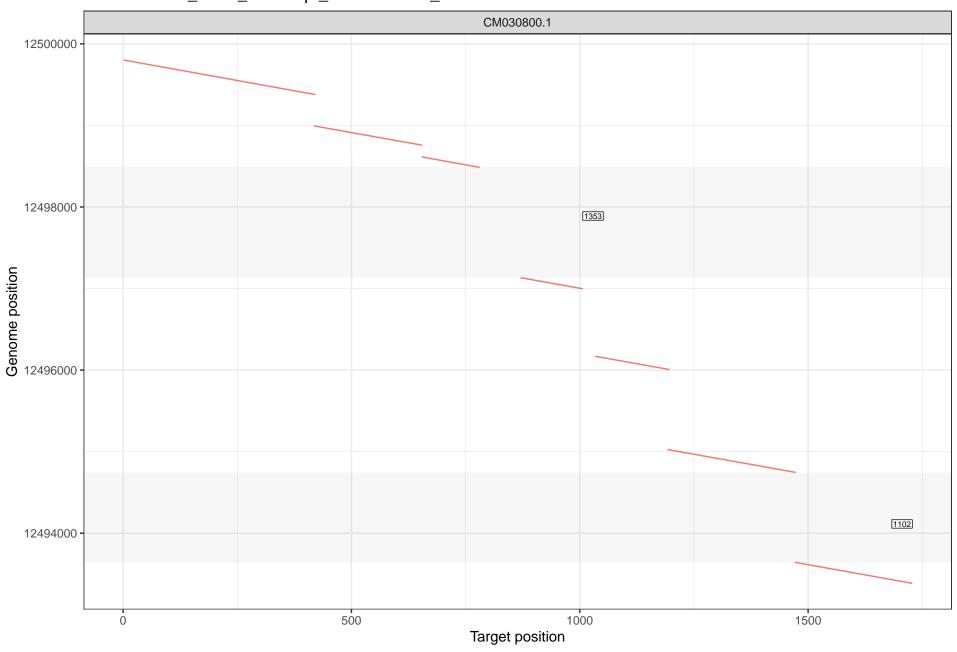
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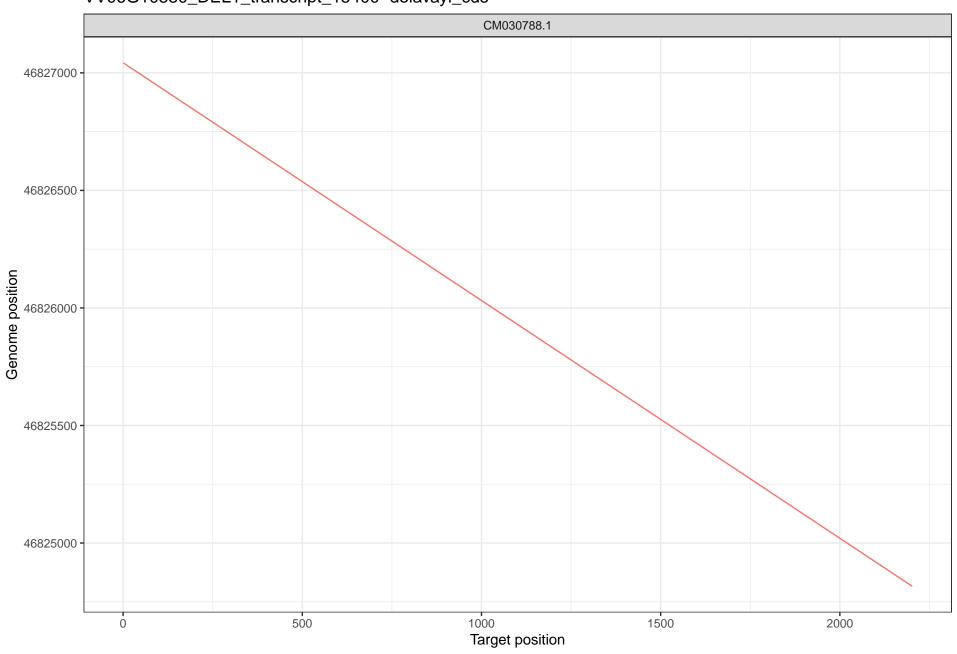
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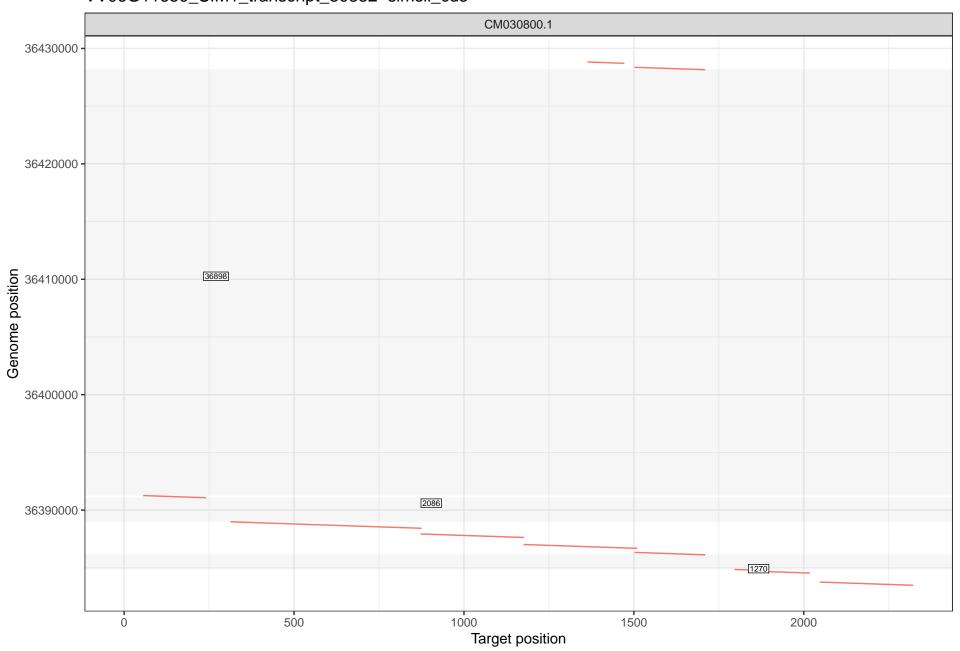
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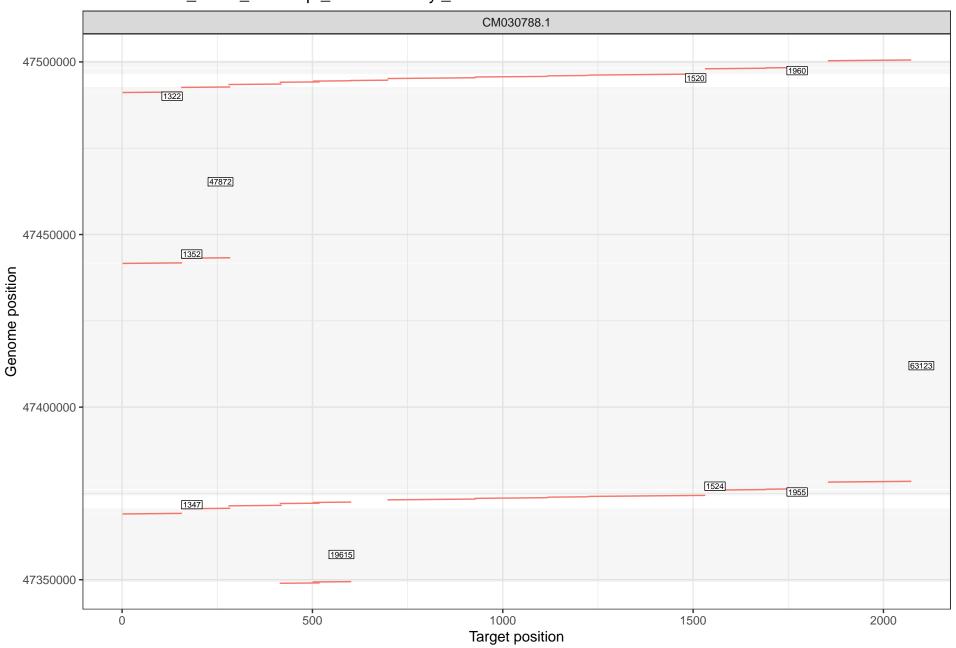
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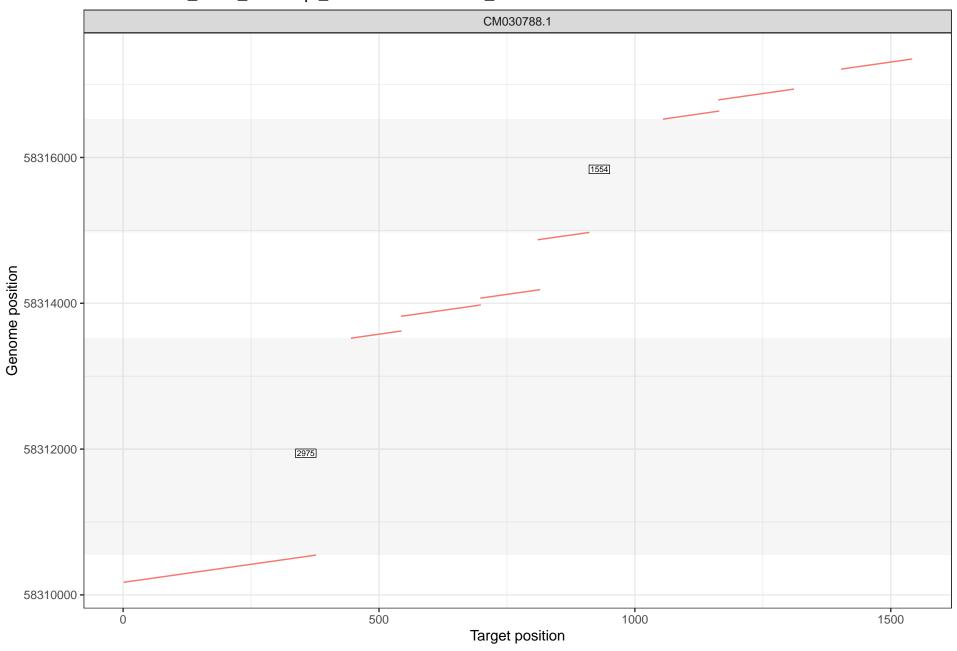


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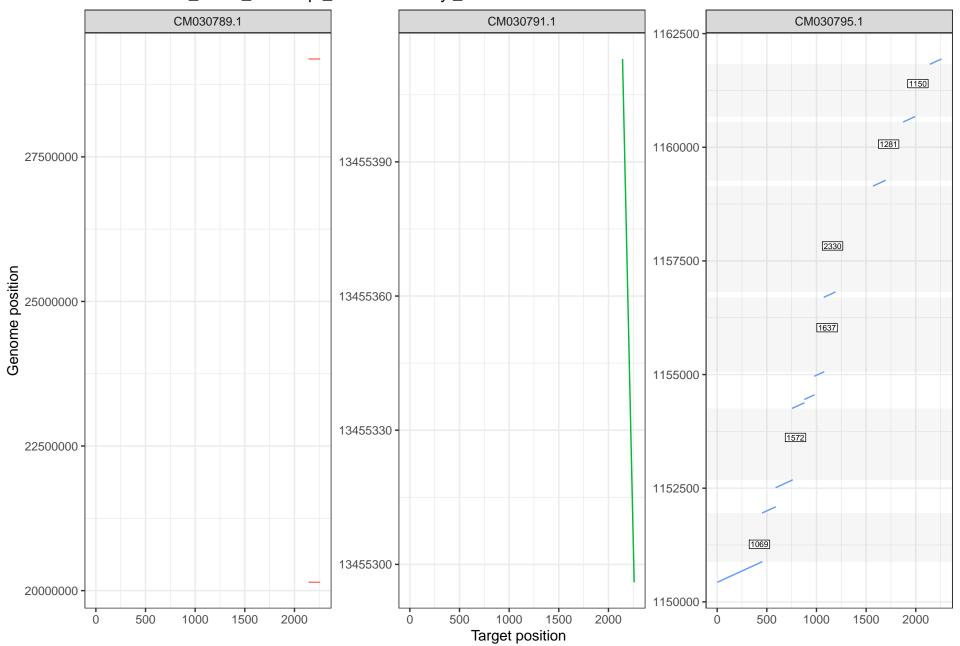


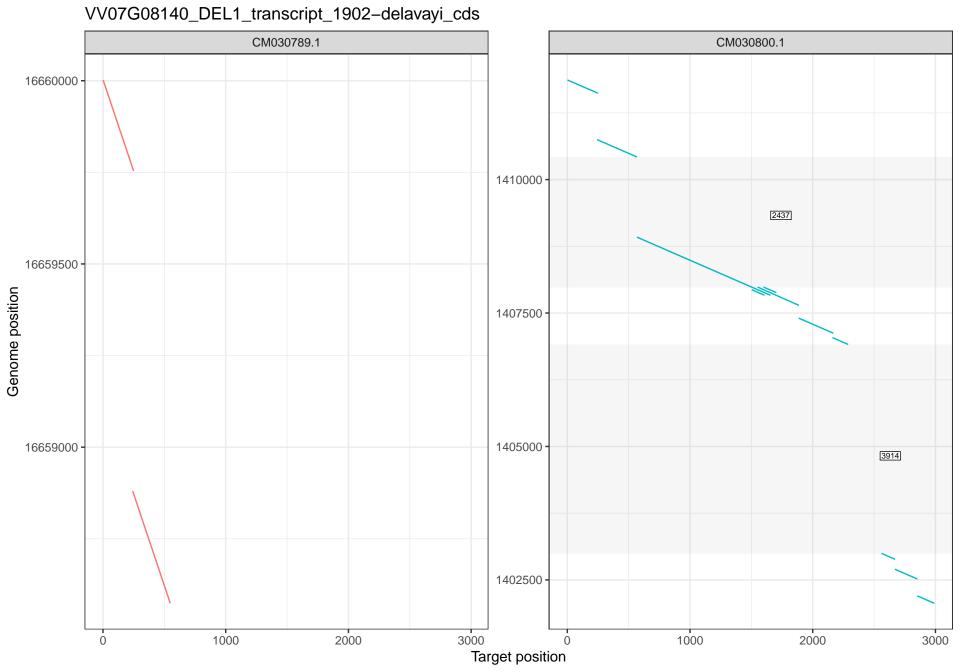
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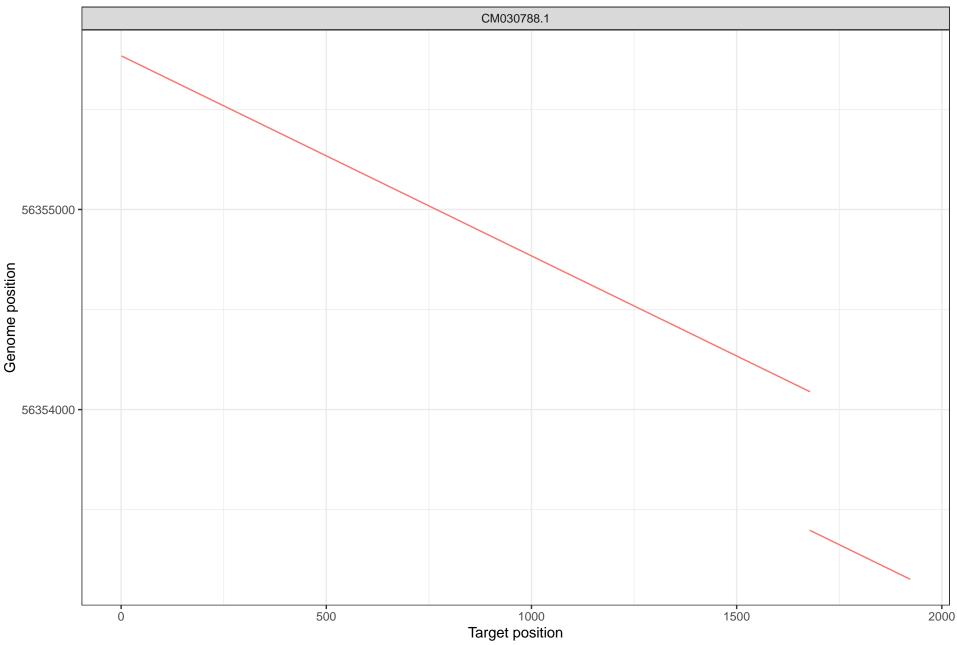


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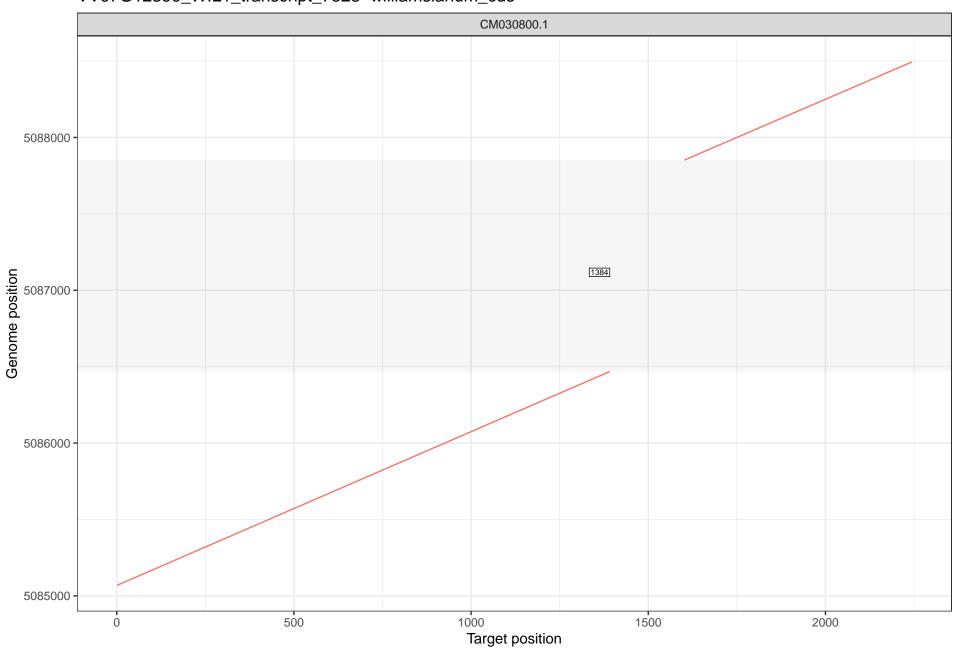




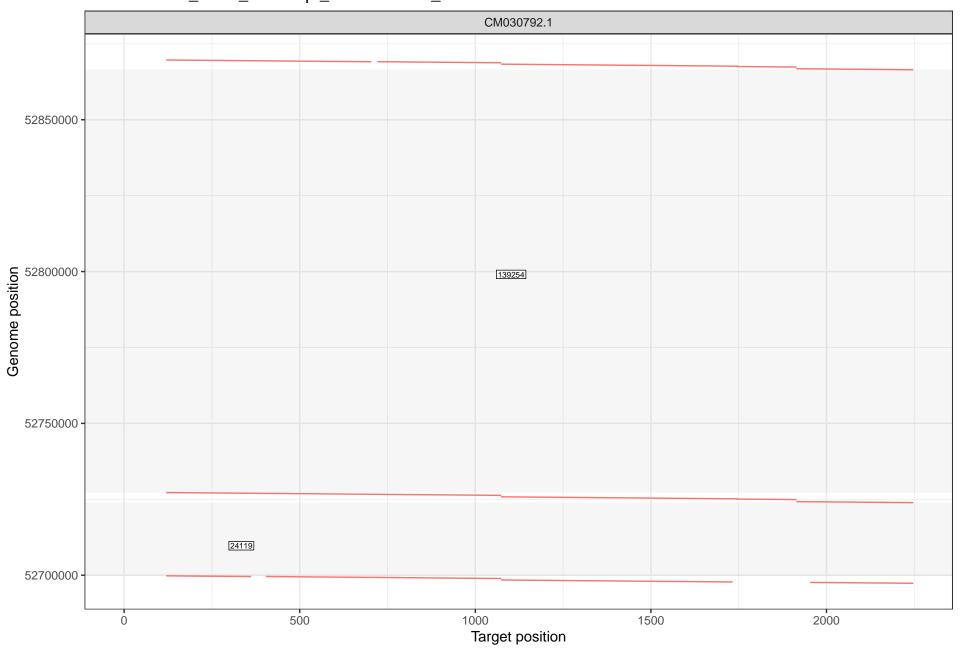
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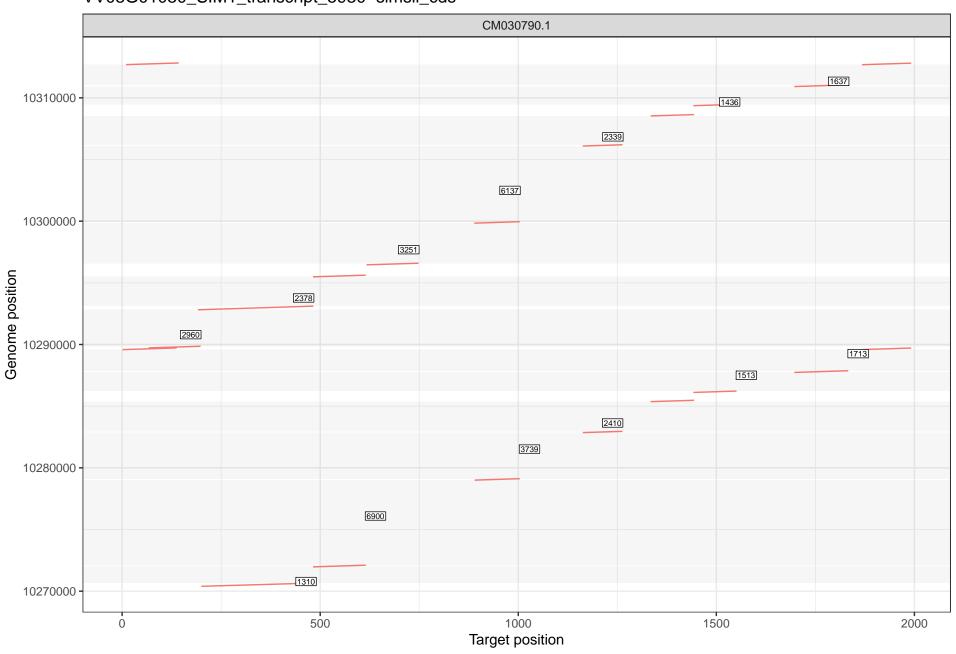
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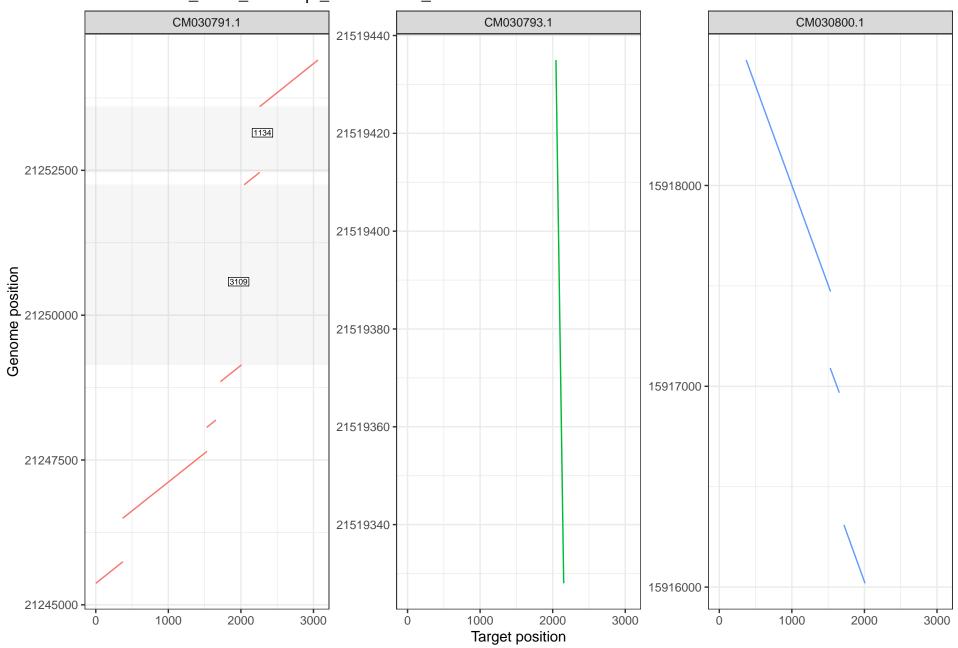
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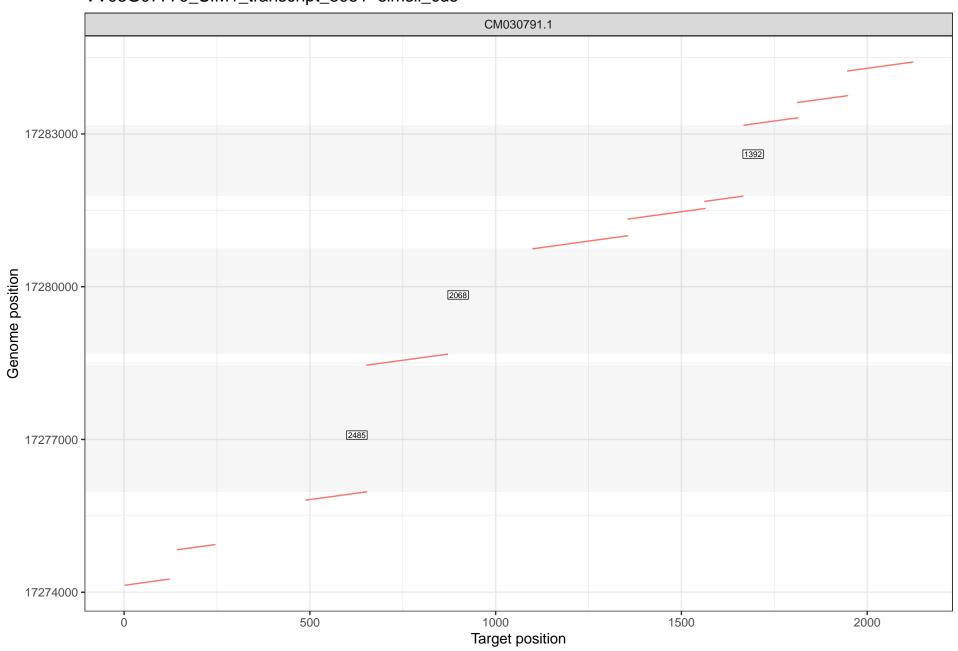


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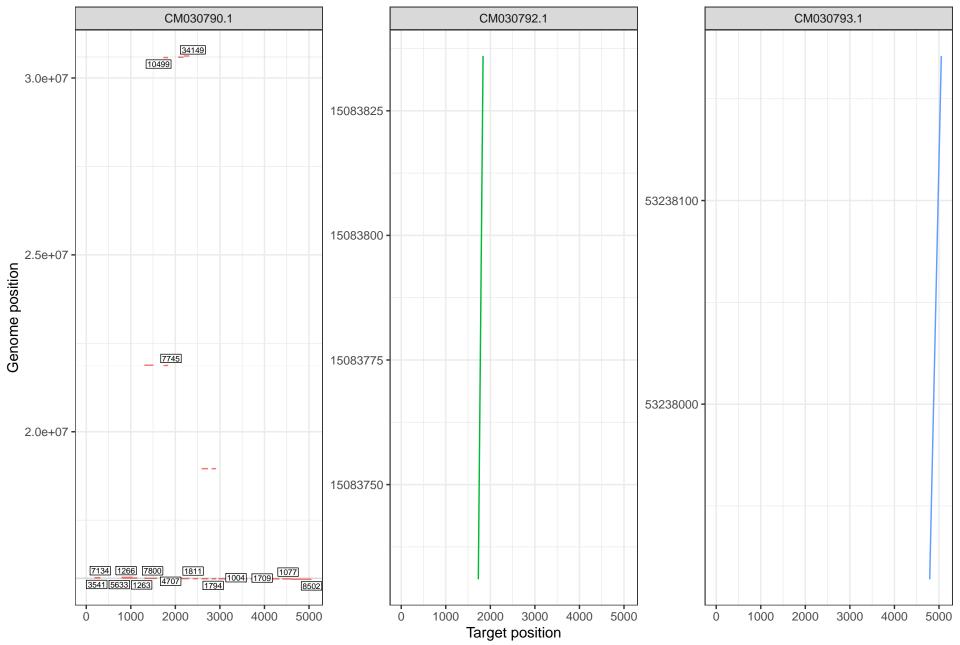


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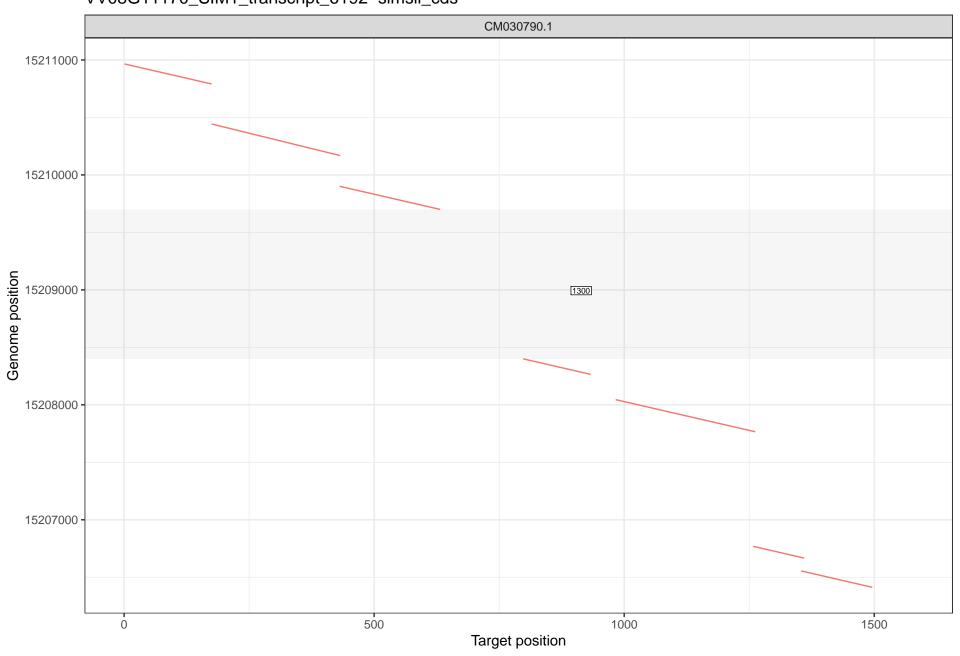
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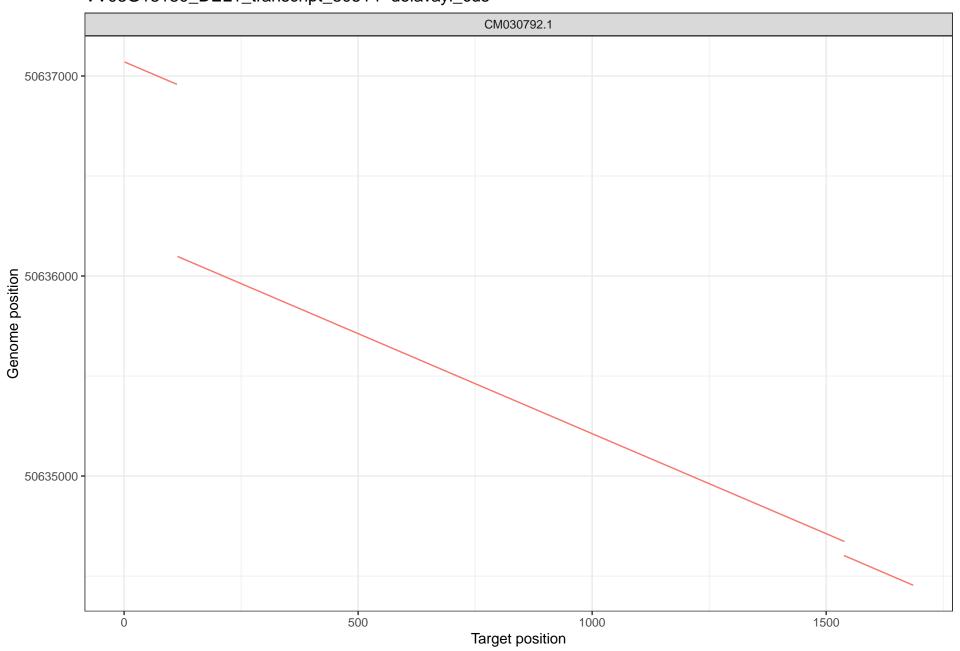
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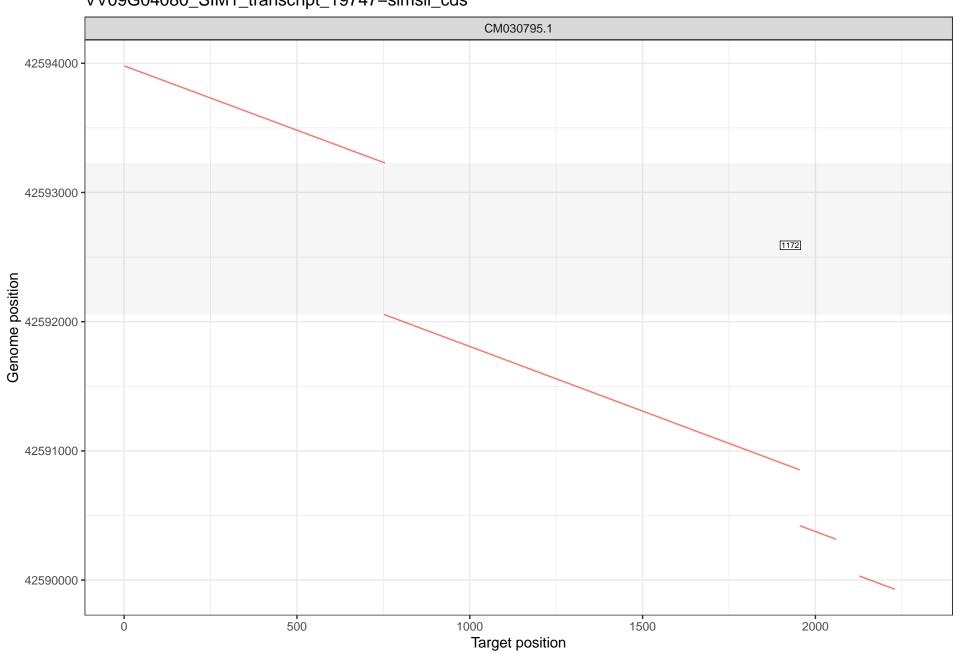


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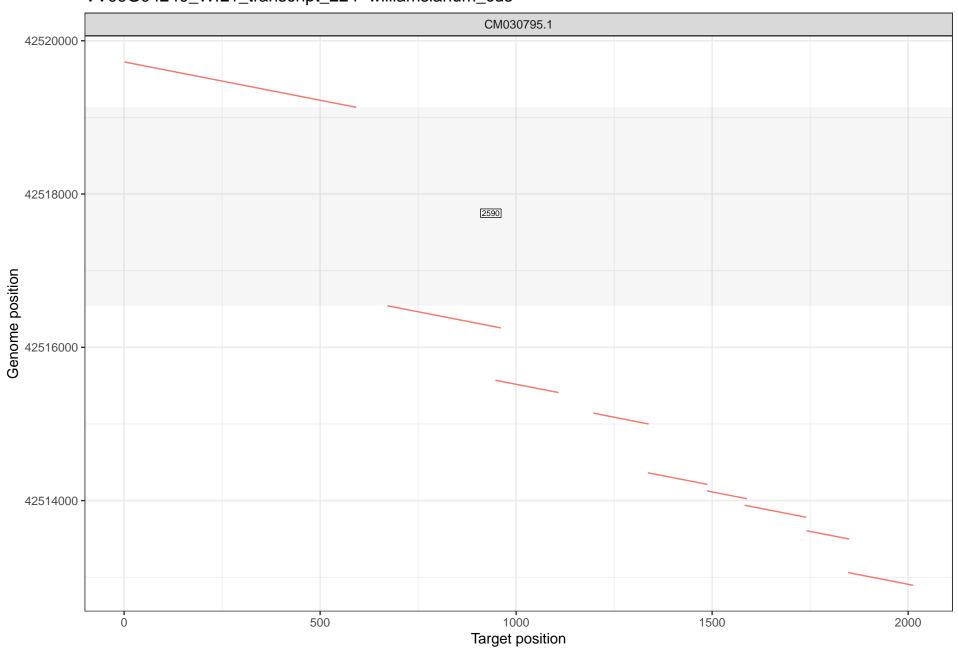


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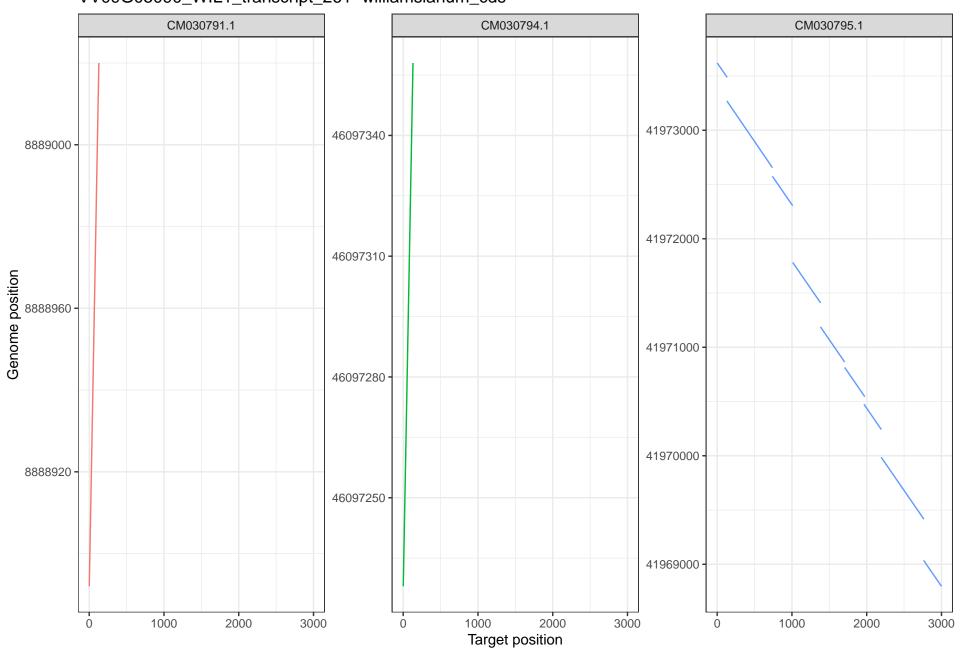
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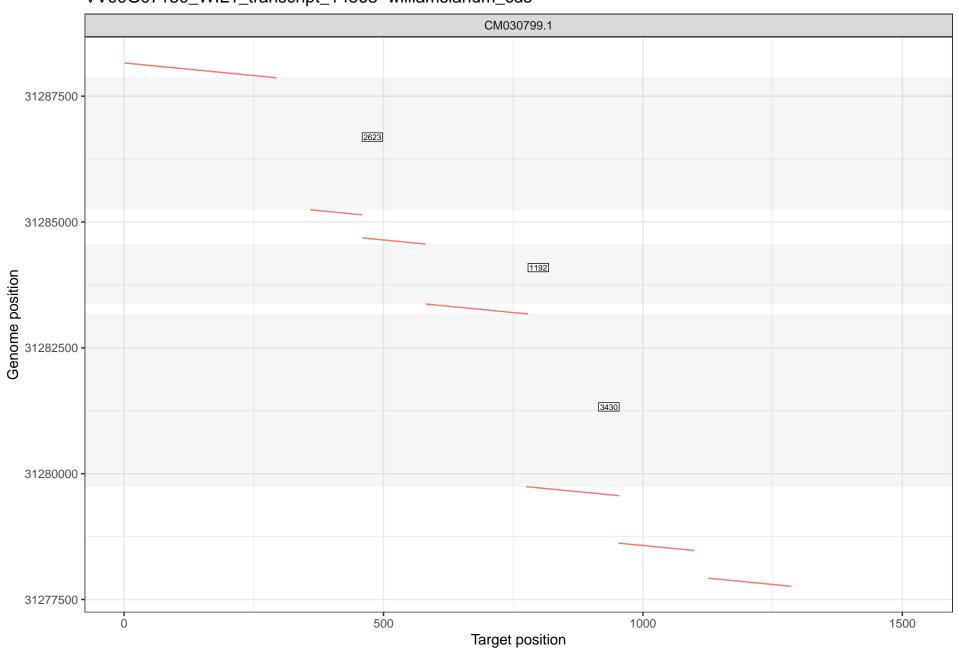
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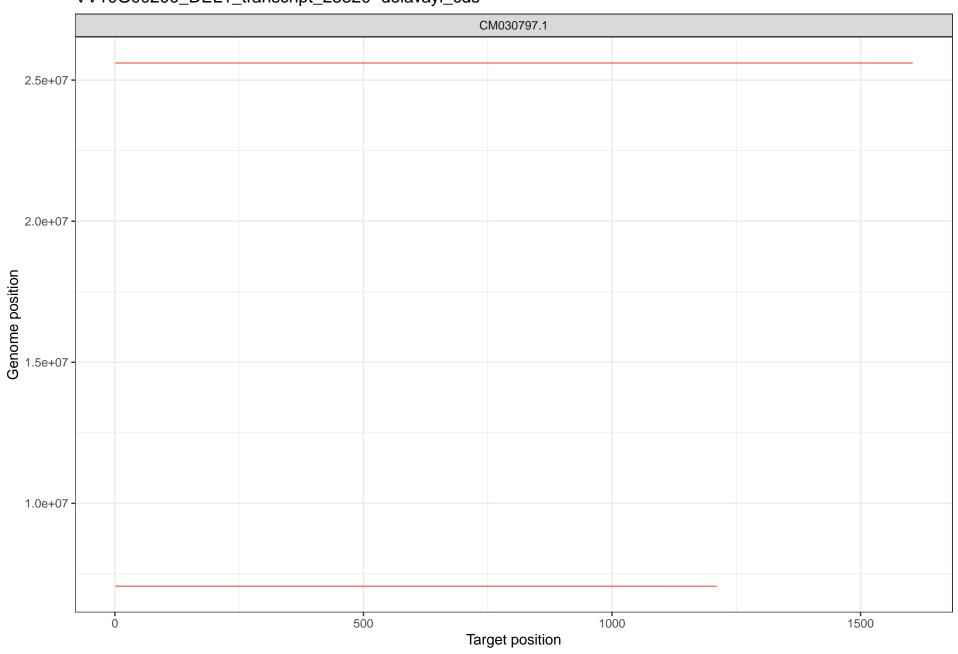
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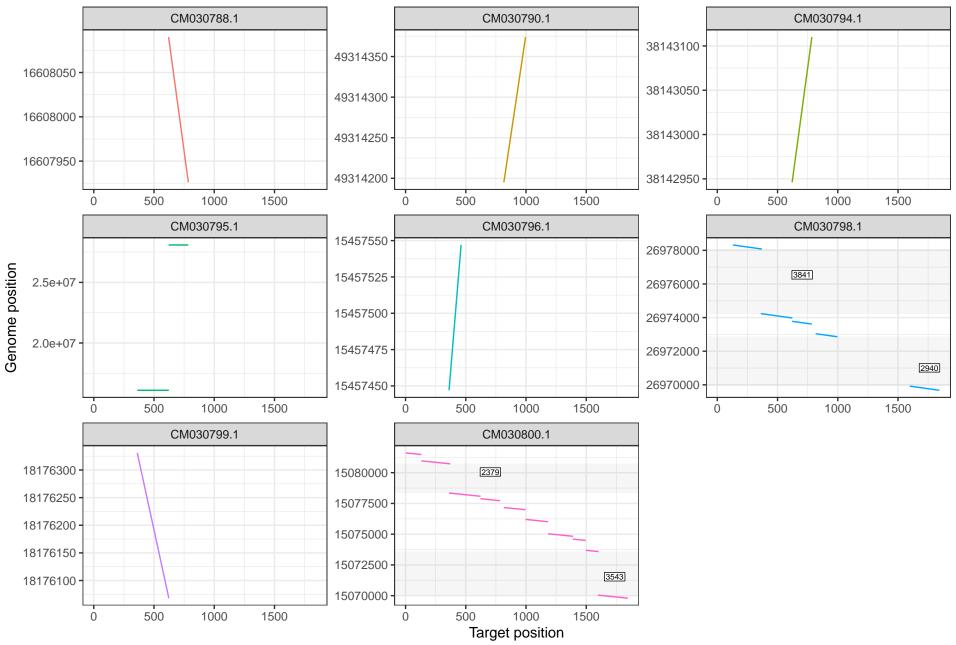
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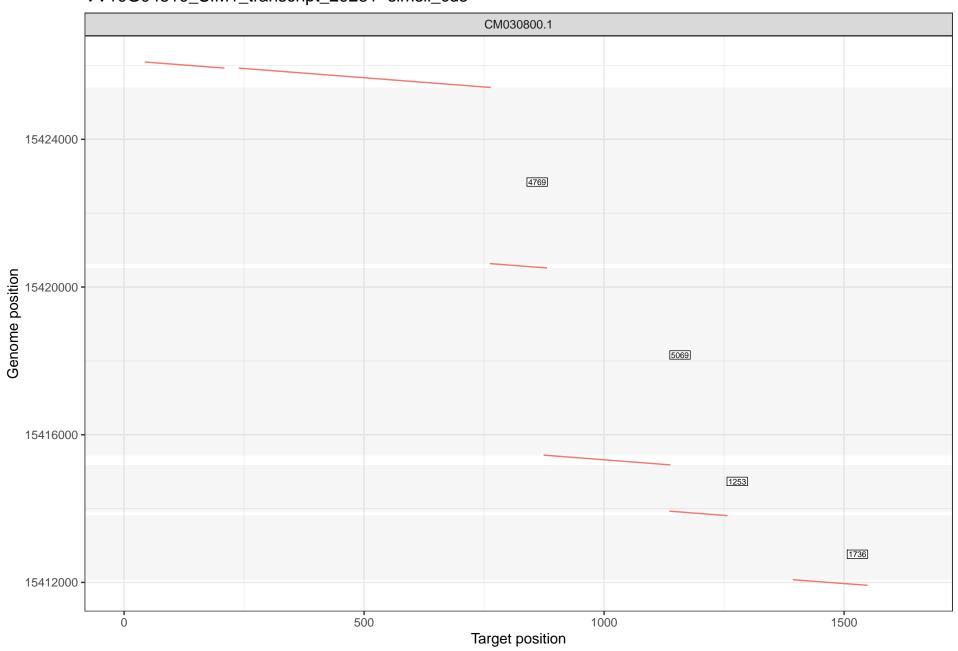
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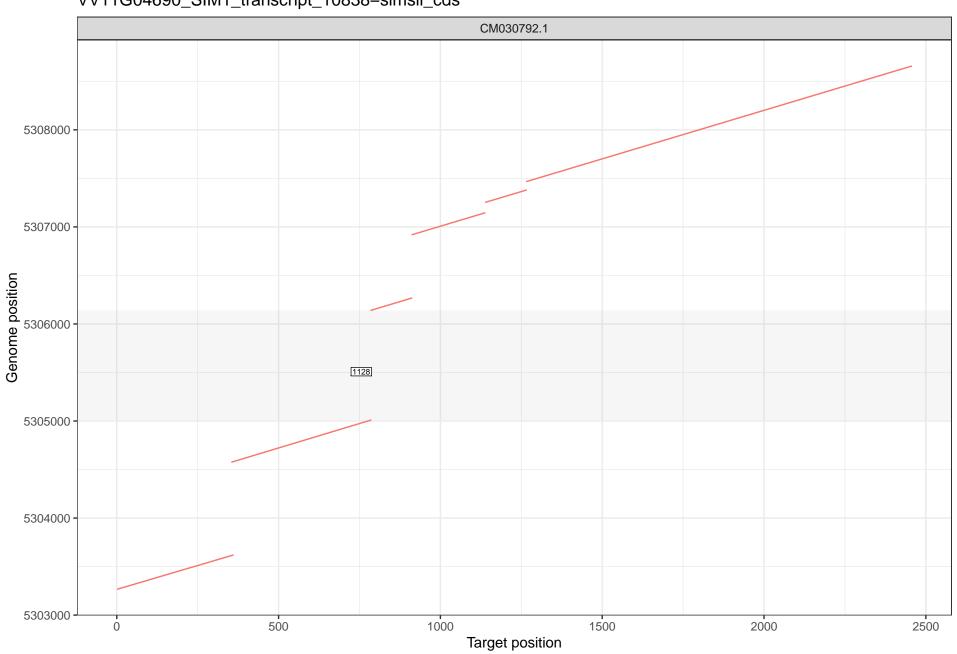
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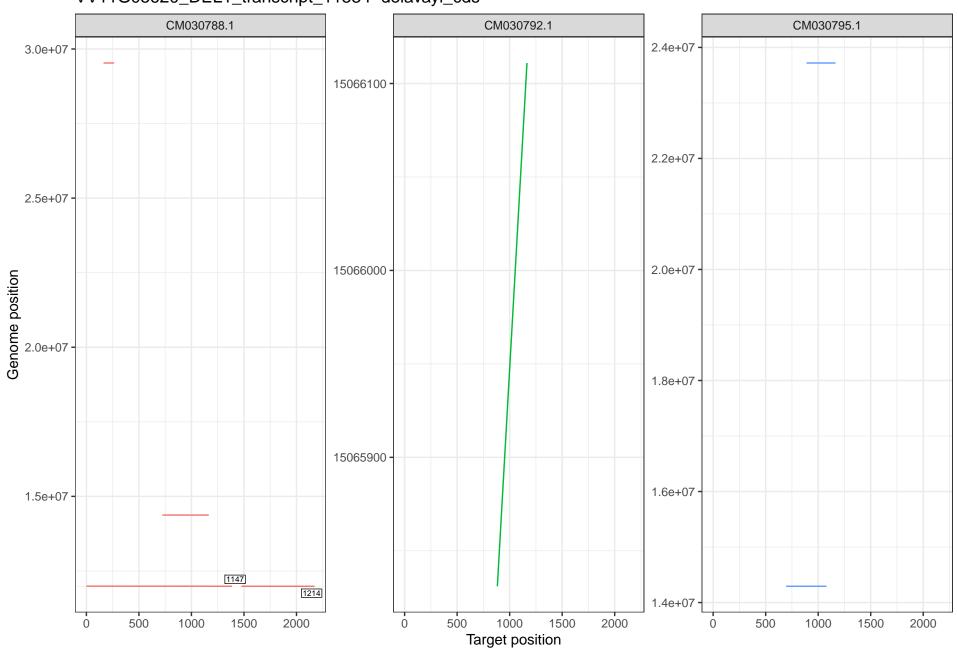
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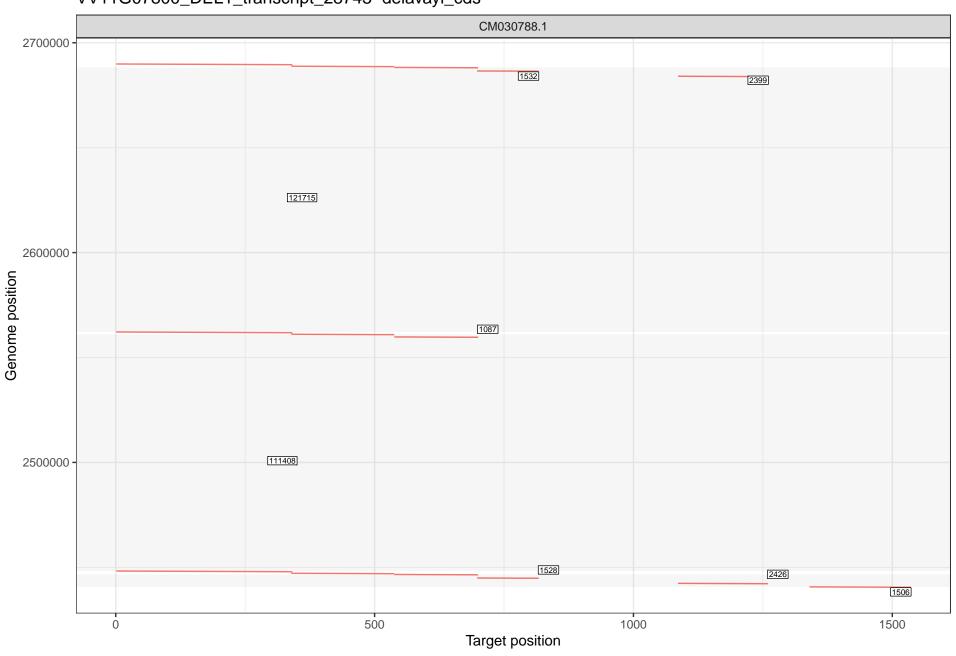
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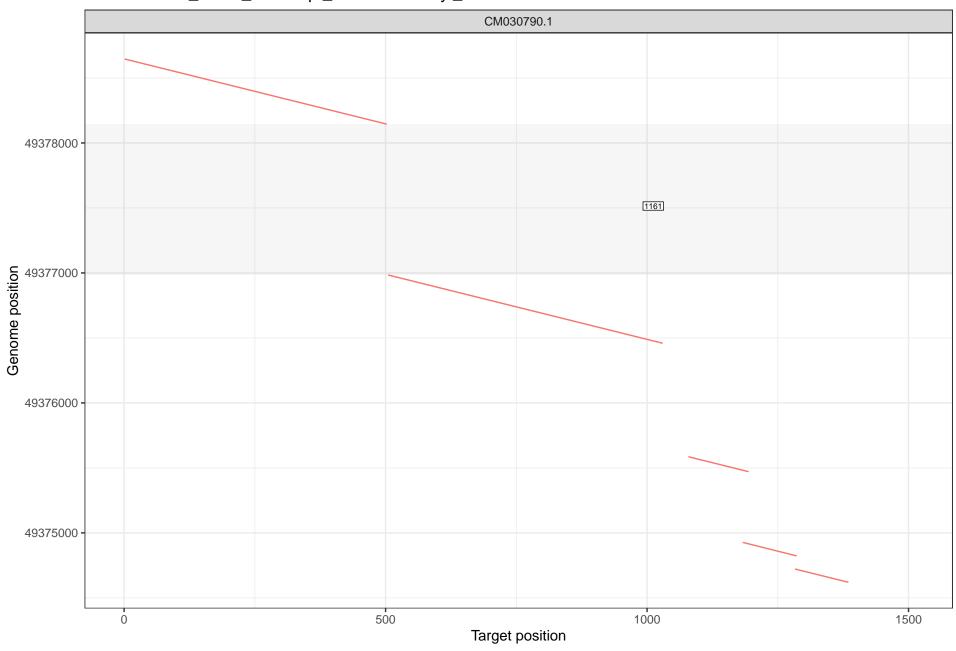


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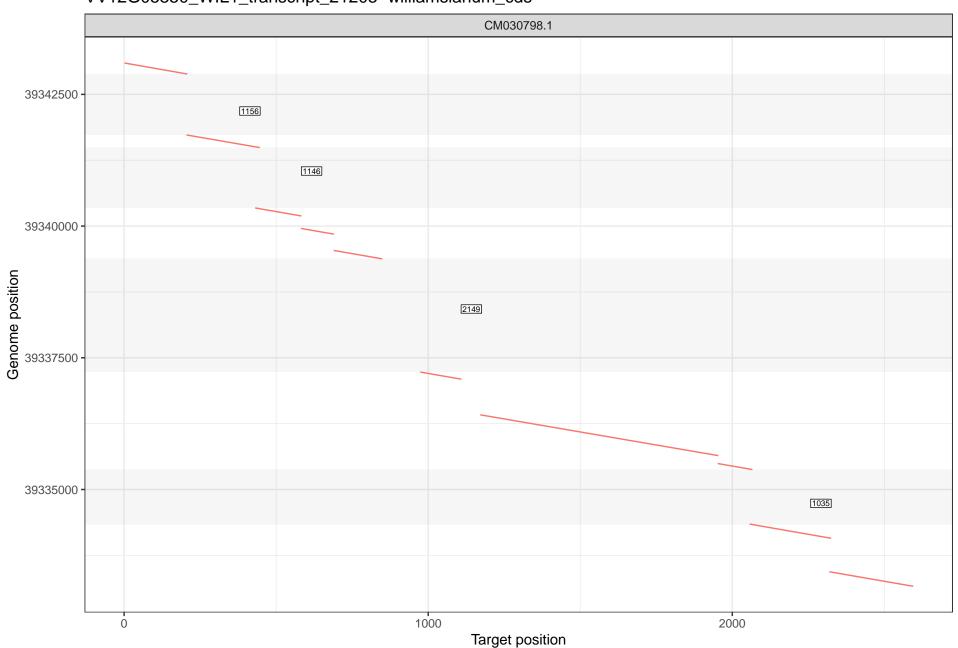


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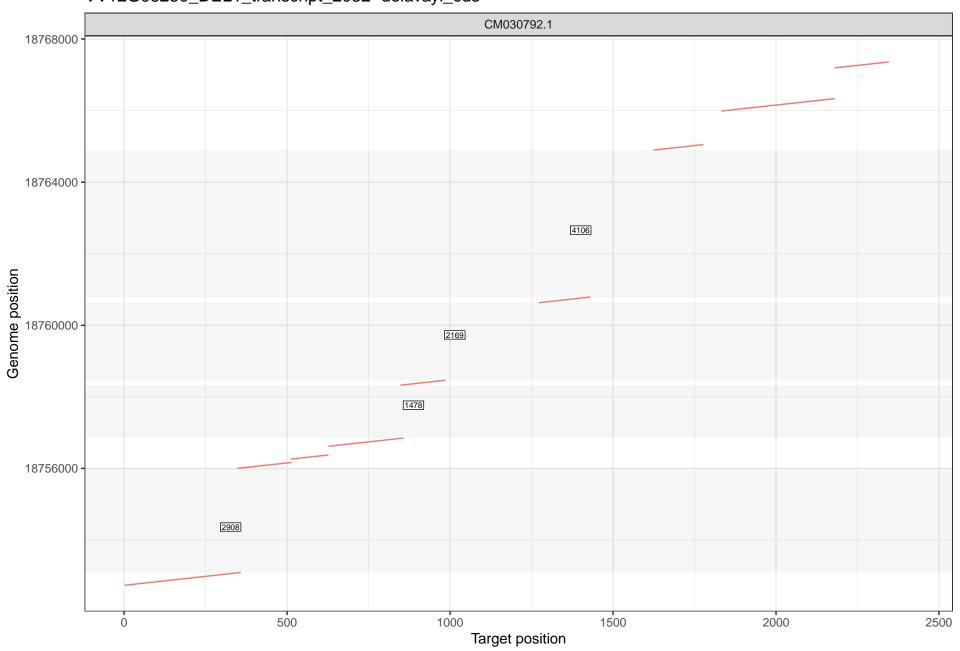
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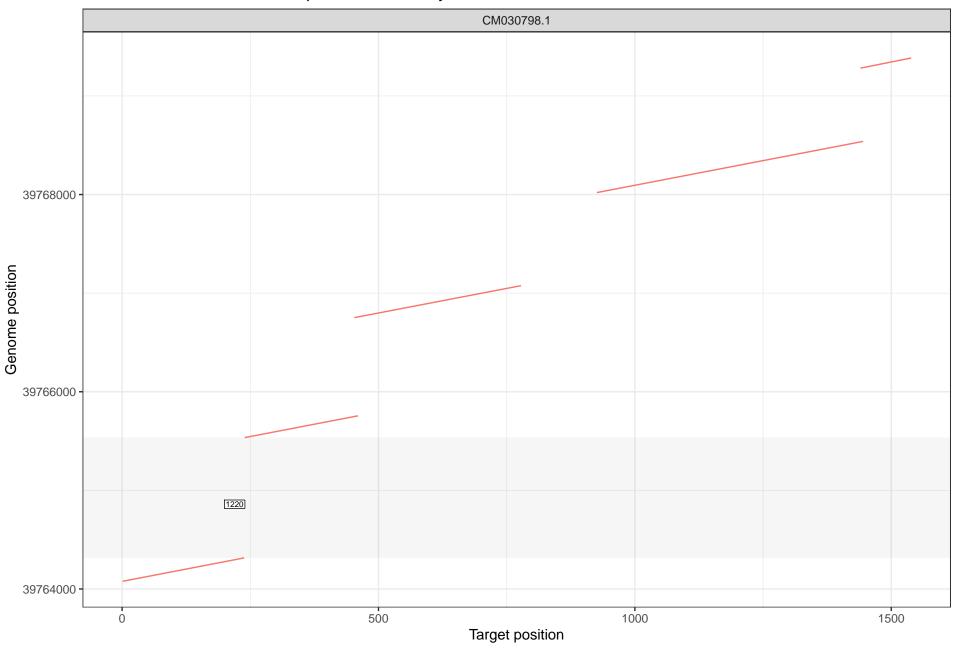
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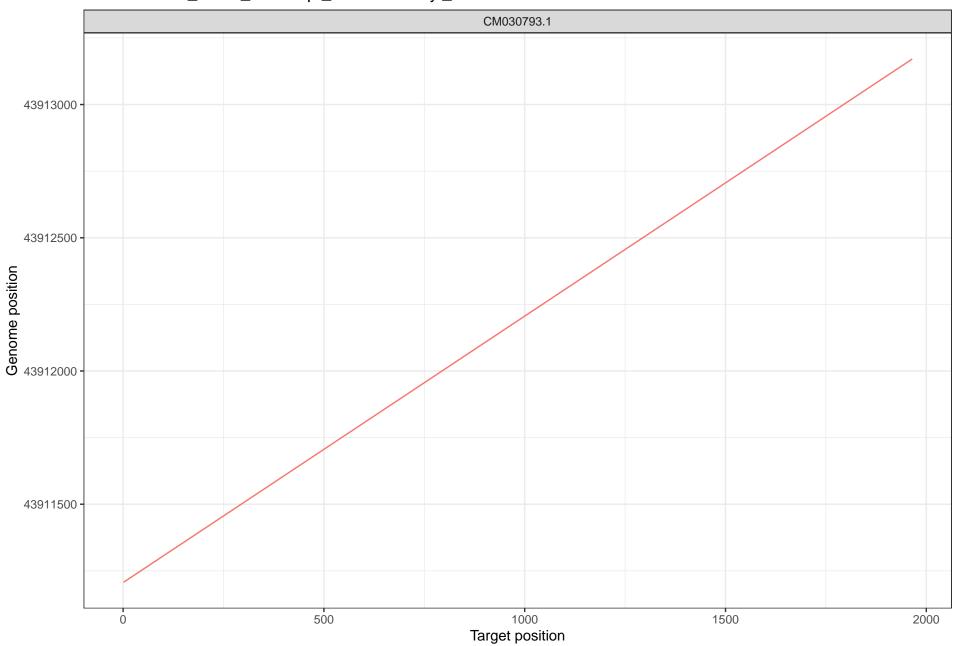
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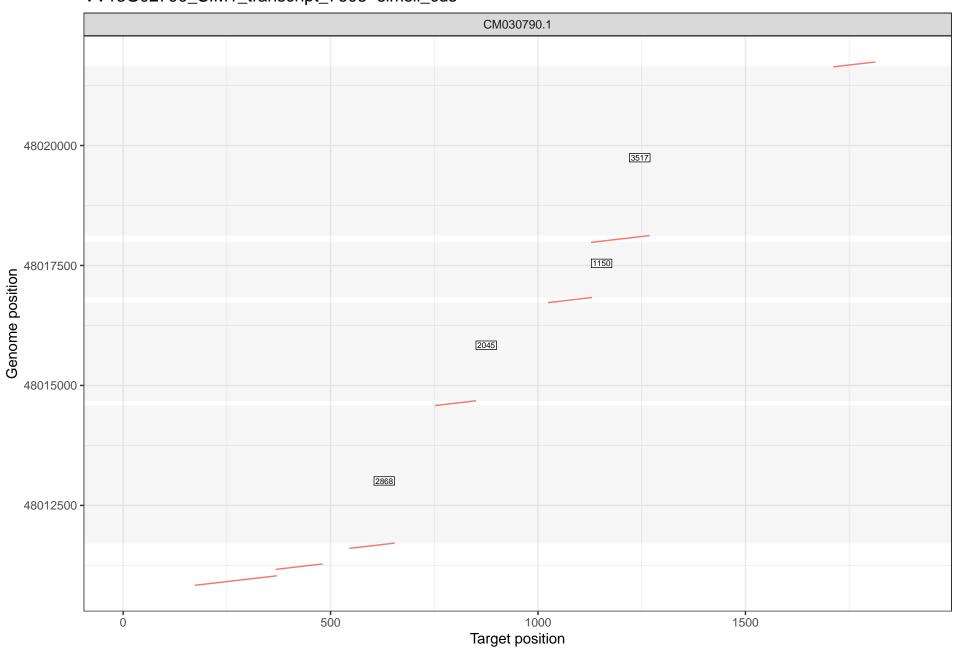
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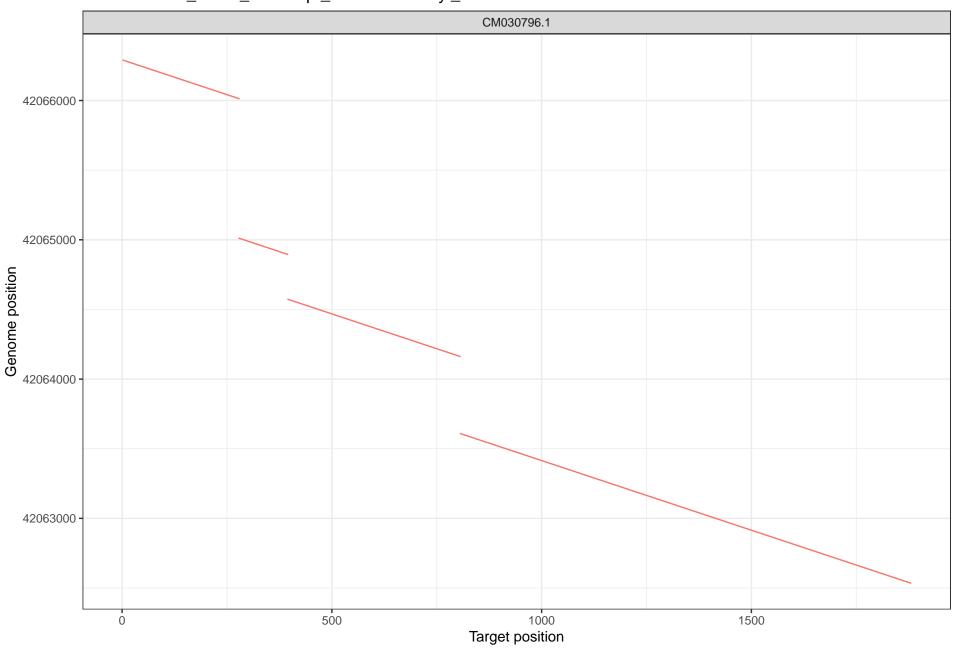
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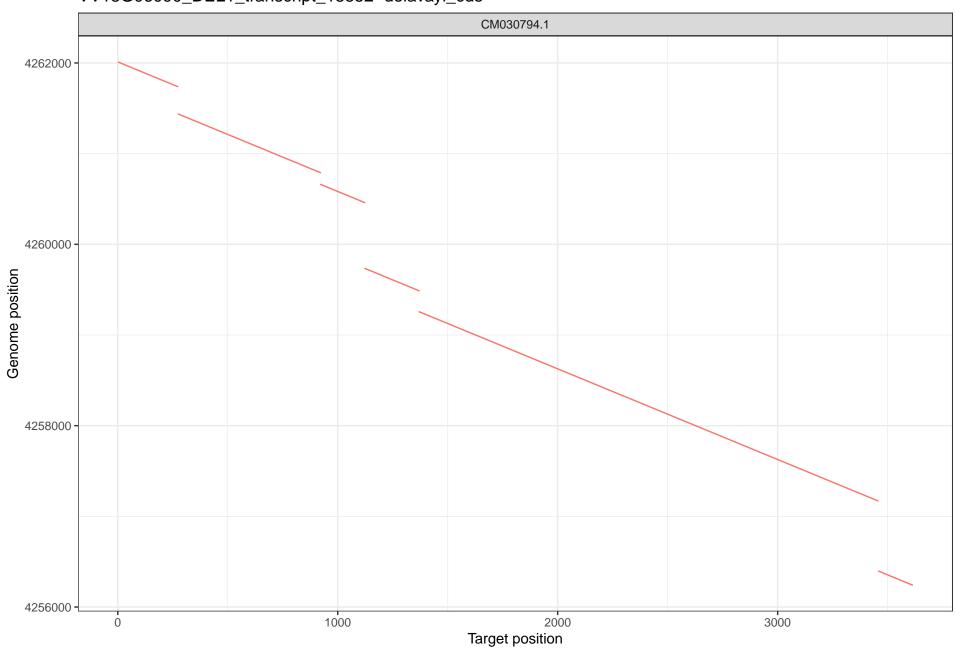
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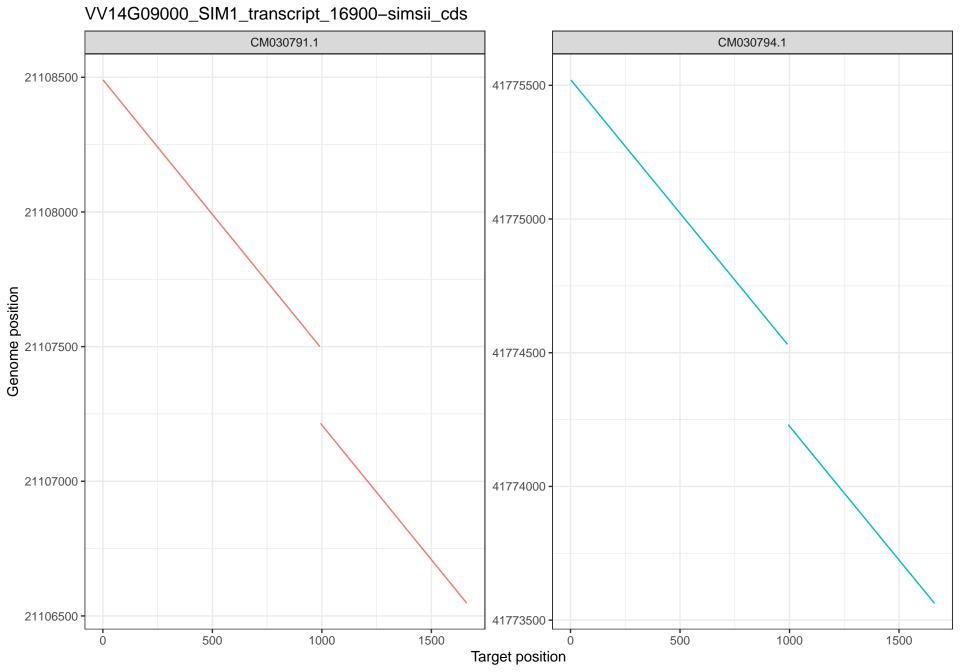


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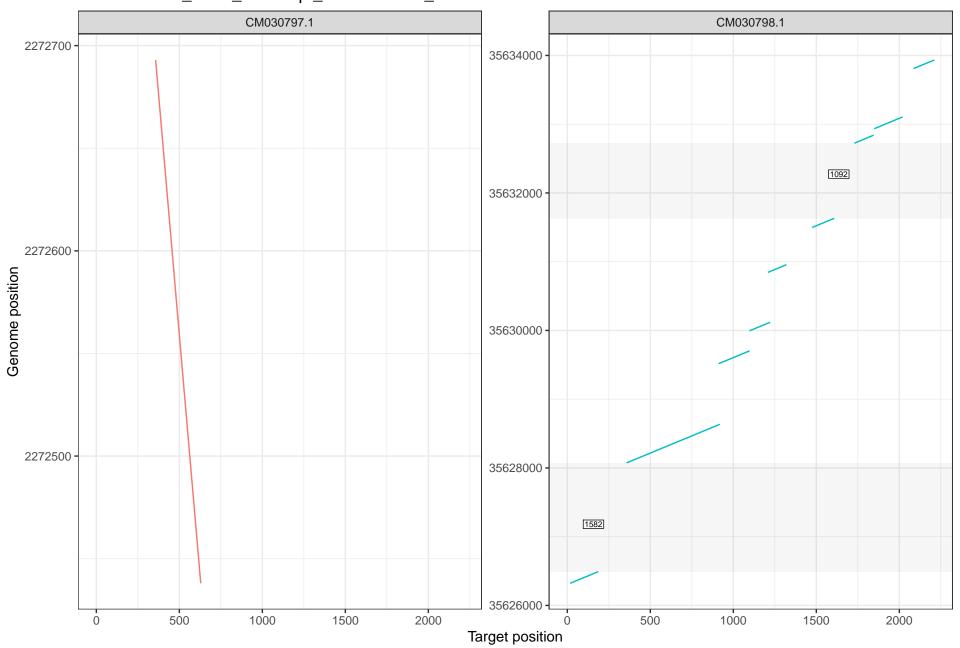


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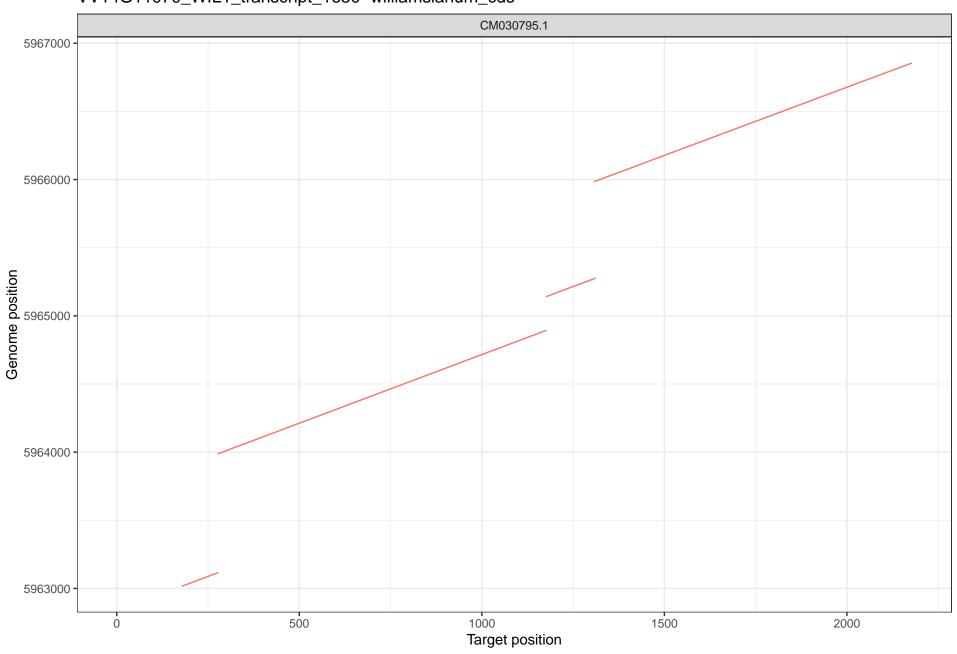
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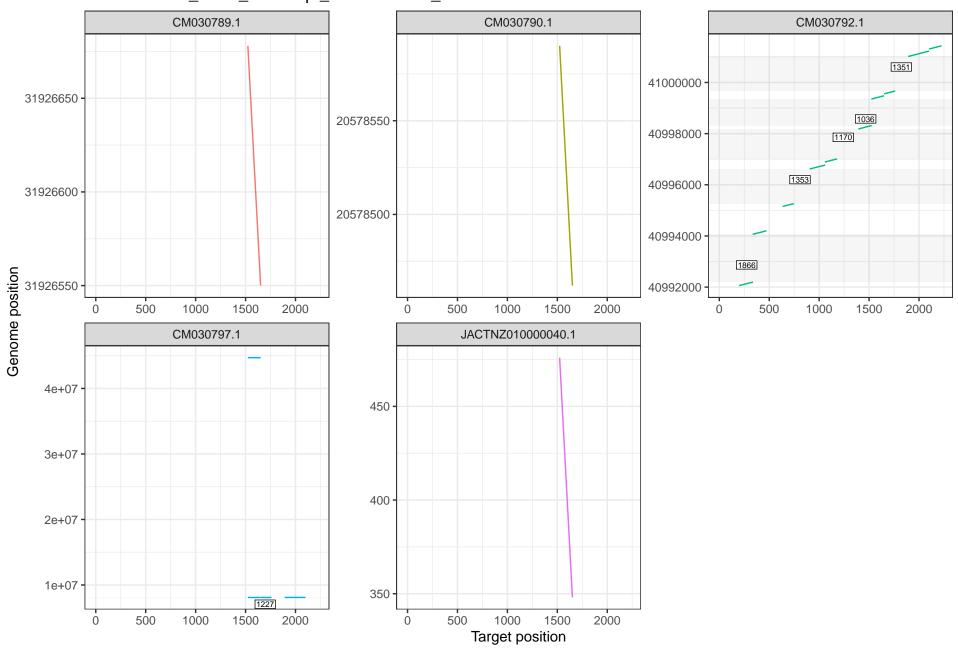
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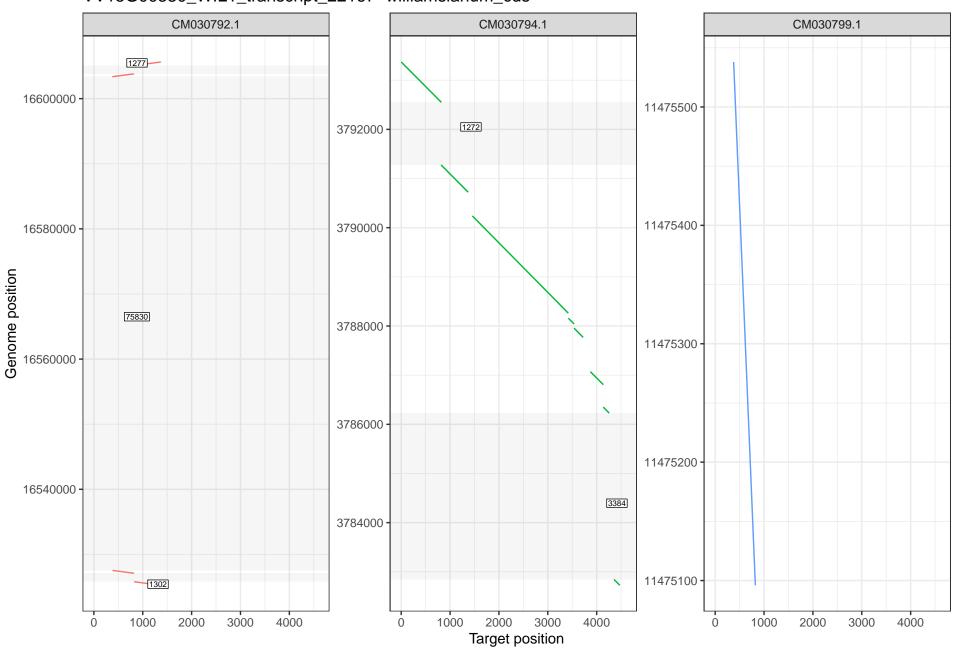
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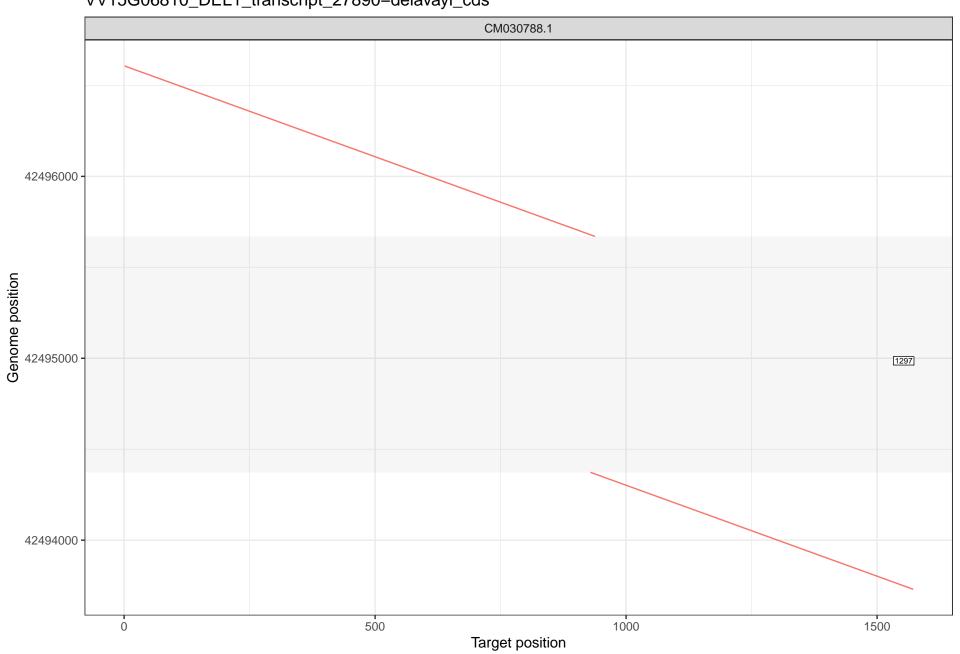
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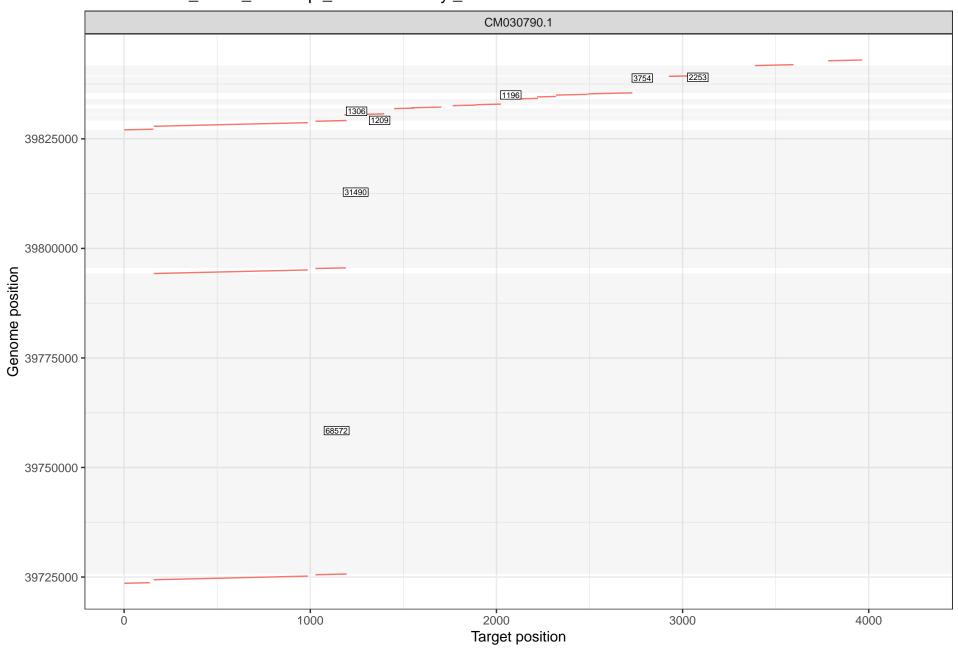
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VV16G01560\_DEL1\_transcript\_31096-delavayi\_cds



VV16G04850\_SIM1\_transcript\_22076-simsii\_cds CM030788.1 CM030789.1 CM030791.1 CM030790.1 5e+07 4e+07 6e+07 3015 3530 4e+07 -3e+07 4e+07 3e+07 4e+07 2e+07 2e+07 3e+07 2e+07 1e+07 1e+07 2e+07 0e+00 500 1000 1500 500 1000 1500 500 1000 1500 500 1000 1500 CM030792.1 CM030793.1 CM030794.1 CM030795.1 15500000 5e+07 5e+07 4e+07 15250000 -2.1e+07 4e+07 3e+07 3e+07 15000000 -1.8e+07 966879 2e+07 2e+07 14750000 Genome position 1.5e+07 1e+07 -802321 1e+07 14500000 500 1000 1500 500 1000 1500 500 1000 1500 500 1000 1500 0 CM030796.1 CM030797.1 CM030798.1 CM030799.1 5e+07 1695 7676 285740 1698 2846 4e+07 4e+07 -4e+07 3e+07 -2e+07 3e+07 3e+07 2e+07 2e+07 3354 2e+07 1e+07 1e+07 -1e+07 -0e+00 0e+00 955741 500 500 500 1000 1500 1000 1500 1000 1500 500 1000 1500 0 CM030800.1 JACTNZ010000036.1 3.2e + 071406350 -3.0e+07 1406300 2.8e+07 1406250 -2.6e+07 1406200 -

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0

1000

1500

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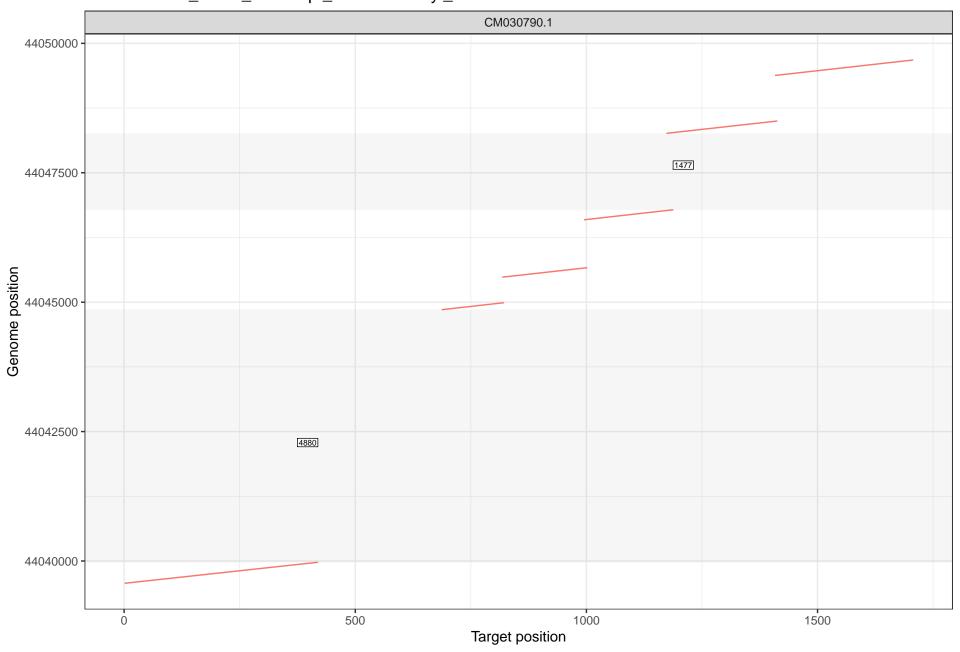
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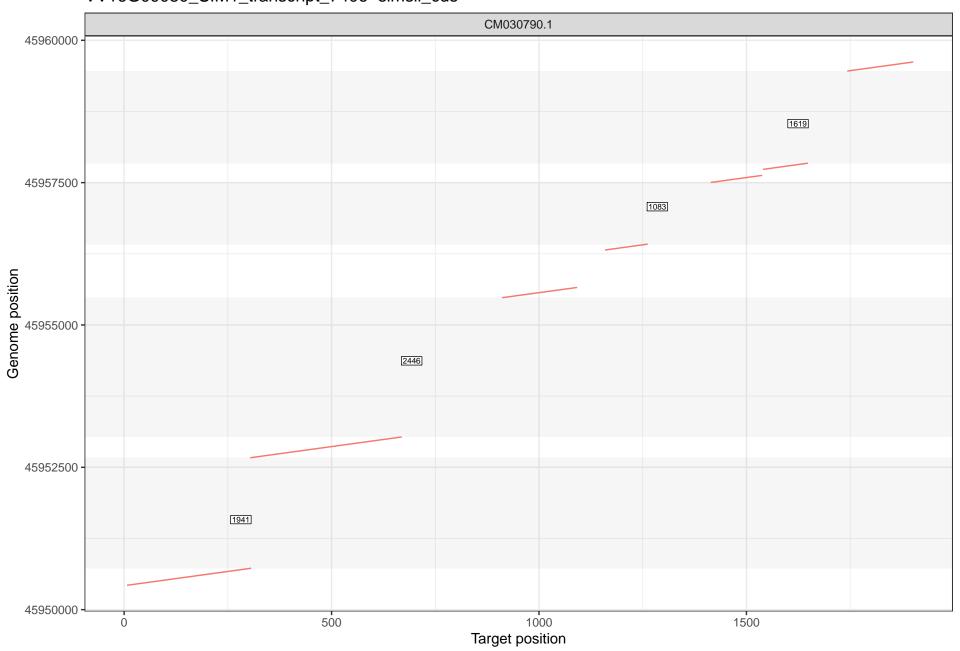
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Target position

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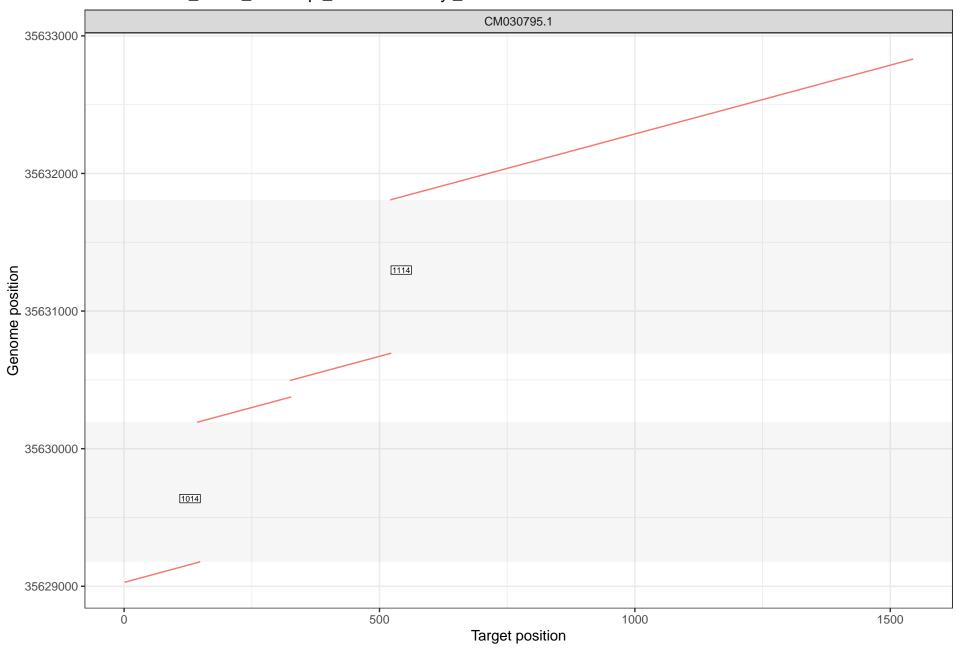


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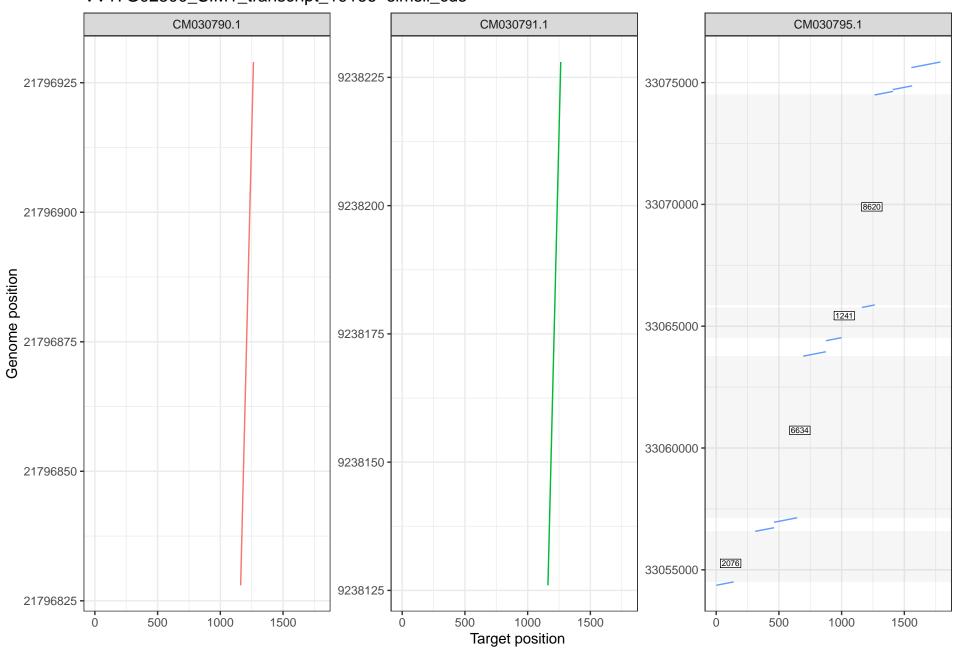
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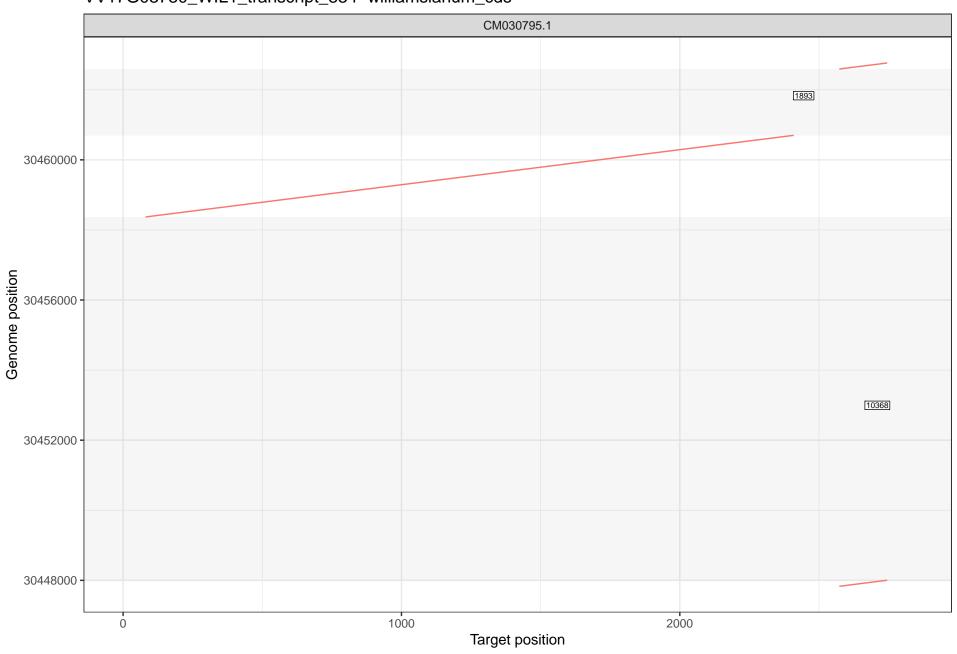


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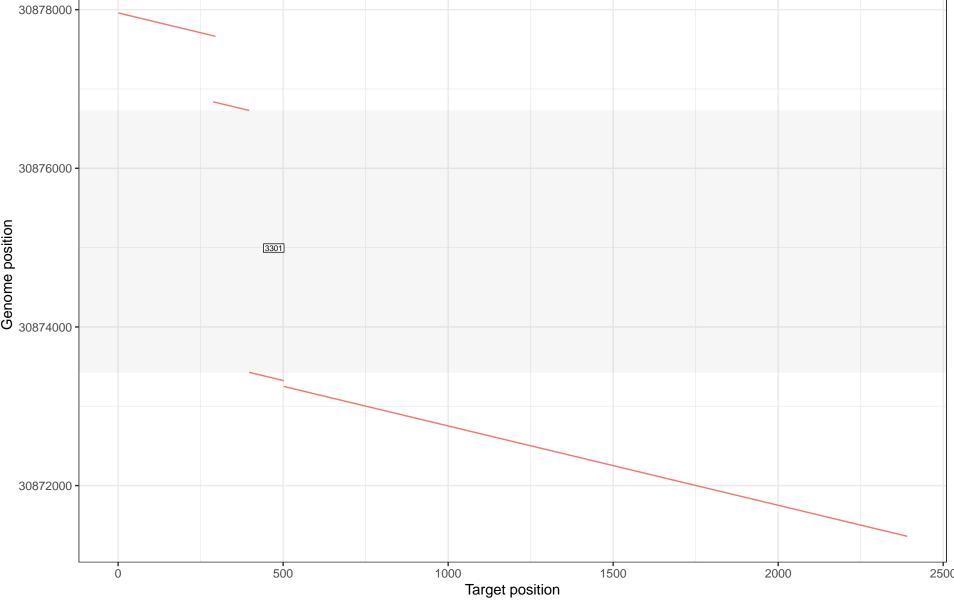
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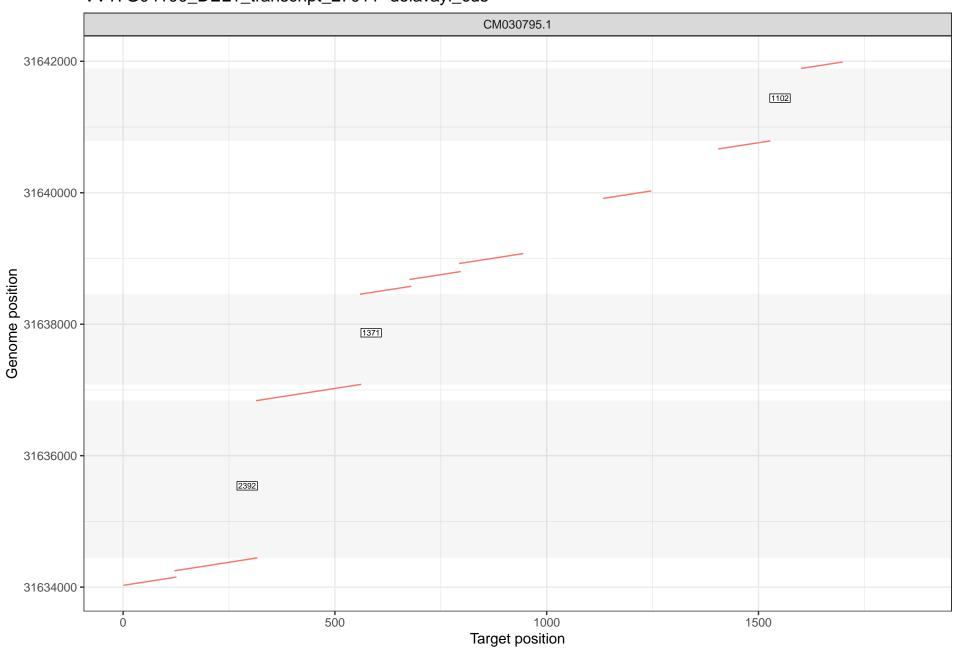
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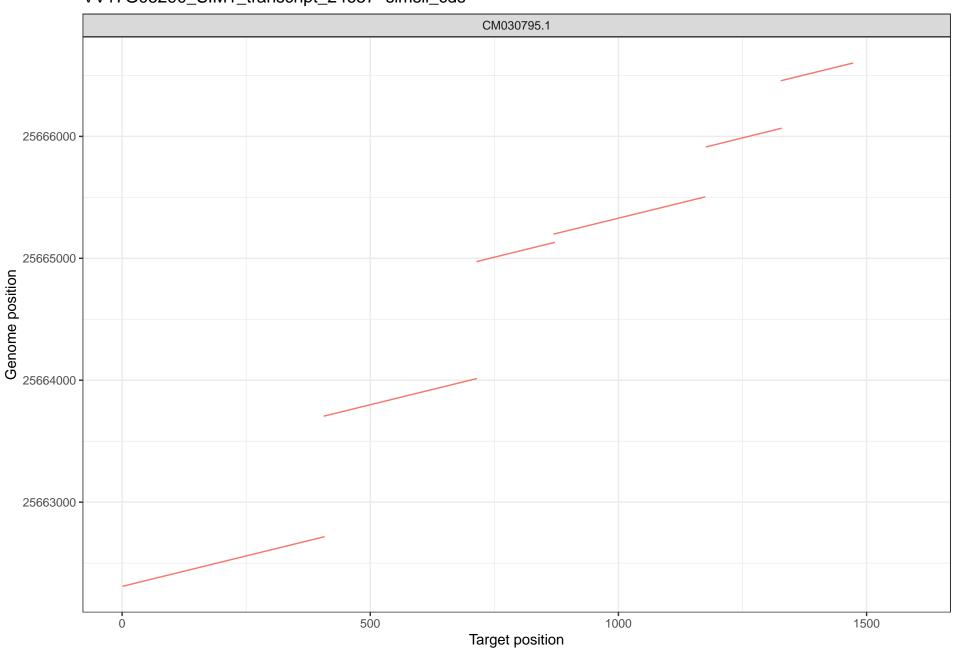
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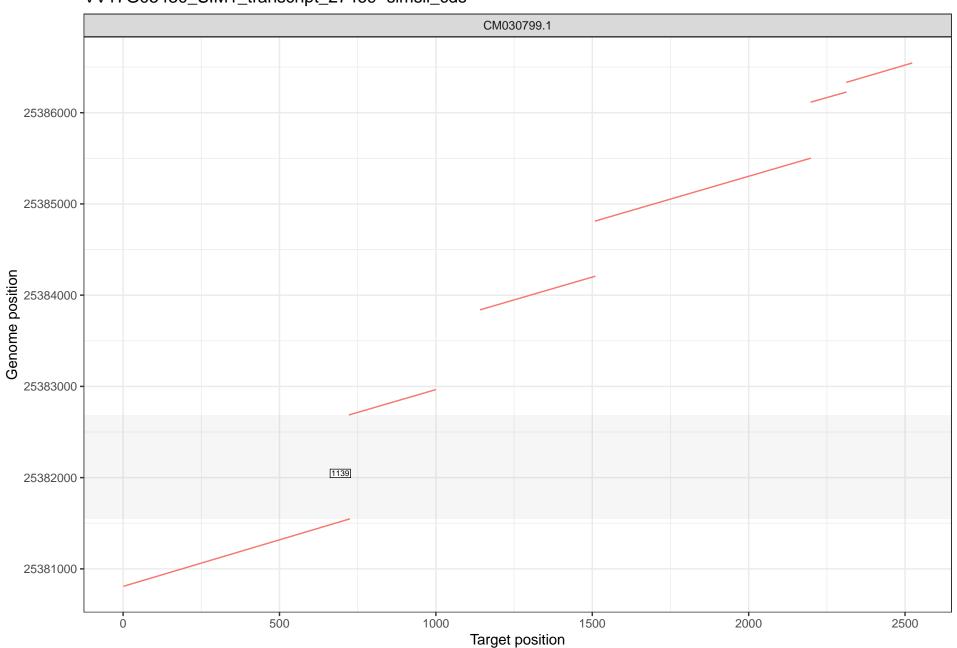
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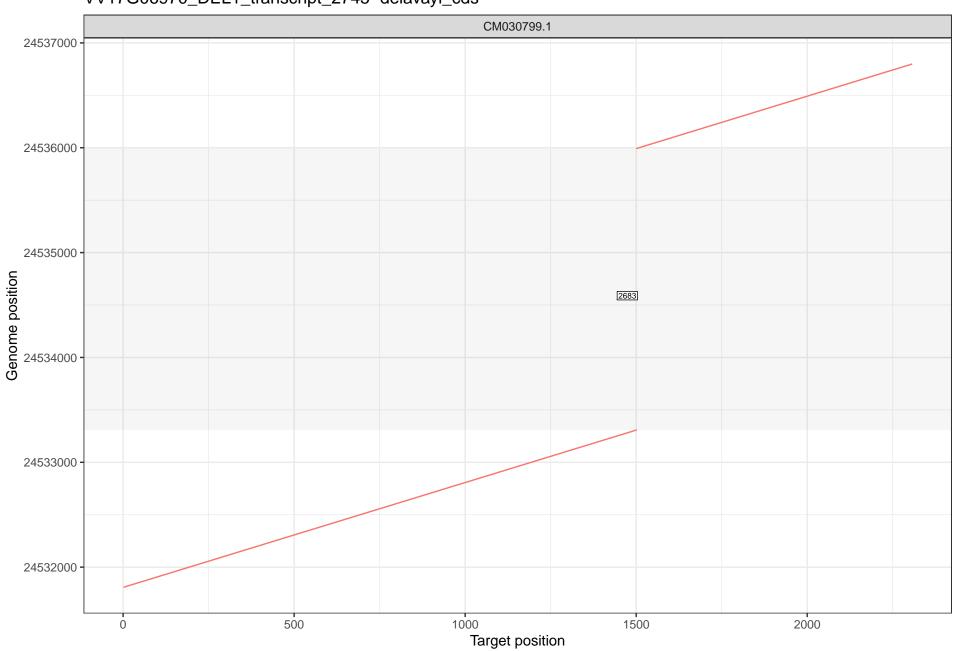
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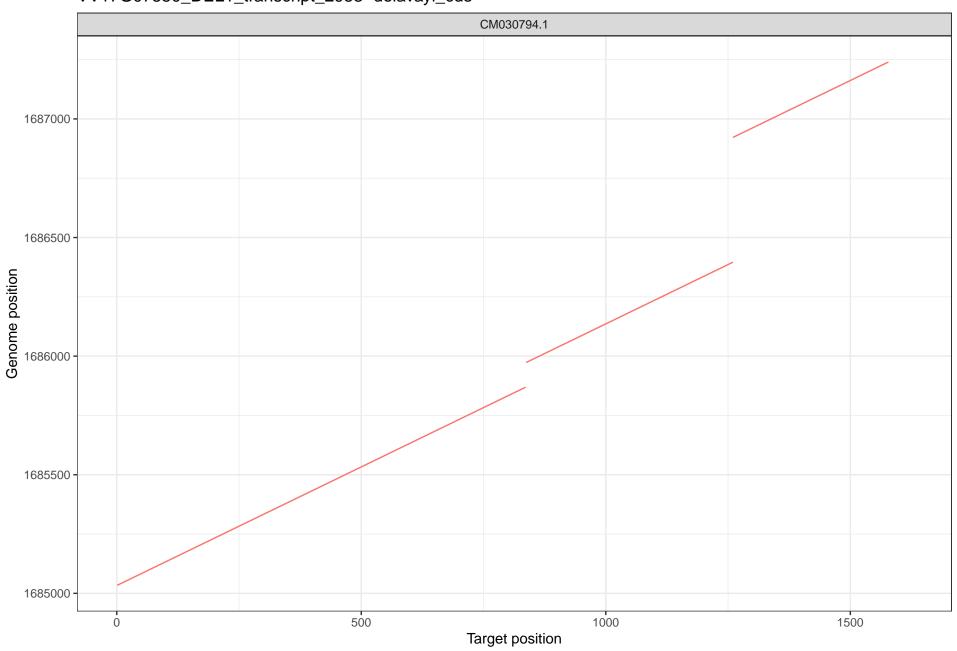
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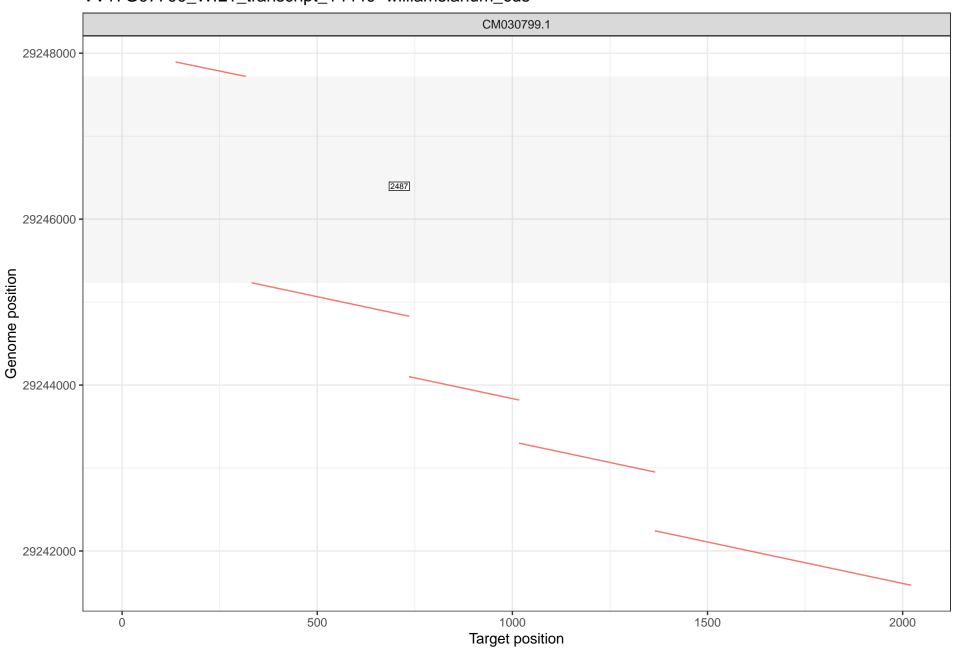
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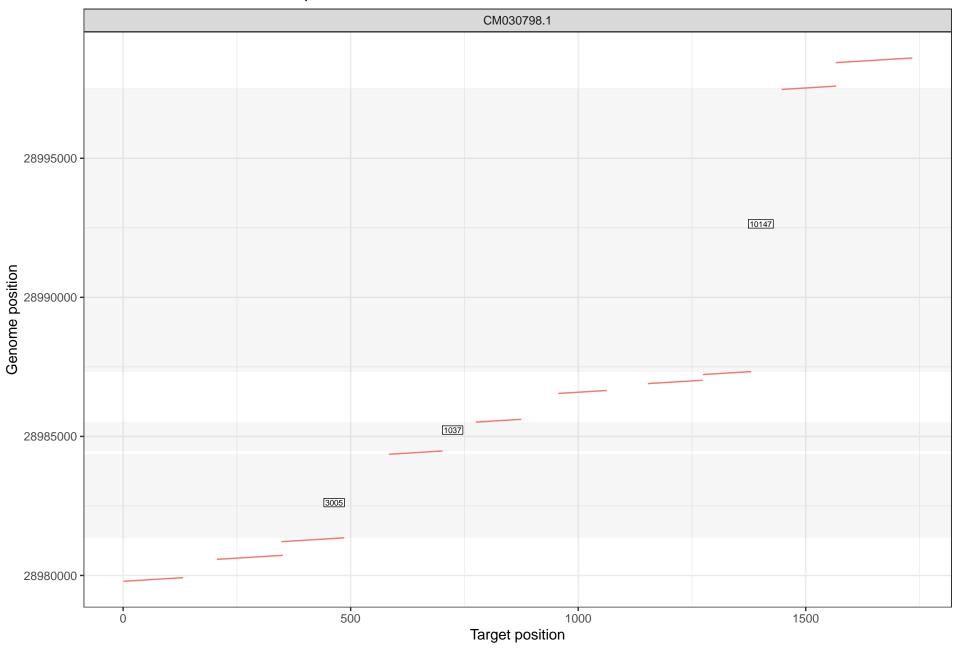
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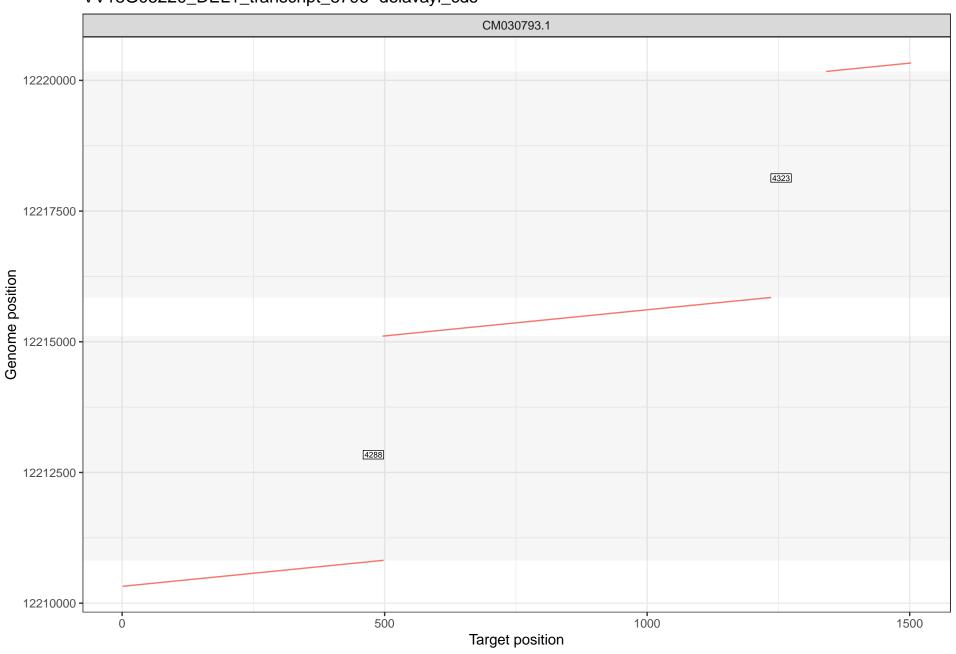
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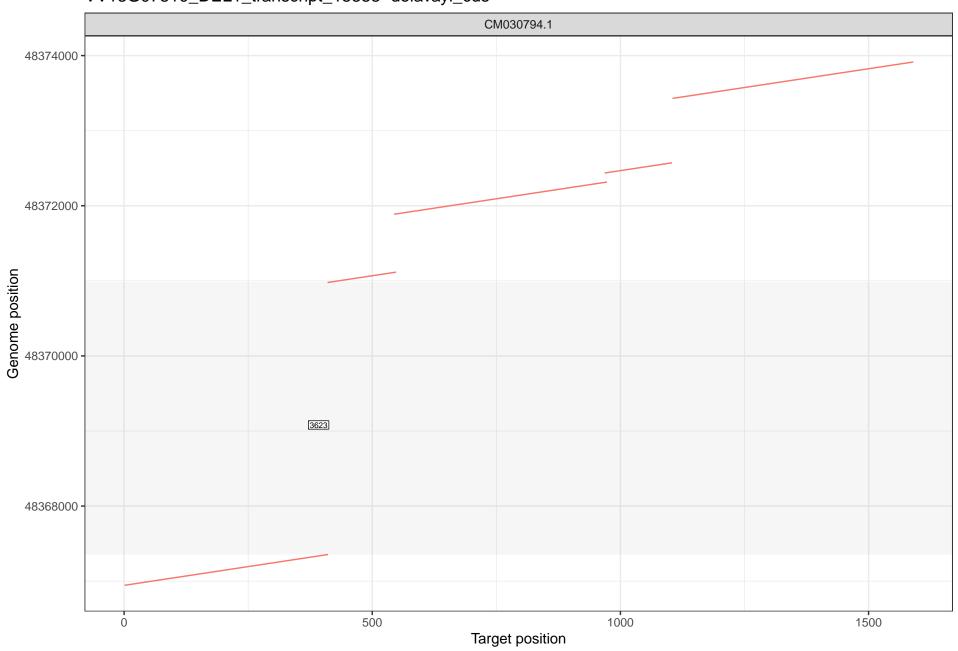
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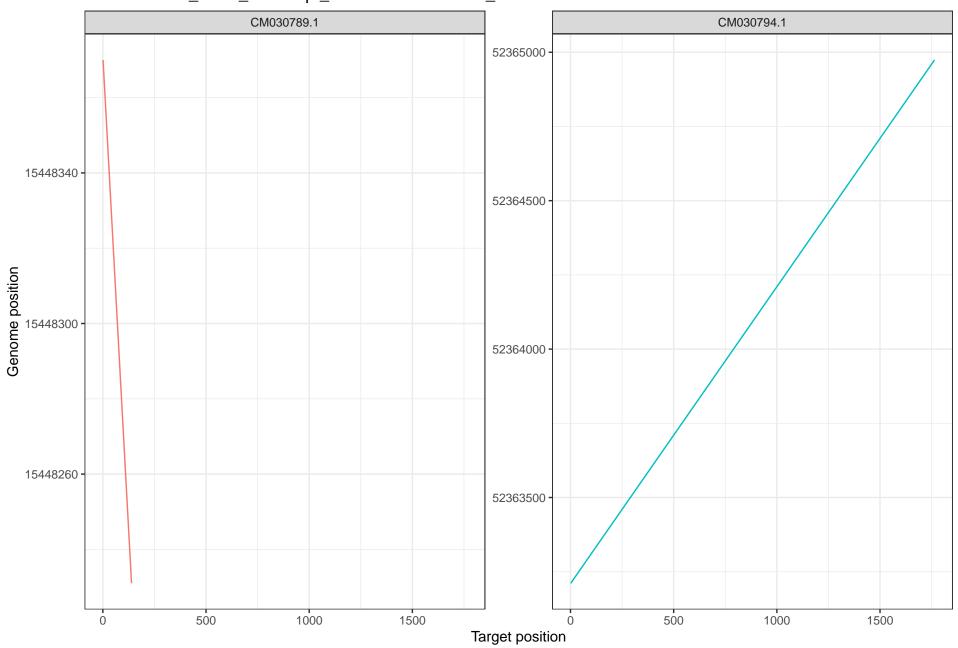


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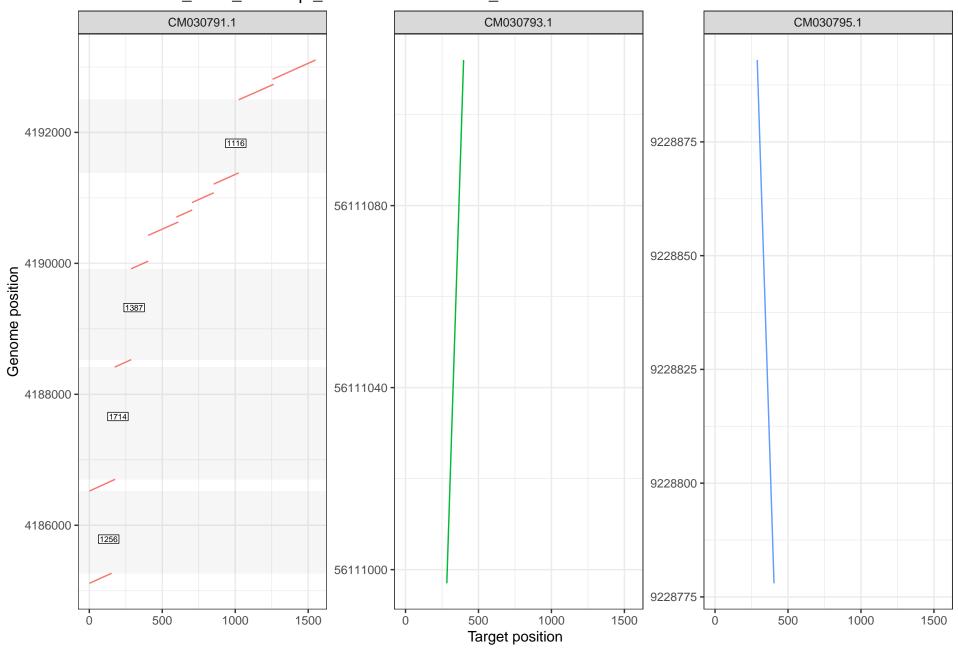


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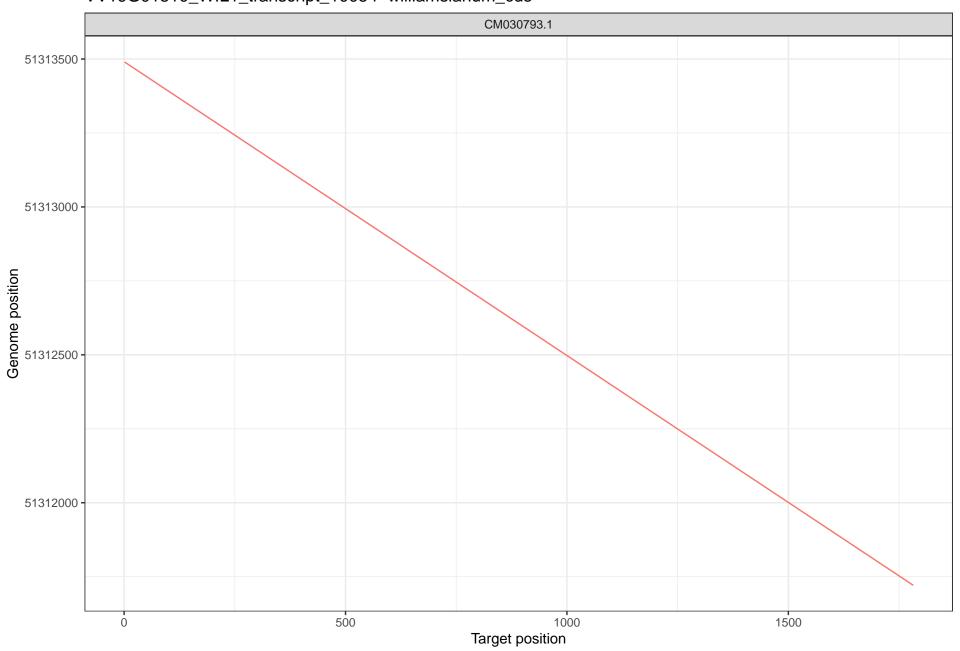
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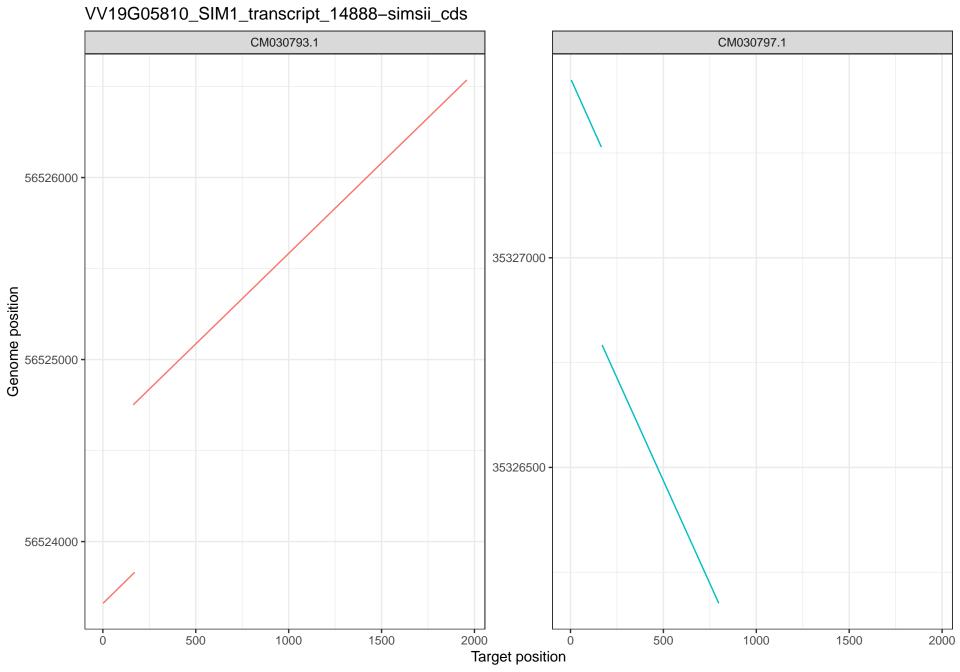
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VV19G06040\_DEL1\_transcript\_3202-delavayi\_cds

