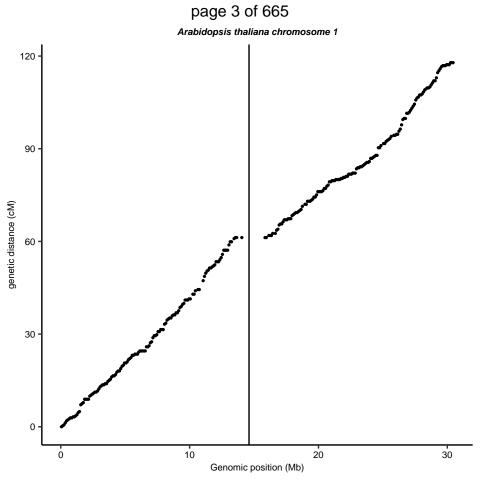
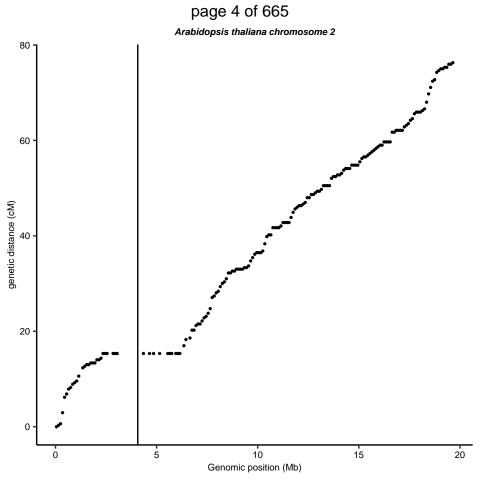
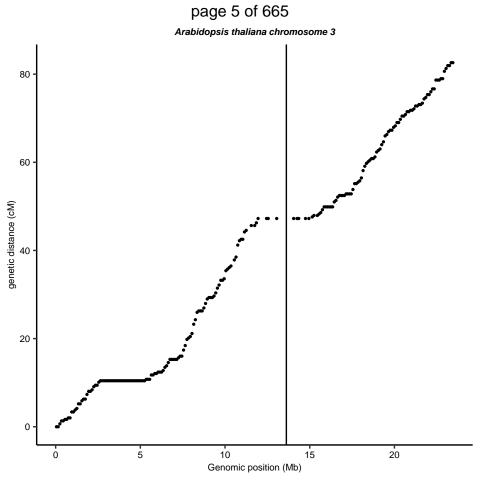
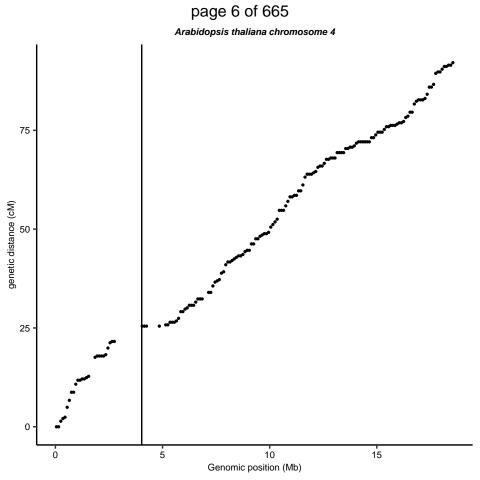
page 1 of 665 Aegilops speltoides chromosome 2S genetic distance (cM) Genomic position (Mb)

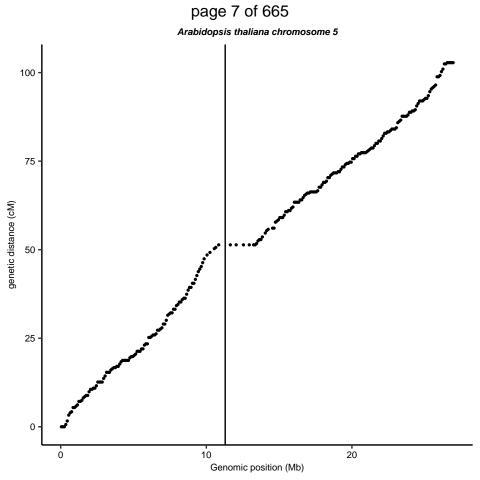
page 2 of 665 Aegilops speltoides chromosome 7S genetic distance (cM) Genomic position (Mb)





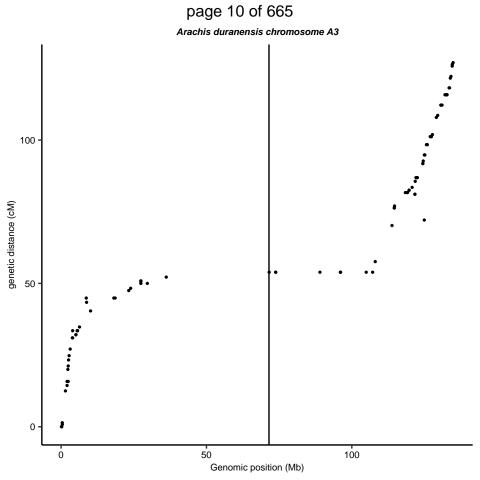






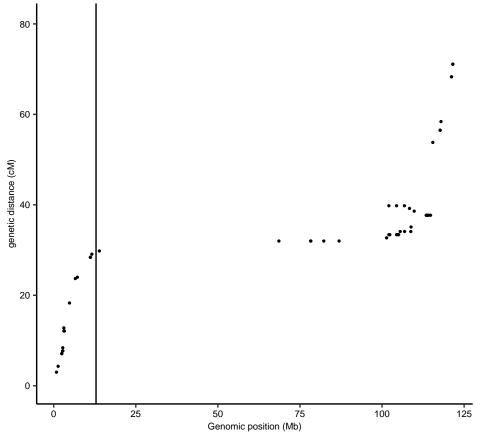
page 8 of 665 Arachis duranensis chromosome A1 genetic distance (cM) Genomic position (Mb)

page 9 of 665 Arachis duranensis chromosome A2 80 60 genetic distance (cM) 20 0 25 75 50 ö Genomic position (Mb)



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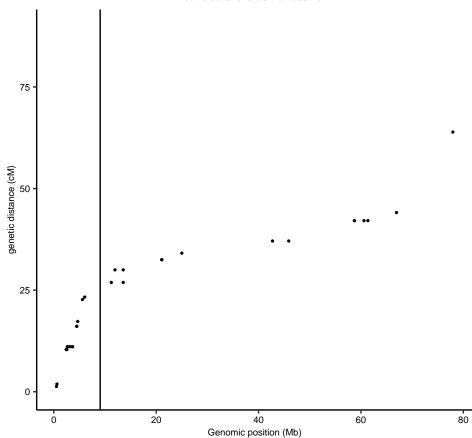
Arachis duranensis chromosome A4



page 12 of 665 Arachis duranensis chromosome A5 100 genetic distance (cM) 50 -30 60 90 Genomic position (Mb)

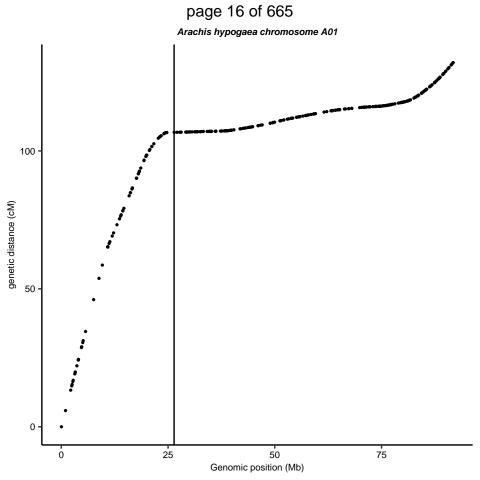
page 13 of 665

Arachis duranensis chromosome A7

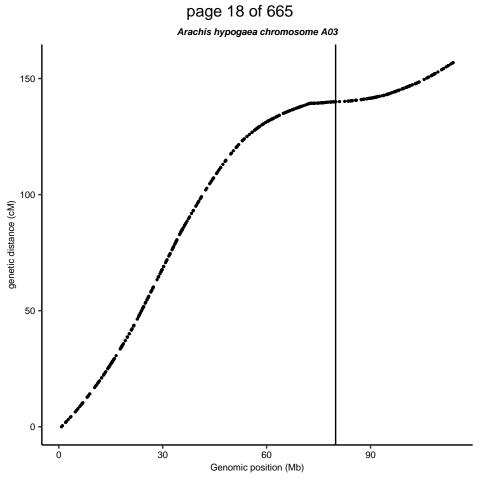


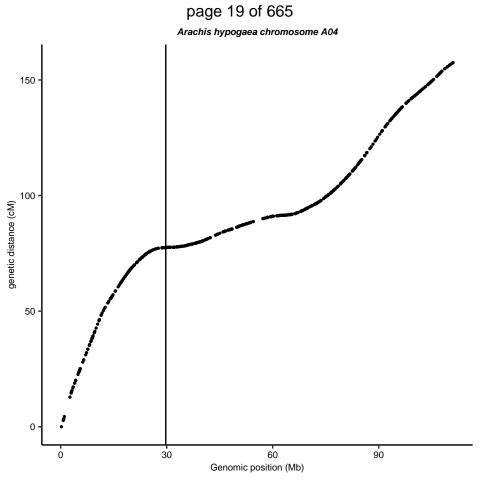
page 14 of 665 Arachis duranensis chromosome A8 genetic distance (cM) Genomic position (Mb)

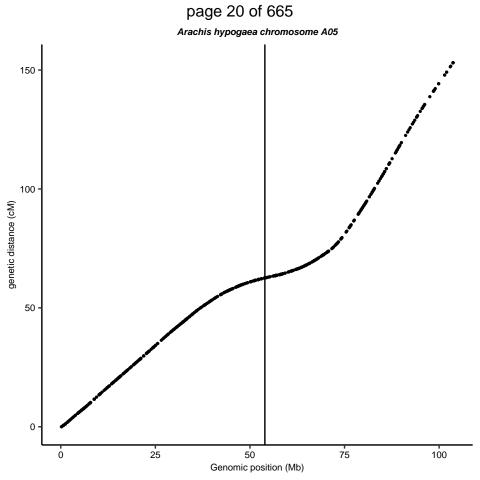
page 15 of 665 Arachis duranensis chromosome A10 genetic distance (cM) Genomic position (Mb)

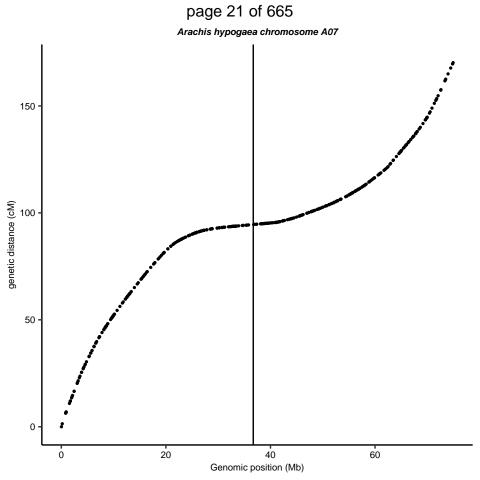


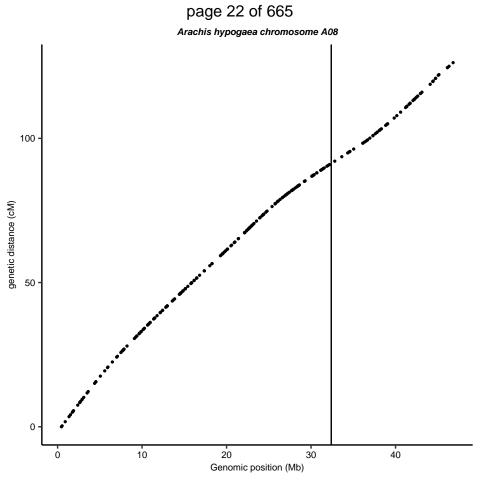
page 17 of 665 Arachis hypogaea chromosome A02 genetic distance (cM) Genomic position (Mb)

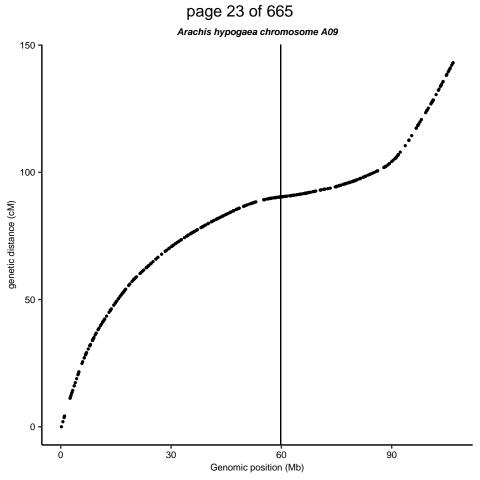


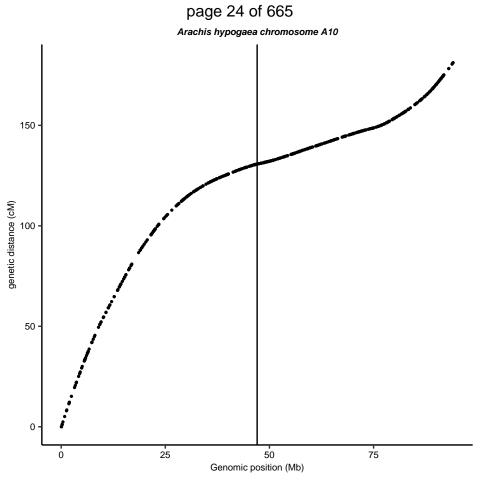


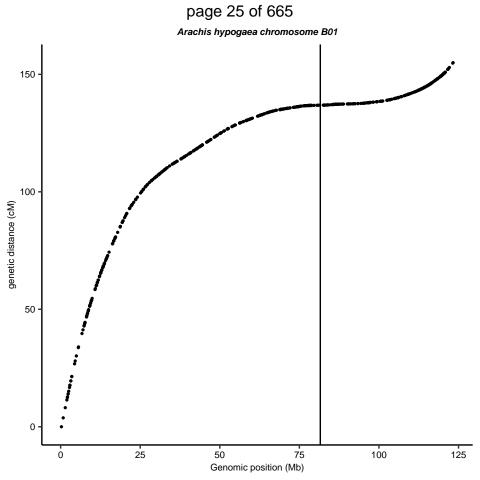


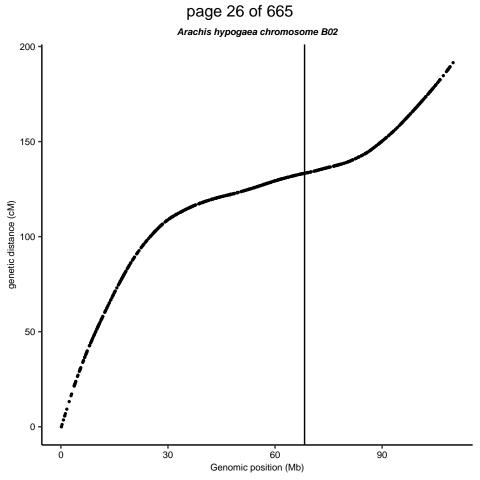


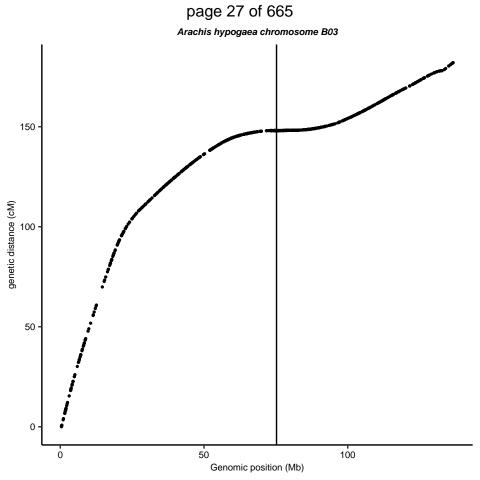


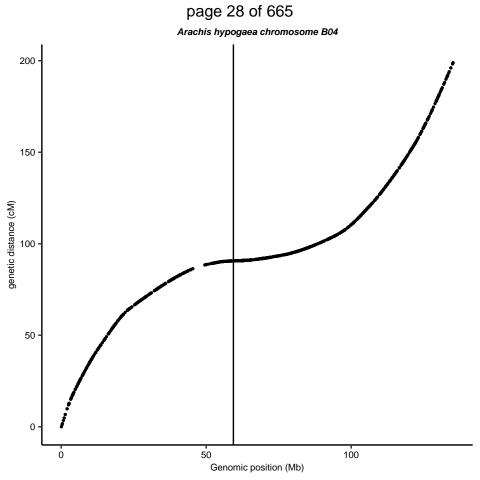


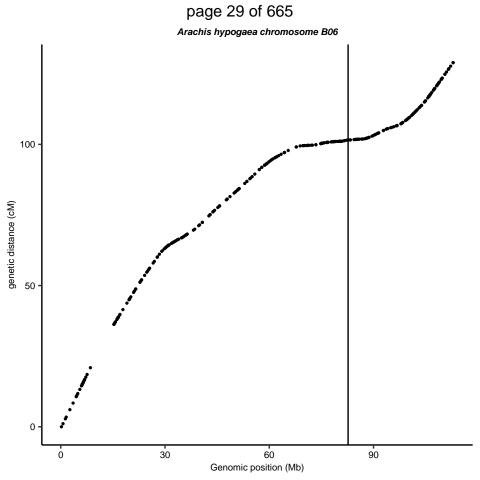


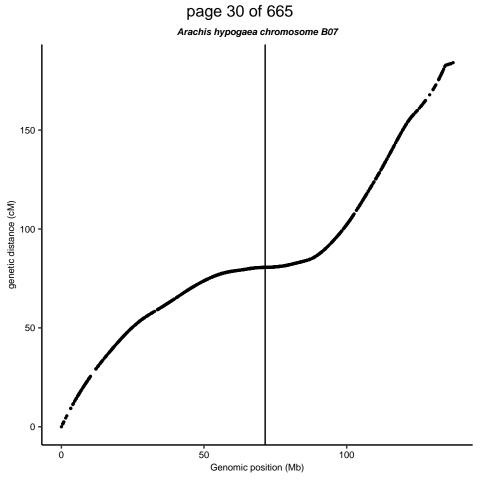


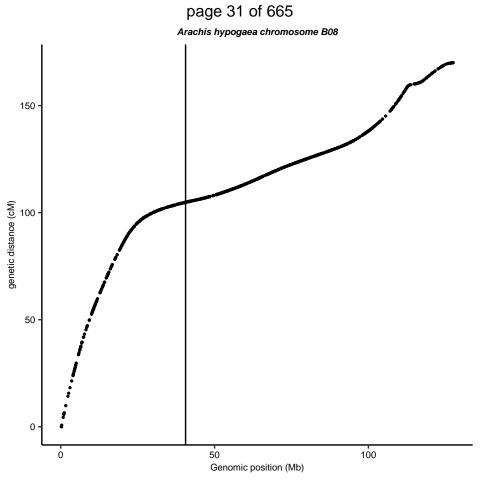


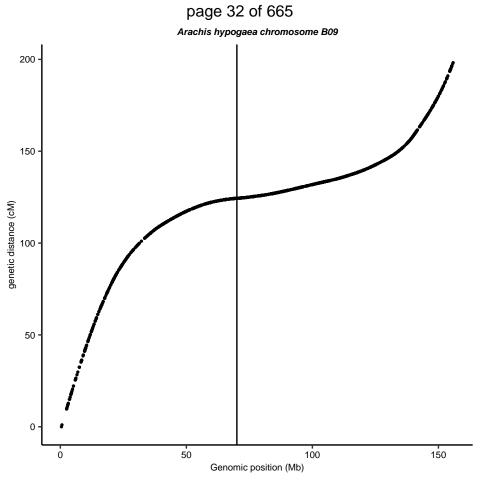


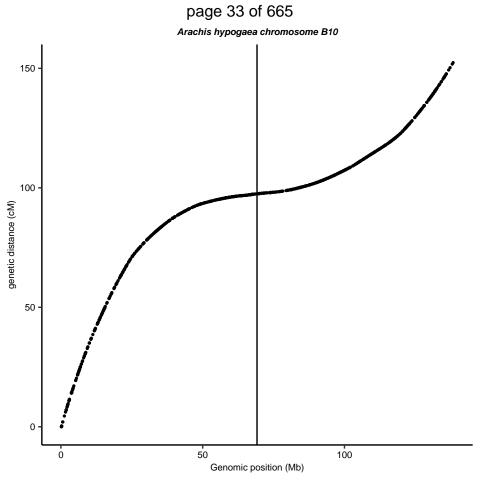


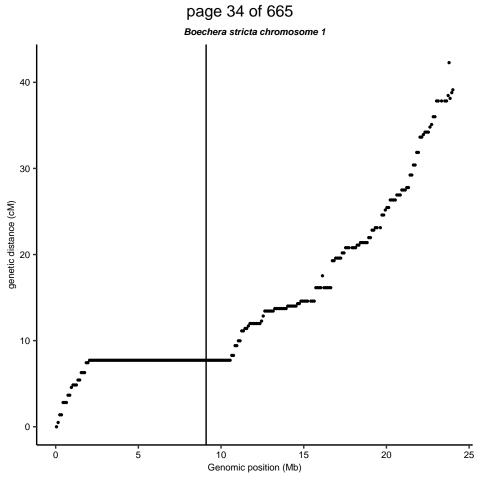


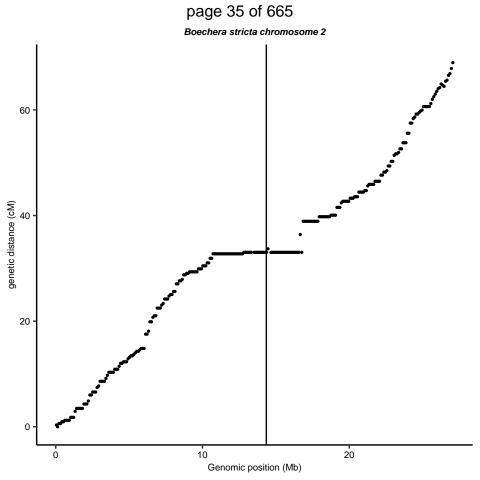


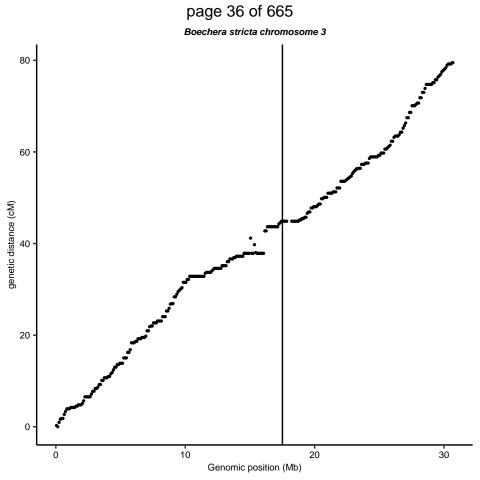


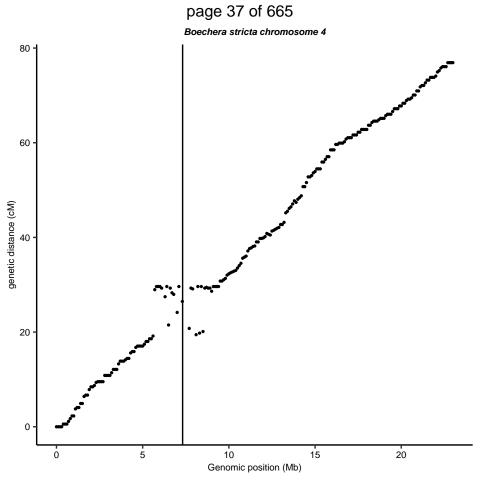


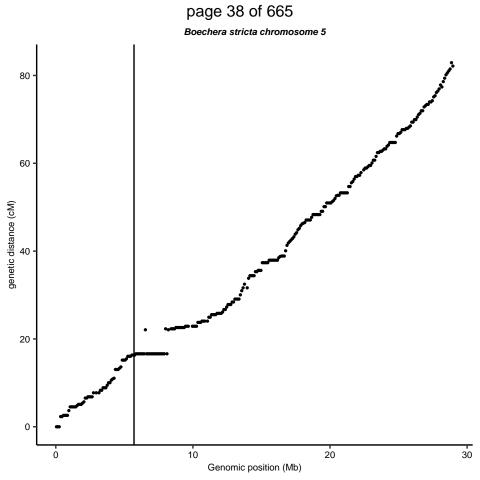


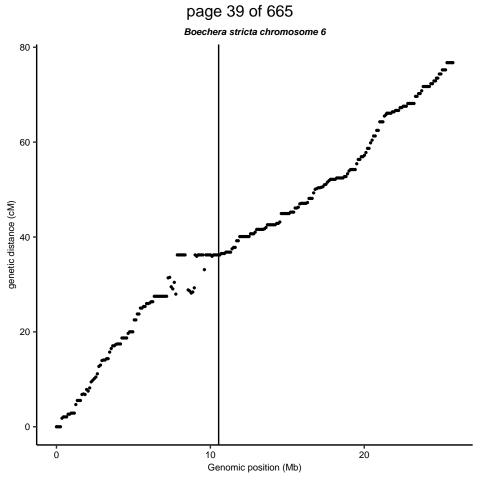


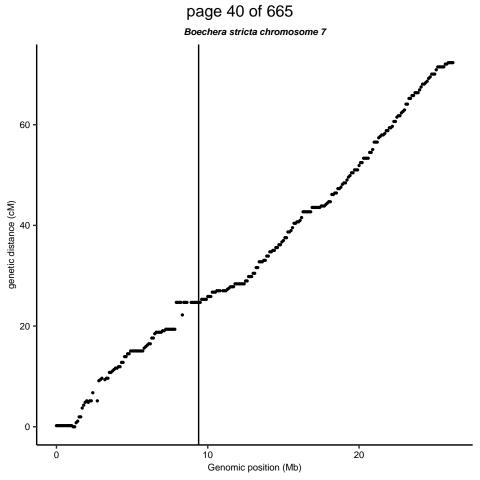


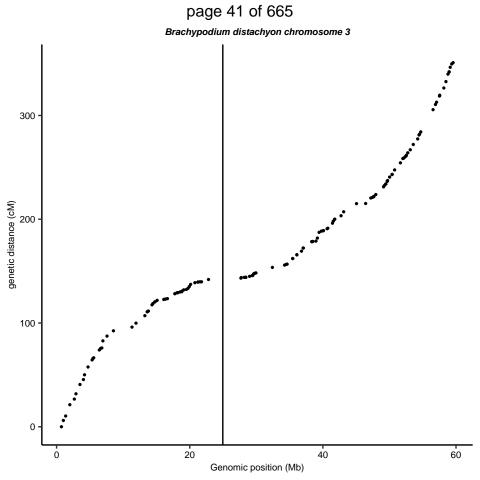






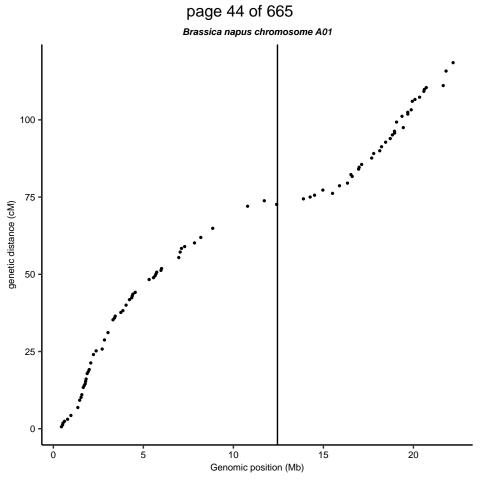




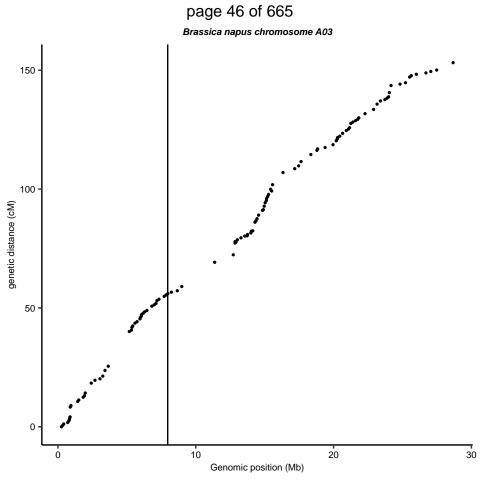


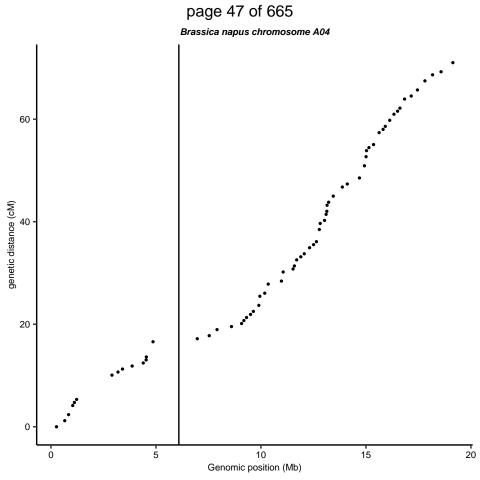
page 42 of 665 Brachypodium distachyon chromosome 4 genetic distance (cM) Genomic position (Mb)

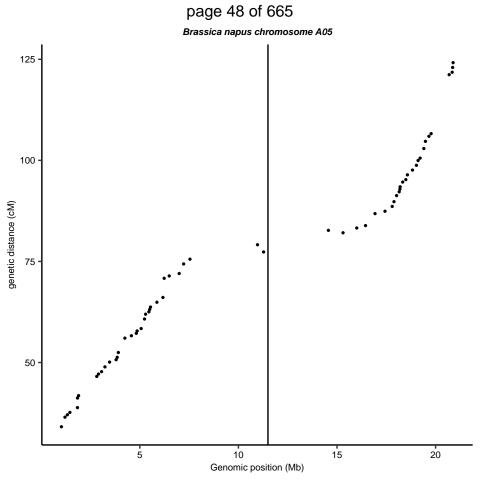
page 43 of 665 Brachypodium distachyon chromosome 5 150 genetic distance (cM) 100 50 0 10 20 Genomic position (Mb)



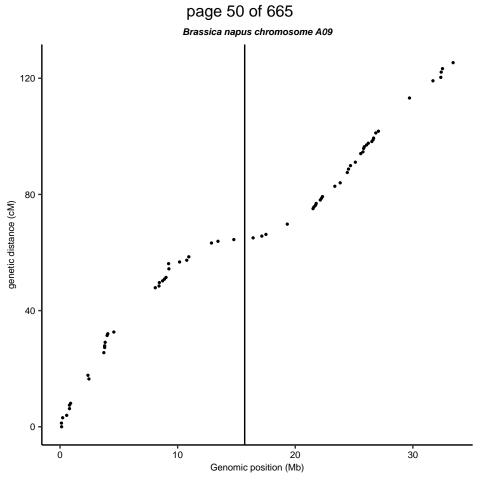
page 45 of 665 Brassica napus chromosome A02 75 genetic distance (cM) Genomic position (Mb)

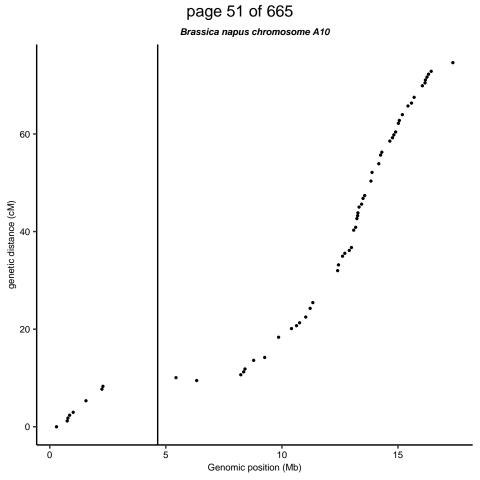


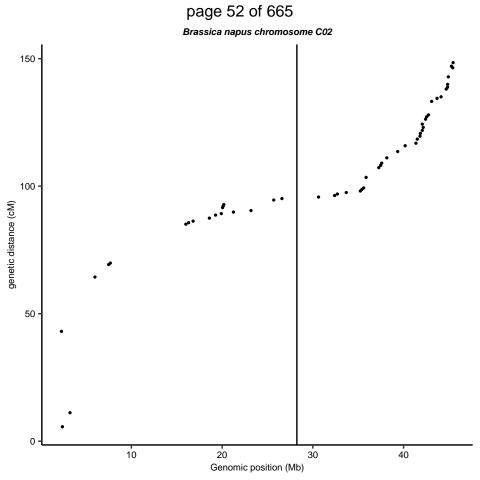


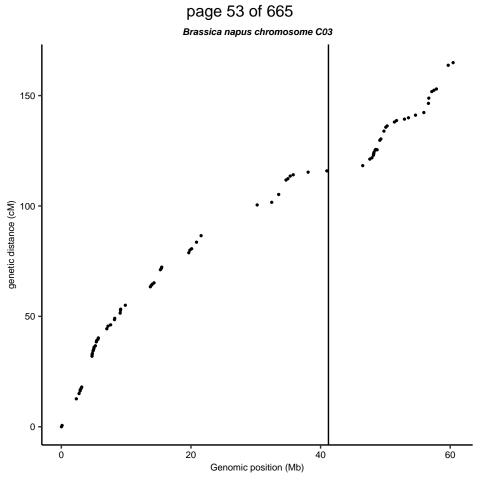


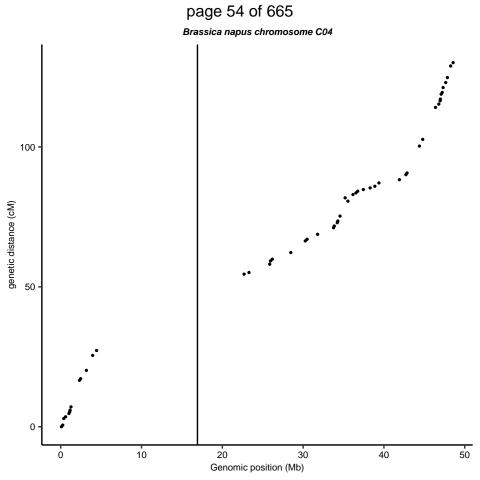
page 49 of 665 Brassica napus chromosome A07 genetic distance (cM) Genomic position (Mb)

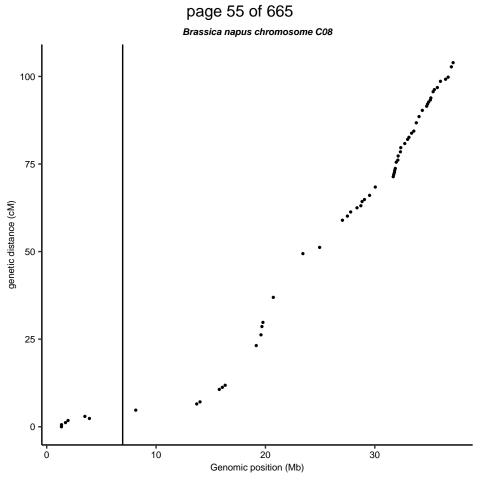


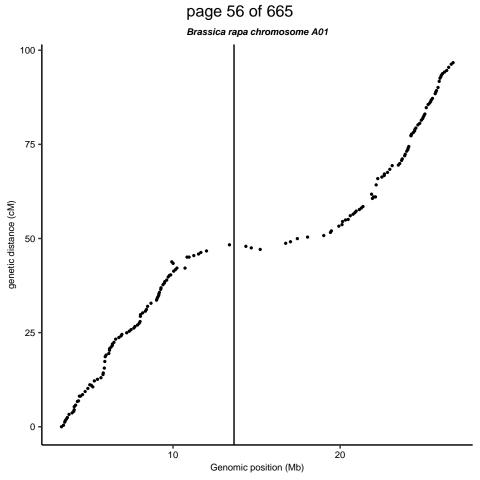


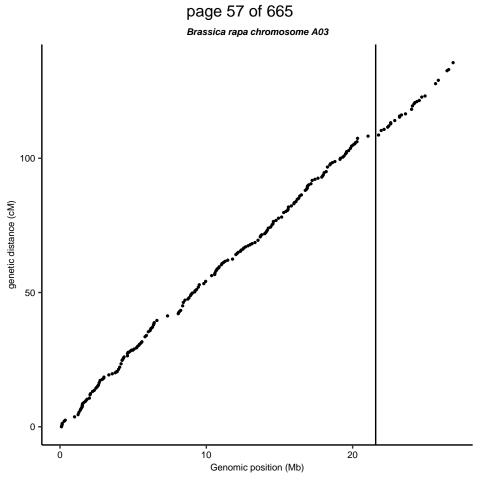


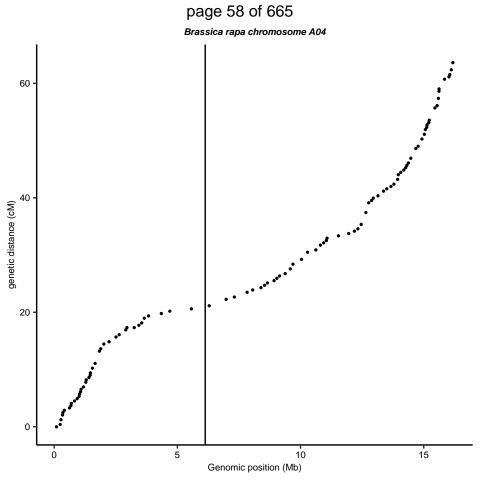


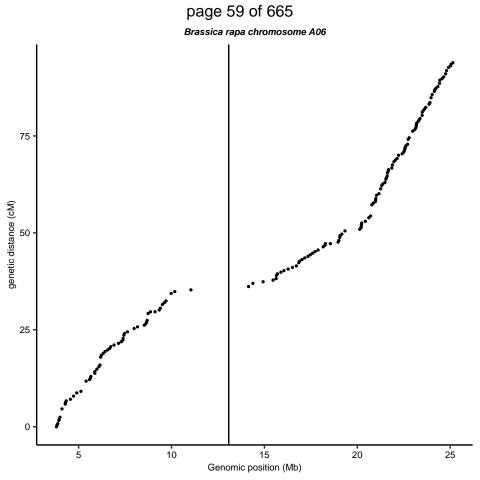


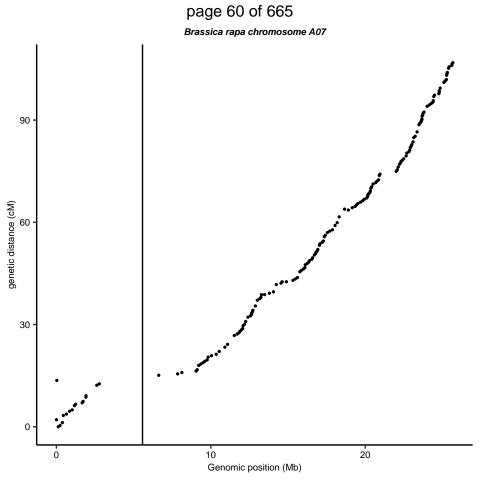


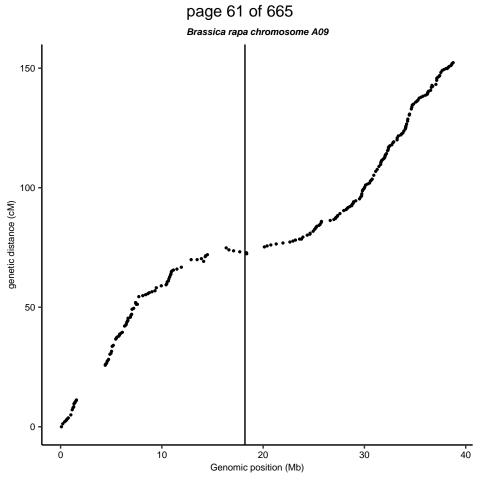


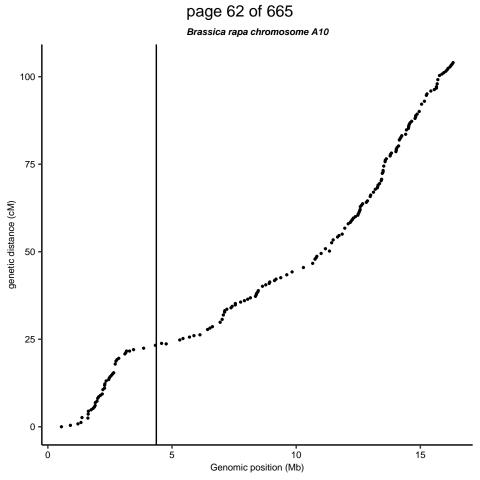


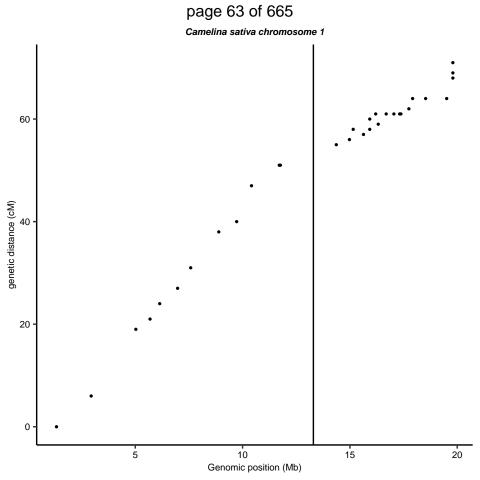


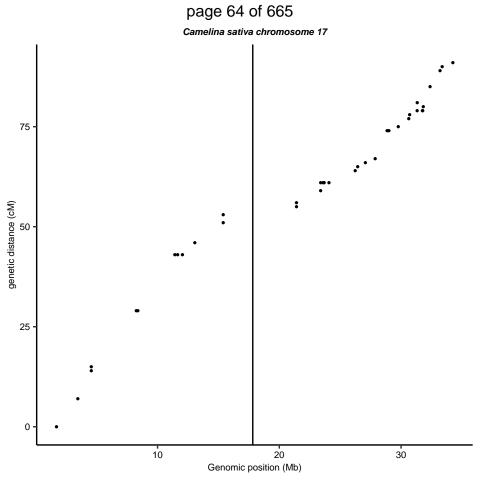


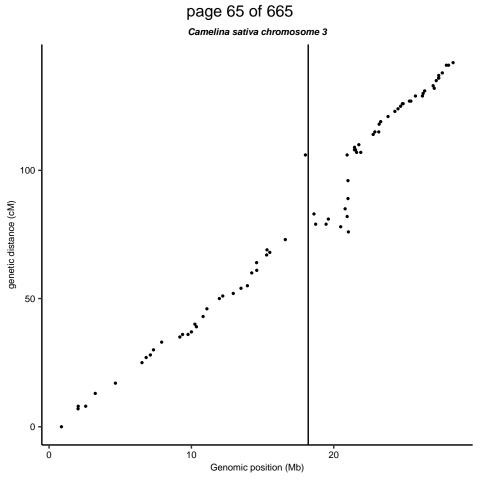


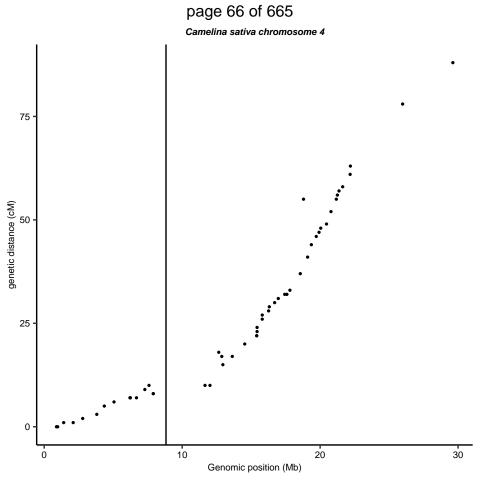


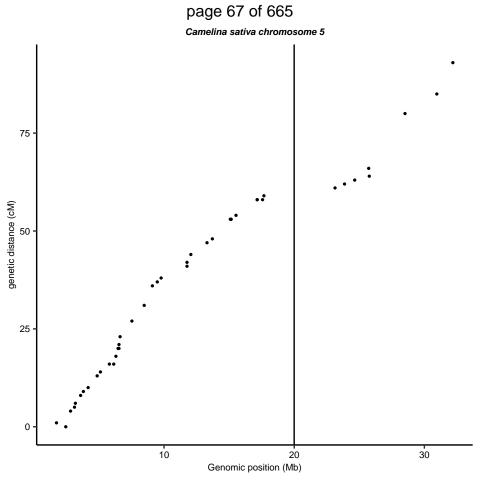


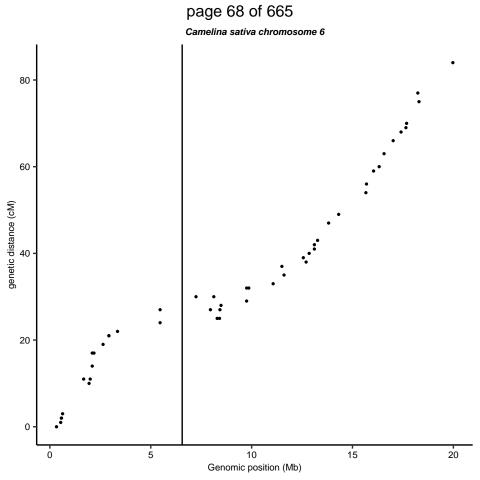


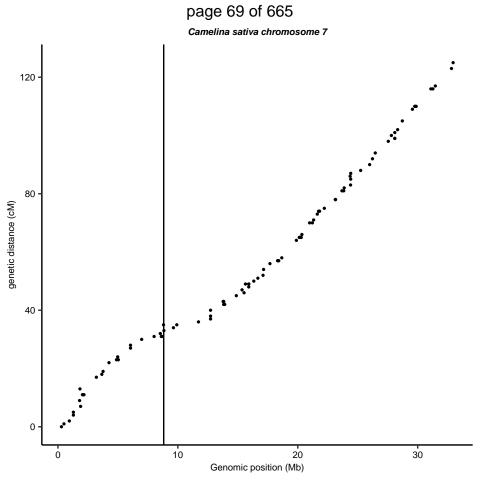


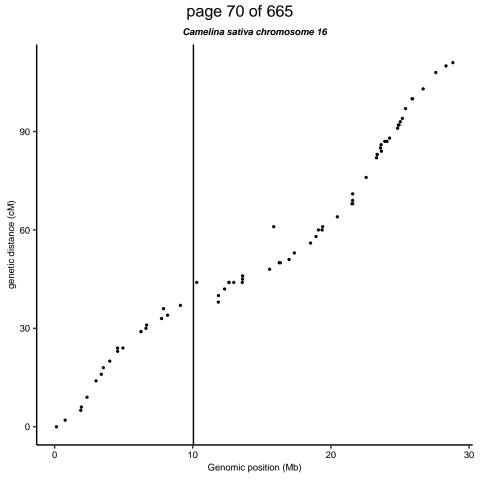


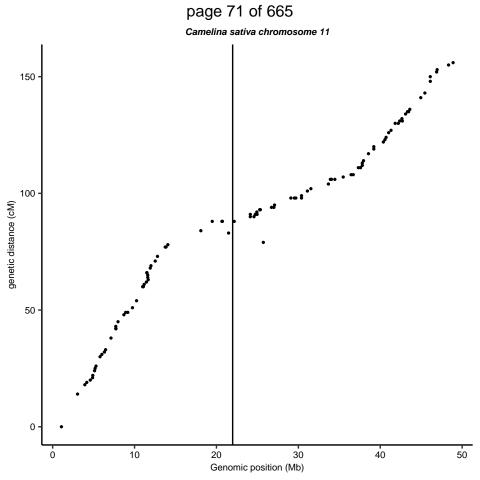


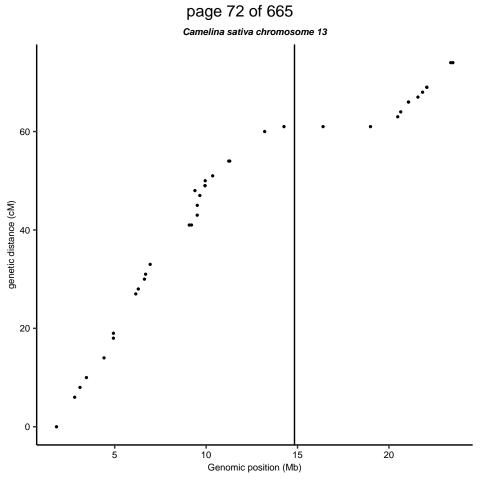


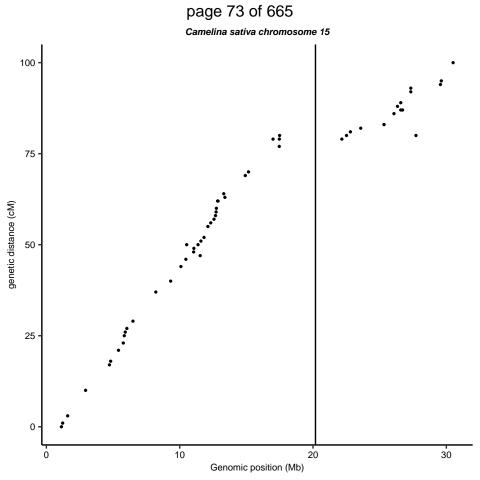




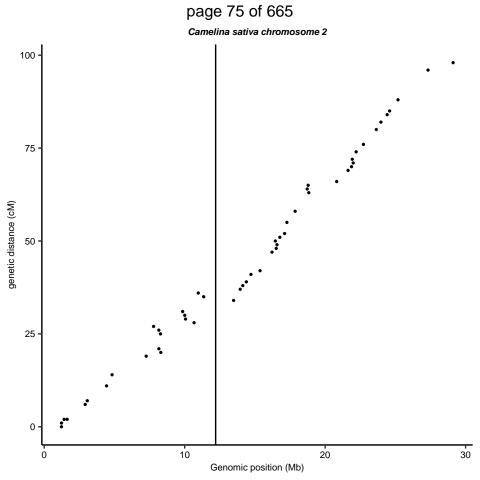




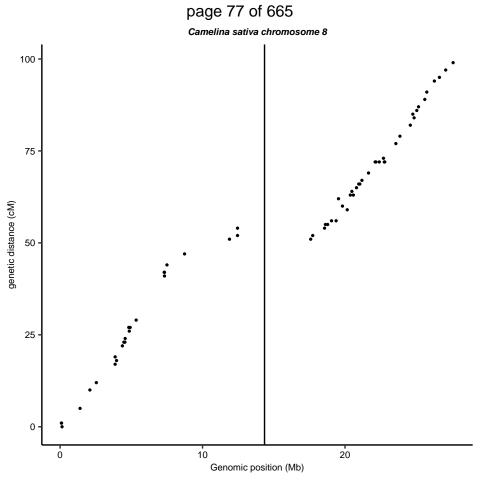


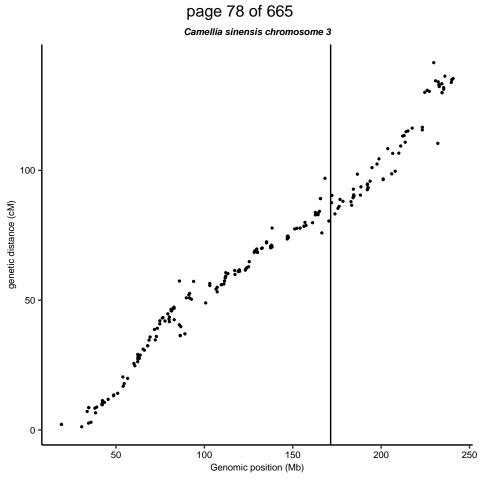


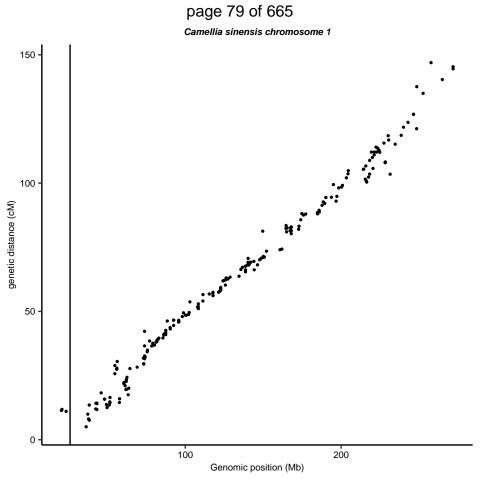
page 74 of 665 Camelina sativa chromosome 19 75 genetic distance (cM) Genomic position (Mb)

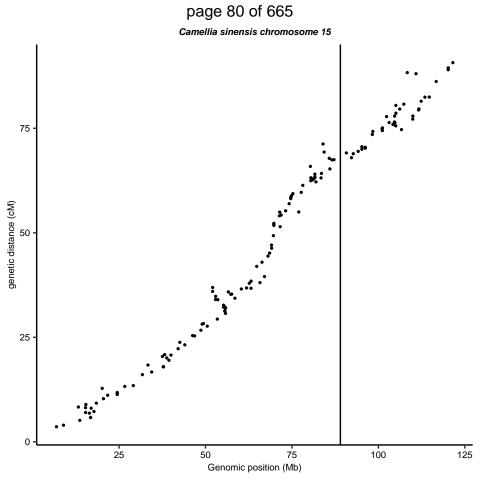


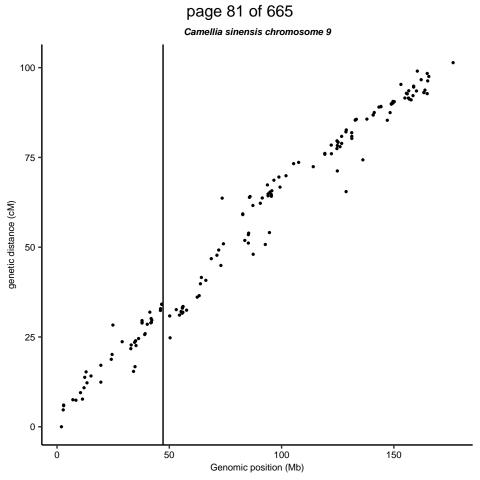
page 76 of 665 Camelina sativa chromosome 20 75 genetic distance (cM) 25 10 20 30 Genomic position (Mb)



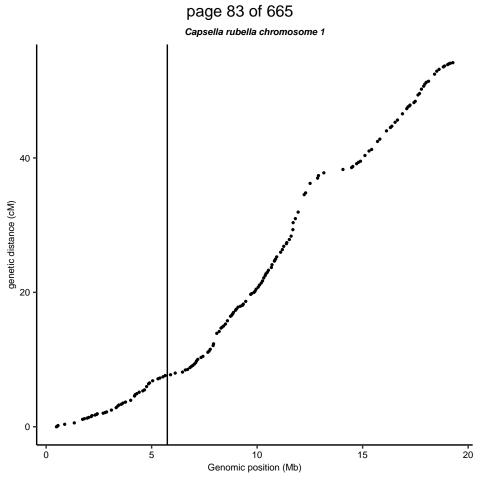


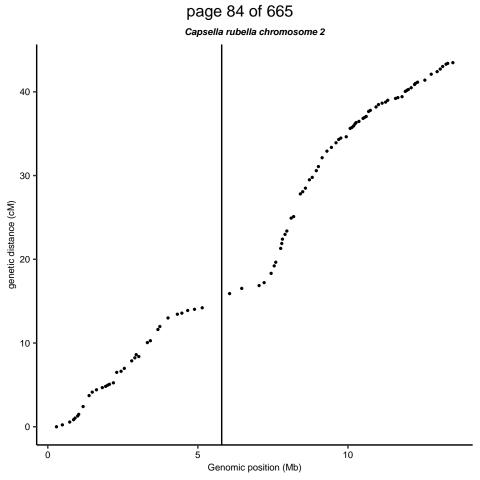


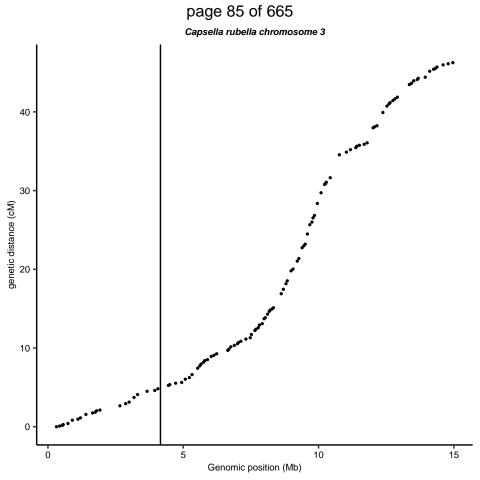


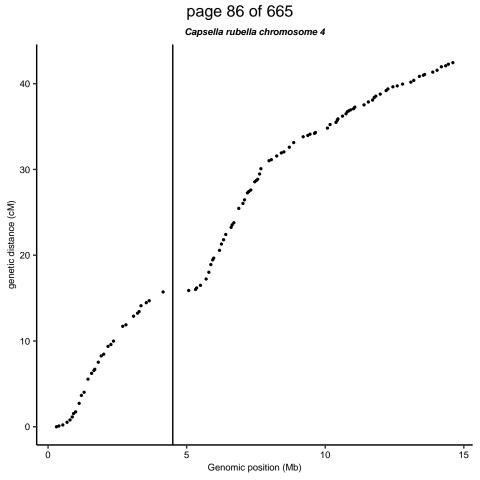


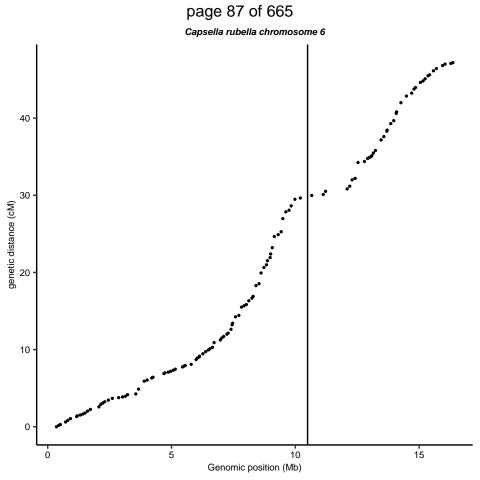
page 82 of 665 Camellia sinensis chromosome 8 90 genetic distance (cM) 30 0 50 150 100 Genomic position (Mb)

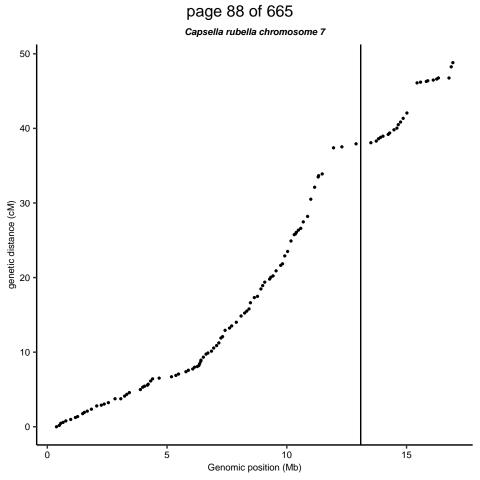


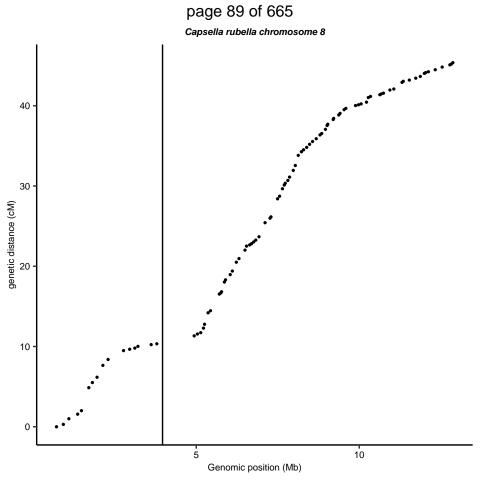


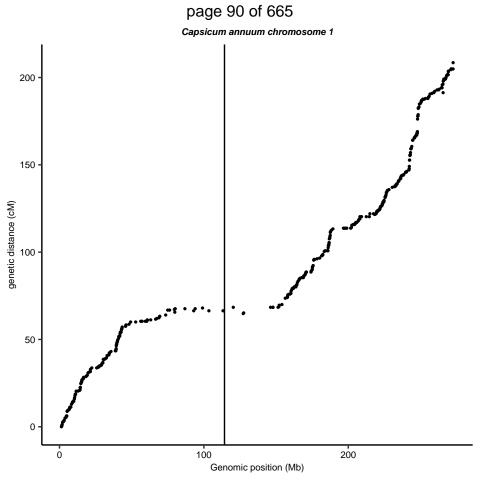


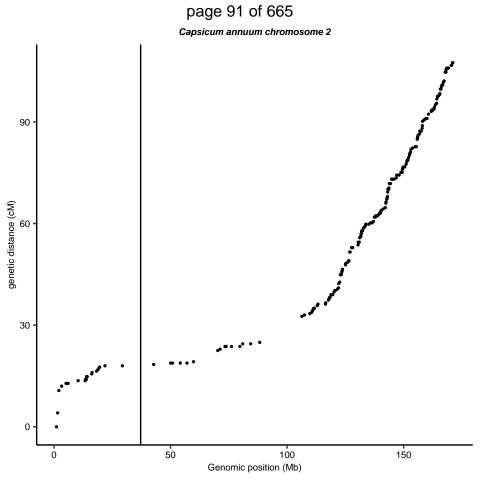


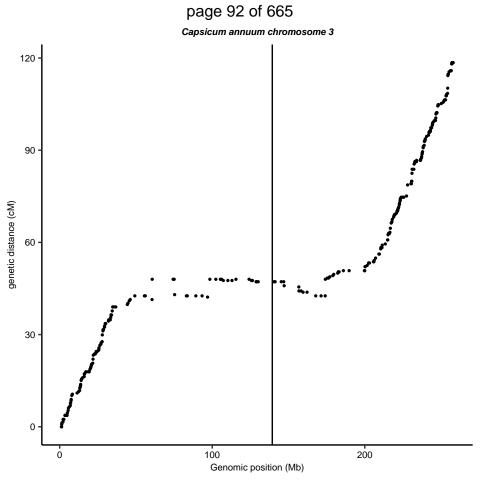


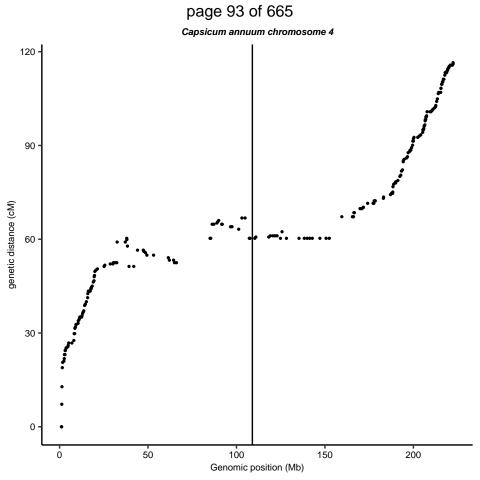


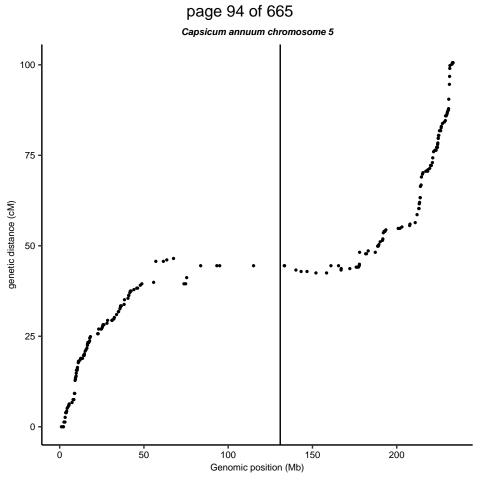


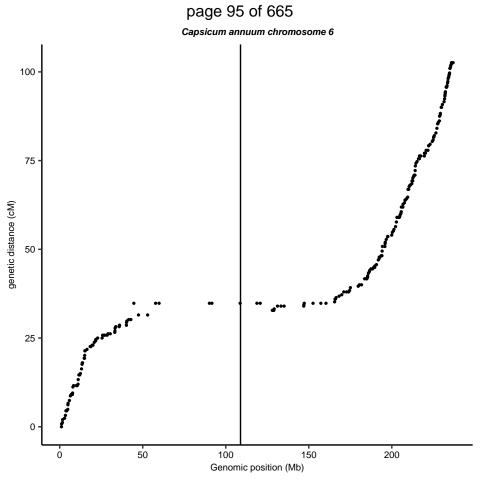


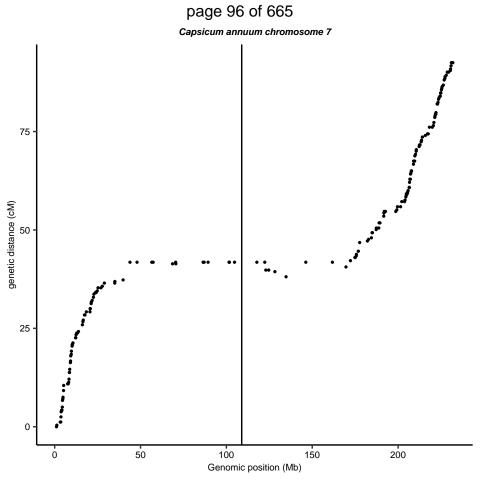




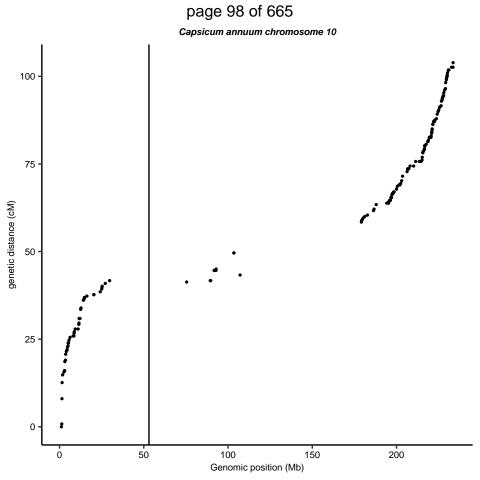


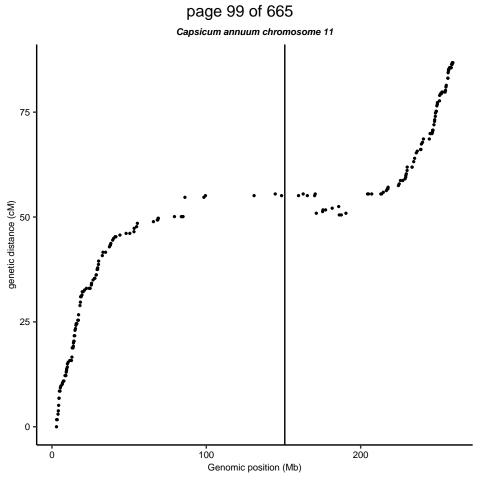


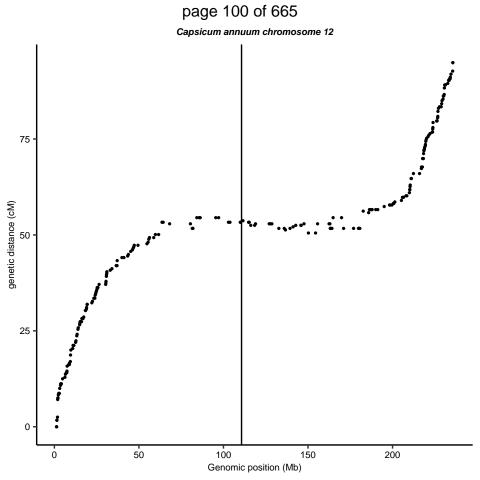


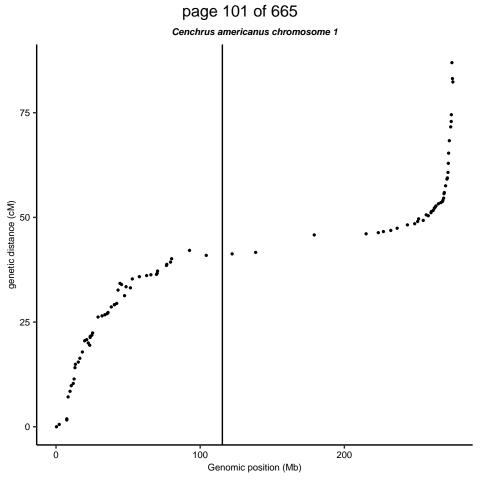


page 97 of 665 Capsicum annuum chromosome 9 genetic distance (cM) Genomic position (Mb)









page 102 of 665 Cenchrus americanus chromosome 2 genetic distance (cM) Ö Genomic position (Mb)

page 103 of 665 Cenchrus americanus chromosome 3 40 genetic distance (cM) 20 0 100 200 ö 300 Genomic position (Mb)

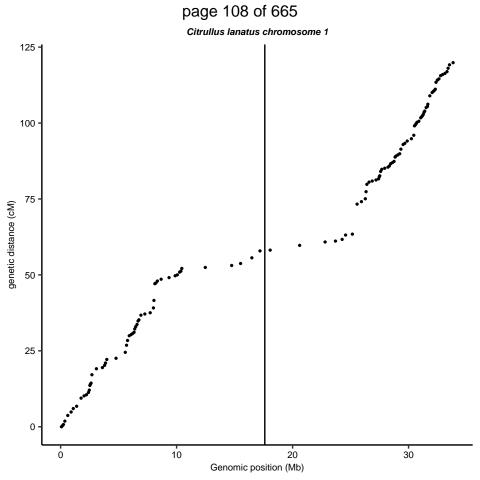
page 104 of 665 Cenchrus americanus chromosome 4 genetic distance (cM) Genomic position (Mb)

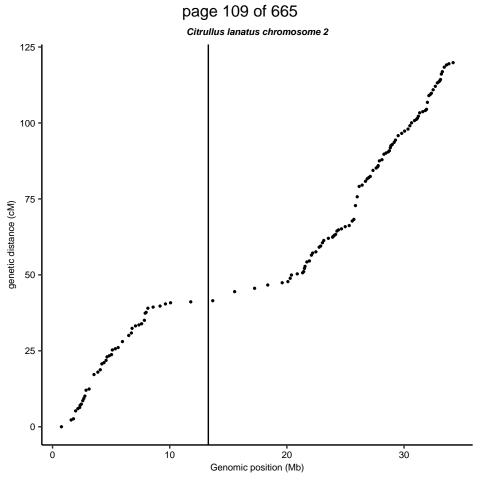
page 105 of 665 Cenchrus americanus chromosome 5 75 genetic distance (cM) 25 0 50 100 150 Genomic position (Mb)

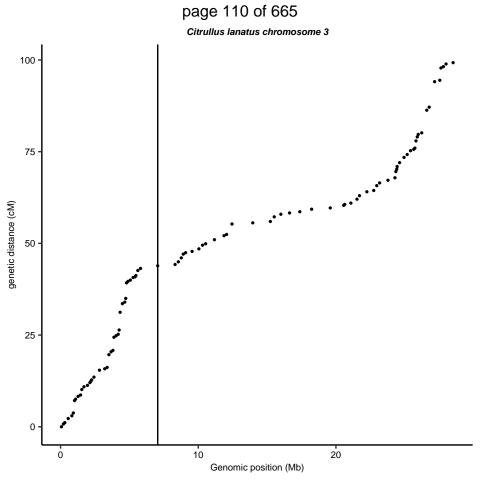
page 106 of 665 Cenchrus americanus chromosome 6 genetic distance (cM)

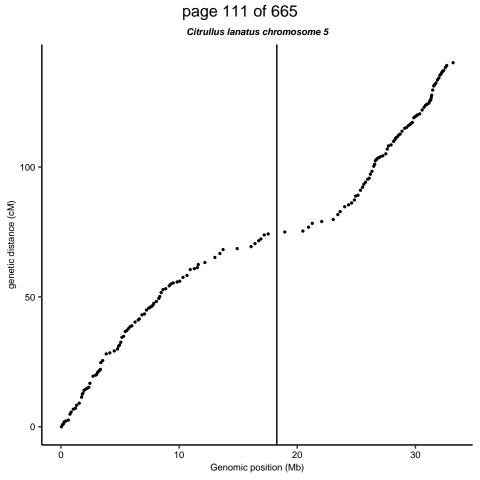
Genomic position (Mb)

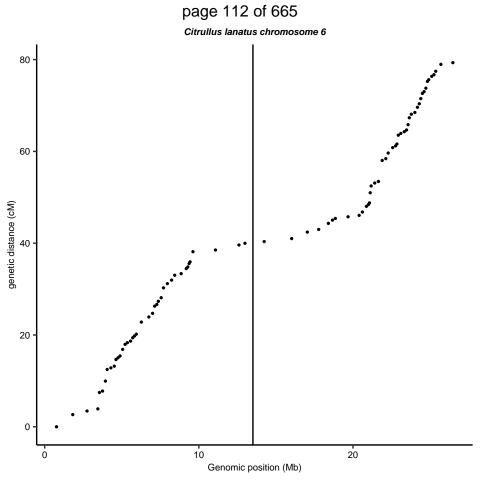
page 107 of 665 Cenchrus americanus chromosome 7 genetic distance (cM) Genomic position (Mb)

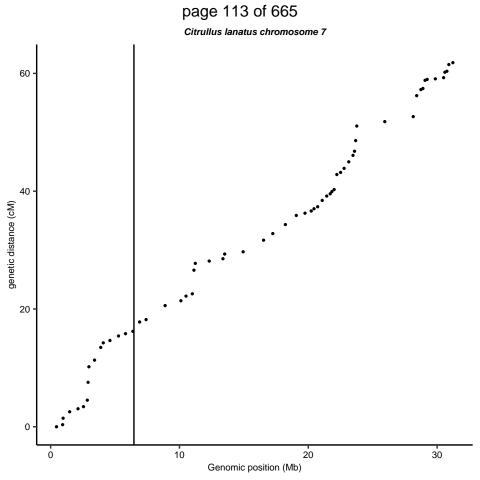


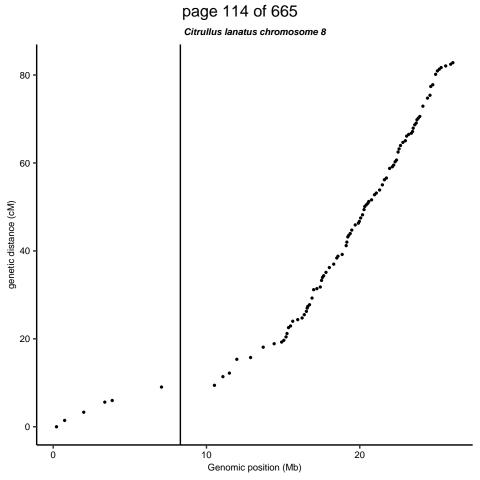


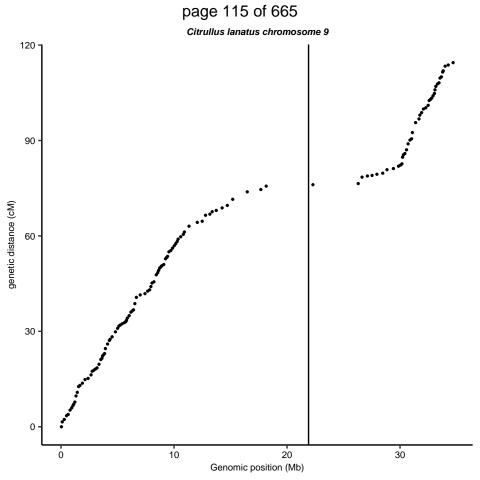


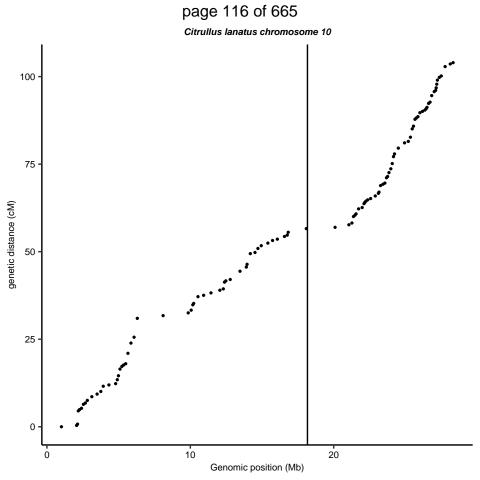


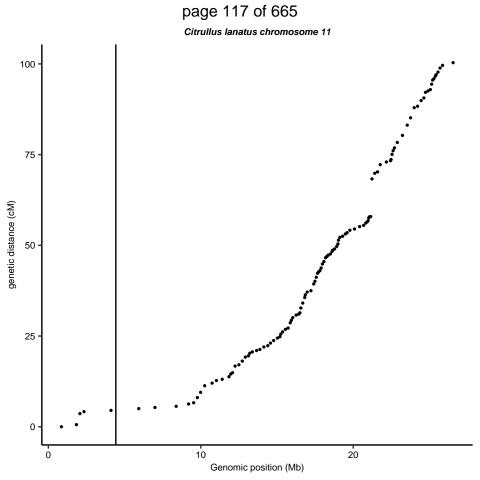


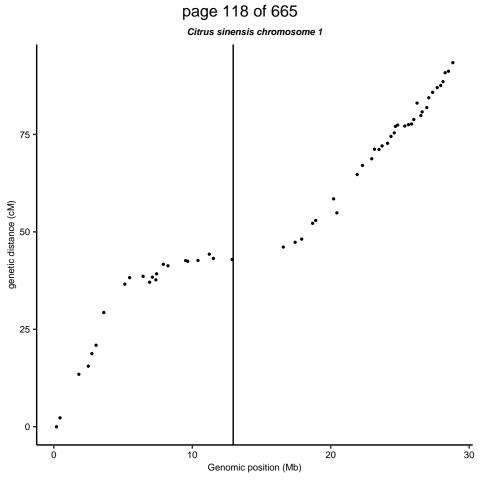




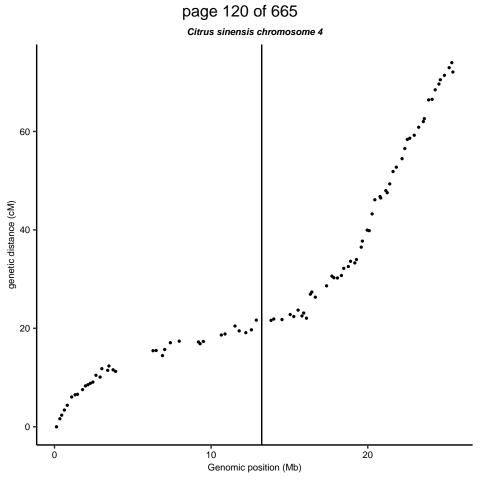


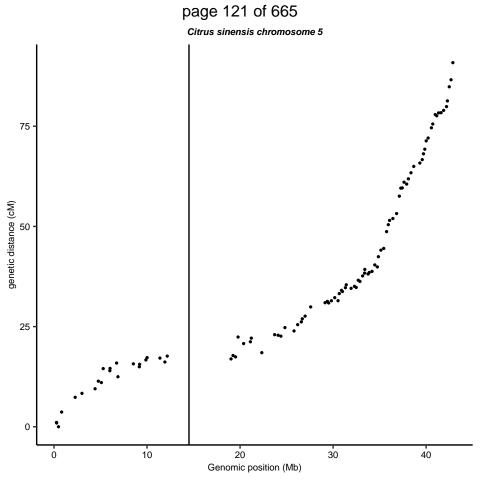




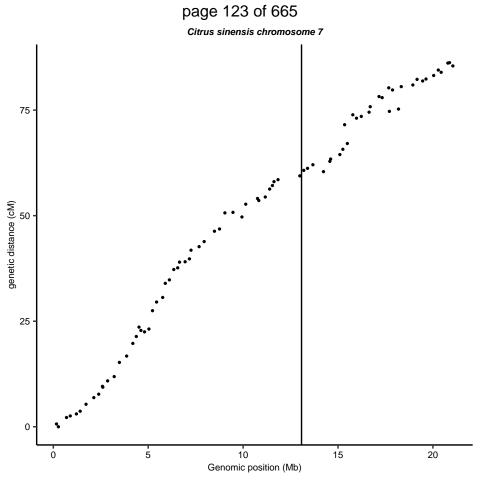


page 119 of 665 Citrus sinensis chromosome 3 75 genetic distance (cM) Genomic position (Mb)





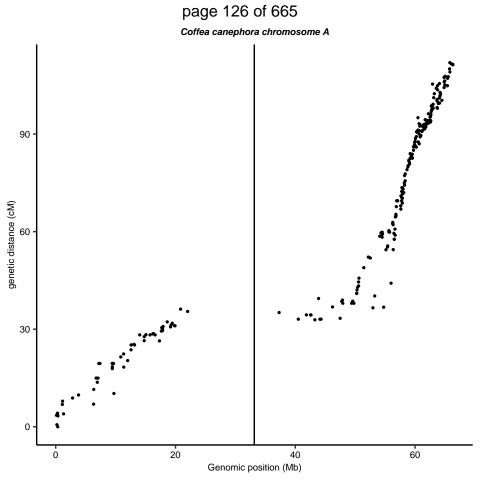
page 122 of 665 Citrus sinensis chromosome 6 genetic distance (cM) Genomic position (Mb)

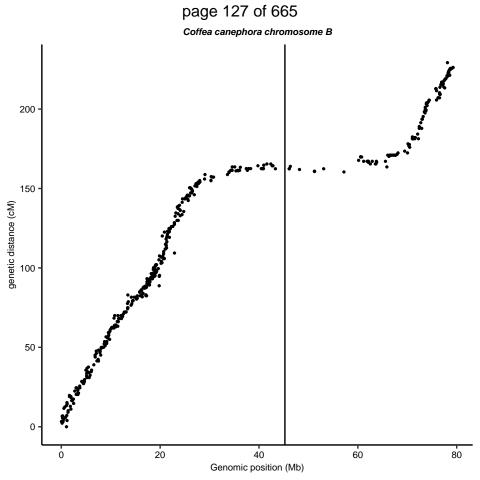


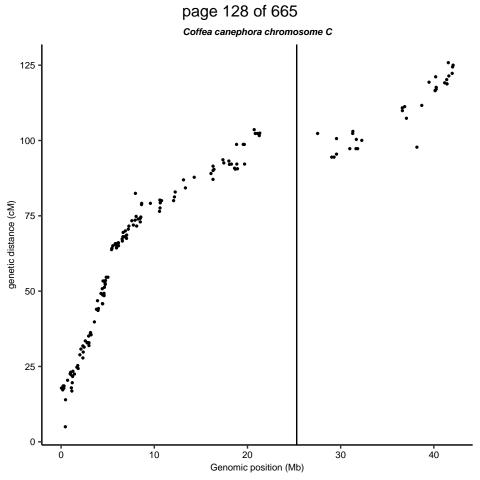
page 124 of 665 Citrus sinensis chromosome 8 genetic distance (cM) Ö Genomic position (Mb)

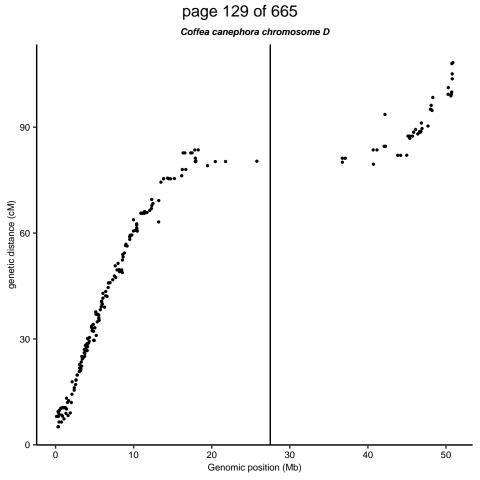
page 125 of 665 Citrus sinensis chromosome 9 60 40 genetic distance (cM) 20 0 -10 20

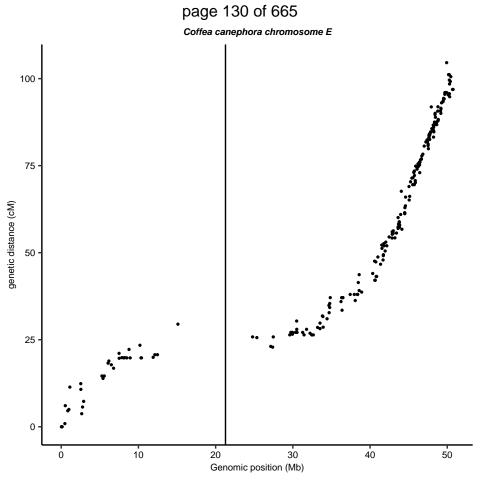
Genomic position (Mb)

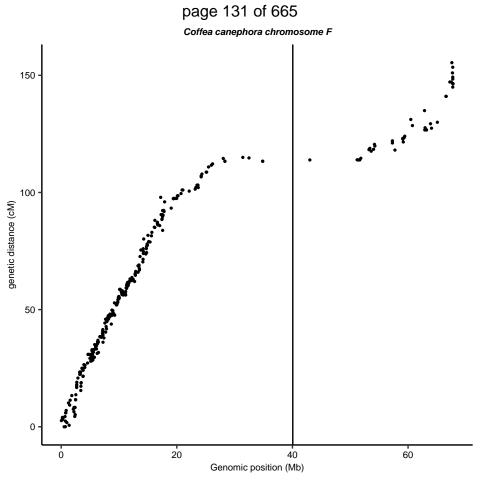


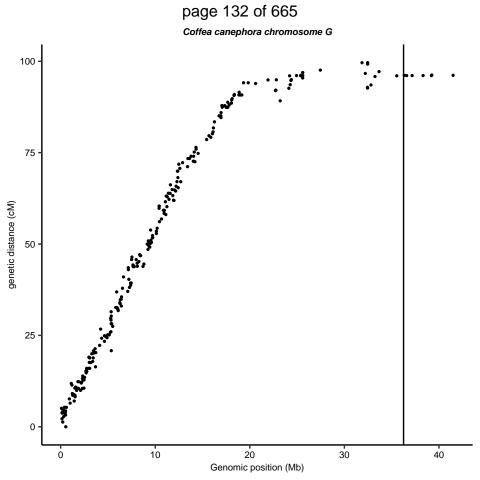


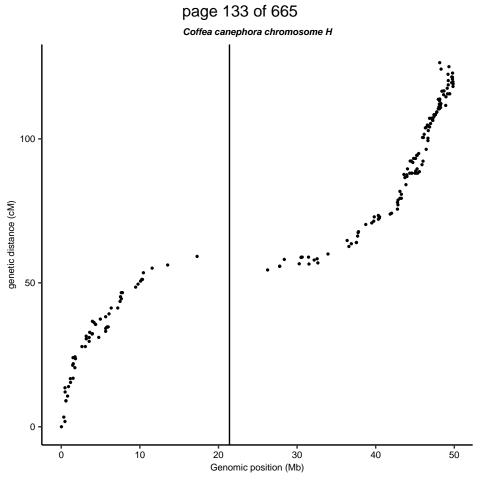


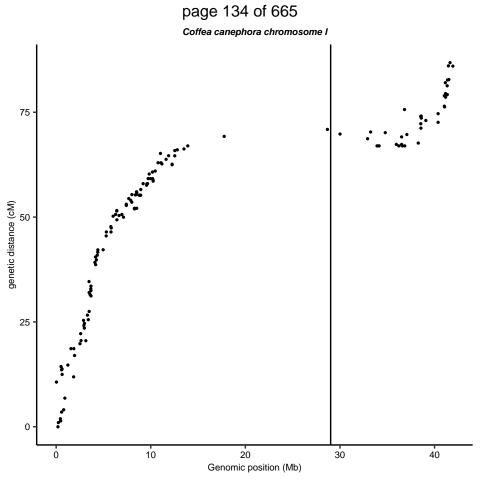




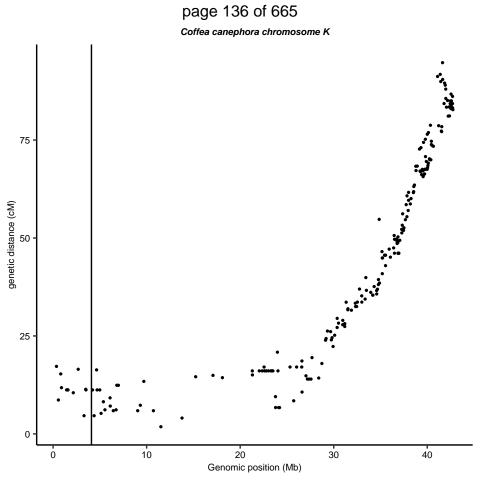






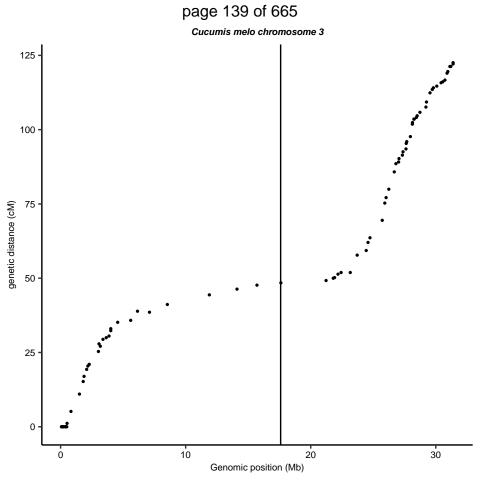


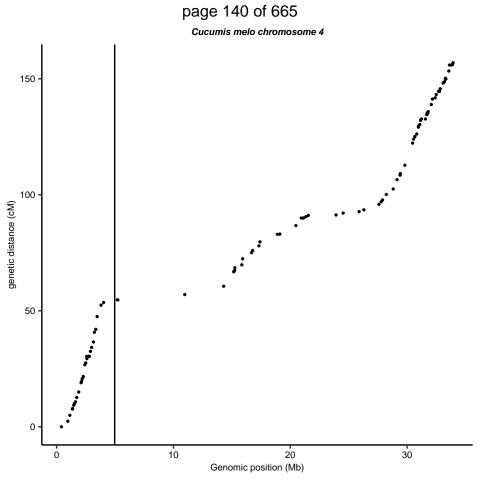
page 135 of 665 Coffea canephora chromosome J genetic distance (cM) Genomic position (Mb)

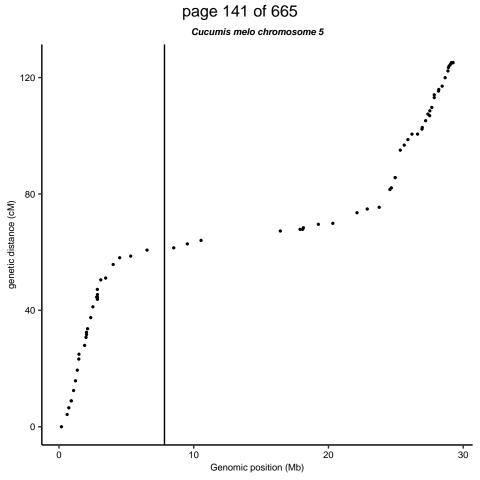


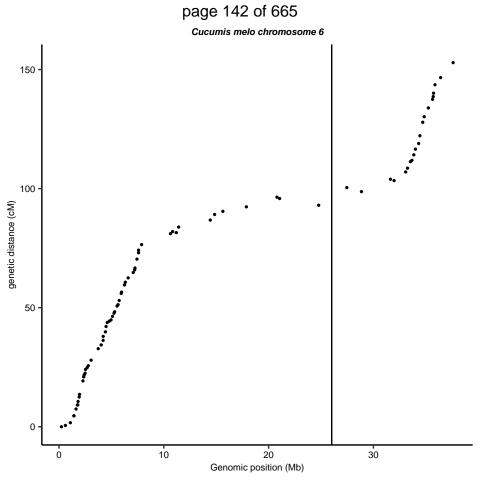
page 137 of 665 Cucumis melo chromosome 1 genetic distance (cM) Genomic position (Mb)

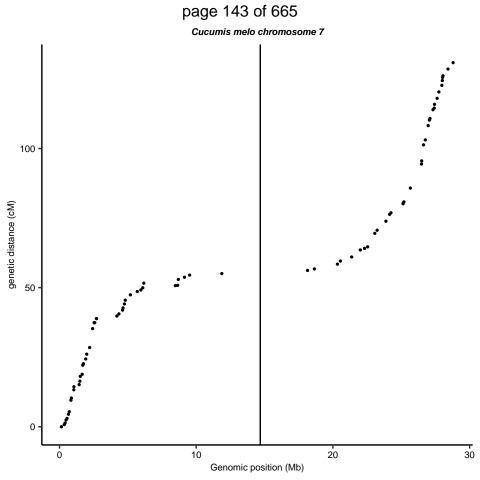
page 138 of 665 Cucumis melo chromosome 2 100 genetic distance (cM) 50 10 20 Genomic position (Mb)







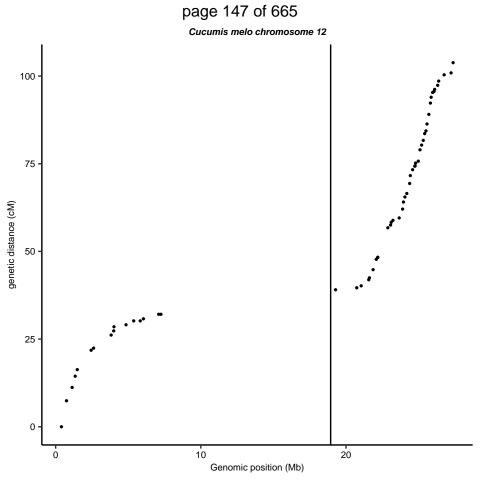


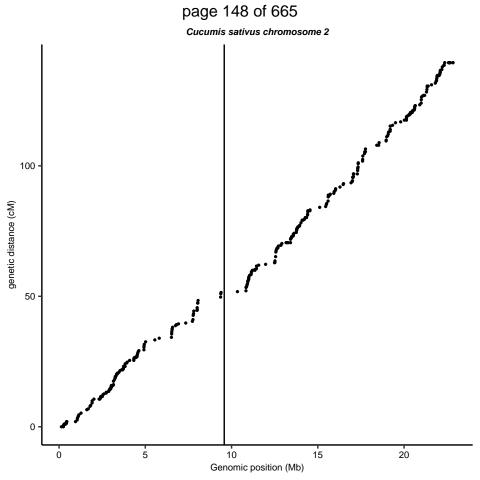


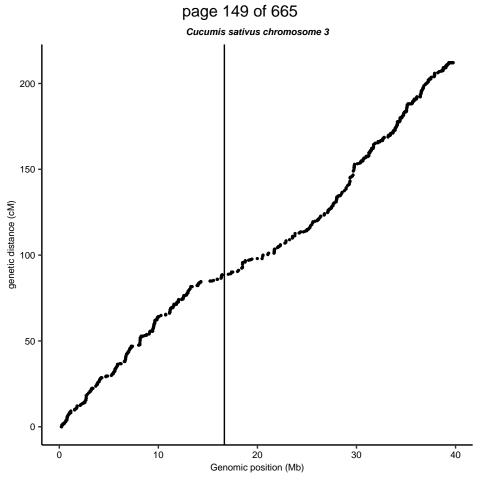
page 144 of 665 Cucumis melo chromosome 8 100 genetic distance (cM) 50 0 10 20 30 Genomic position (Mb)

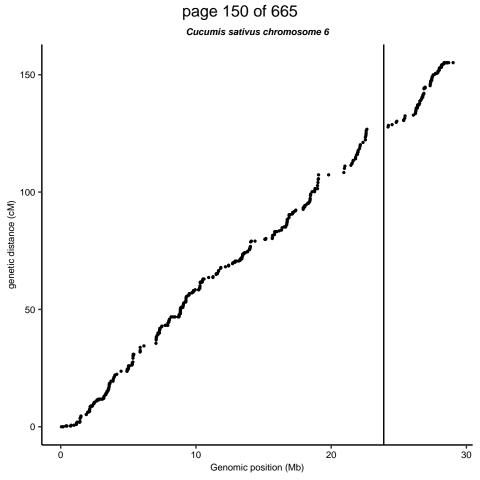
page 145 of 665 Cucumis melo chromosome 9 90 genetic distance (cM) 30 0 10 15 20 25 Ö Genomic position (Mb)

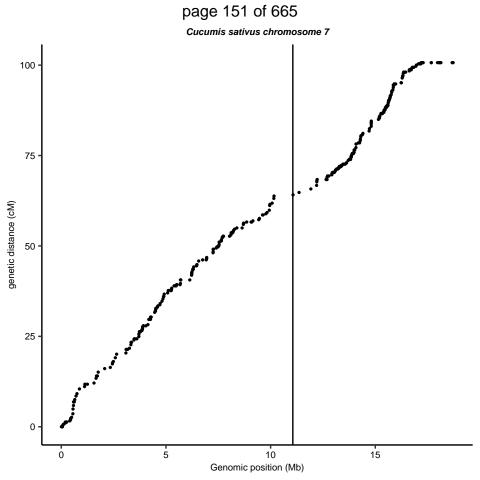
page 146 of 665 Cucumis melo chromosome 10 100 75 genetic distance (cM) 50 25 0 10 15 20 Genomic position (Mb)

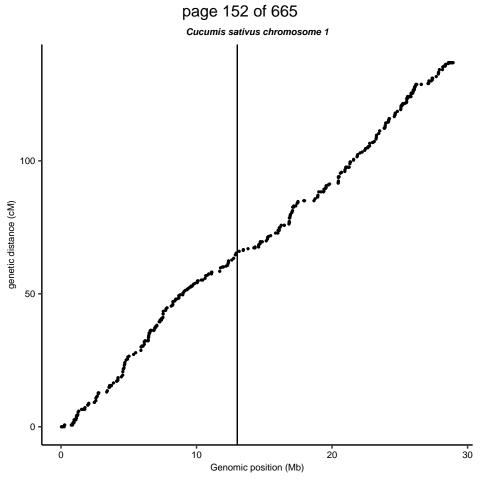


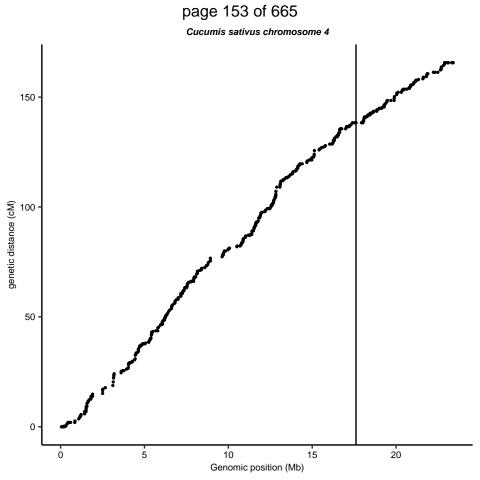


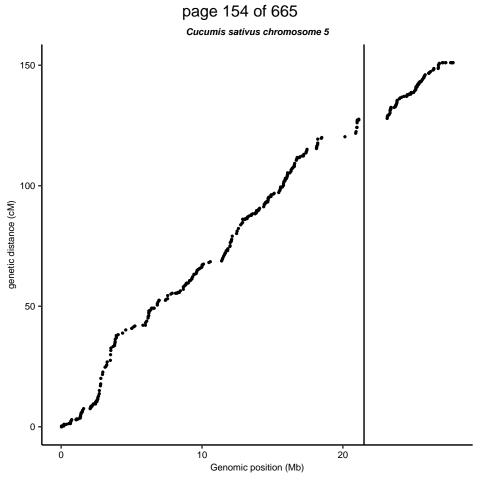


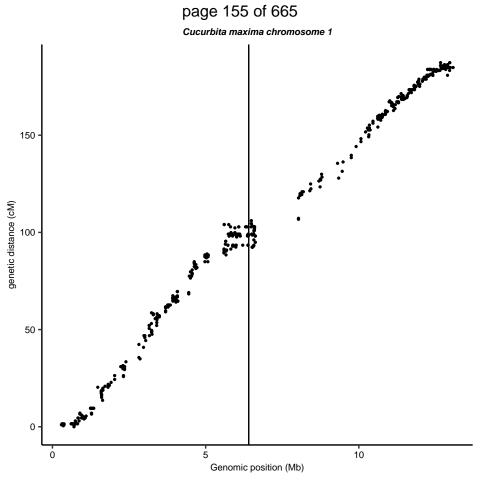


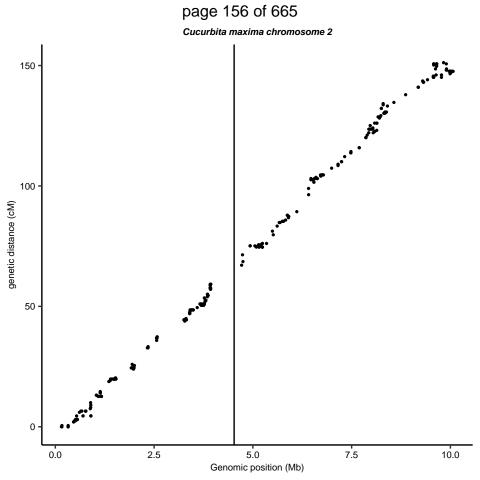


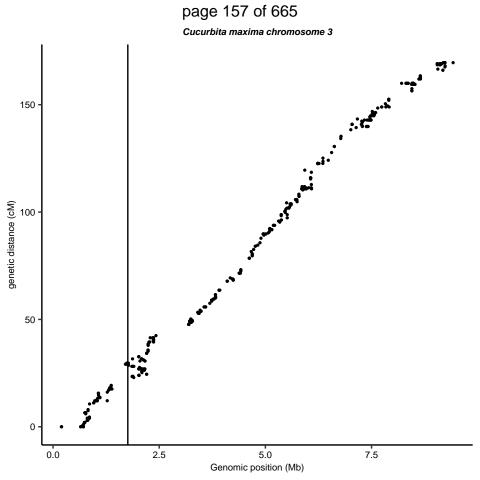


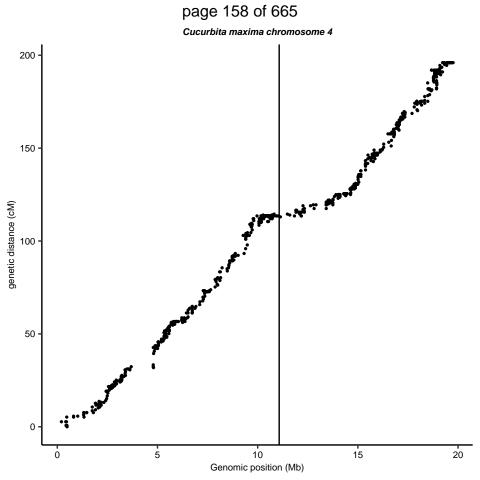


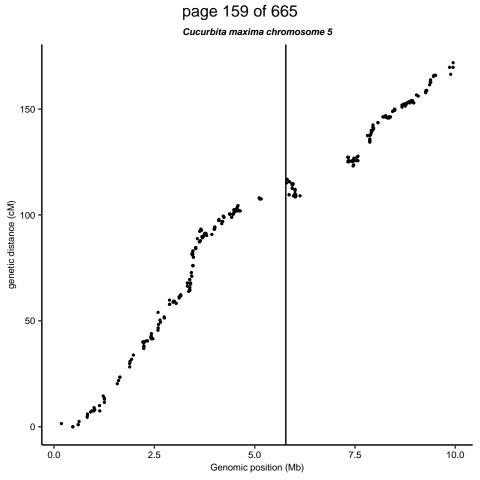


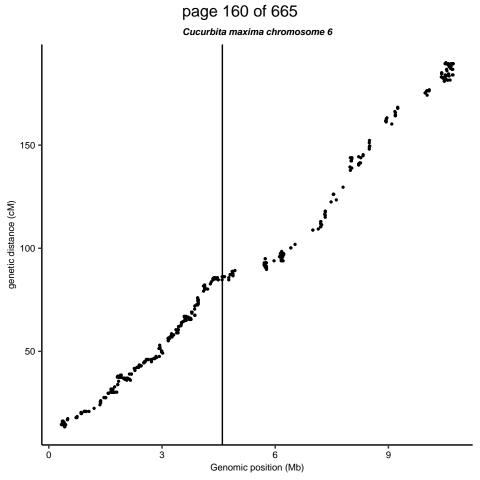


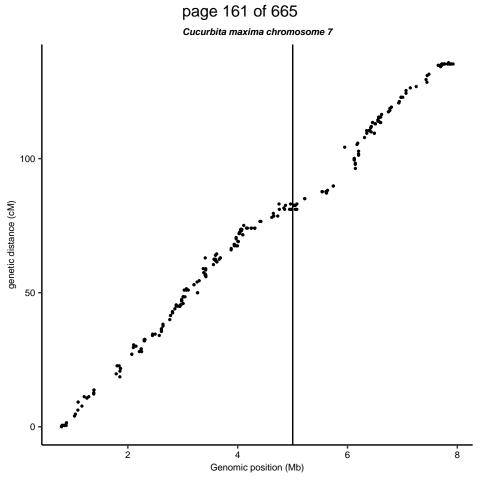


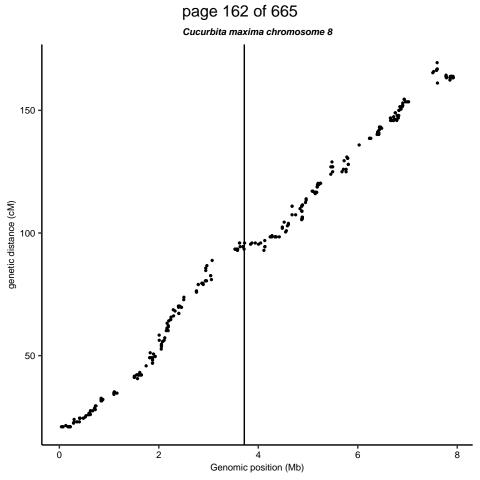


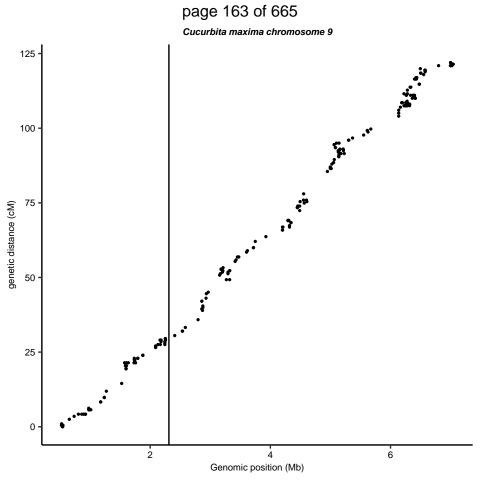


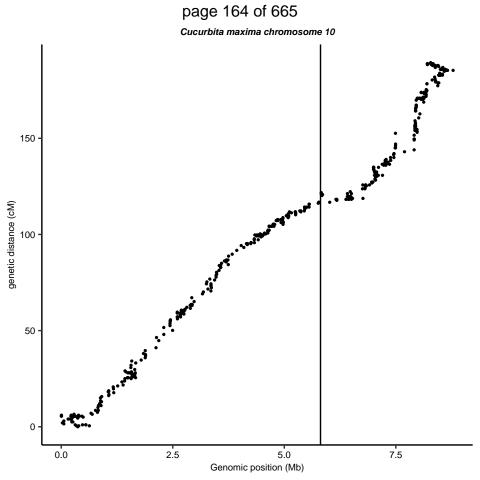


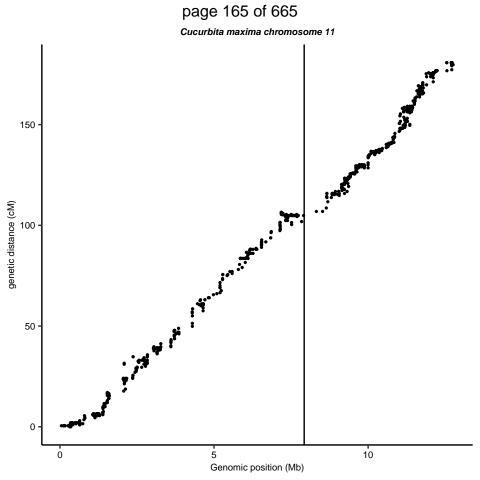


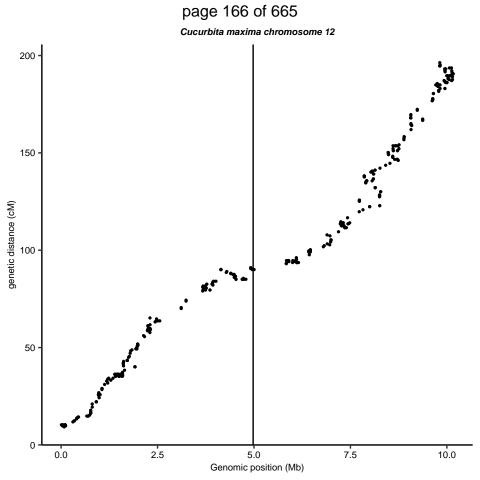


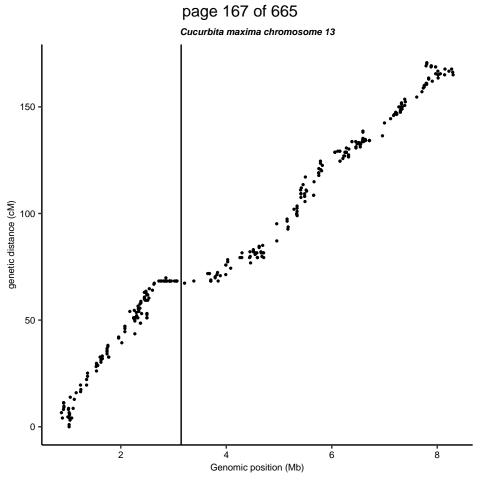


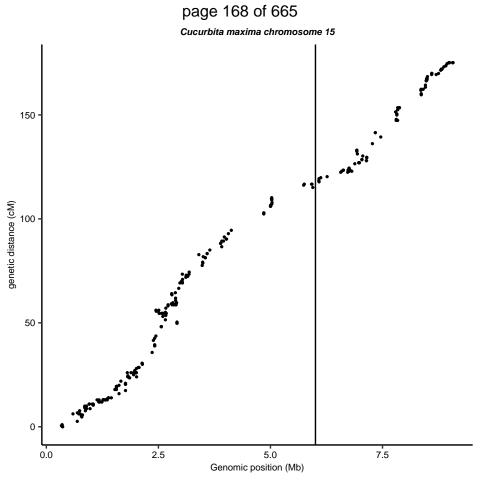


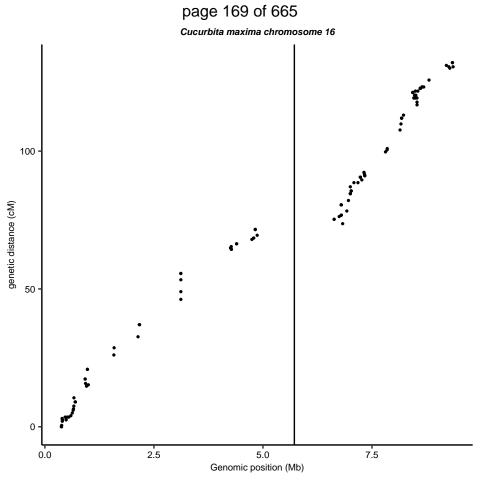


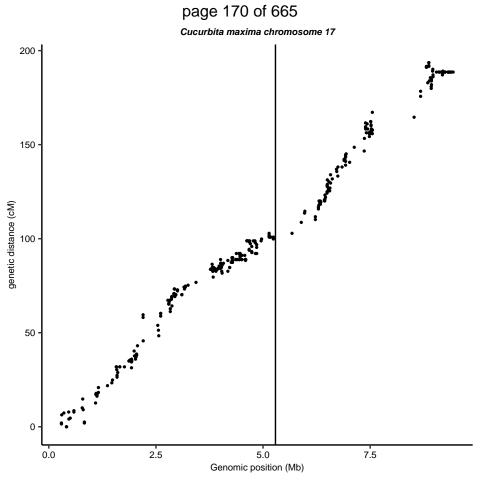


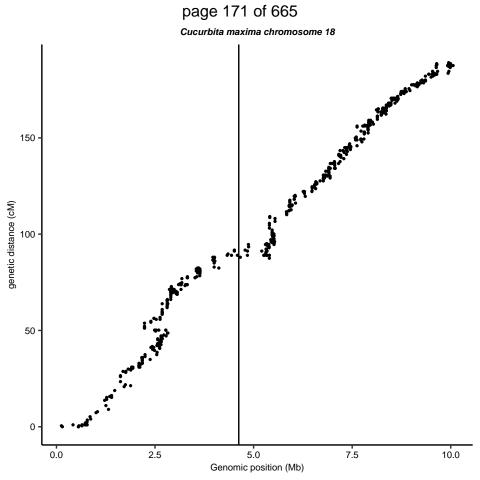


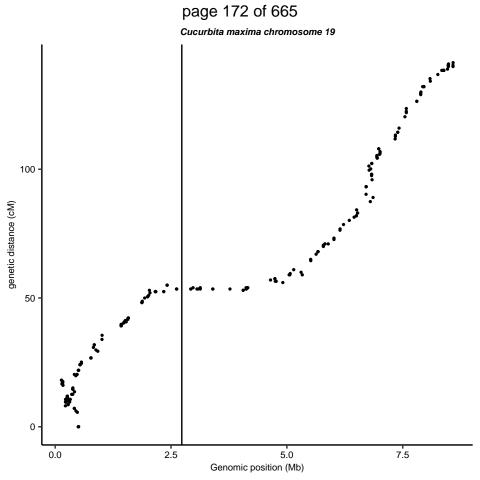


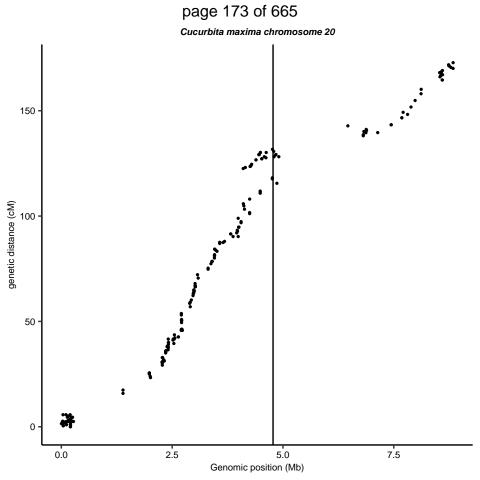


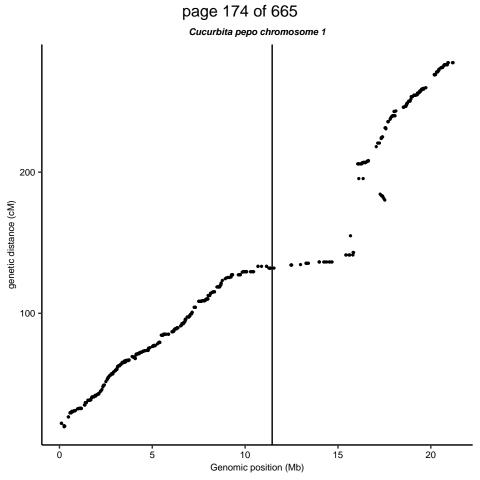


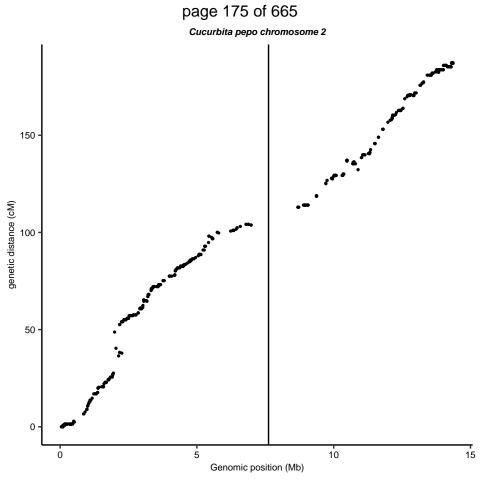


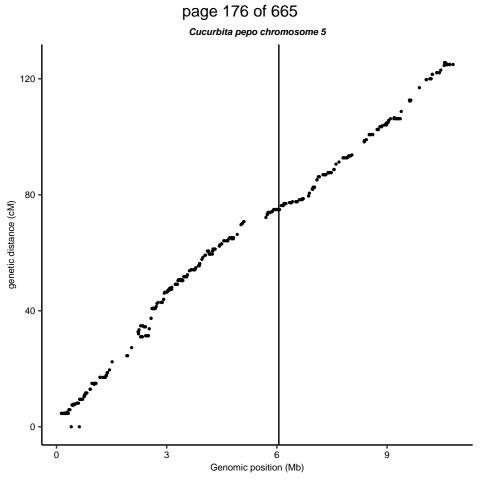


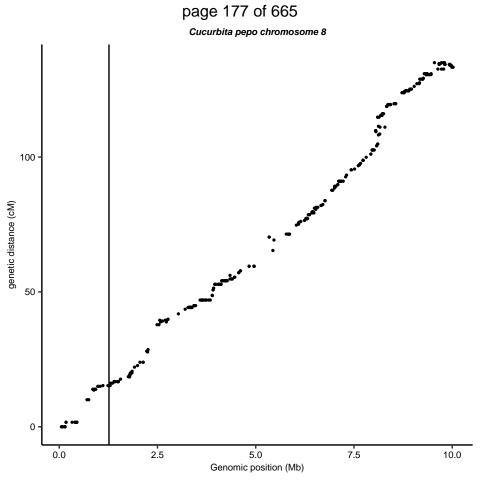




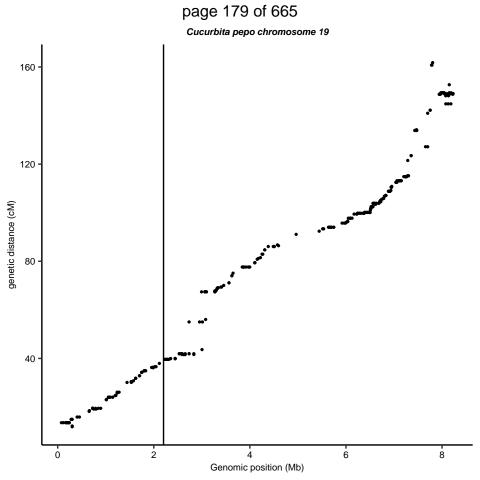


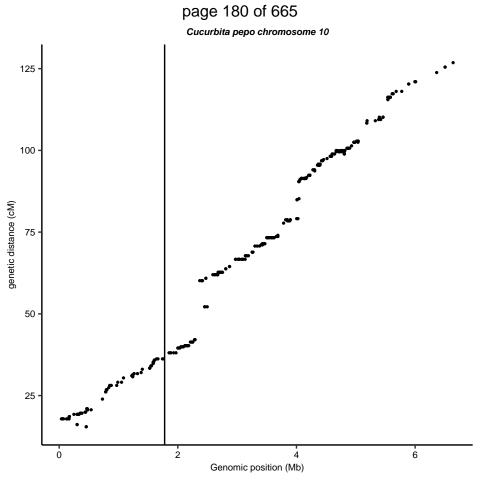


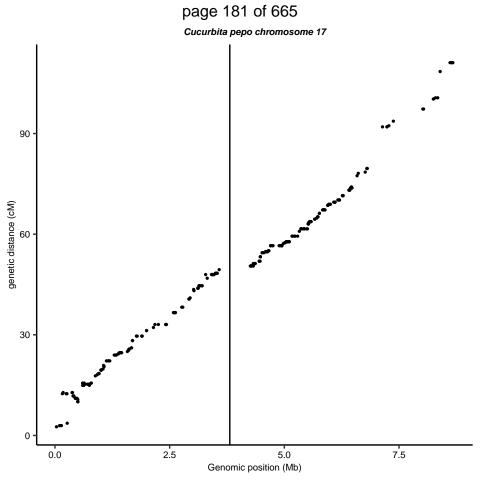




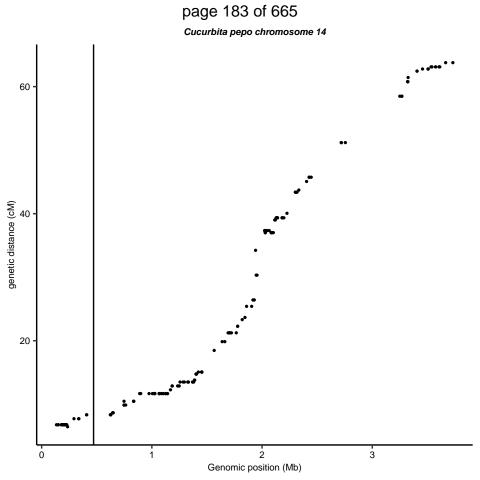
page 178 of 665 Cucurbita pepo chromosome 9 150 100 genetic distance (cM) 50 0 2.5 7.5 0.0 10.0 5.0 Genomic position (Mb)

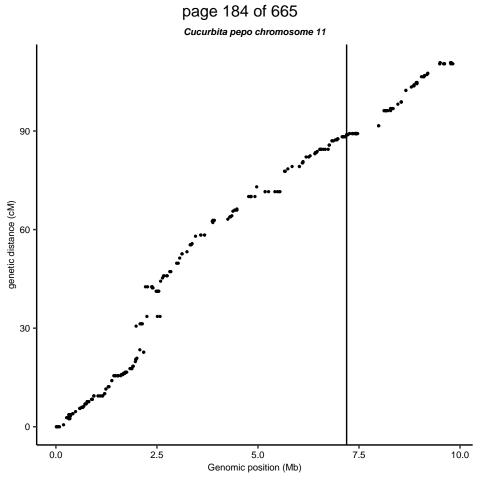


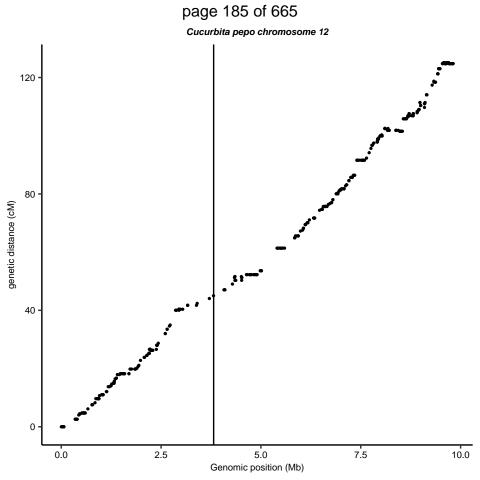


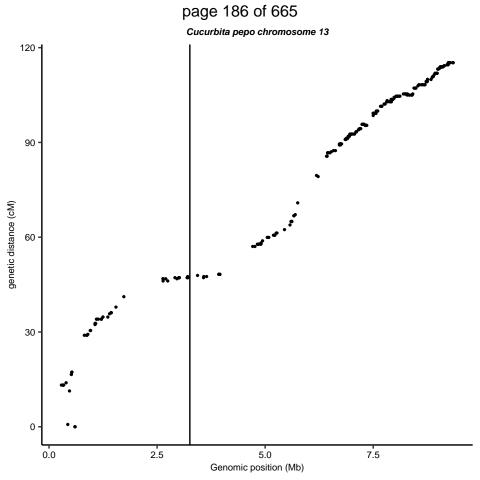


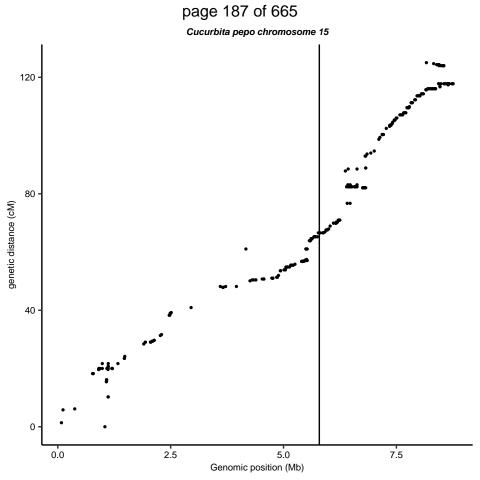
page 182 of 665 Cucurbita pepo chromosome 7 150 -100 genetic distance (cM) 50 0 0.0 2.5 7.5 5.0 10.0 Genomic position (Mb)

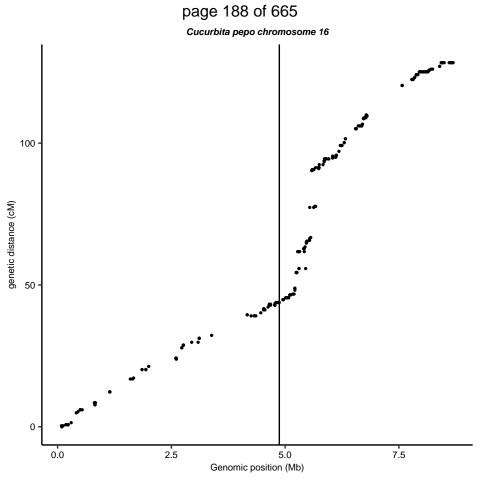


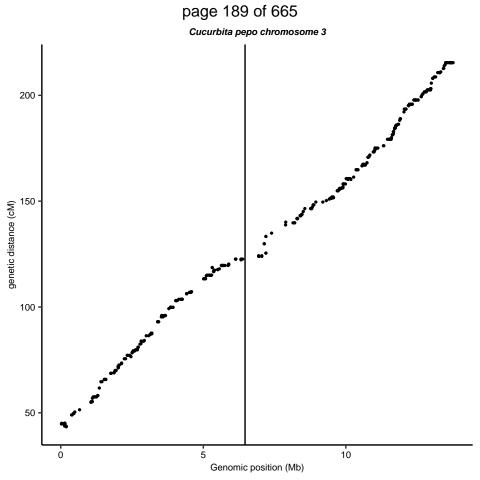


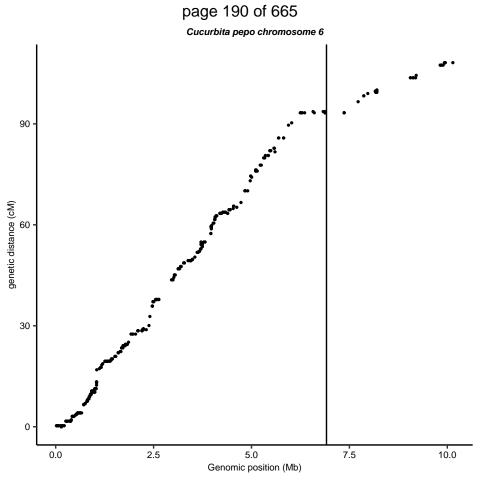


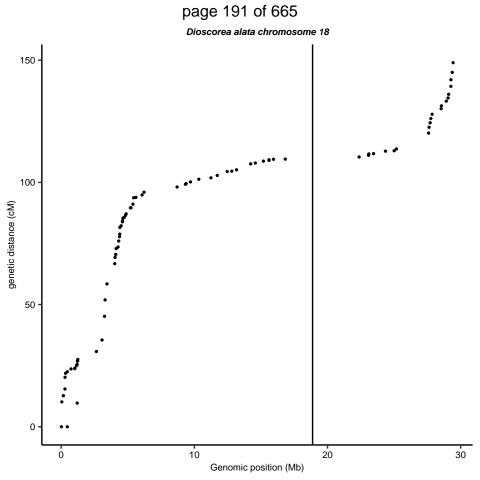






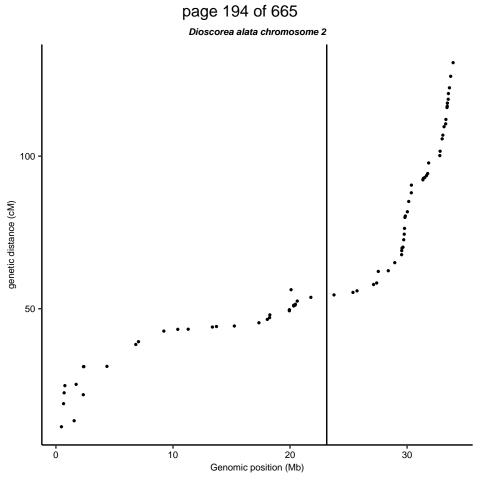






page 192 of 665 Dioscorea alata chromosome 1 125 100 genetic distance (cM) 75 -50 25 10 20 Genomic position (Mb)

page 193 of 665 Dioscorea alata chromosome 11 genetic distance (cM) Genomic position (Mb)



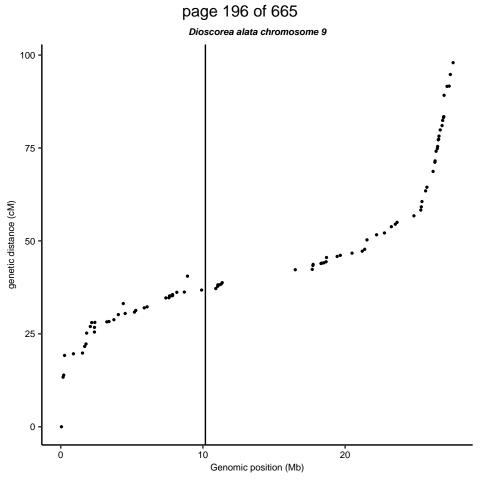
page 195 of 665 Dioscorea alata chromosome 7 150 -100 genetic distance (cM) 50 0 -

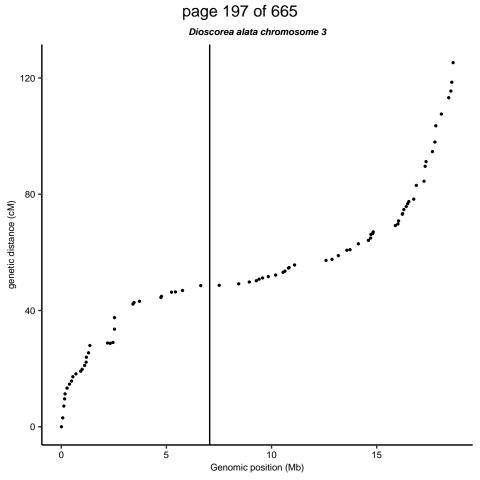
10 Genomic position (Mb)

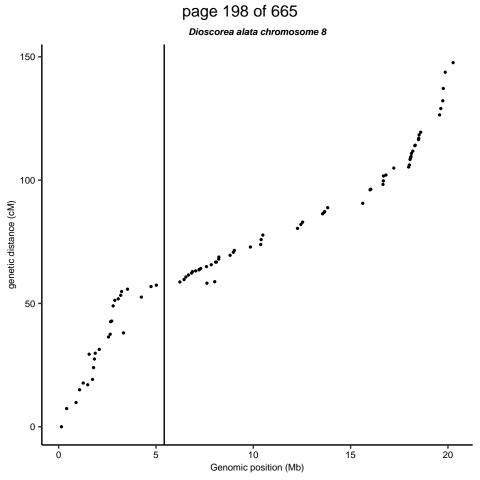
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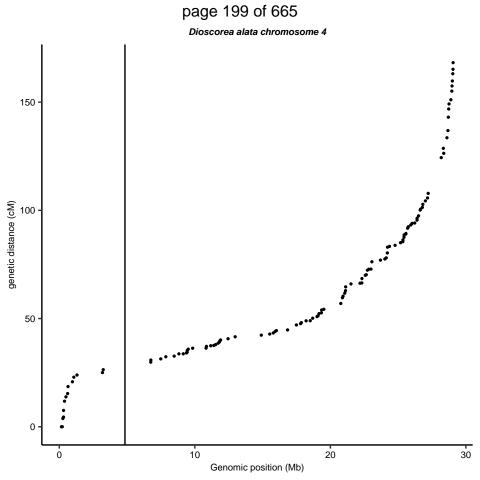
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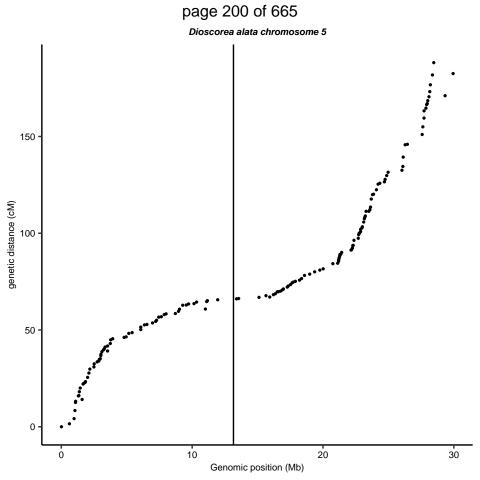
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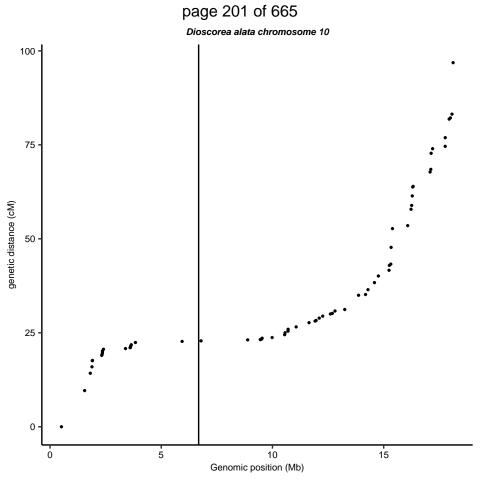


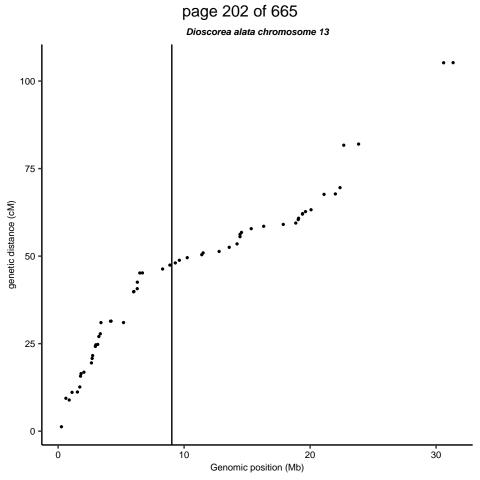




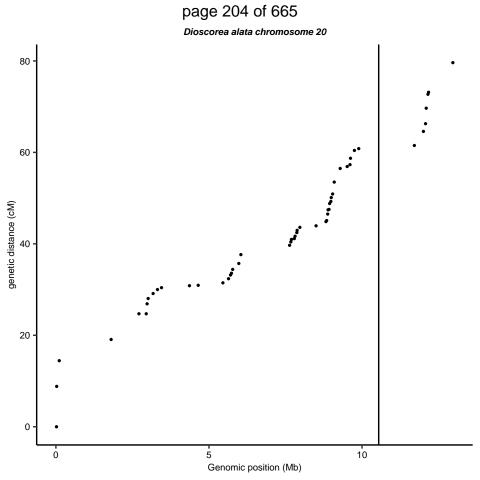


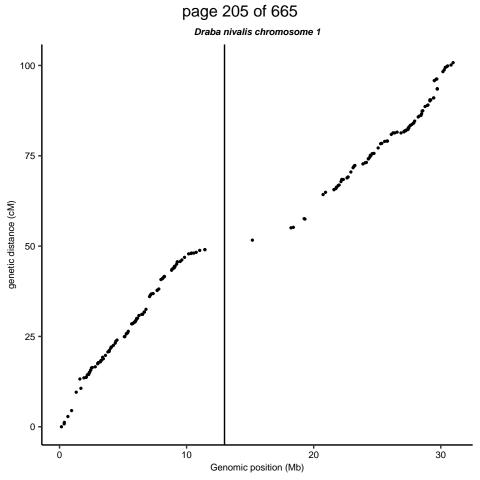


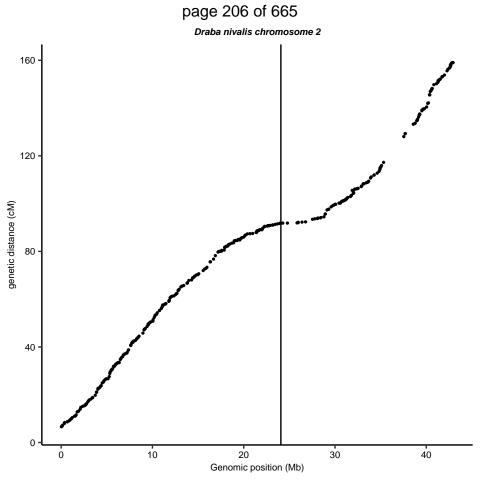


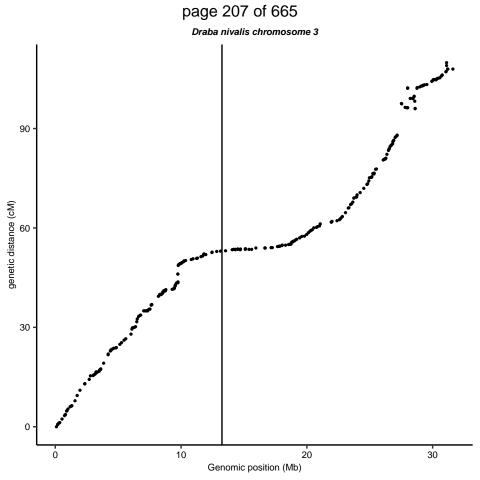


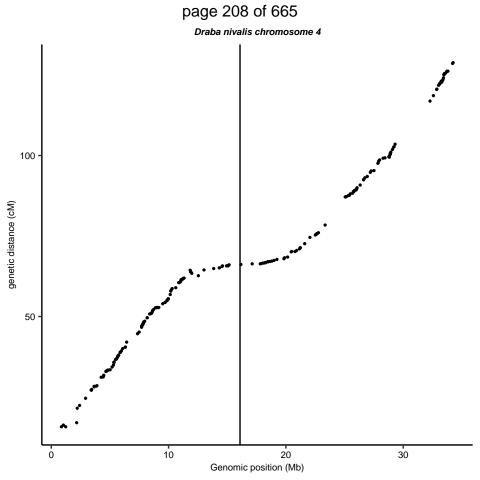
page 203 of 665 Dioscorea alata chromosome 17 150 genetic distance (cM) 100 50 -0 15 20 5 10 Genomic position (Mb)

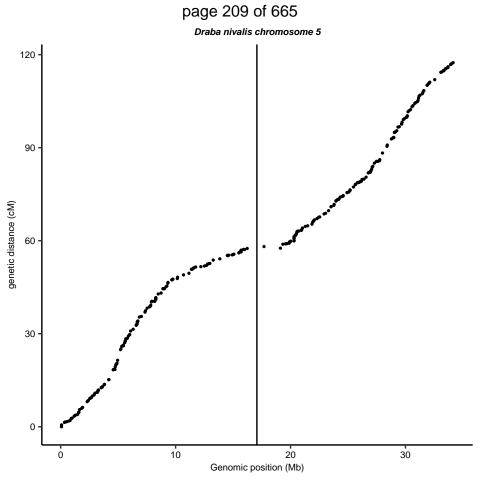


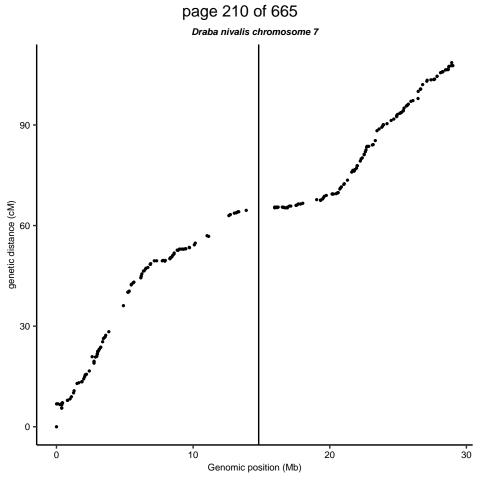


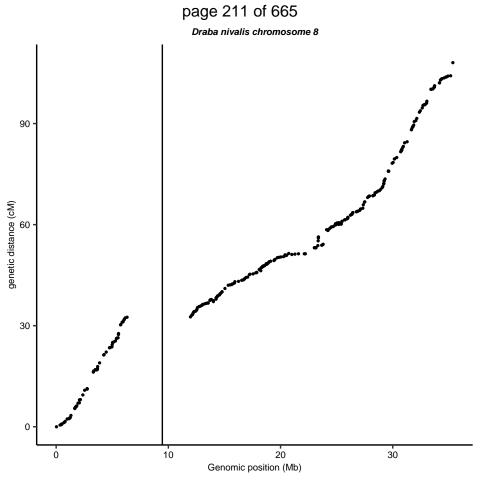


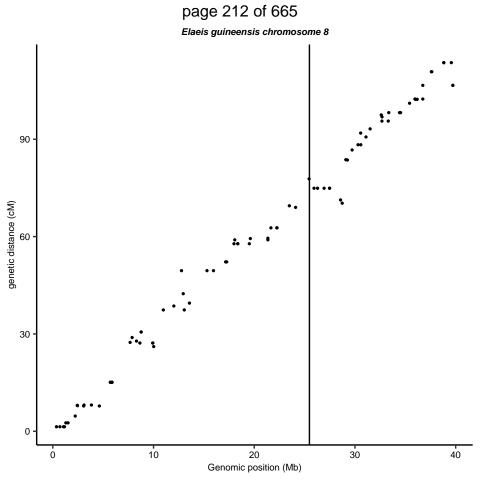


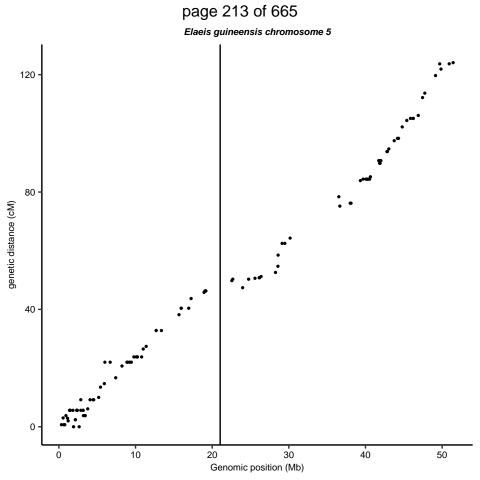




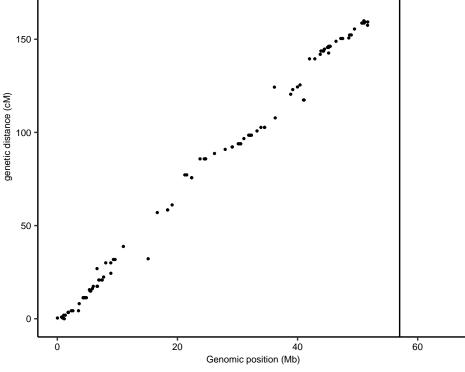


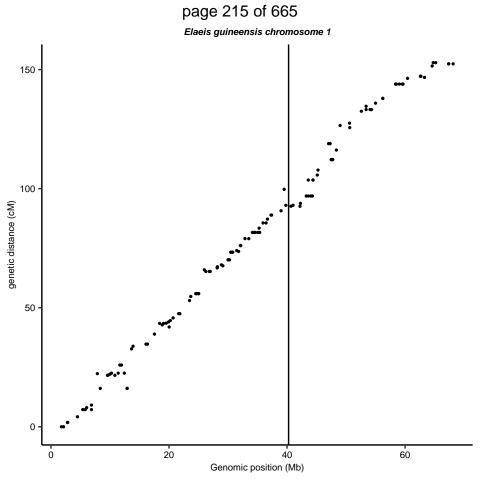


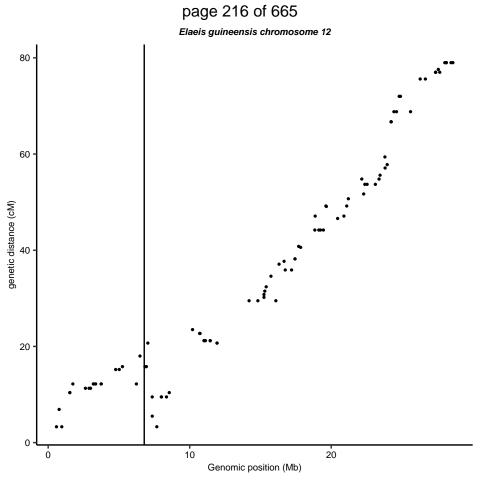




page 214 of 665 Elaeis guineensis chromosome 2 200 -150 genetic distance (cM) 100 50







page 217 of 665 Elaeis guineensis chromosome 10 100 -75 genetic distance (cM) 50 25 10 20 30 Genomic position (Mb)

page 218 of 665 Elaeis guineensis chromosome 14 40 genetic distance (cM) 0

10

Genomic position (Mb)

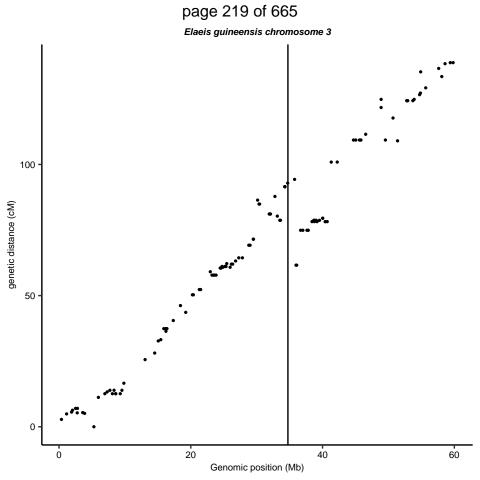
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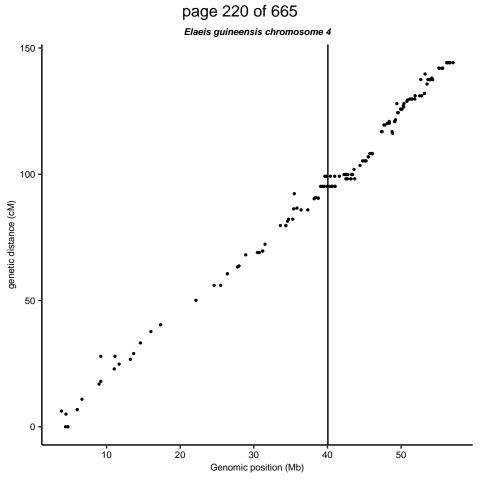
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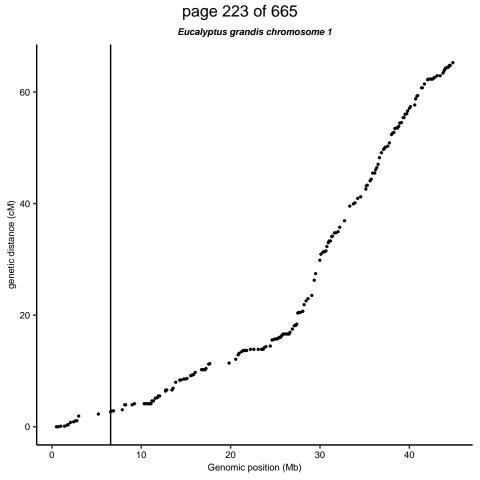
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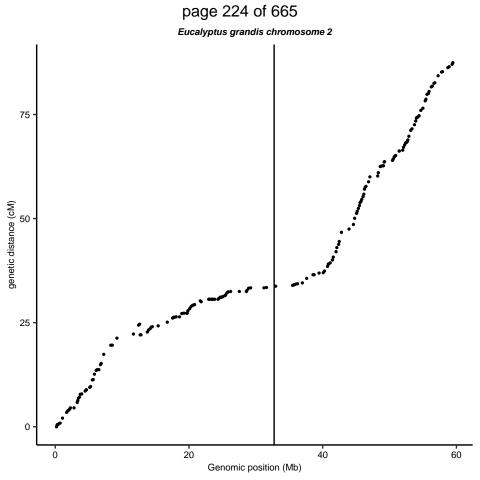


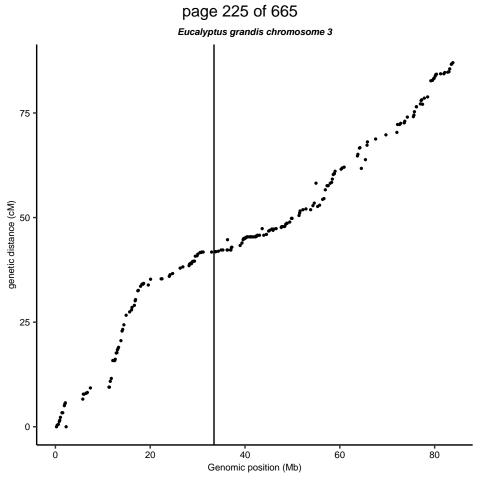


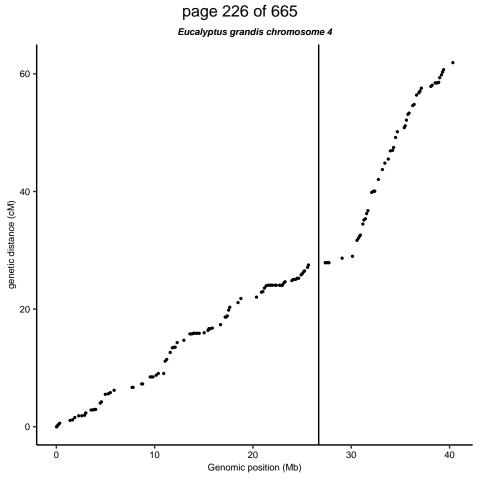
page 221 of 665 Elaeis guineensis chromosome 6 100 genetic distance (cM) 50 10 30 20 40 Genomic position (Mb)

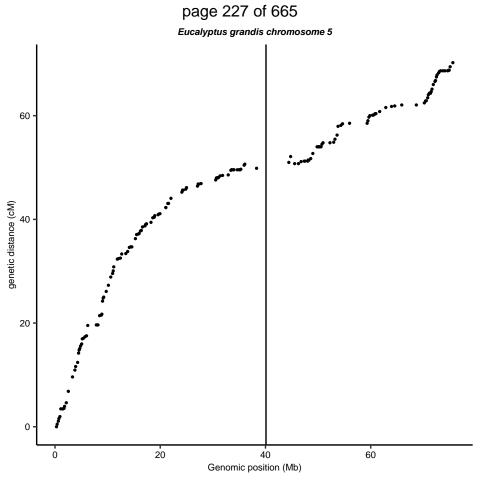
page 222 of 665 Elaeis guineensis chromosome 7 125 -genetic distance (cM) Genomic position (Mb)

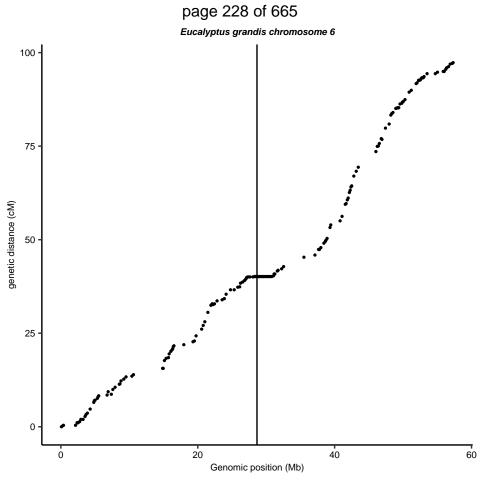


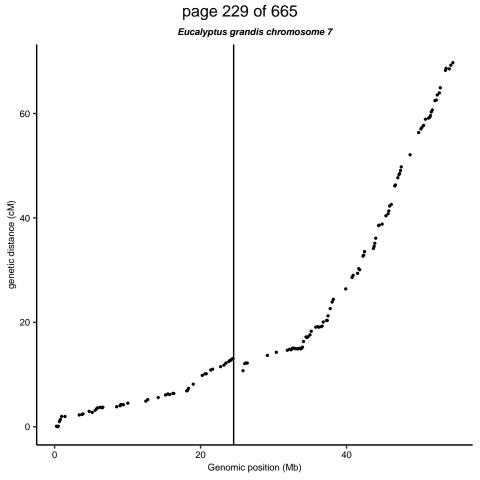


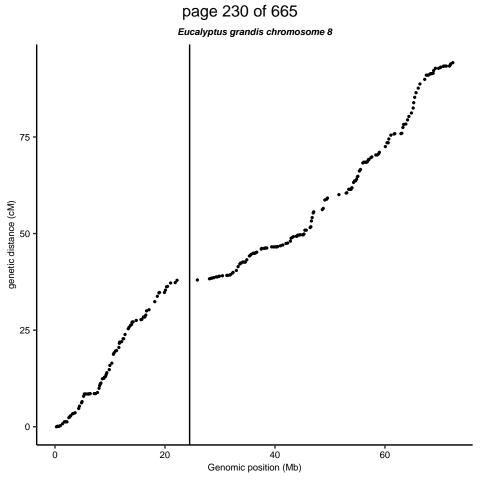


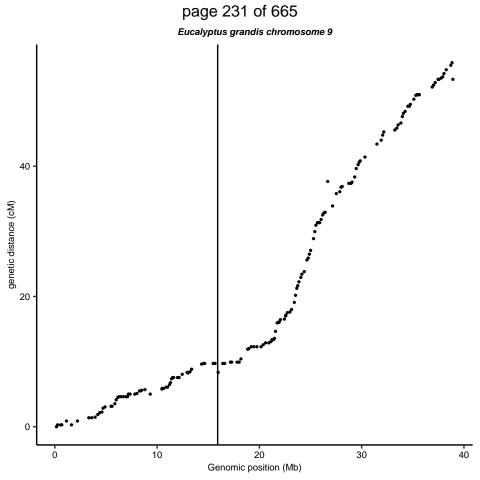


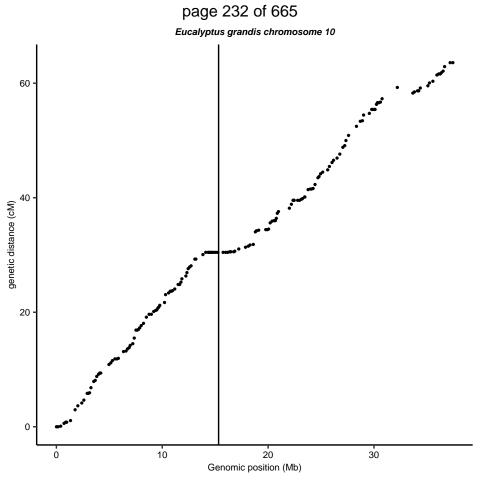


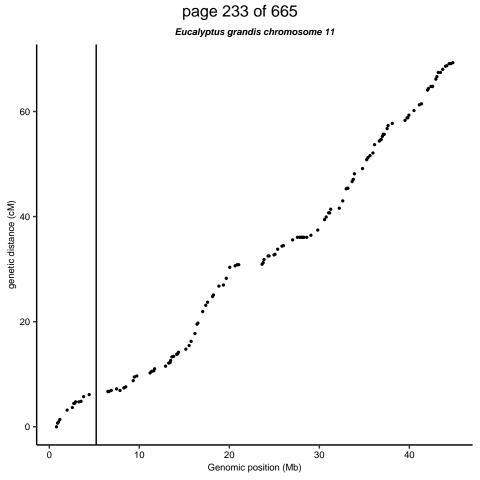


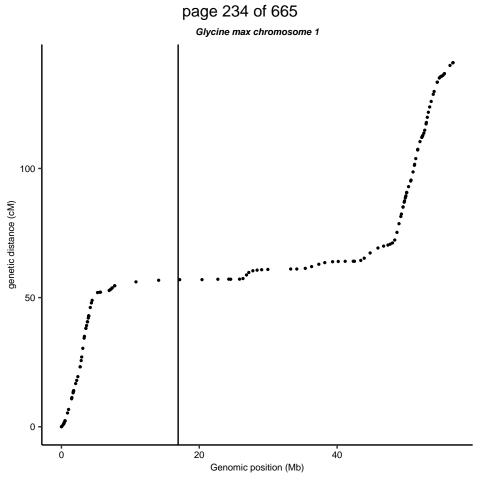


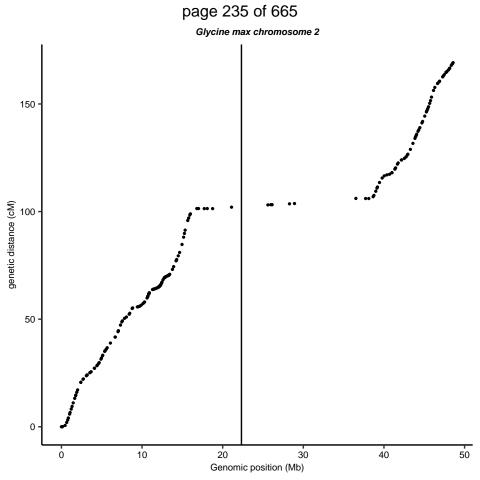


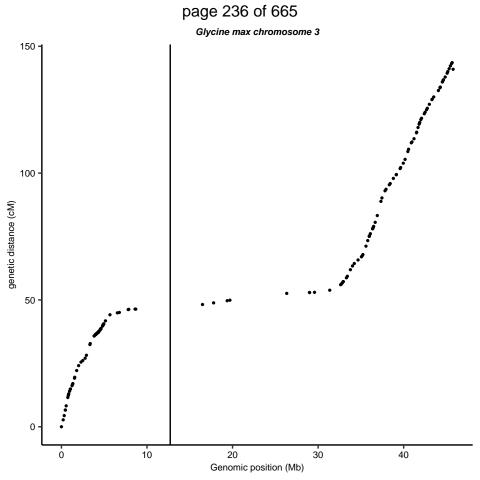


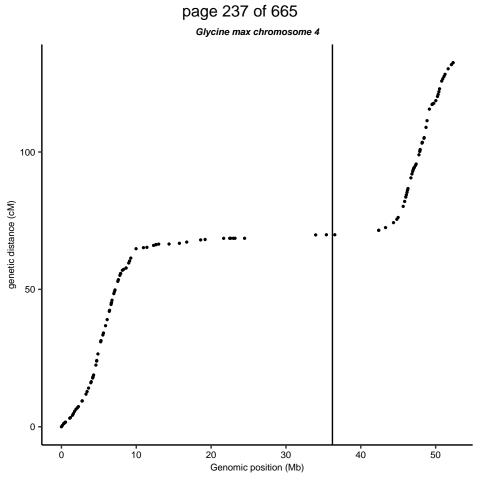


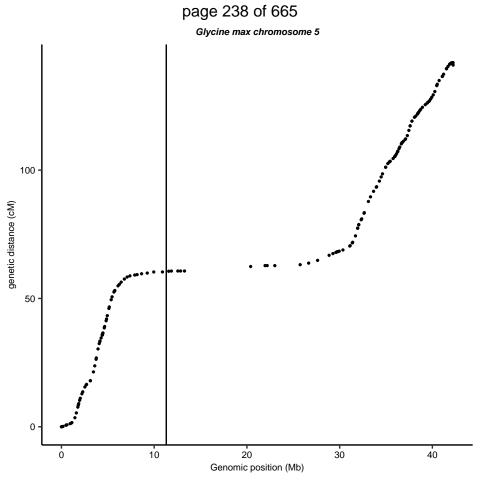


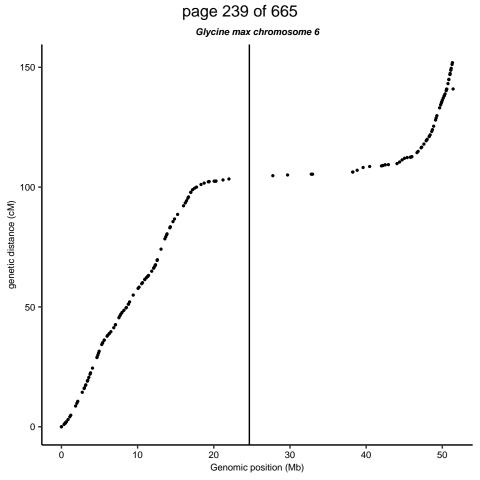


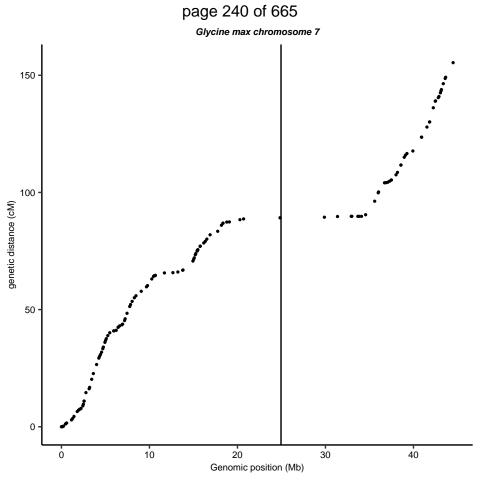


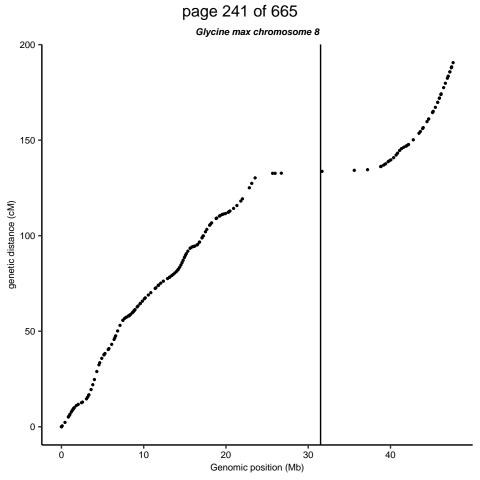


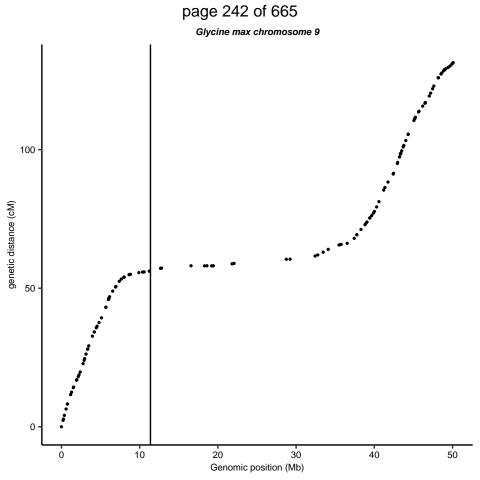


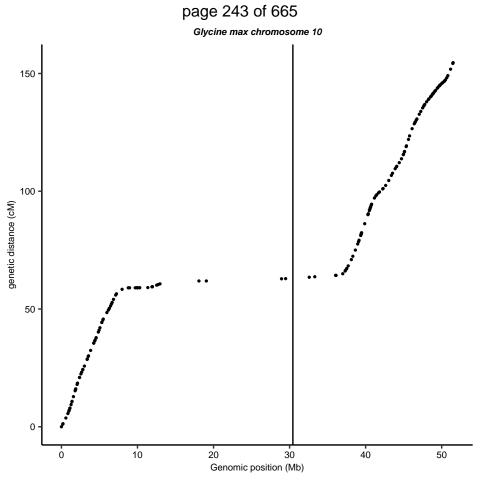


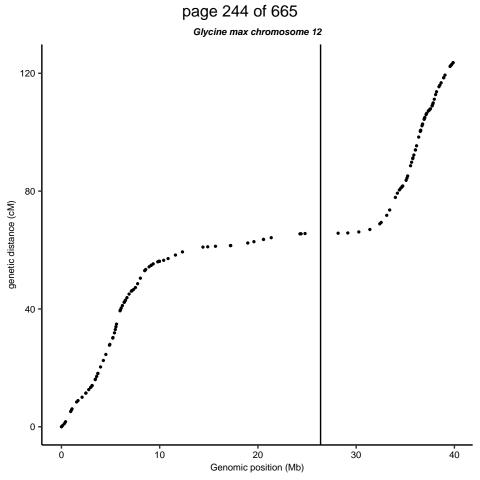


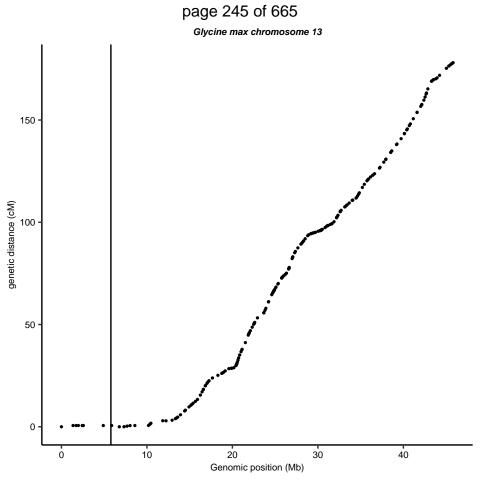


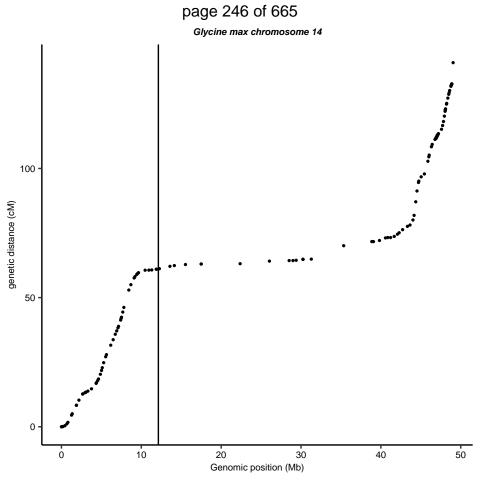


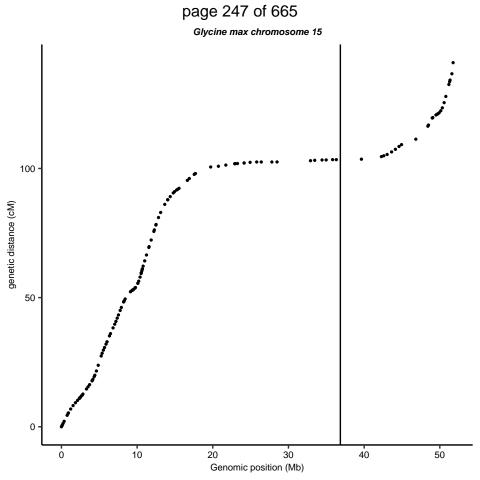


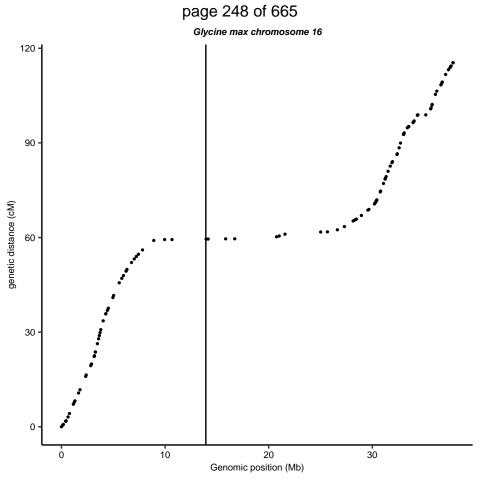


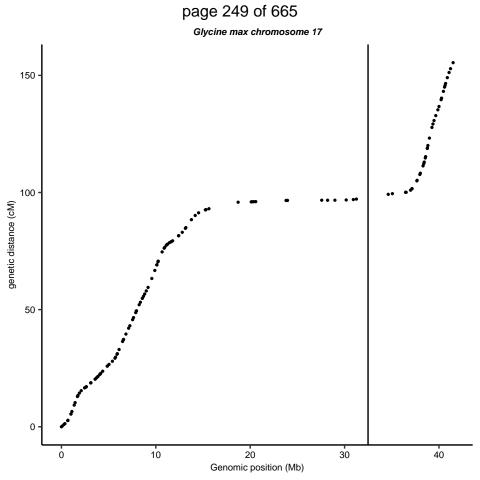


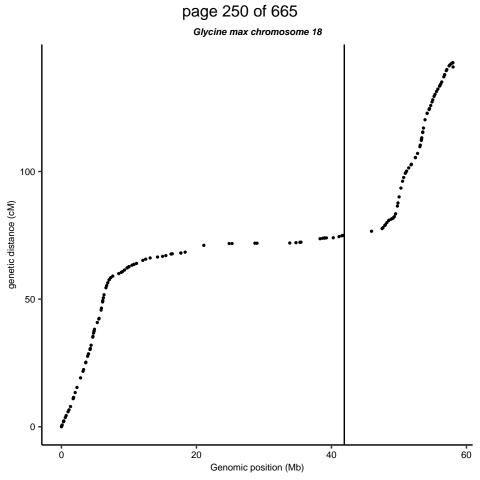


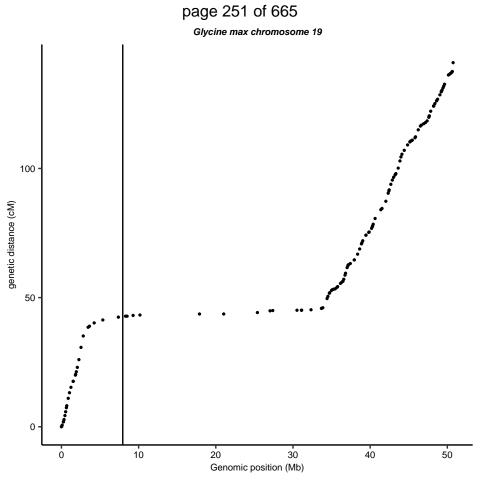


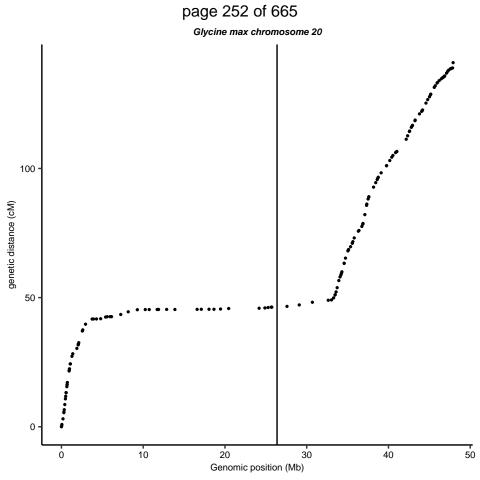






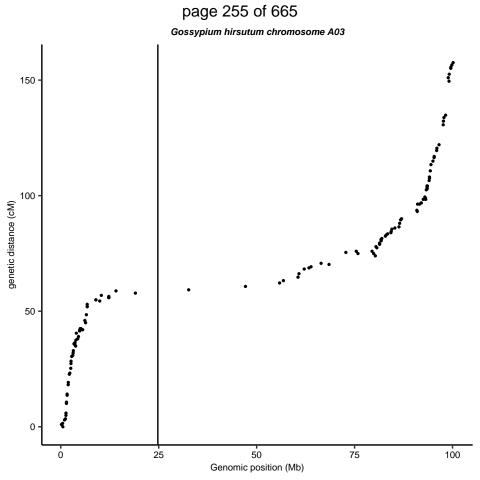




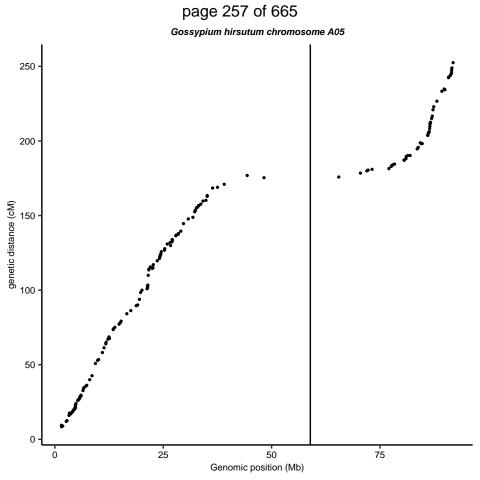


page 253 of 665 Gossypium hirsutum chromosome A01 genetic distance (cM) Genomic position (Mb)

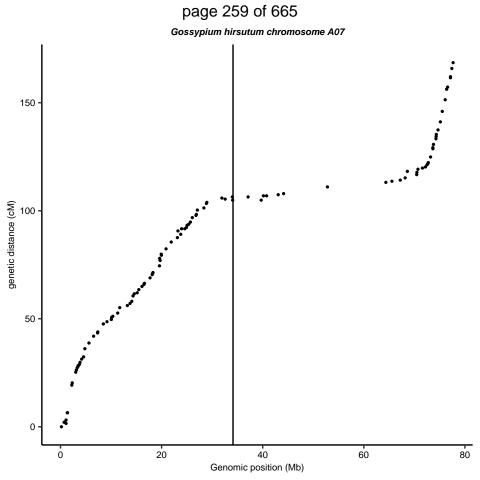
page 254 of 665 Gossypium hirsutum chromosome A02 genetic distance (cM) Genomic position (Mb)



page 256 of 665 Gossypium hirsutum chromosome A04 100 75 genetic distance (cM) 50 25 0 20 60 40 Genomic position (Mb)



page 258 of 665 Gossypium hirsutum chromosome A06 genetic distance (cM) Genomic position (Mb)



page 260 of 665 Gossypium hirsutum chromosome A08 genetic distance (cM) Genomic position (Mb)

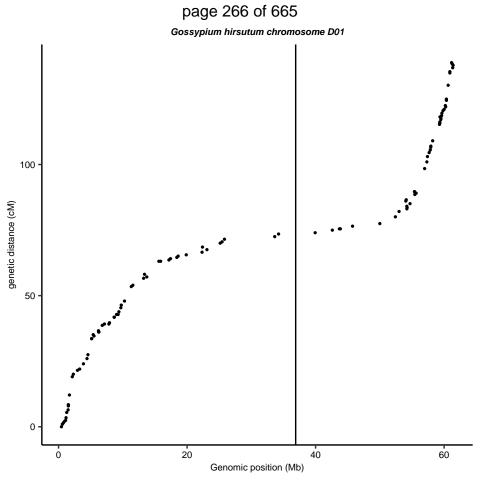
page 261 of 665 Gossypium hirsutum chromosome A09 150 genetic distance (cM) 100 -50 60 20 40 Genomic position (Mb)

page 262 of 665 Gossypium hirsutum chromosome A10 150 100 genetic distance (cM) 50 0 25 75 50 100 Genomic position (Mb)

page 263 of 665 Gossypium hirsutum chromosome A11 genetic distance (cM) 5 Genomic position (Mb)

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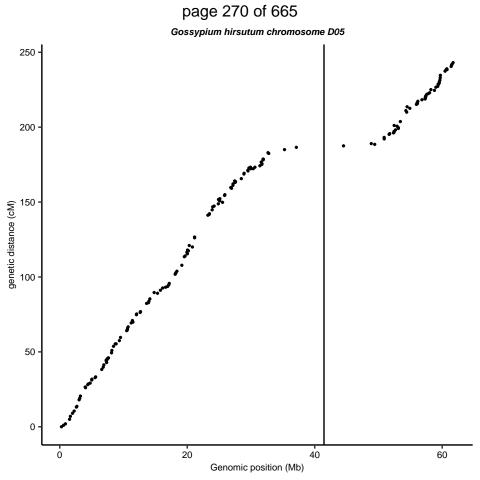
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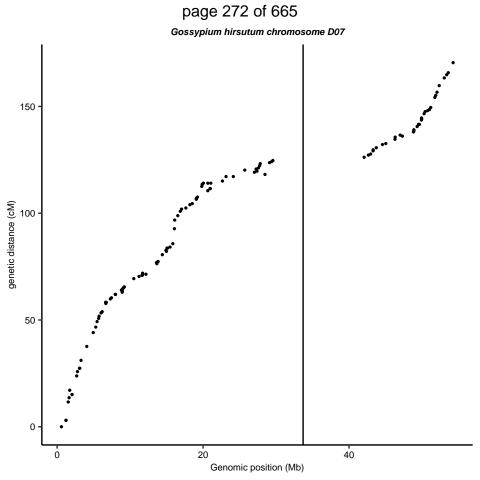
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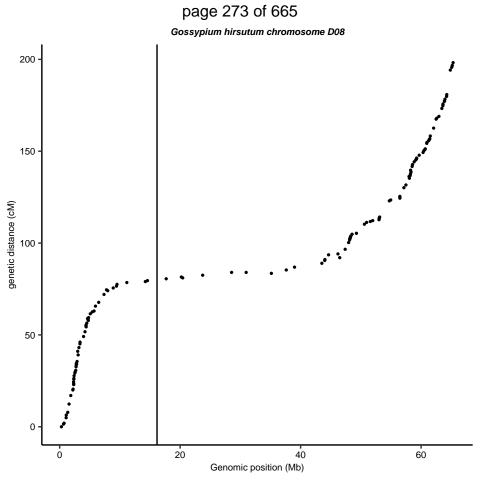
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page 269 of 665 Gossypium hirsutum chromosome D04 genetic distance (cM) Genomic position (Mb)

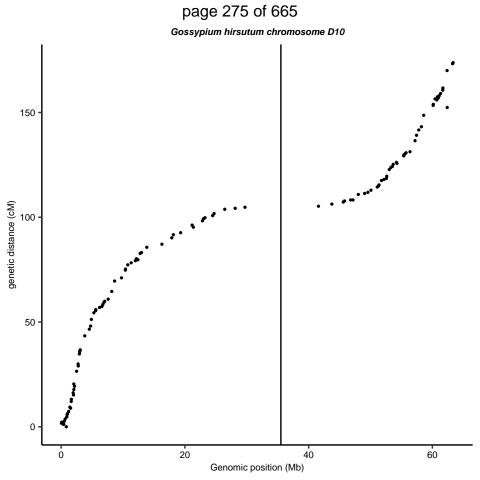


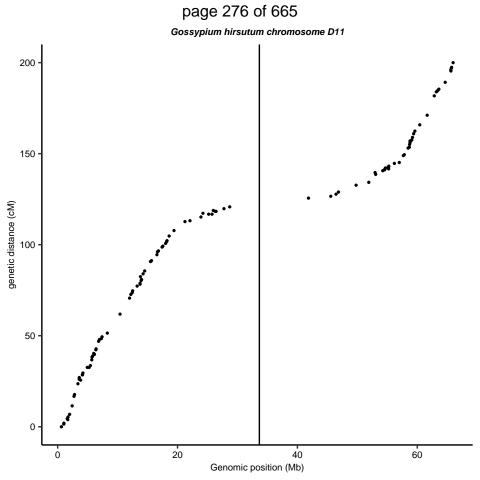
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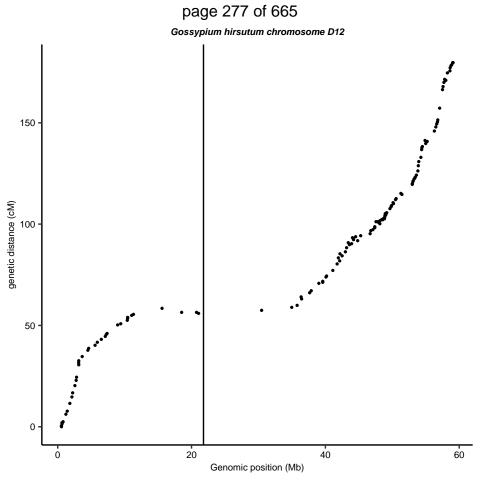


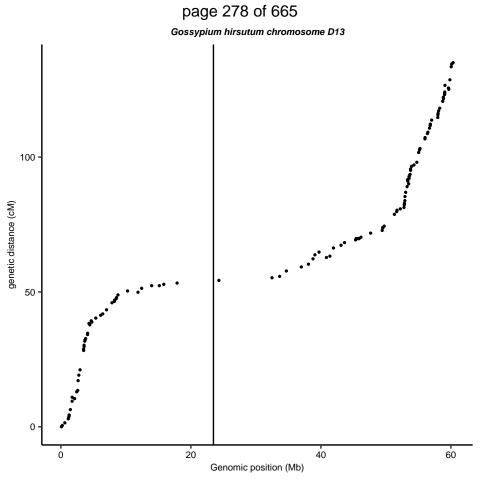


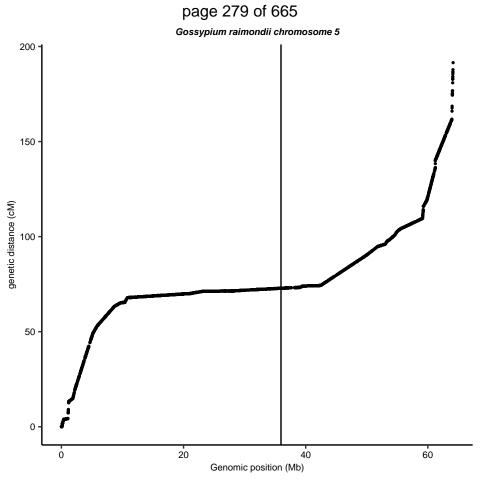
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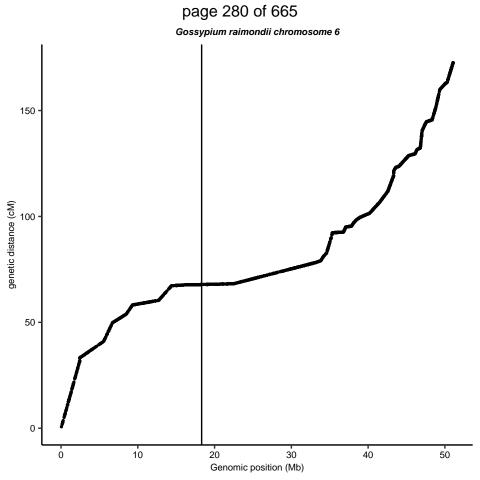


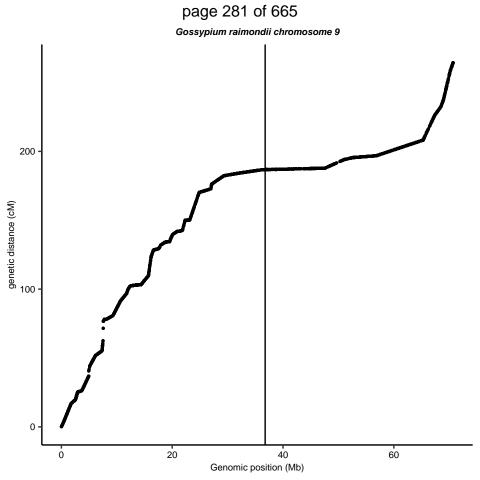


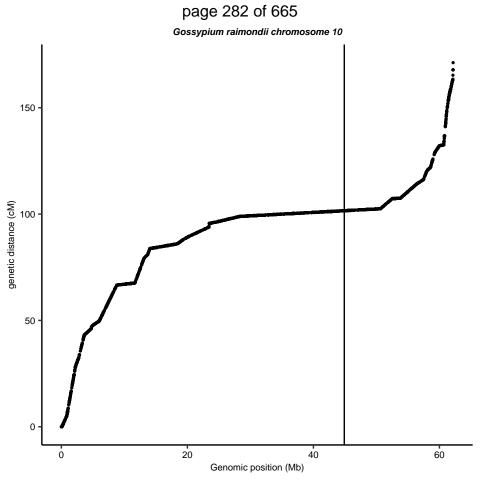


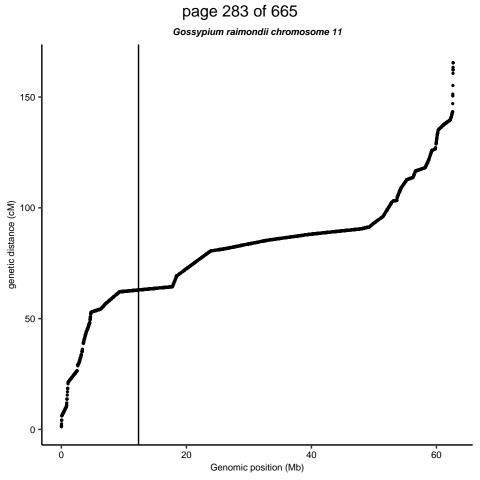


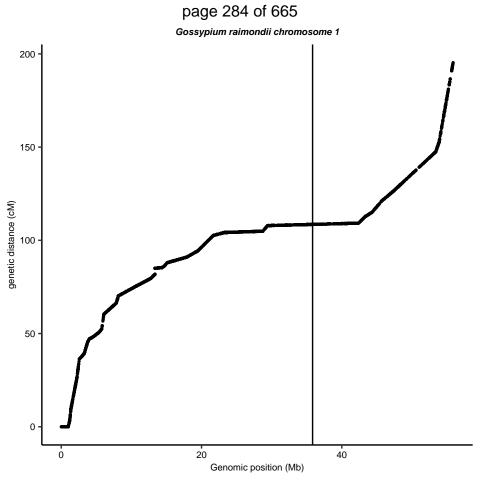


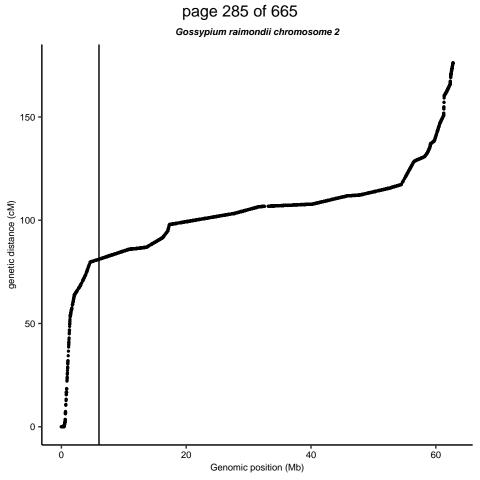


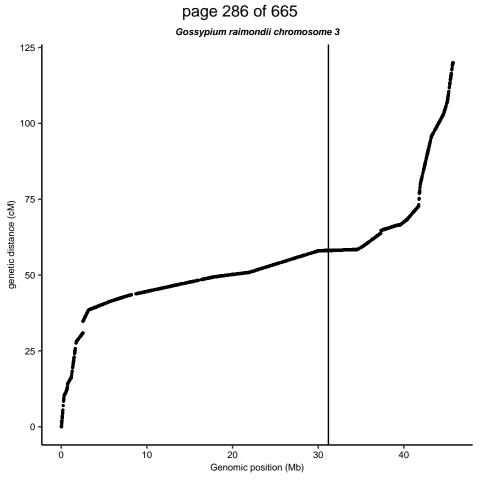


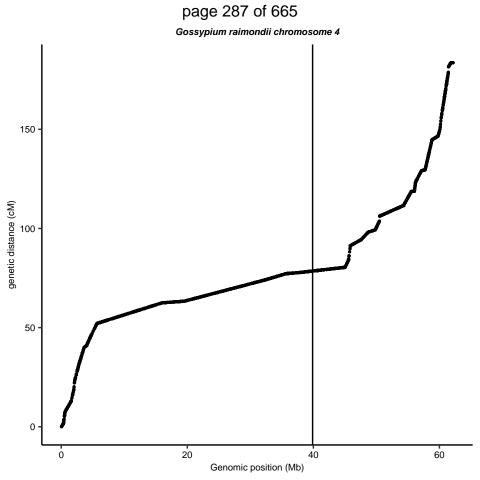


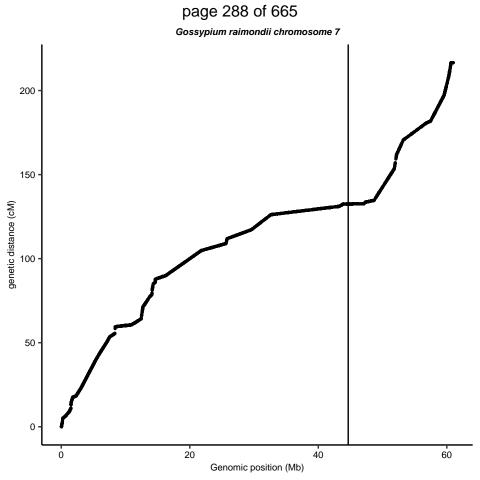


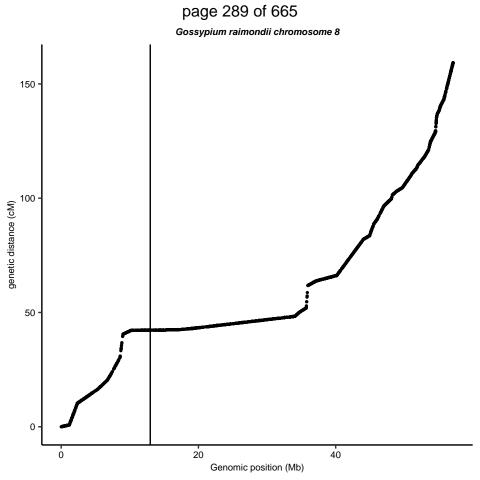


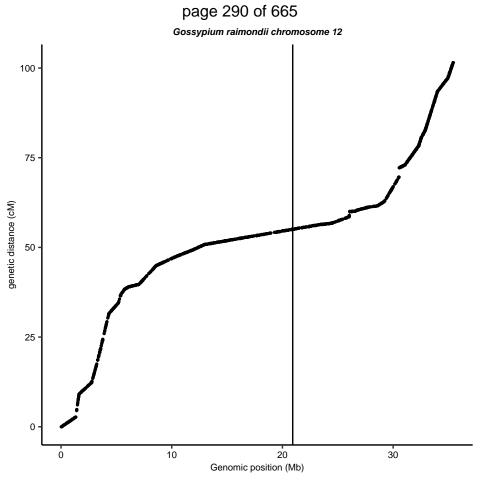


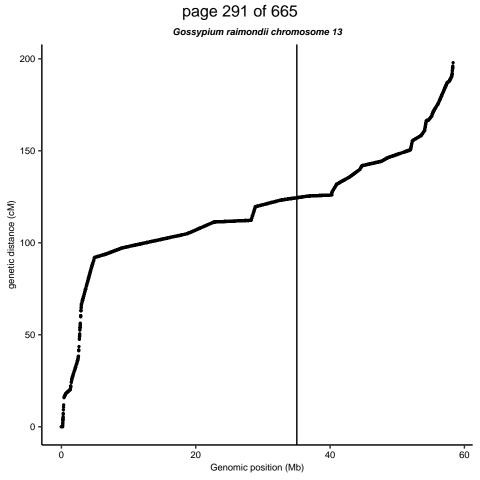


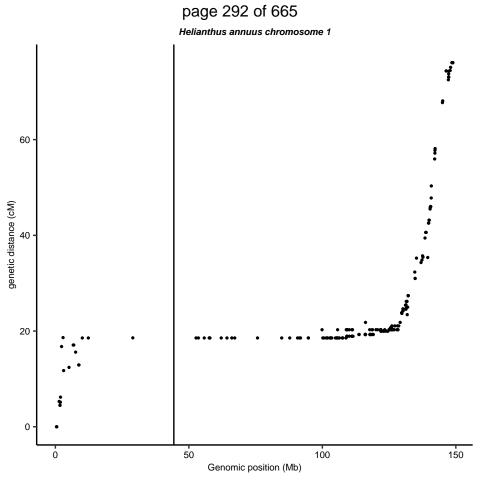


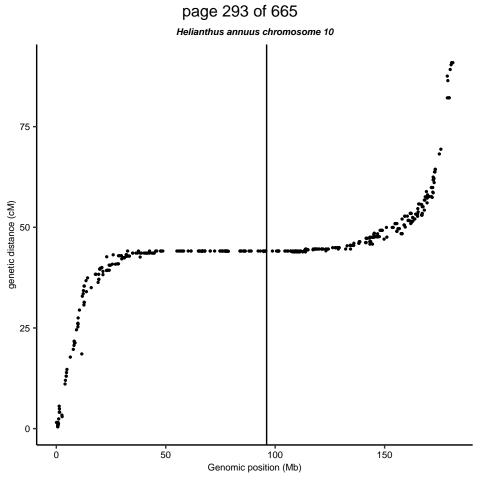


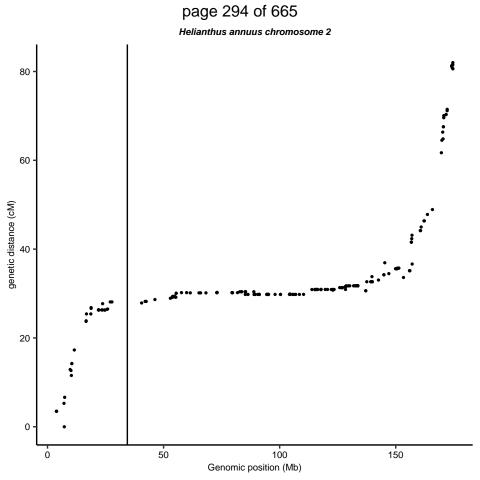


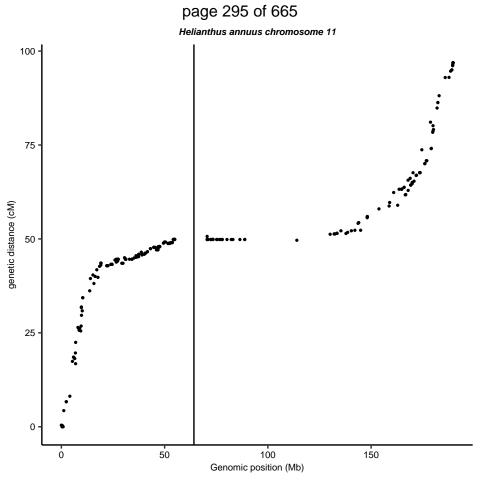


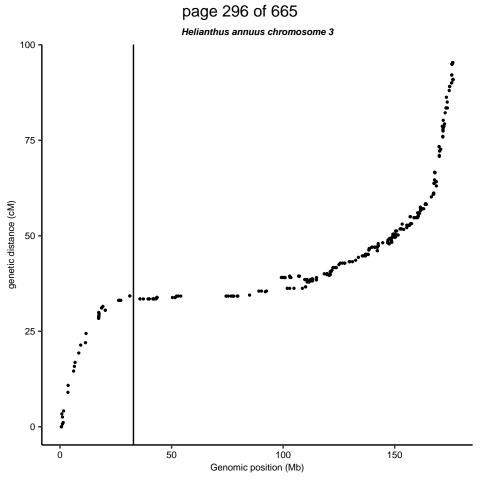


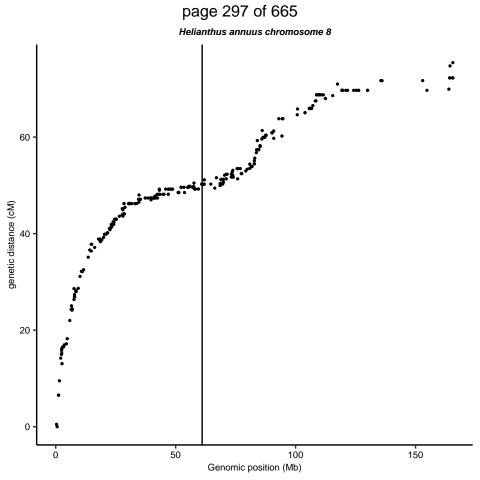




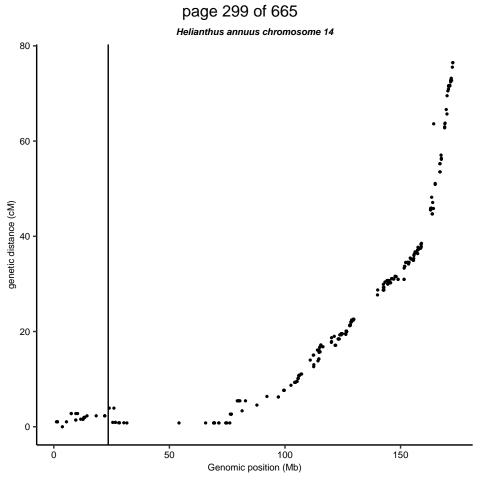


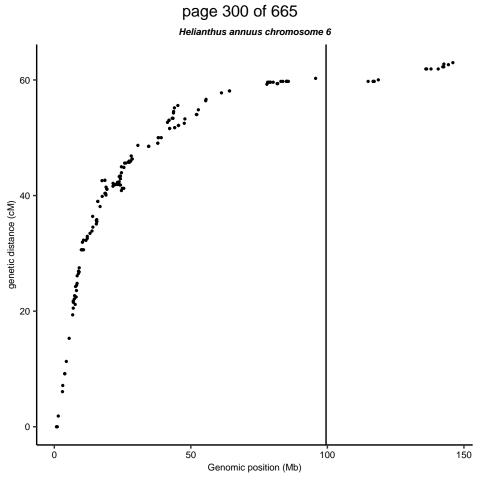


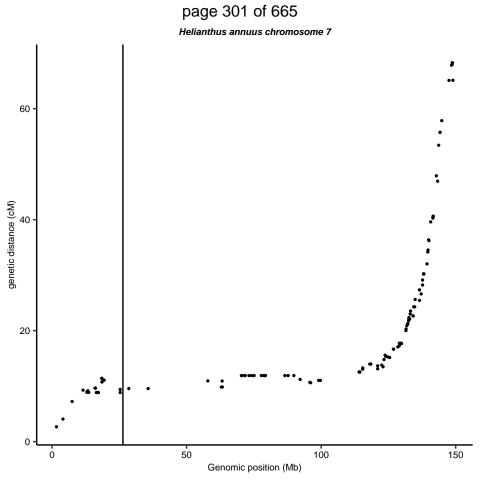


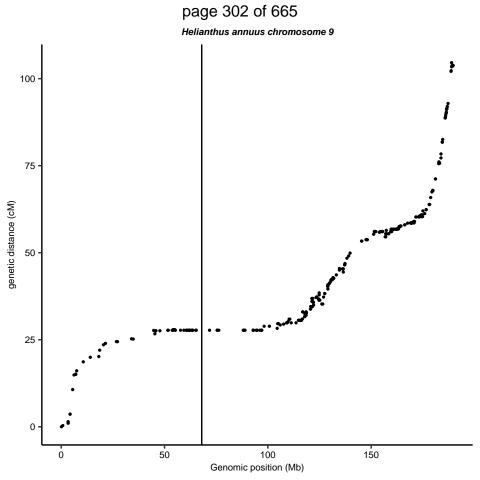


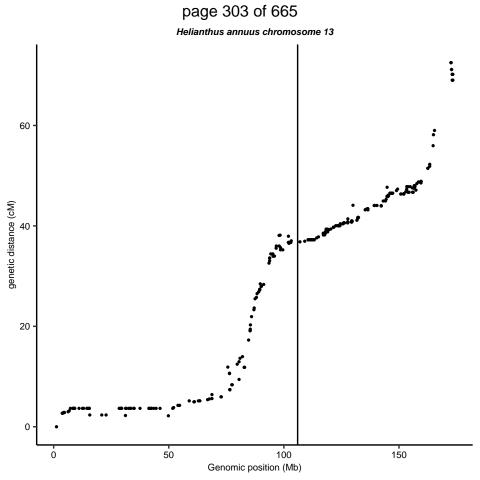
page 298 of 665 Helianthus annuus chromosome 4 100 75 genetic distance (cM) 50 25 50 150 100 200 Genomic position (Mb)

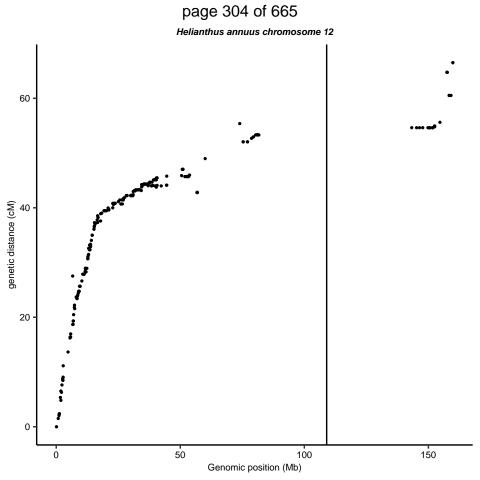


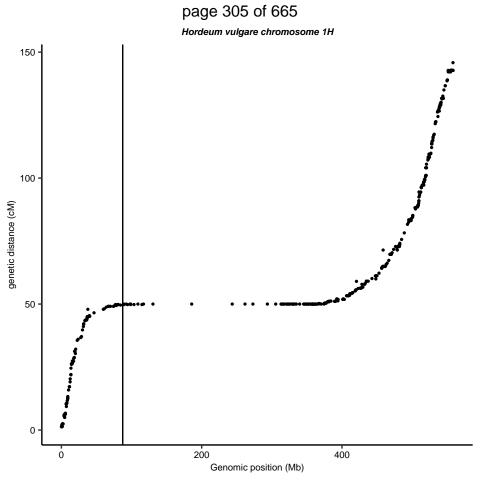


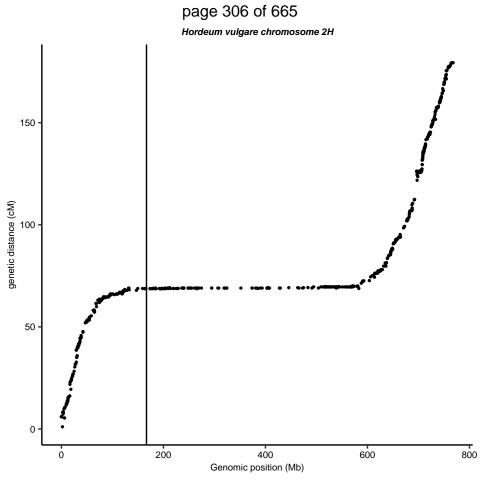


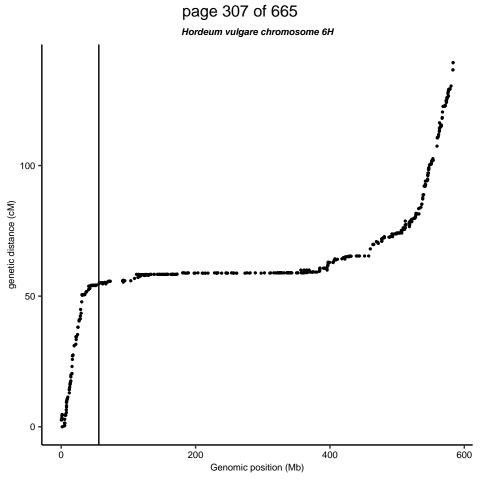


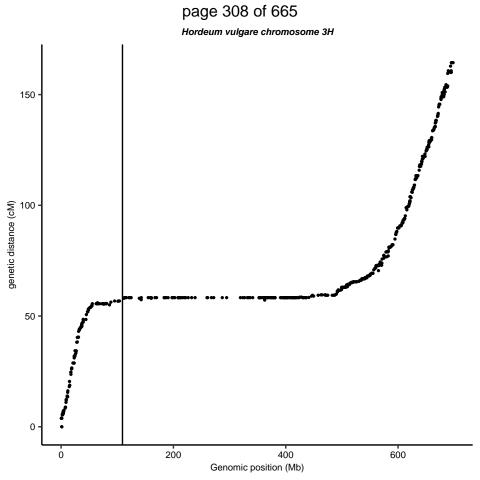


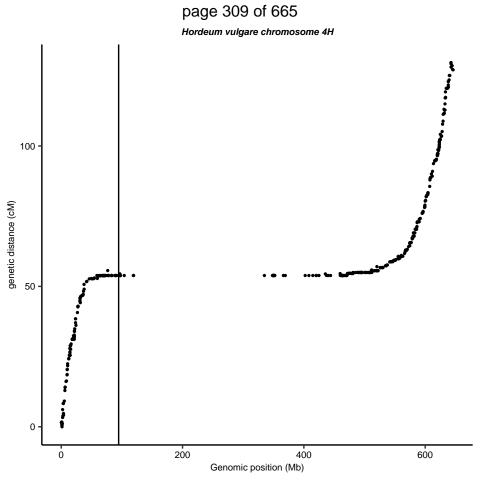


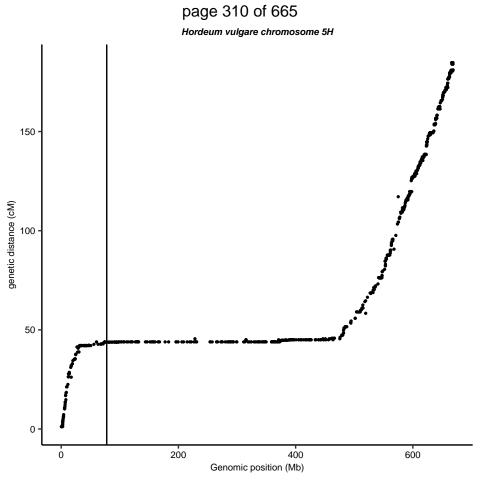


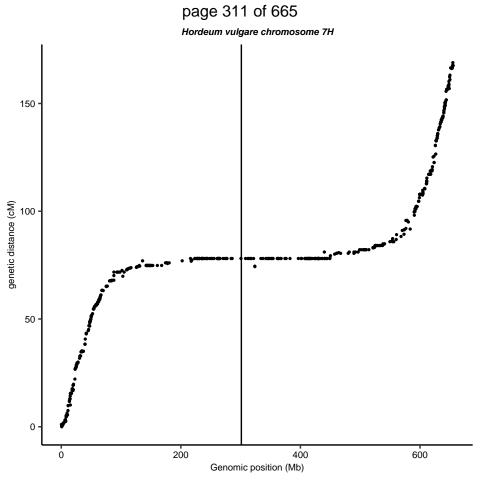


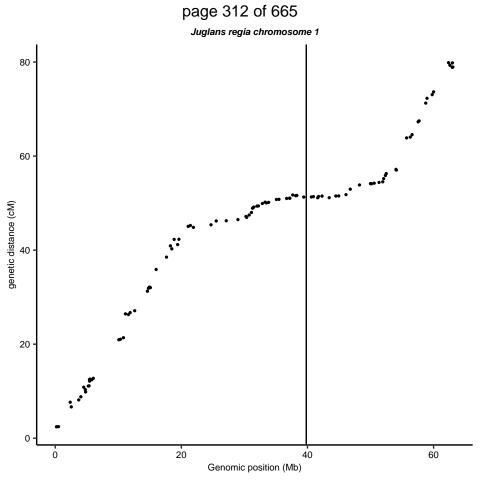


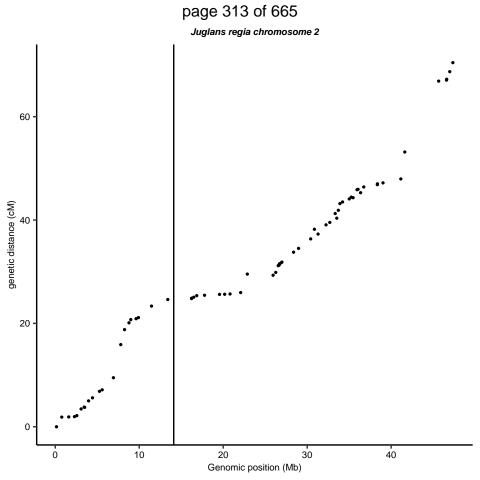


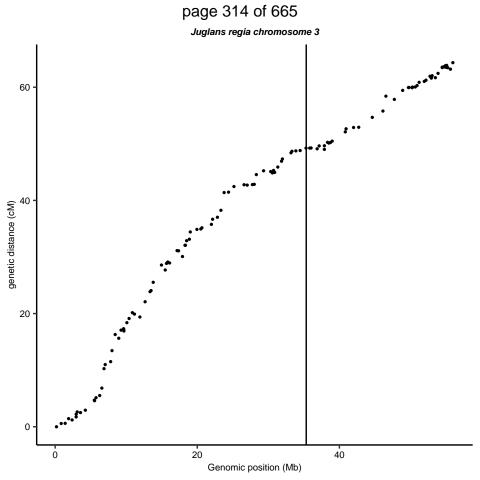


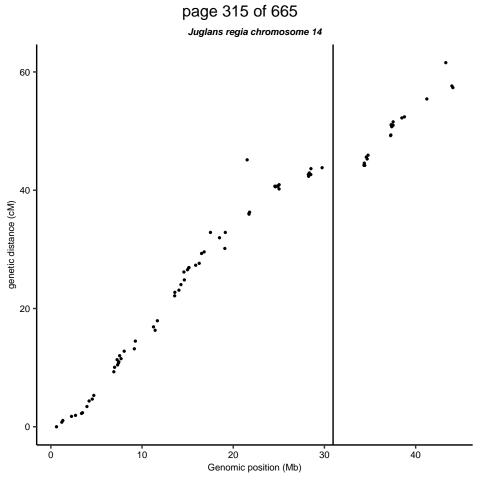


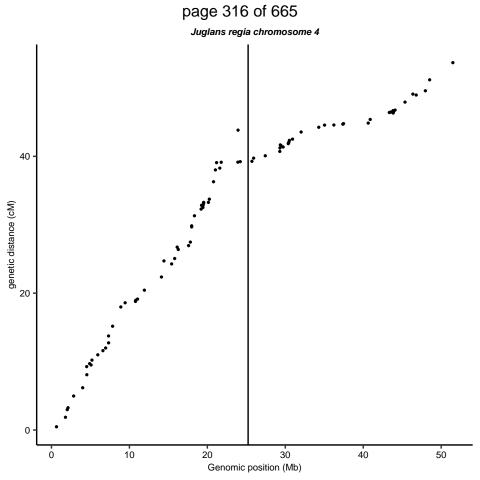




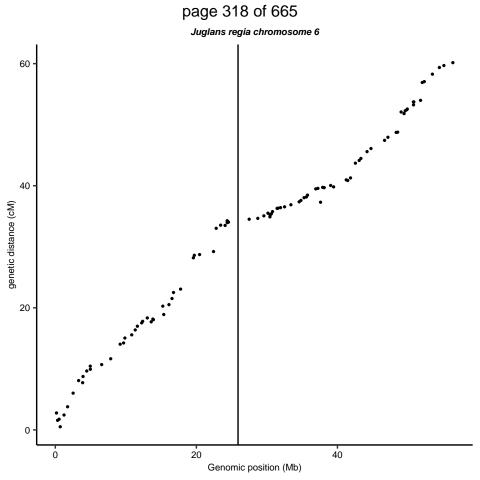


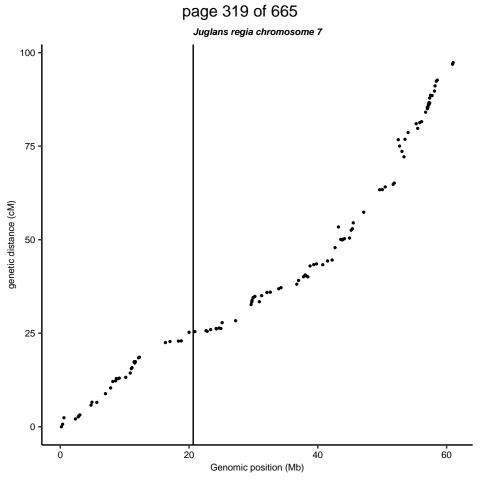


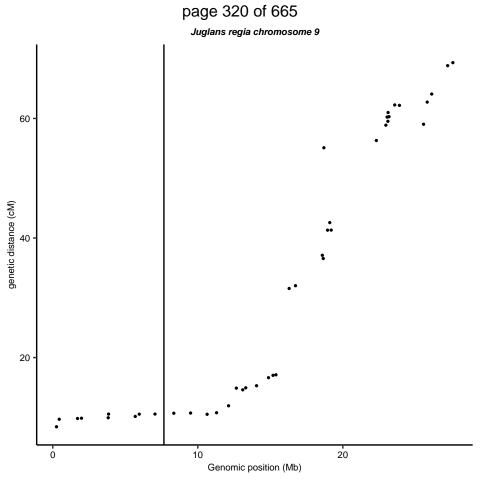


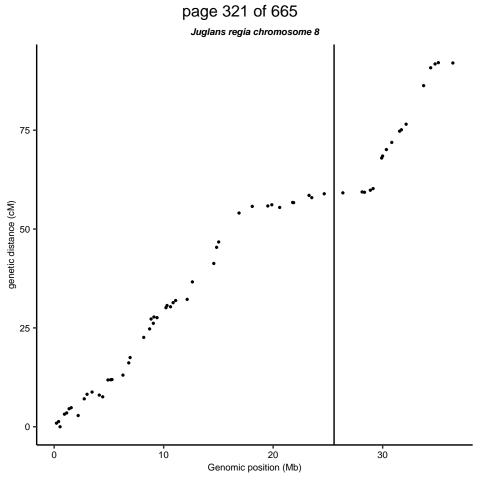


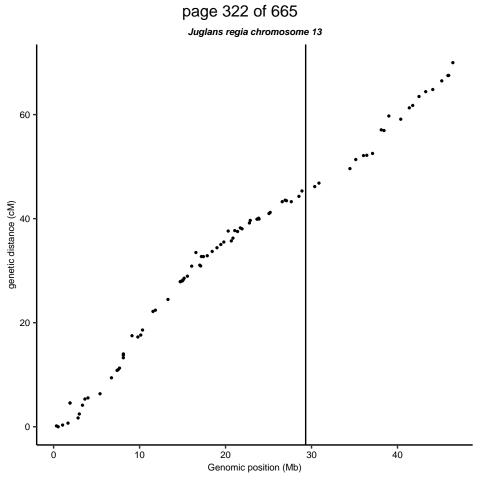
page 317 of 665 Juglans regia chromosome 5 genetic distance (cM) Genomic position (Mb)

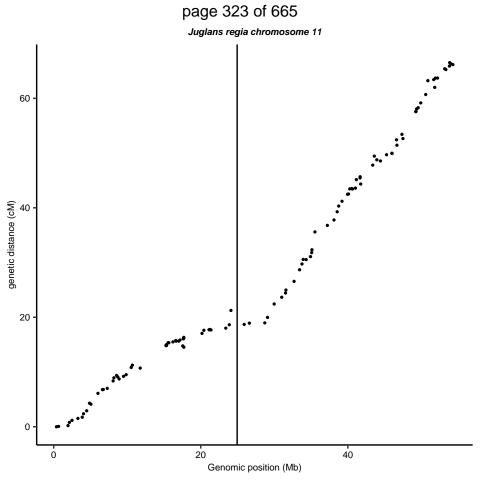


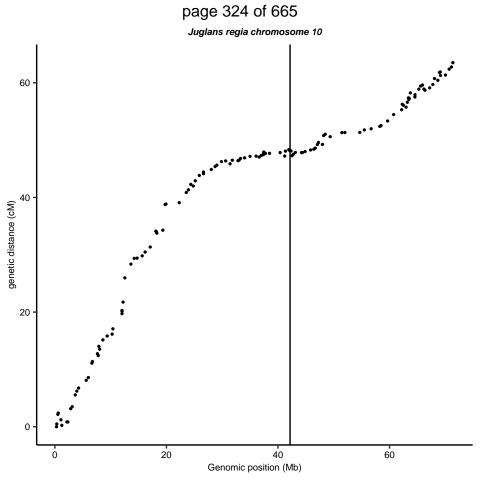


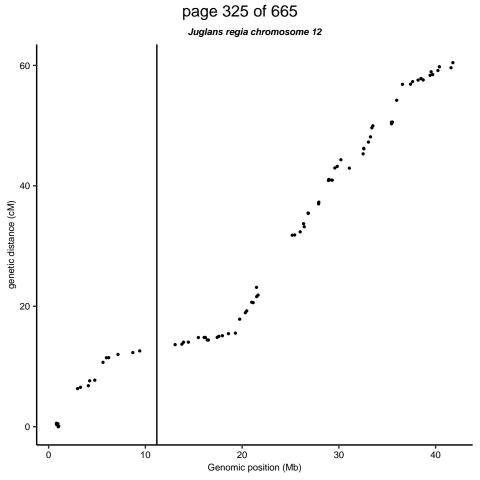


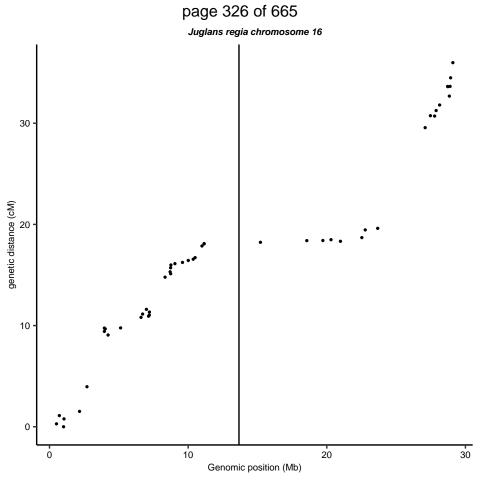


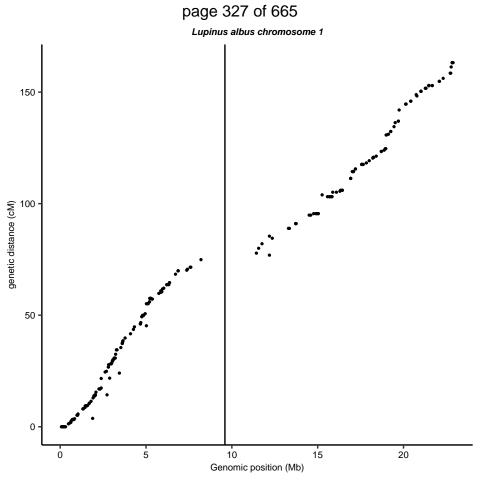


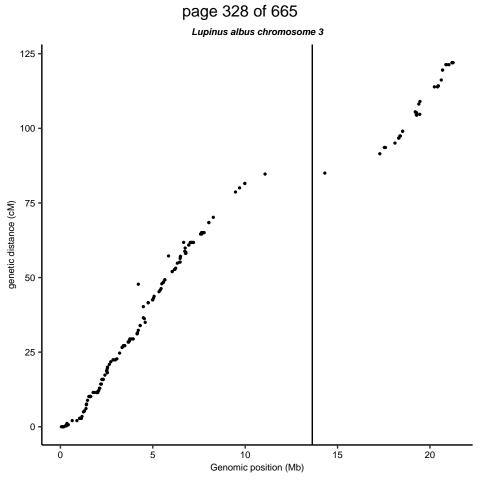


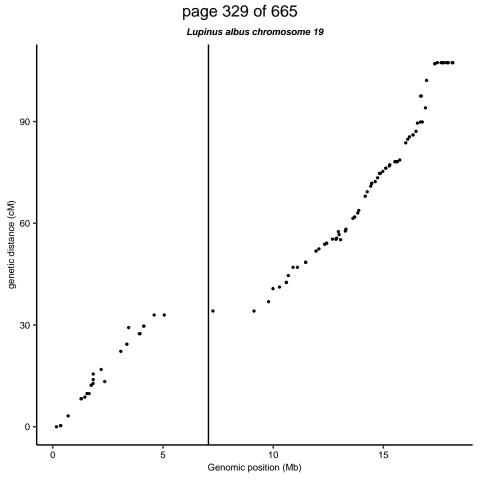


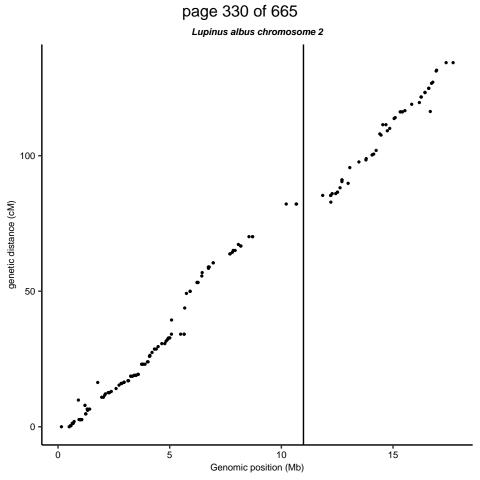


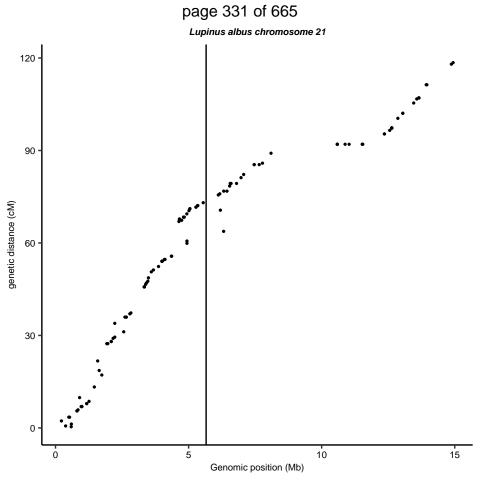


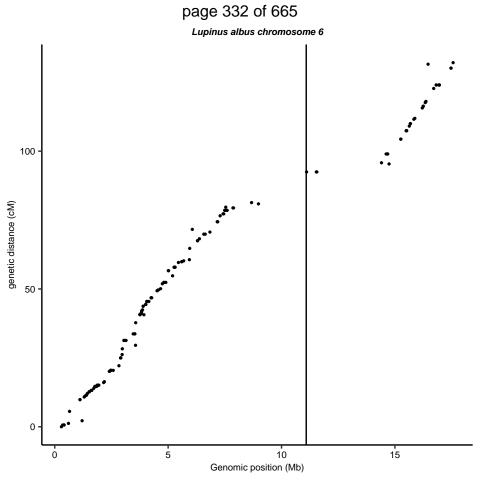


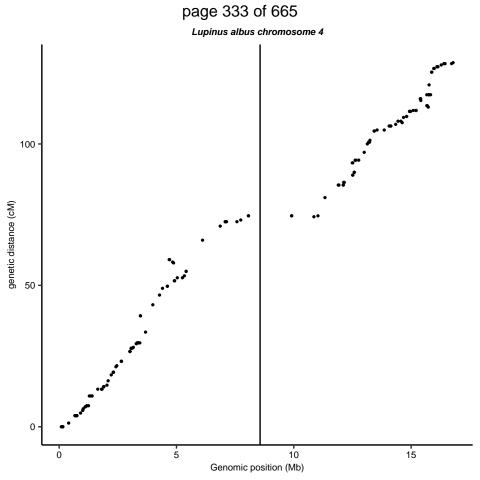


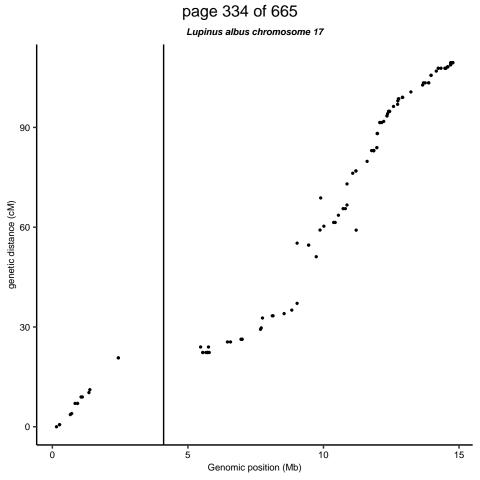


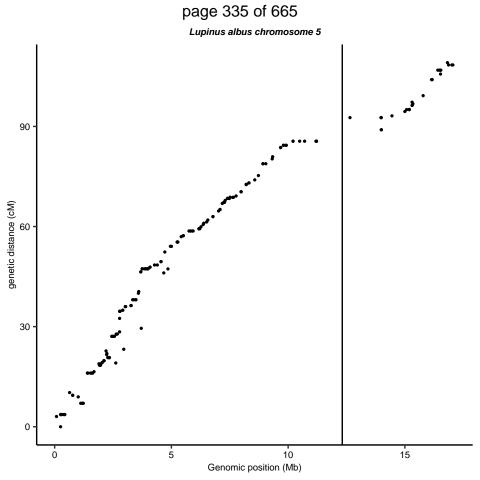


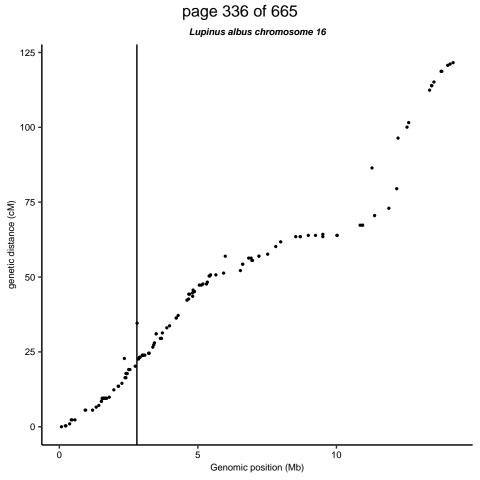


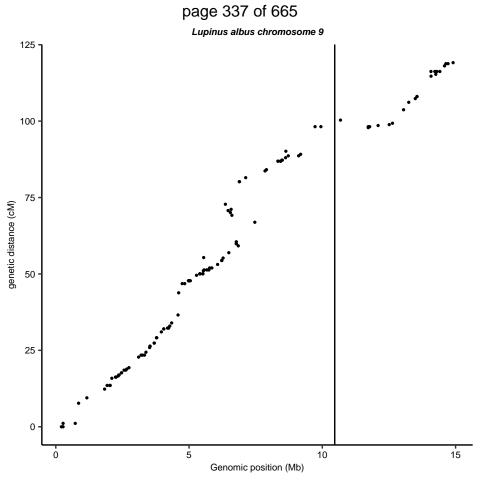


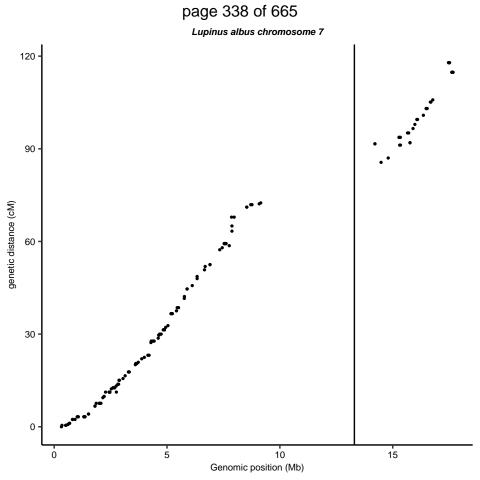


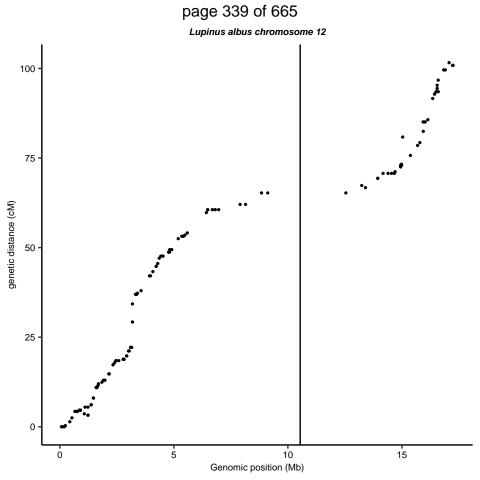


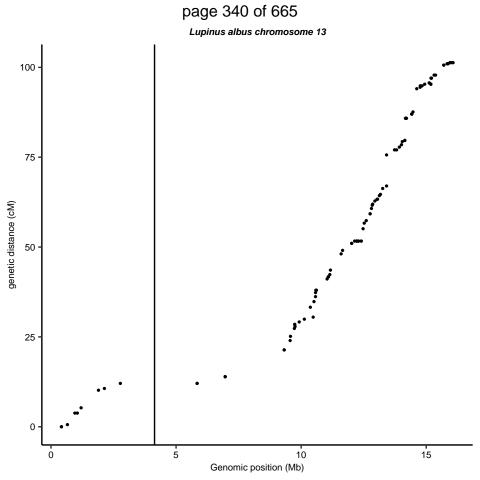


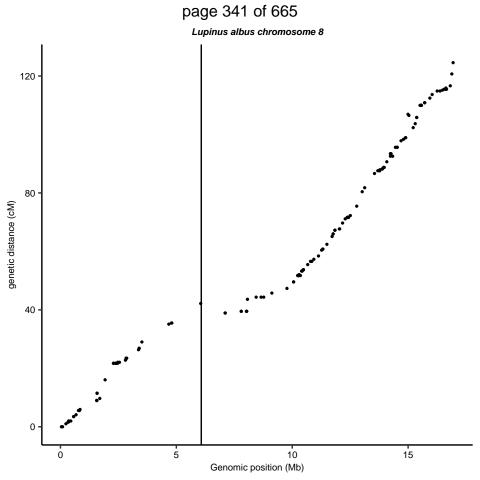


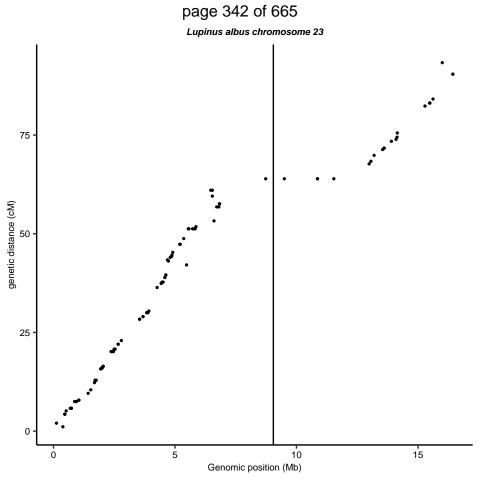


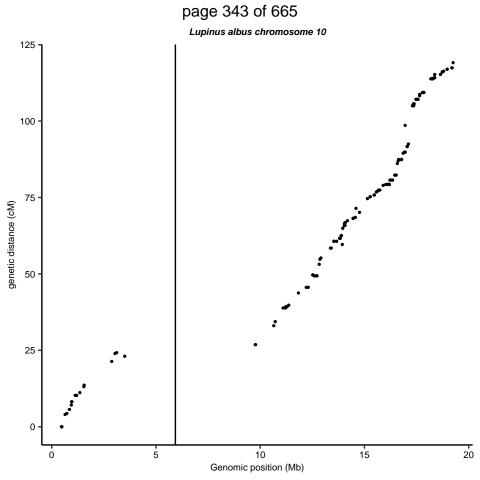


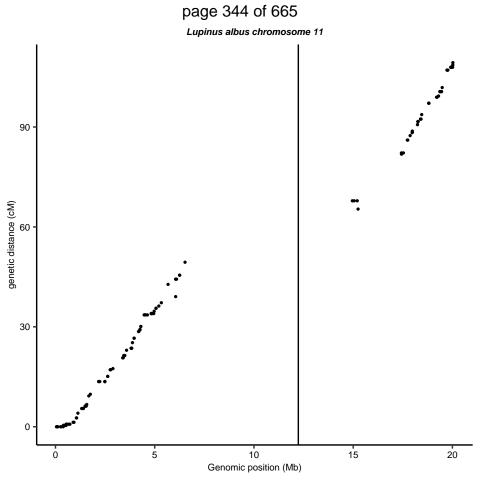


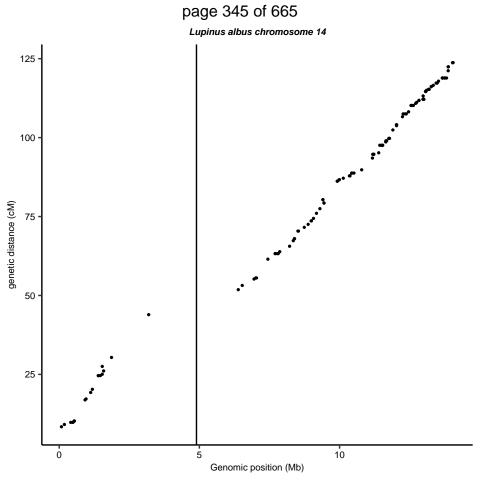






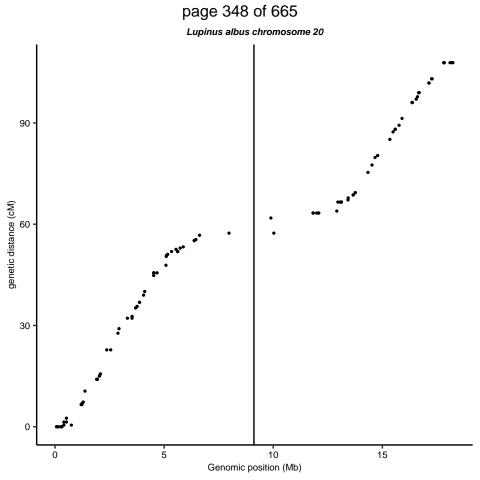


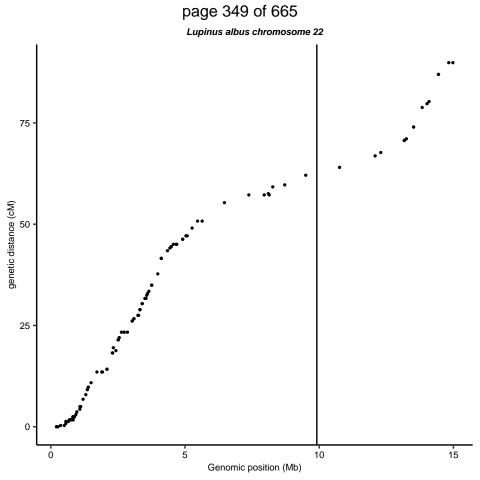


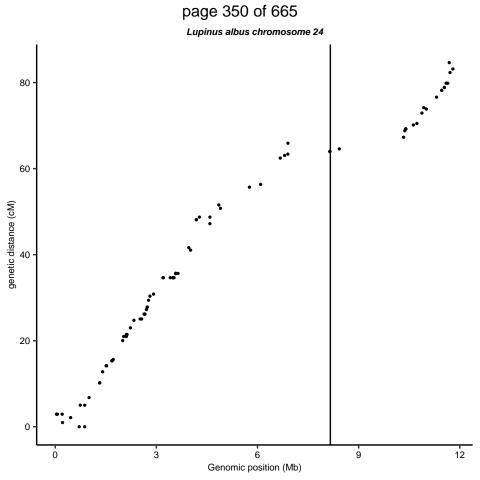


page 346 of 665 Lupinus albus chromosome 15 120 80 genetic distance (cM) 40 -0 15 5 10 20 Genomic position (Mb)

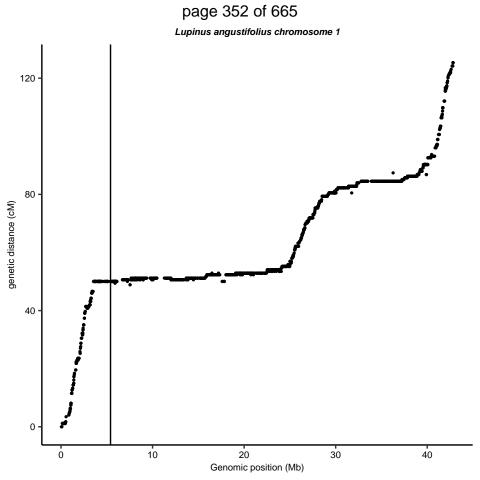
page 347 of 665 Lupinus albus chromosome 18 genetic distance (cM) Genomic position (Mb)

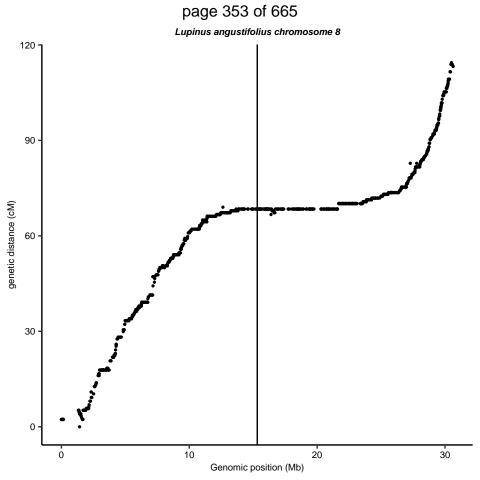


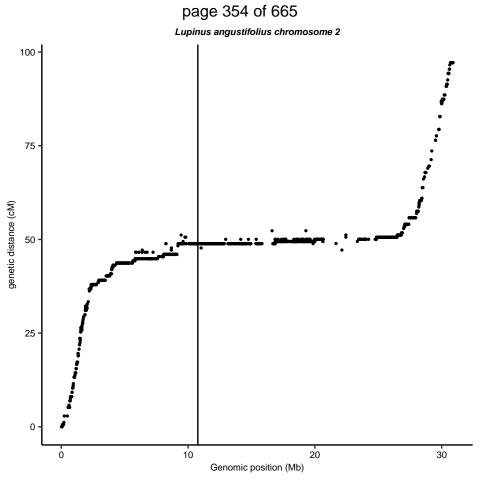


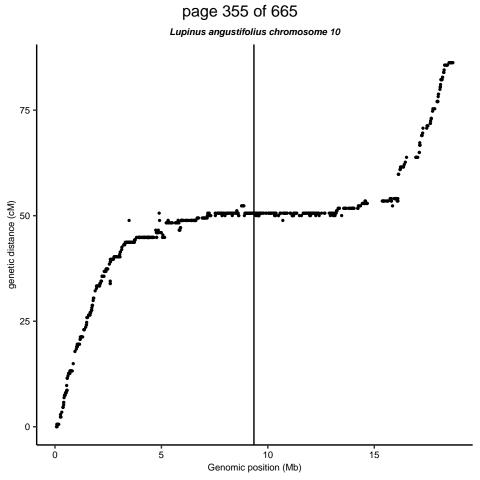


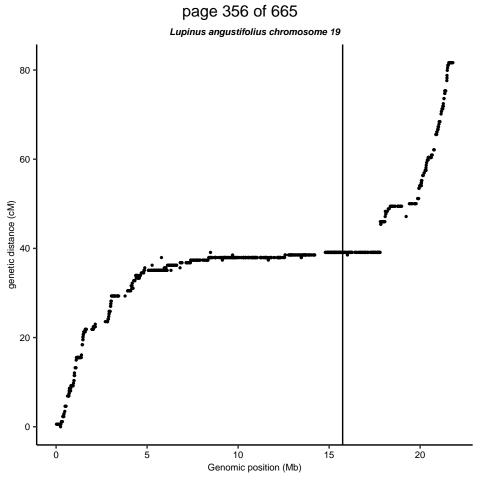
page 351 of 665 Lupinus albus chromosome 25 125 -100 75 genetic distance (cM) 50 25 0 15 5 10 Genomic position (Mb)

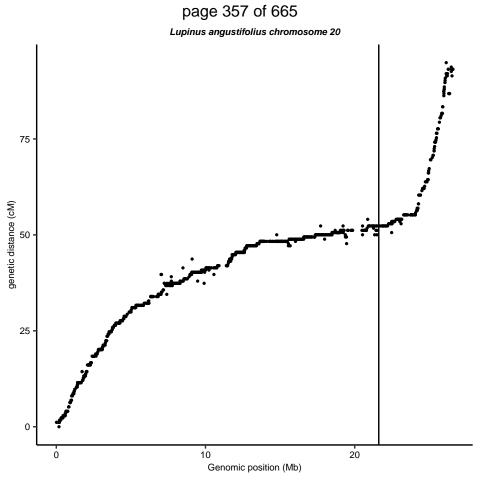


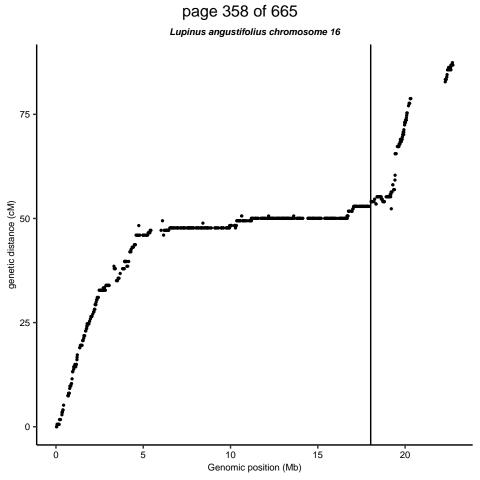


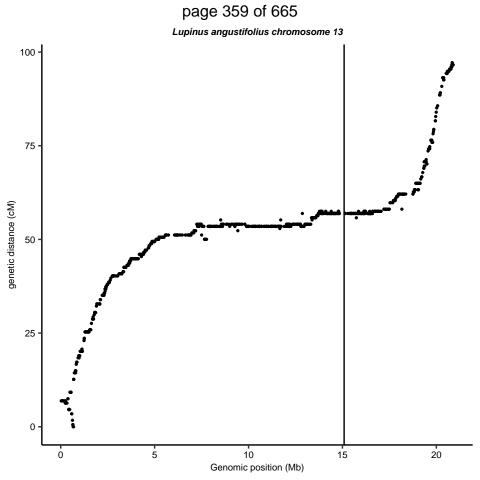


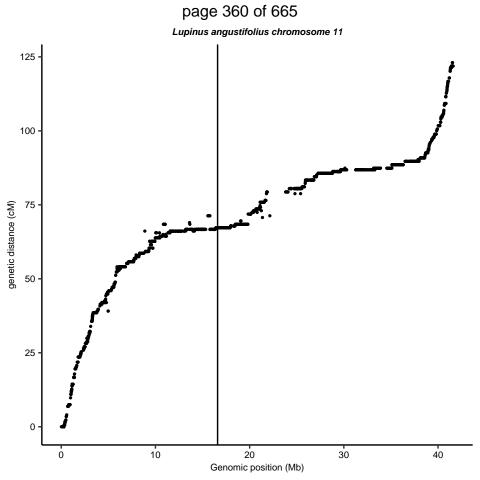


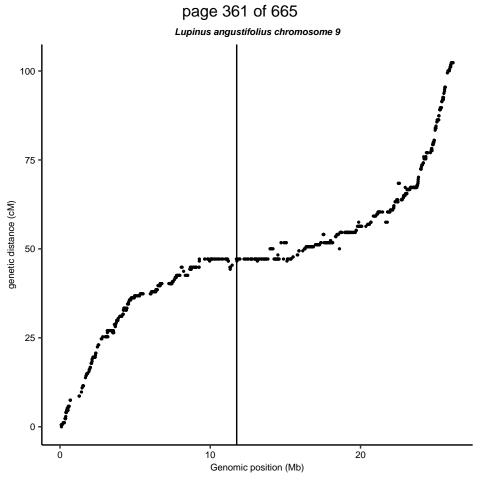


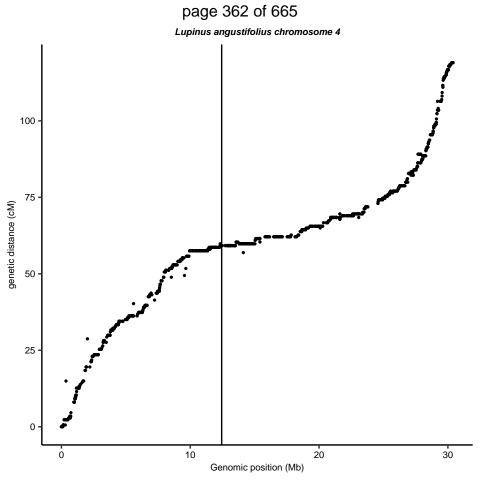


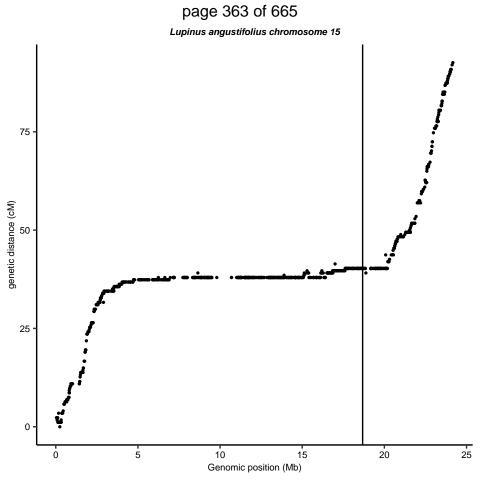


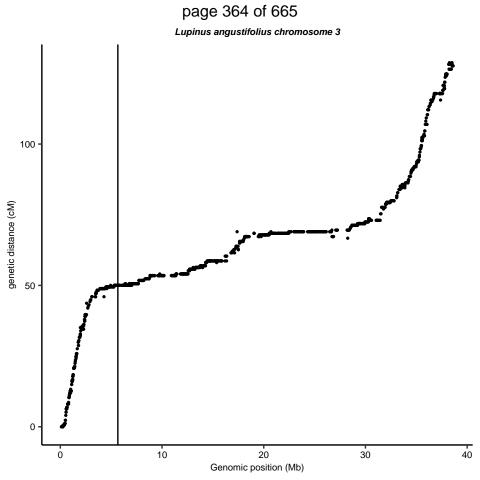


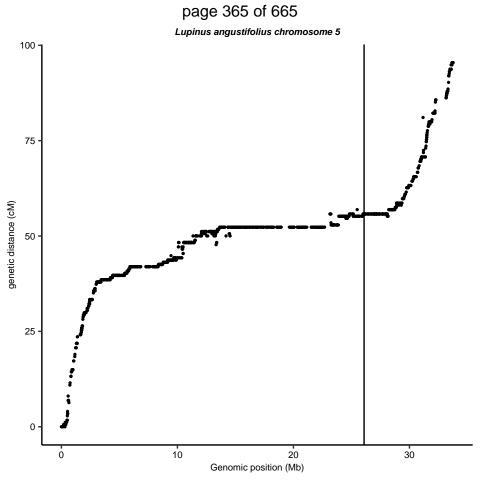


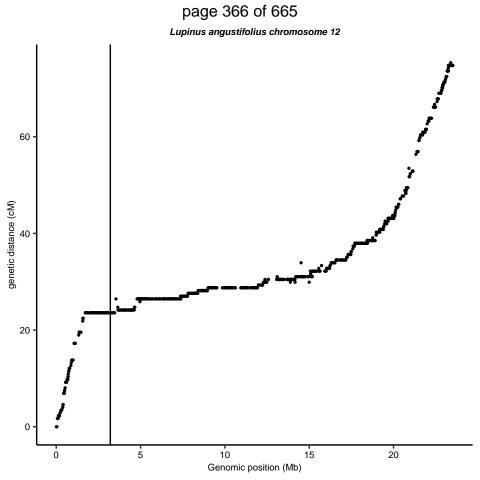


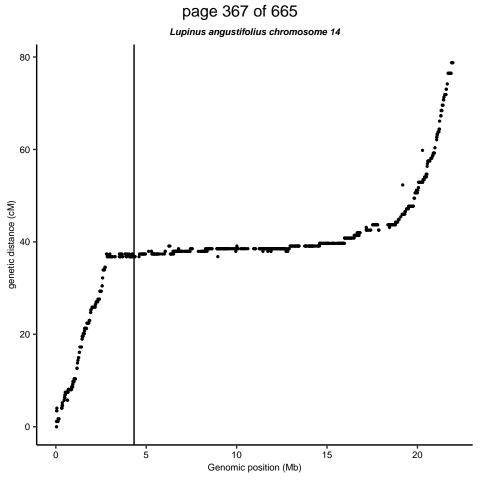


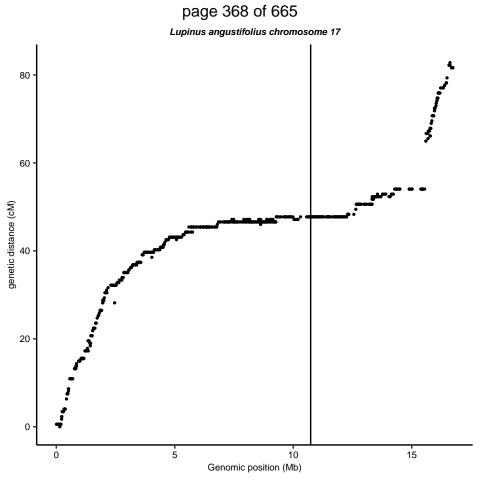


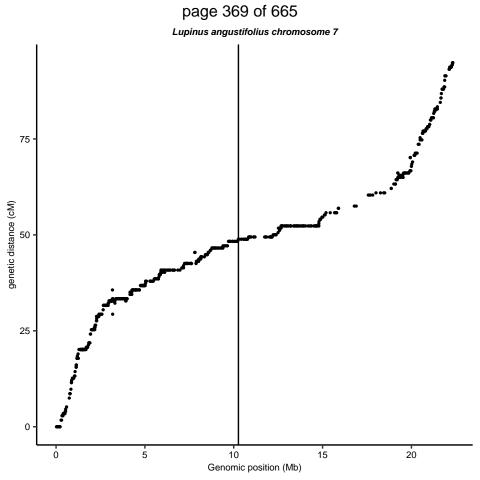


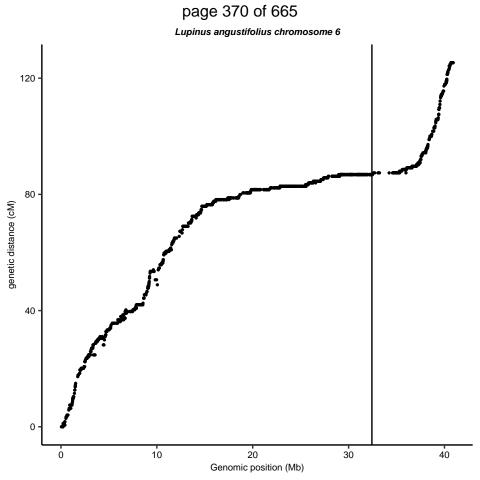


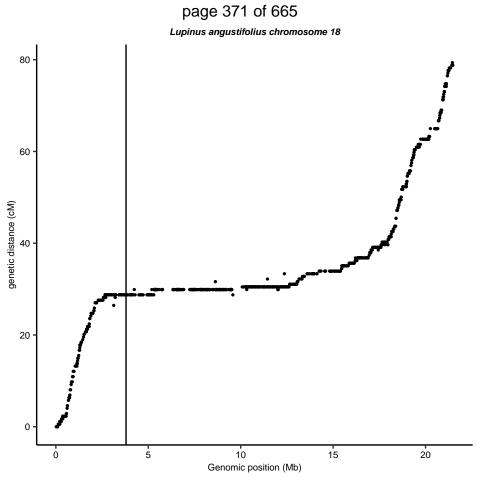


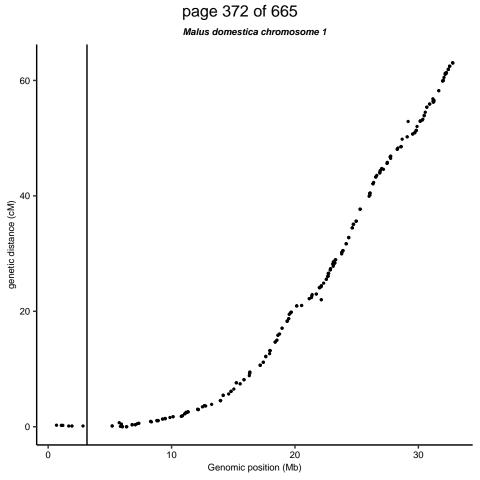


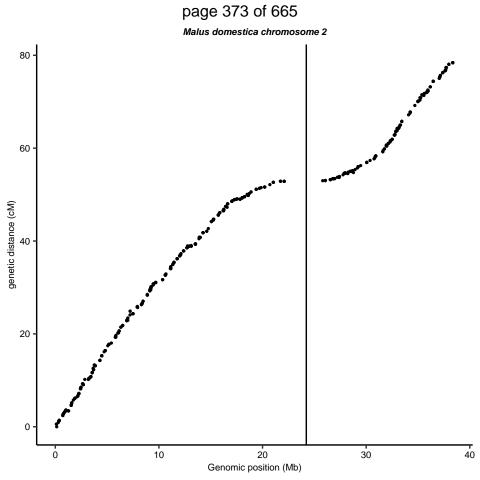


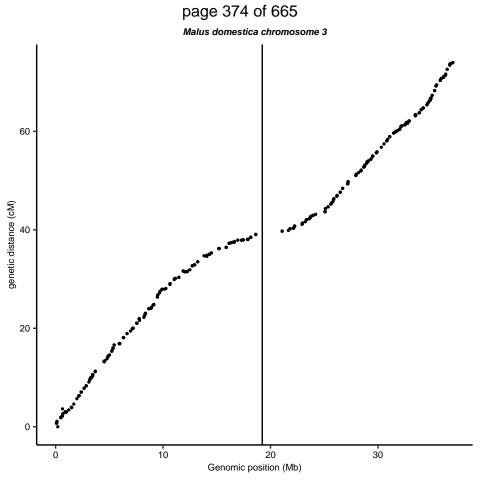


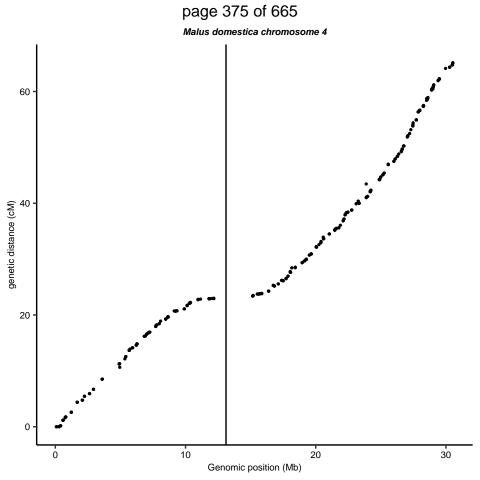


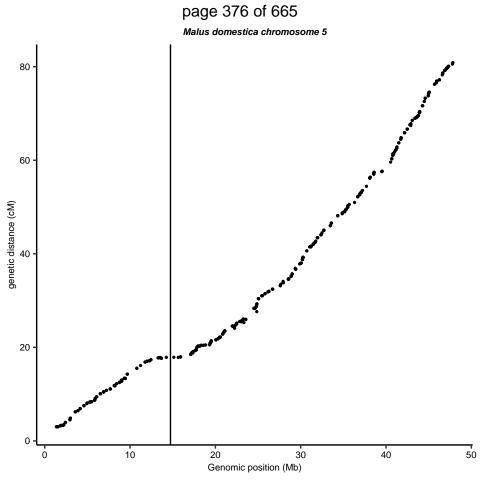


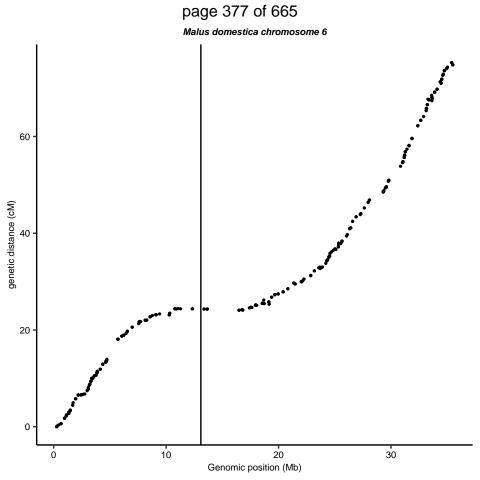


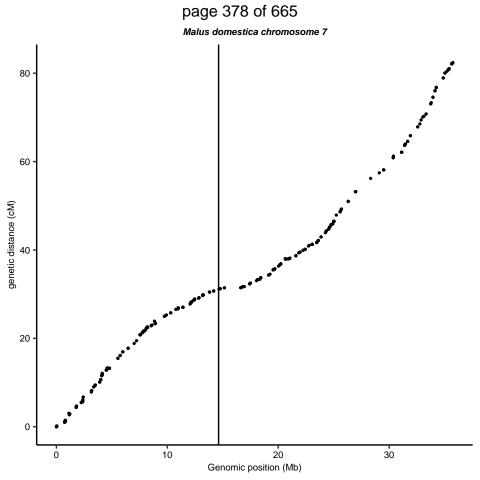


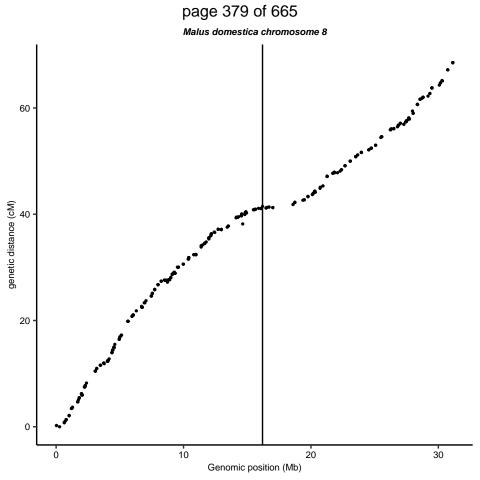


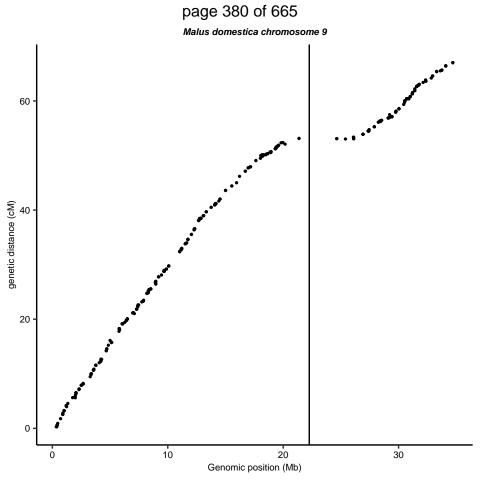


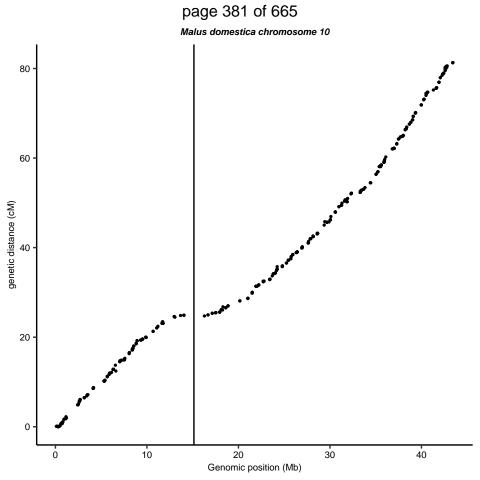


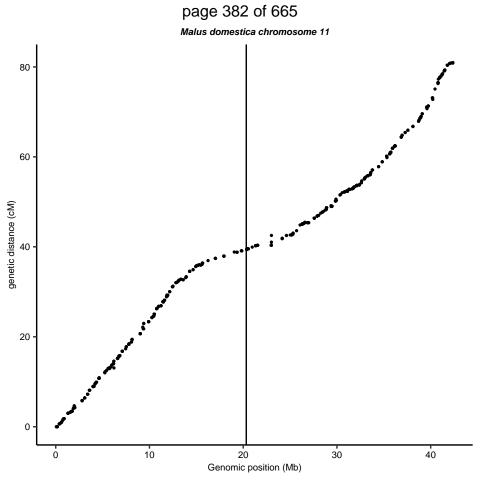


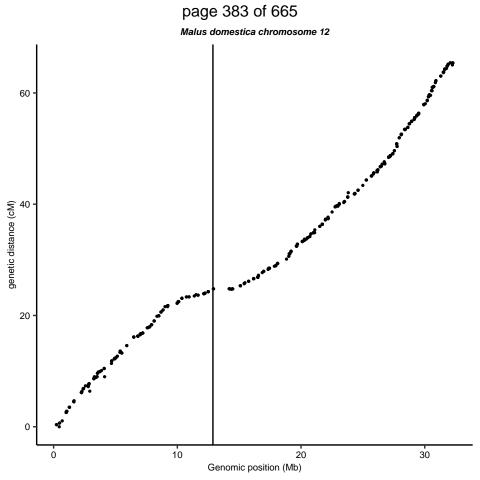


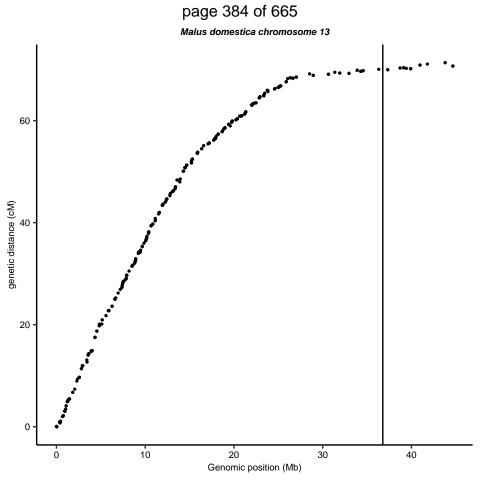


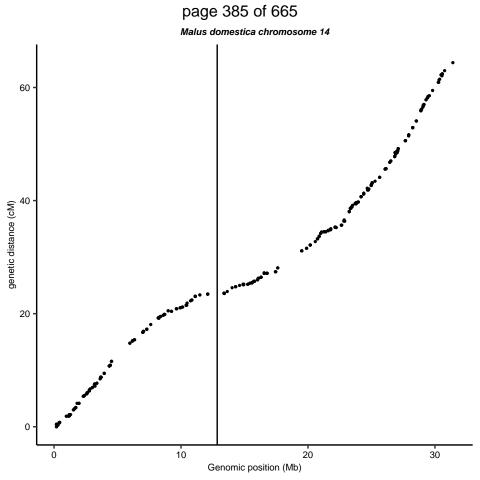


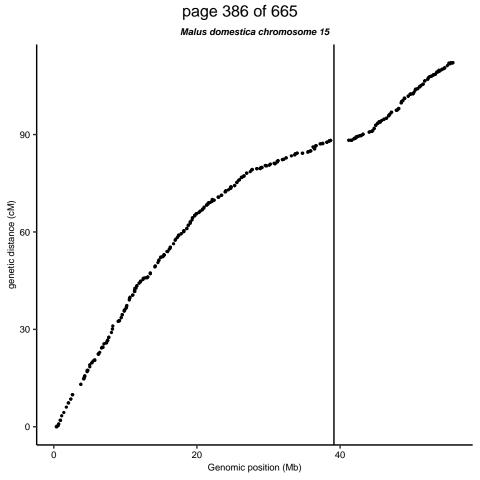


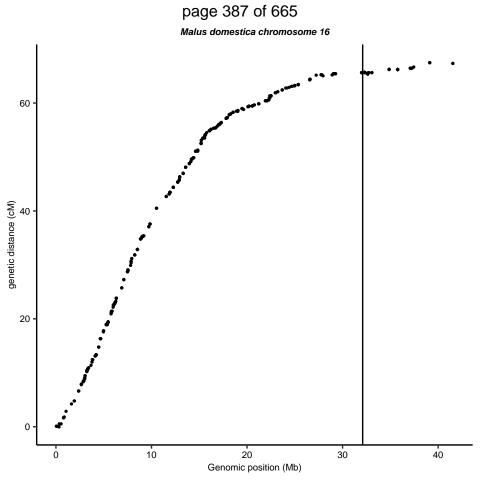


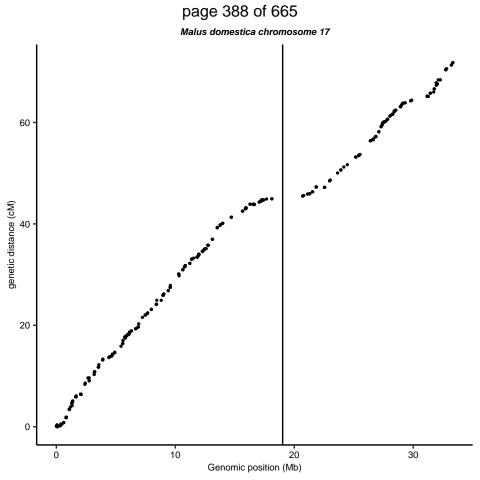


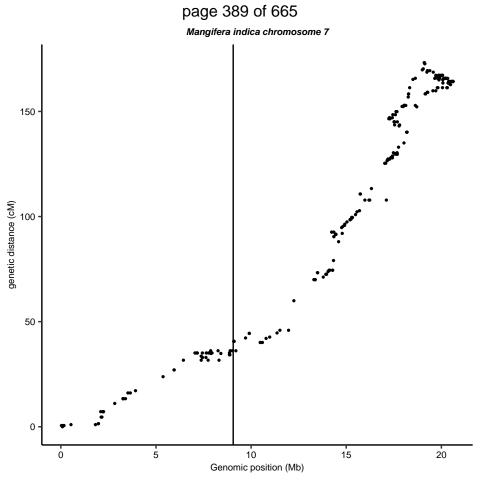


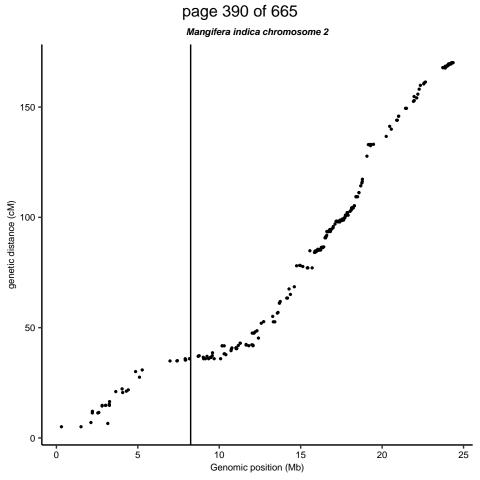


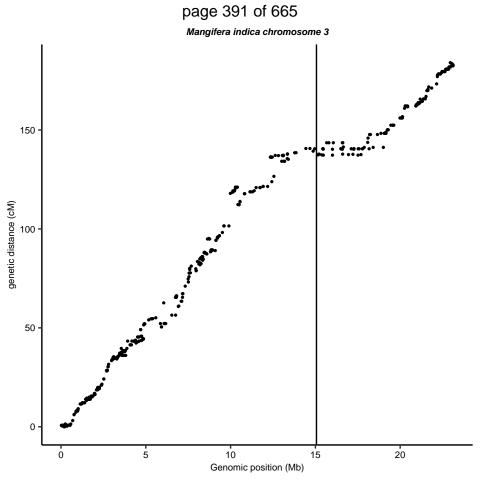


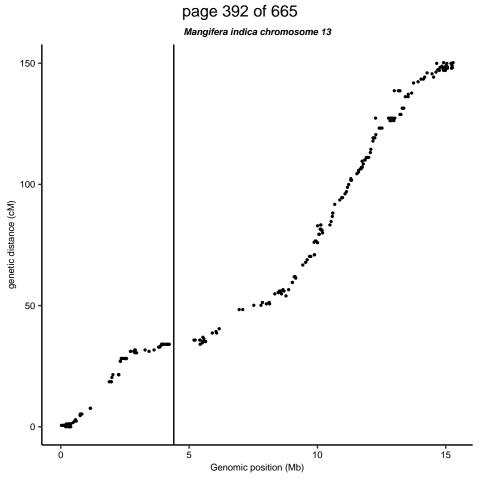


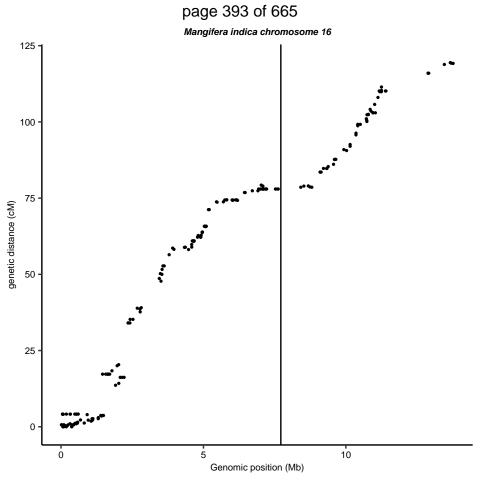


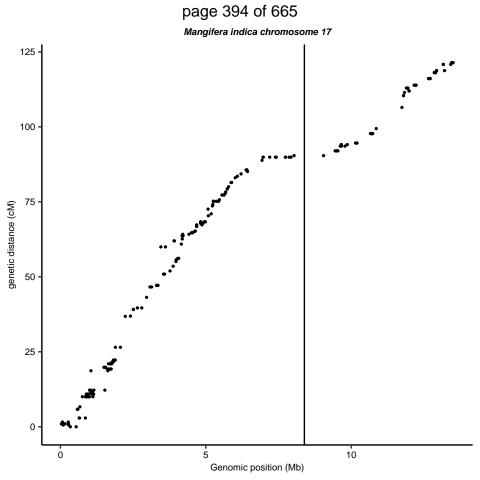


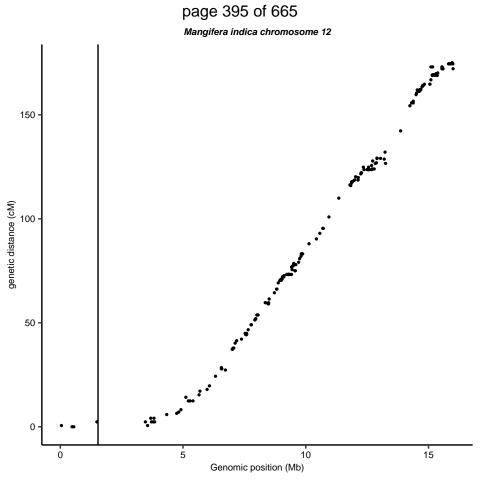


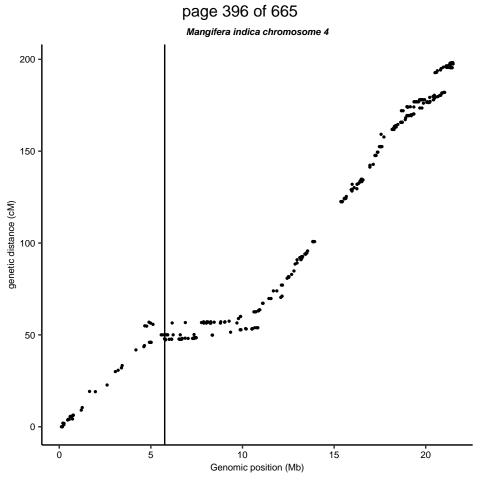


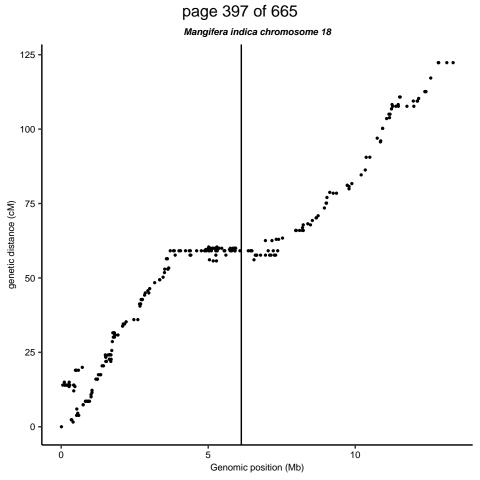


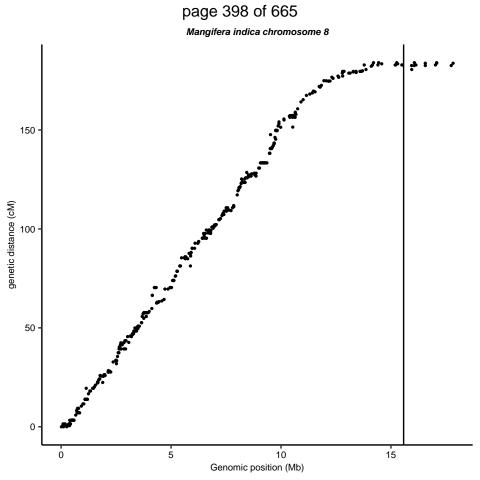


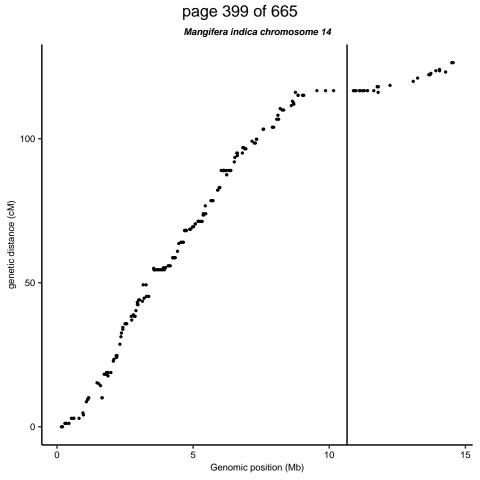


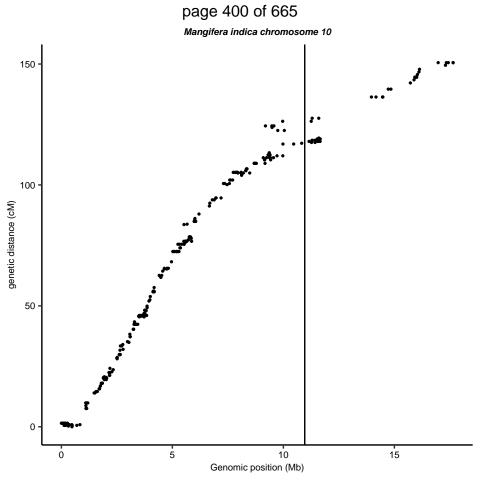


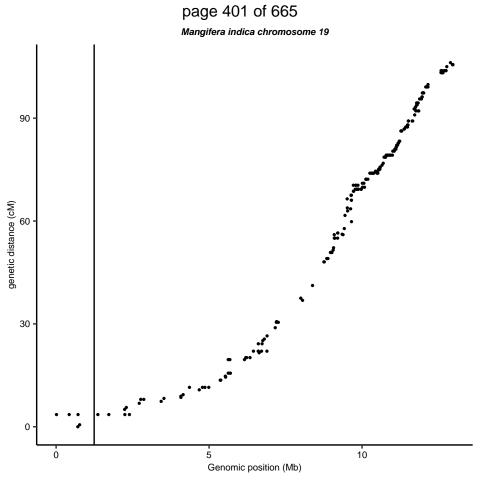


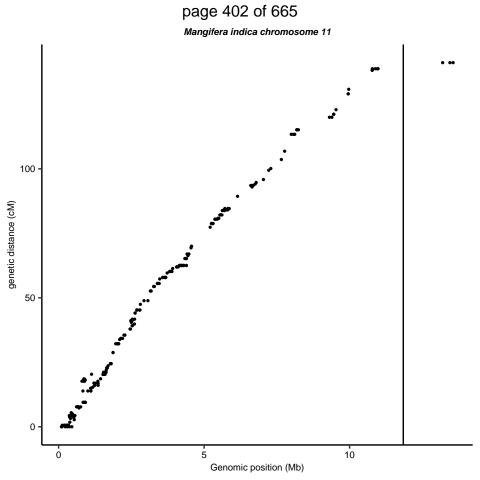


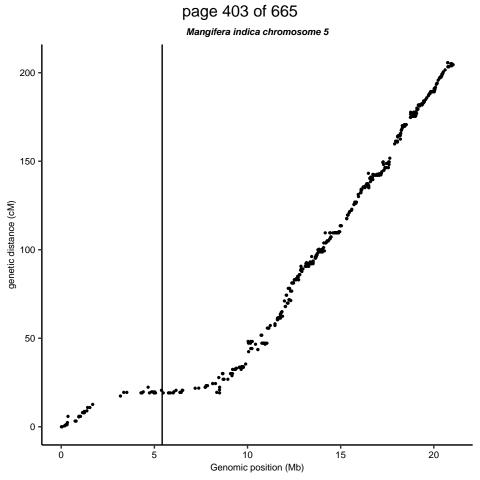


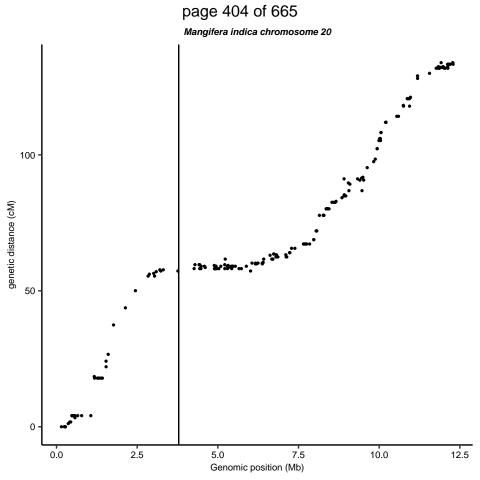


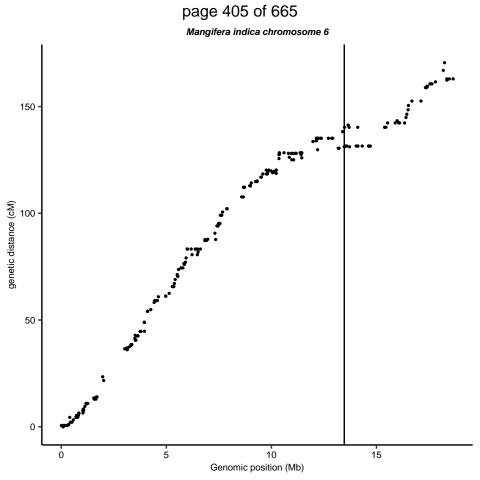


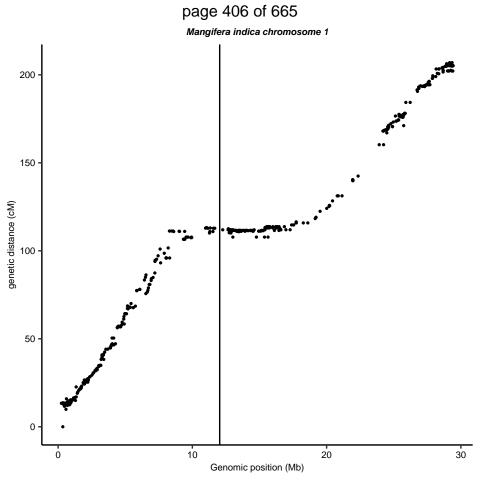


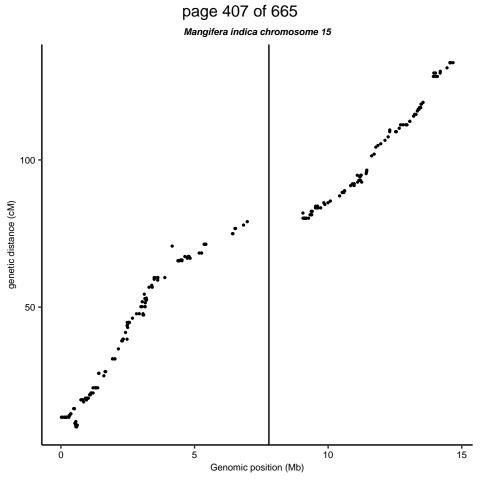


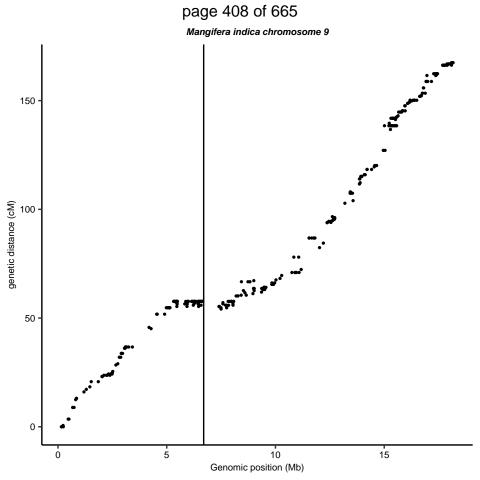


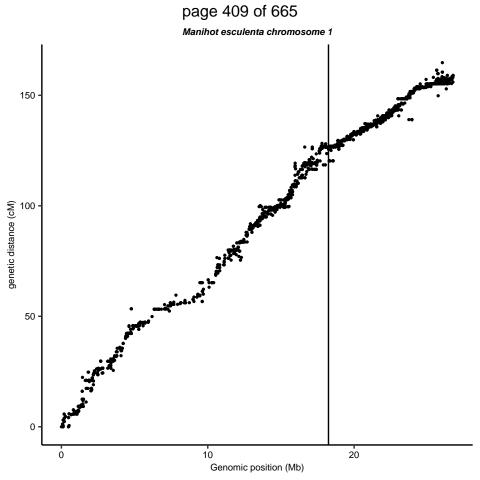


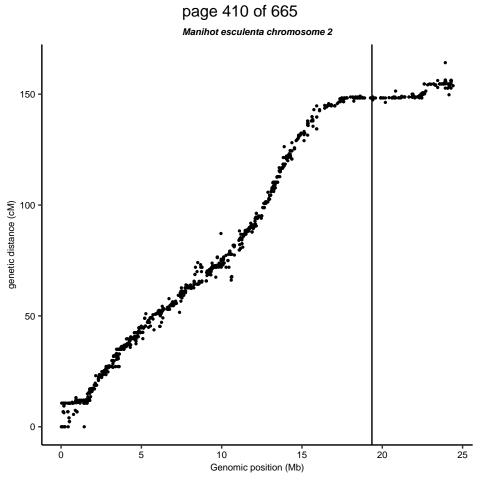


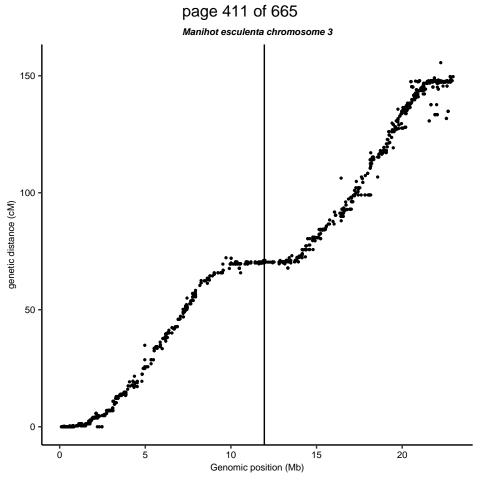


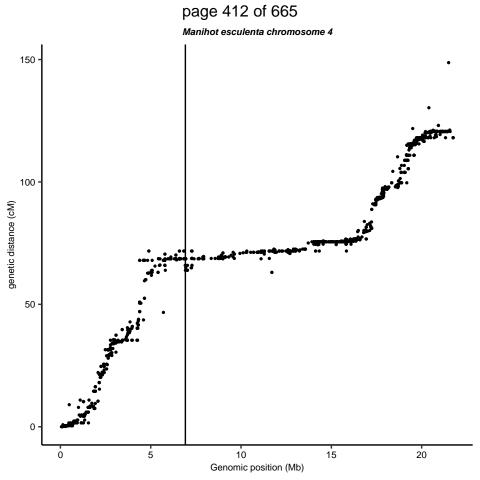


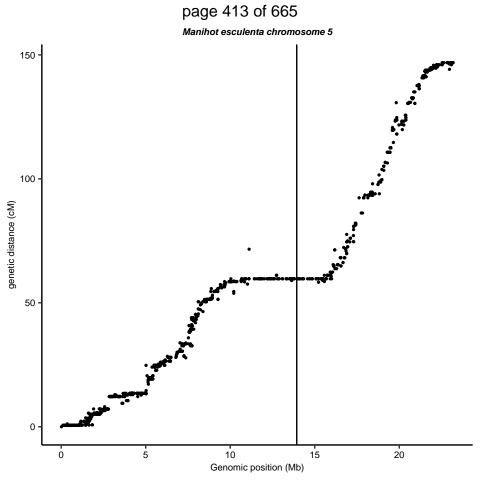


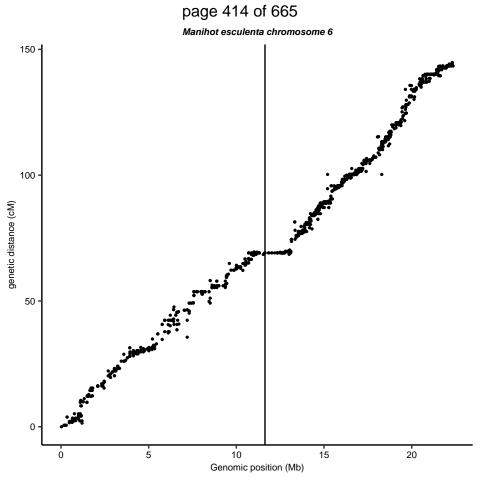


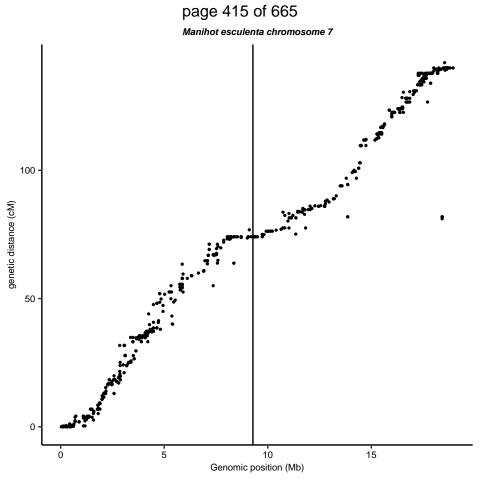


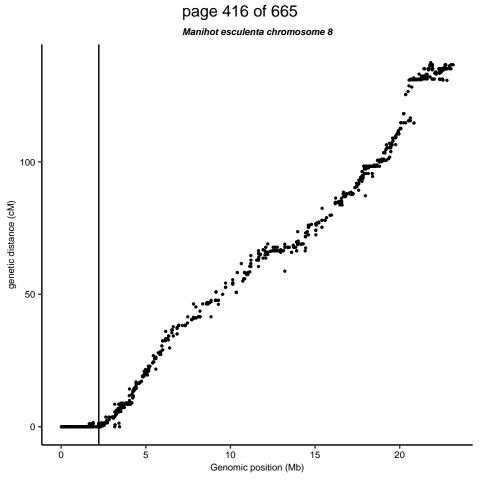


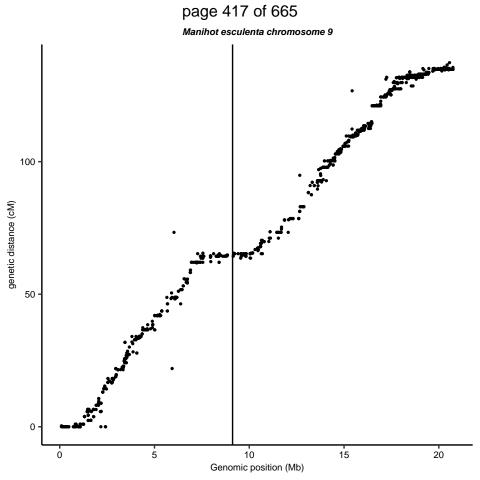


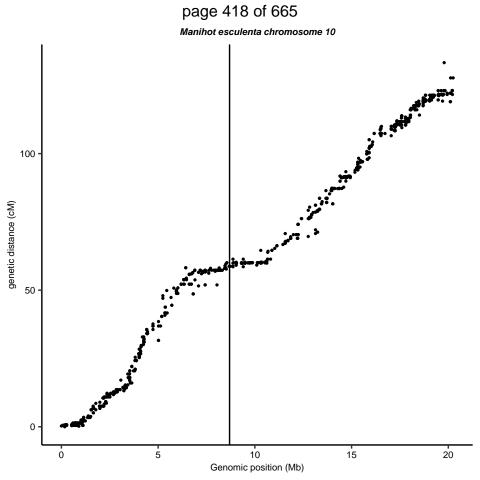


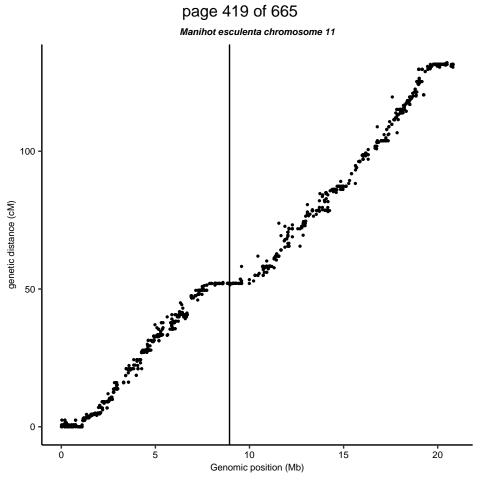


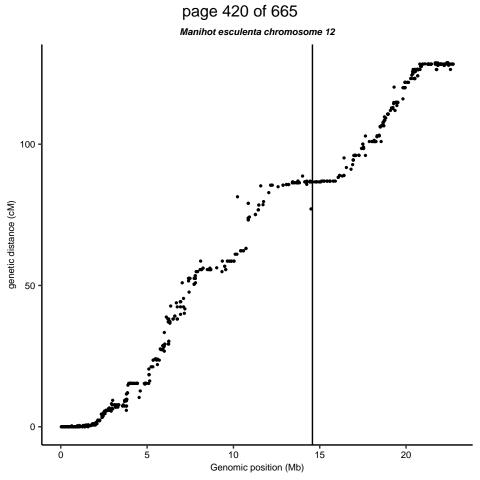


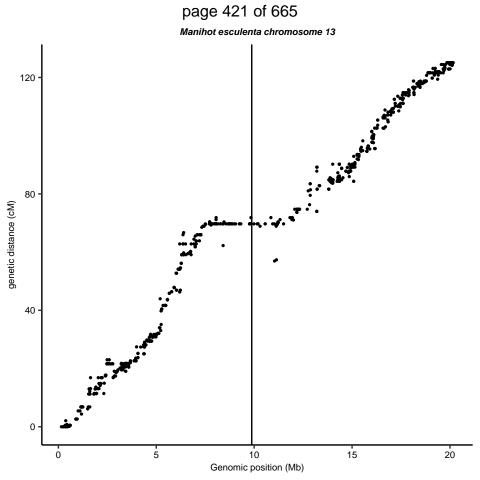


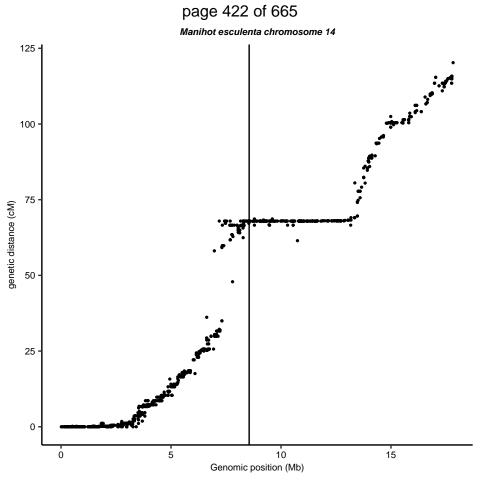


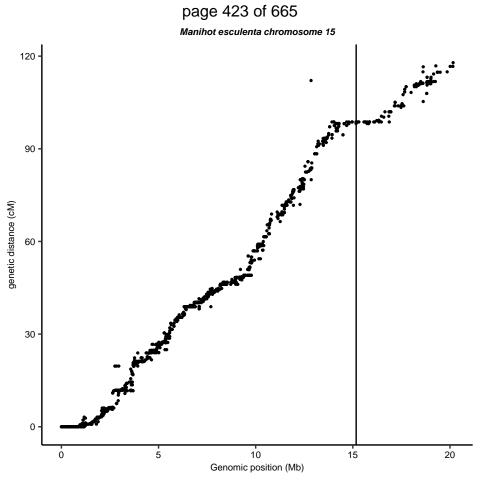


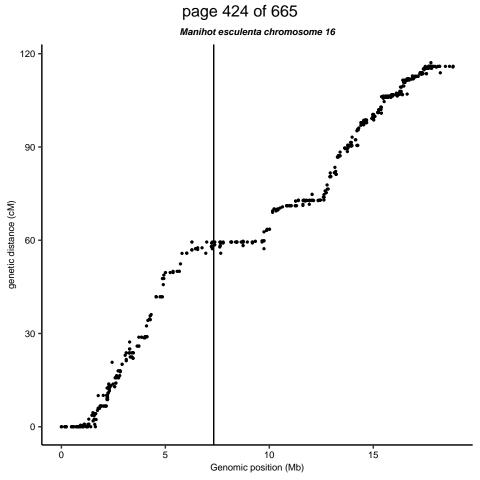


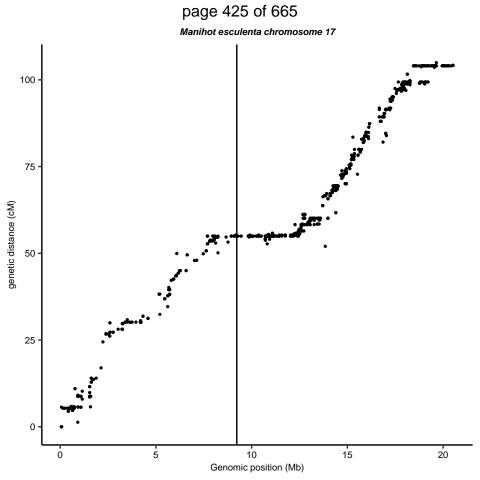


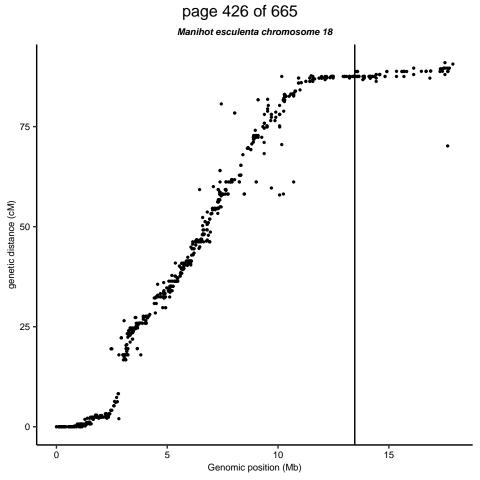


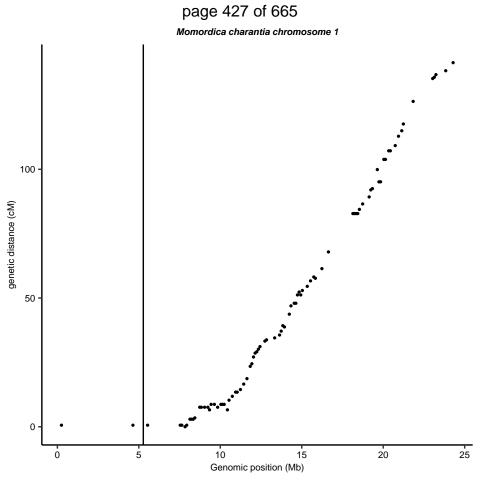






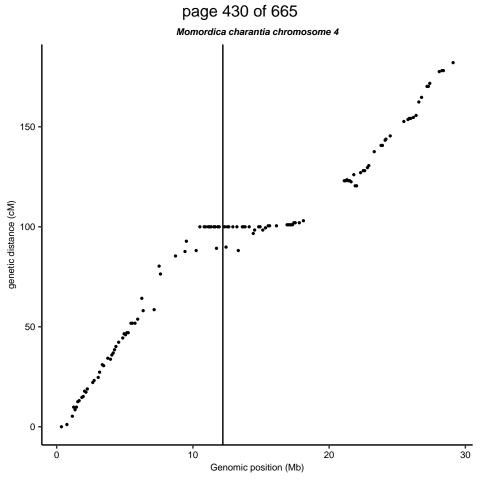




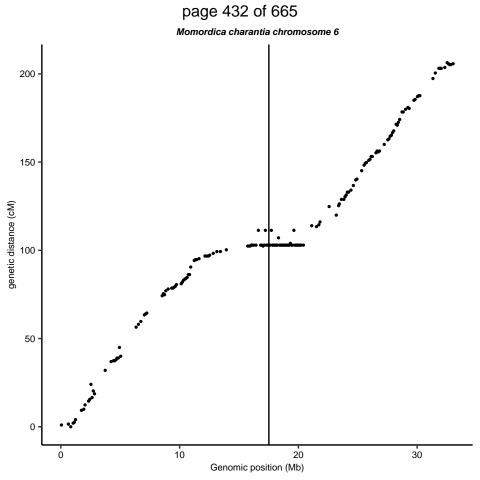


page 428 of 665 Momordica charantia chromosome 2 genetic distance (cM) Genomic position (Mb)

page 429 of 665 Momordica charantia chromosome 3 100 75 genetic distance (cM) 50 25 0 10 15 20 Genomic position (Mb)

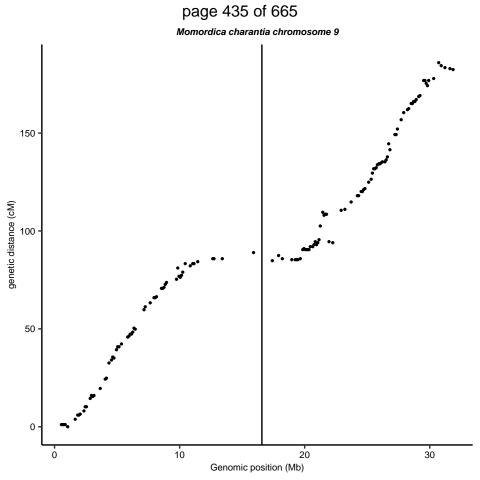


page 431 of 665 Momordica charantia chromosome 5 genetic distance (cM) Genomic position (Mb)



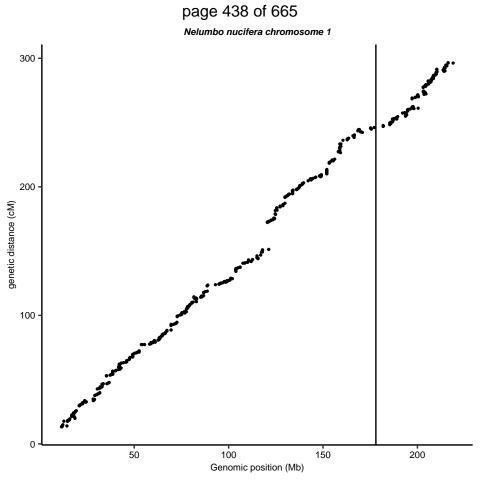
page 433 of 665 Momordica charantia chromosome 7 80 -60 genetic distance (cM) 20 0 20 5 10 15 Genomic position (Mb)

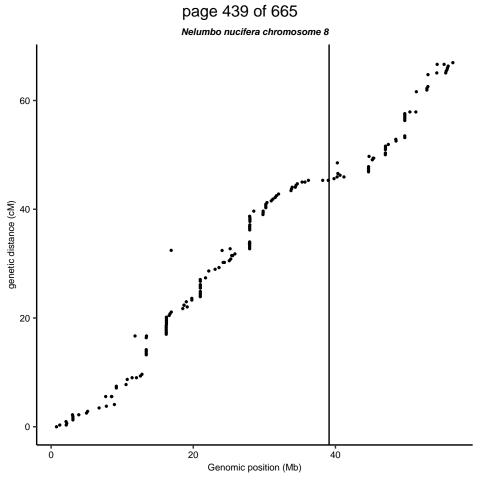
page 434 of 665 Momordica charantia chromosome 8 150 genetic distance (cM) 100 -50 0 10 20 30 Genomic position (Mb)

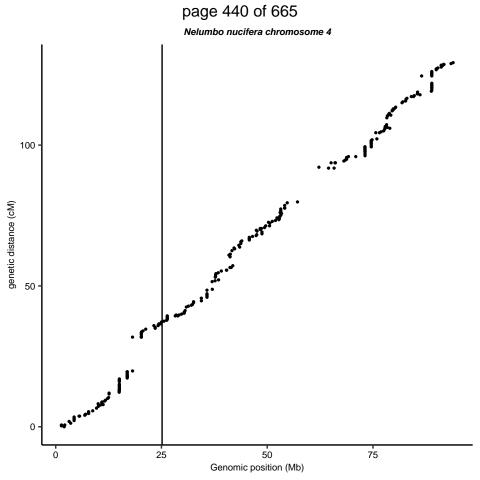


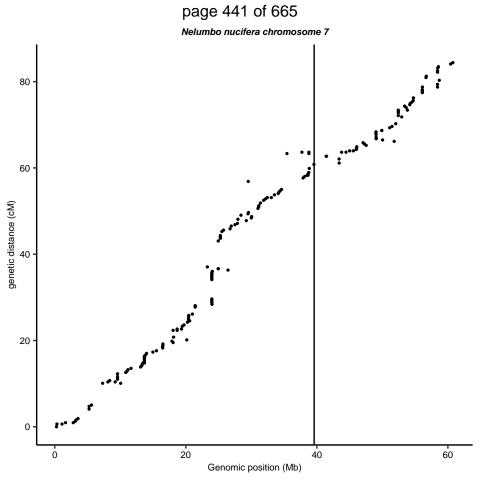
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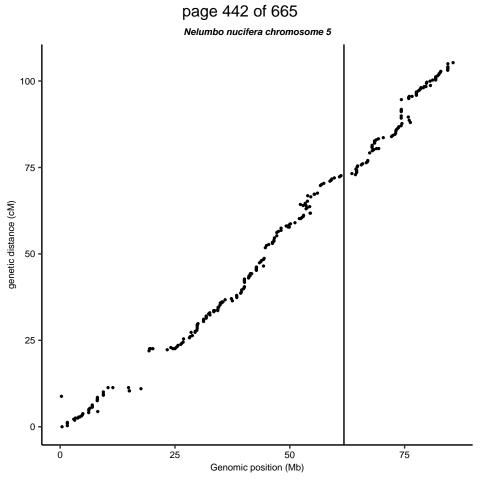
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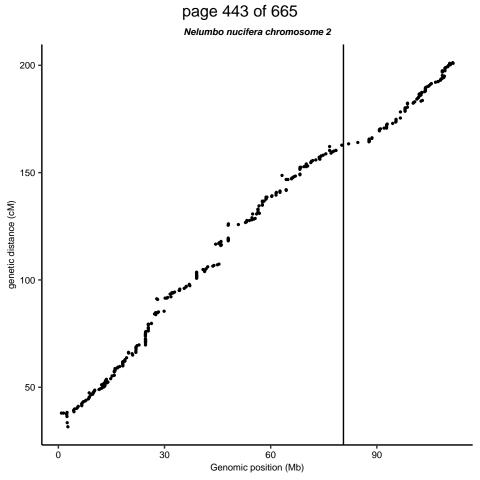


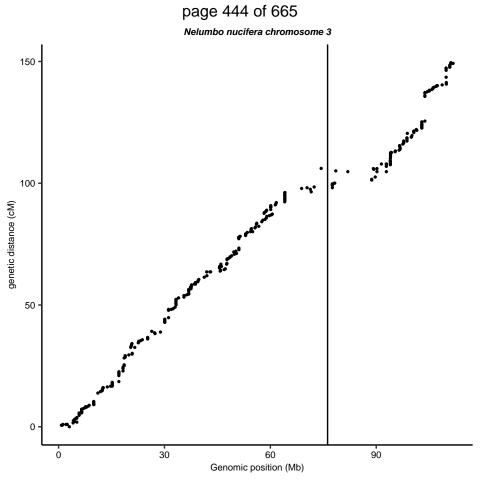


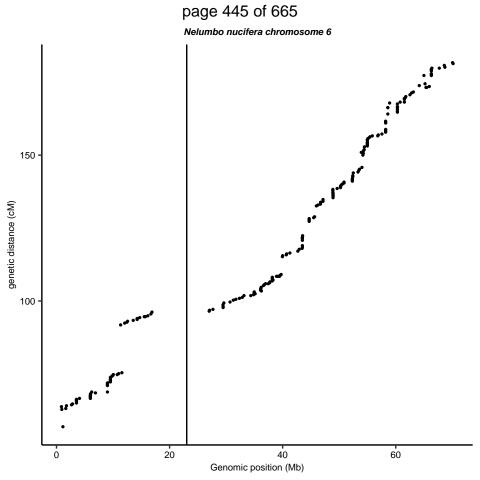


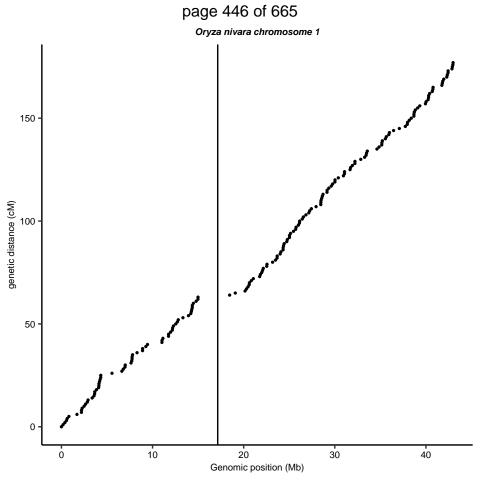


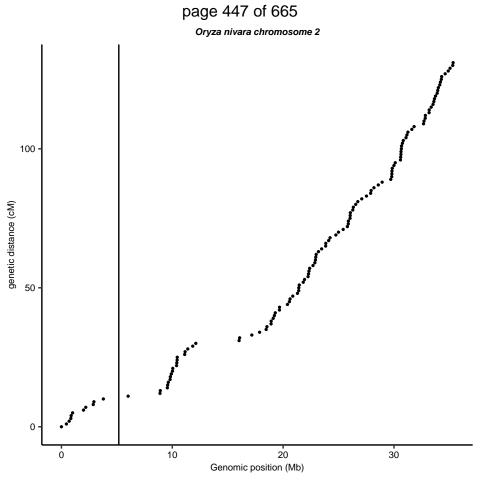


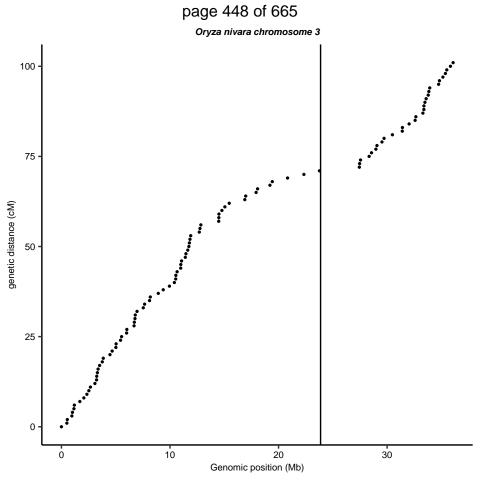


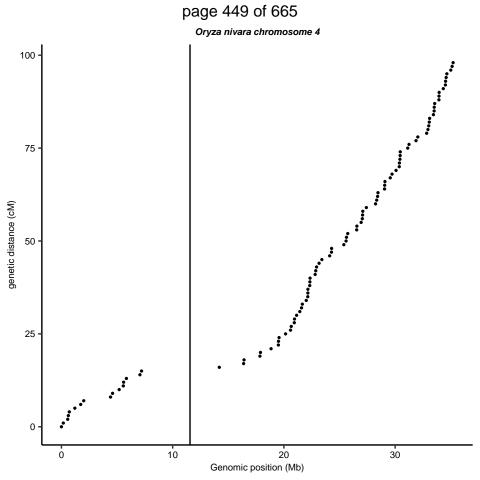


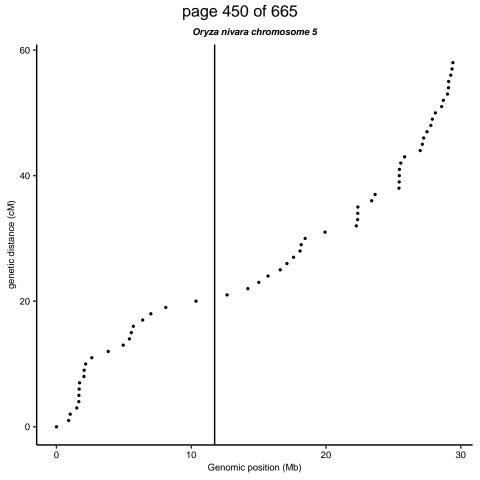


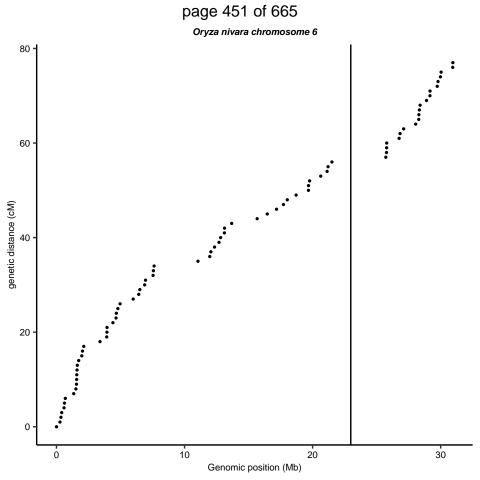


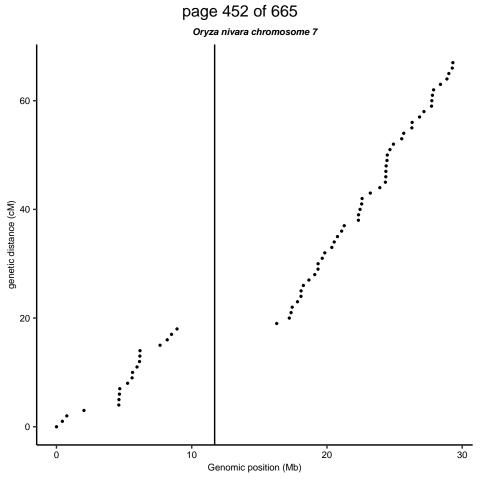


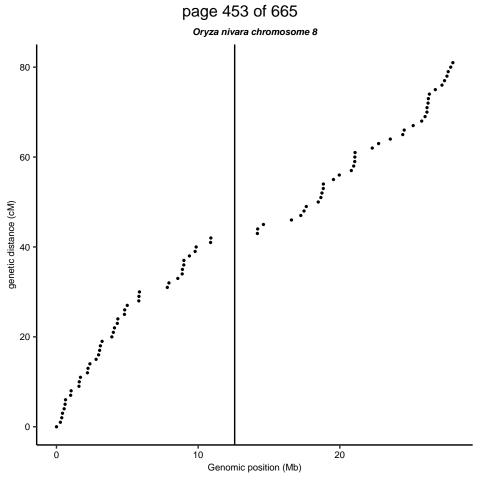


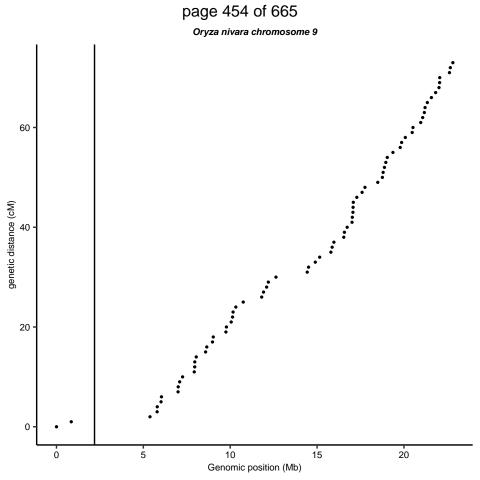


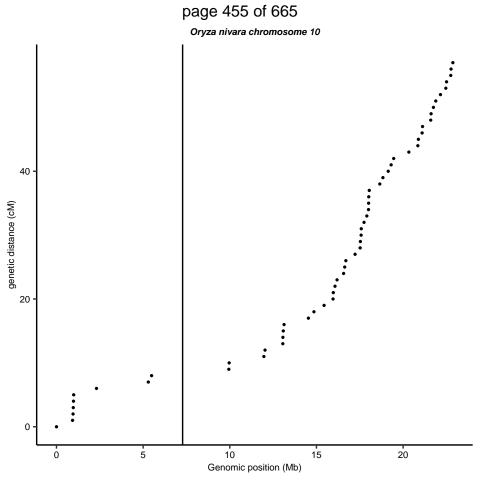


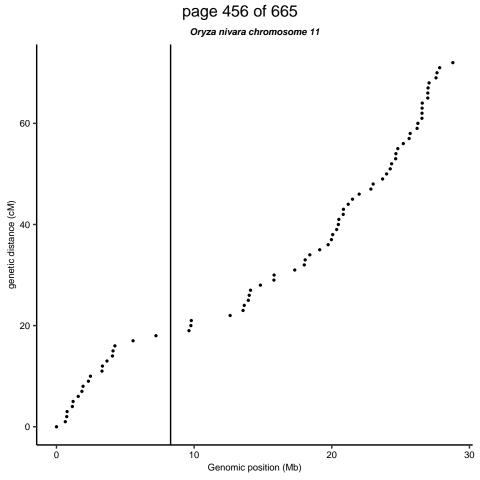


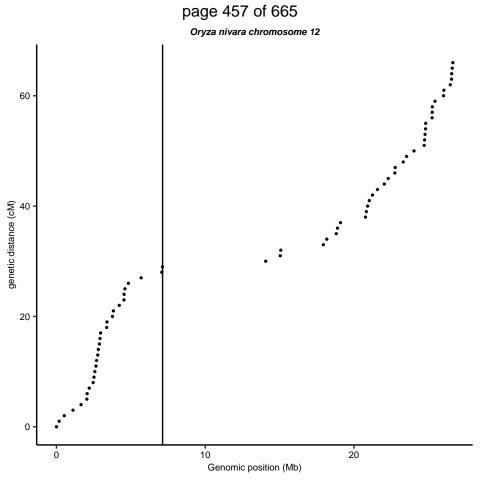


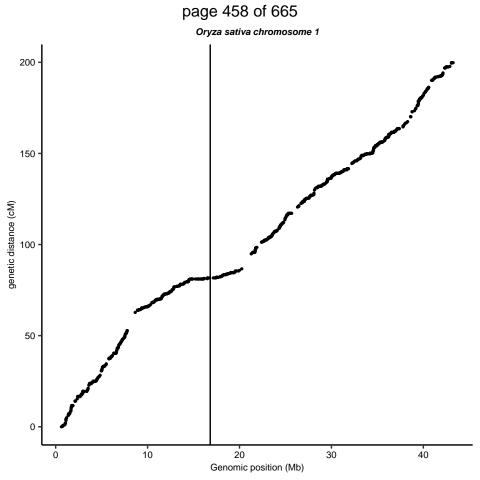


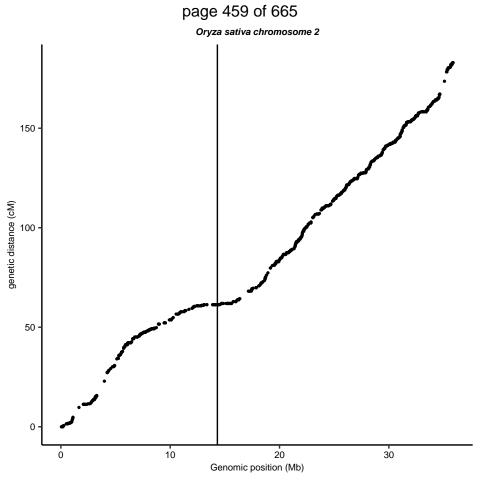


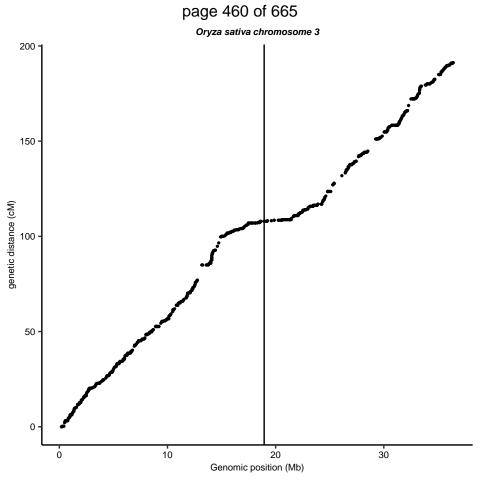


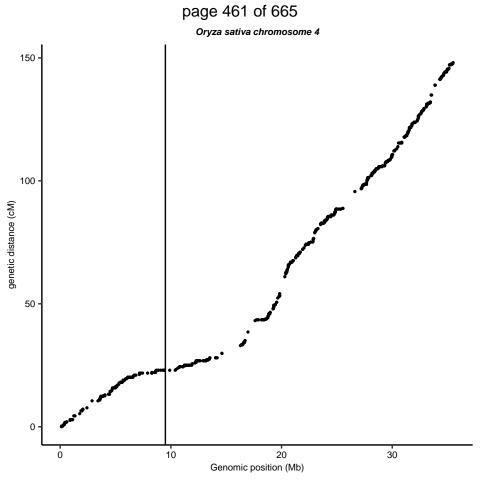


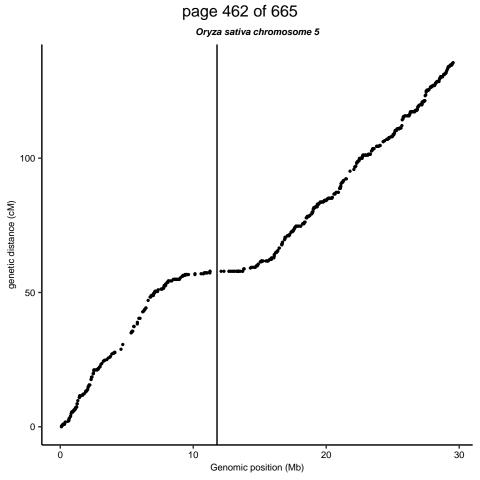


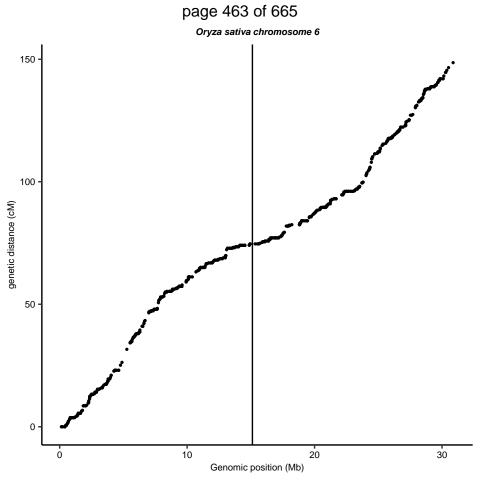


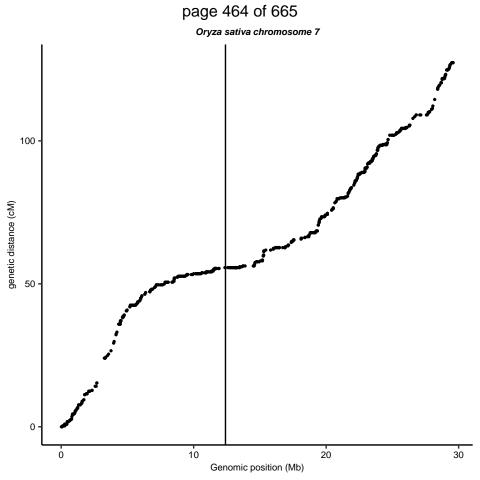


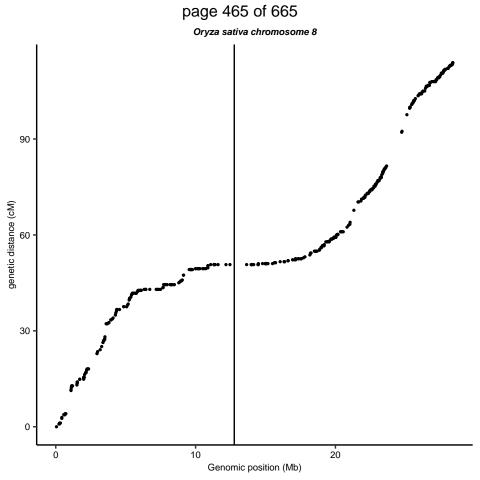


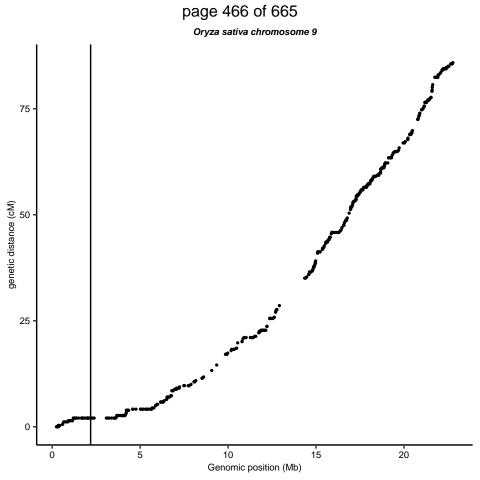


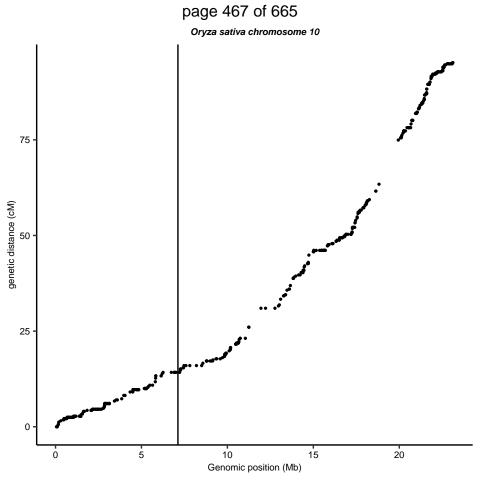


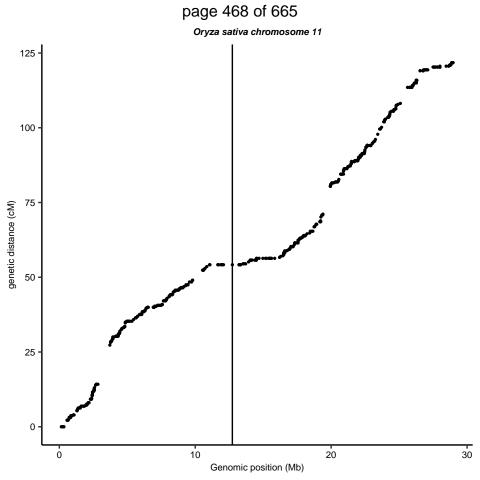


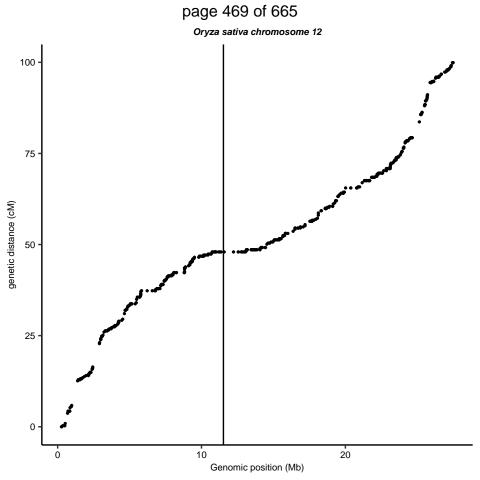


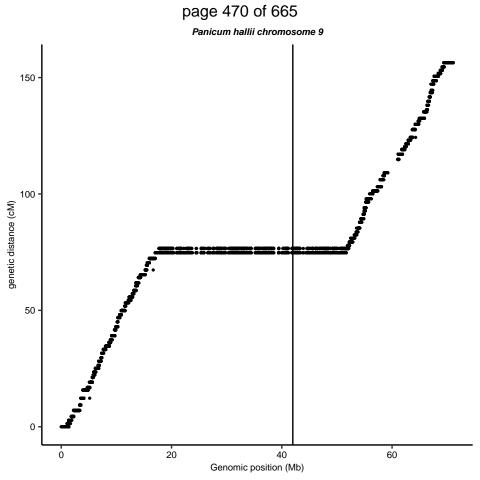


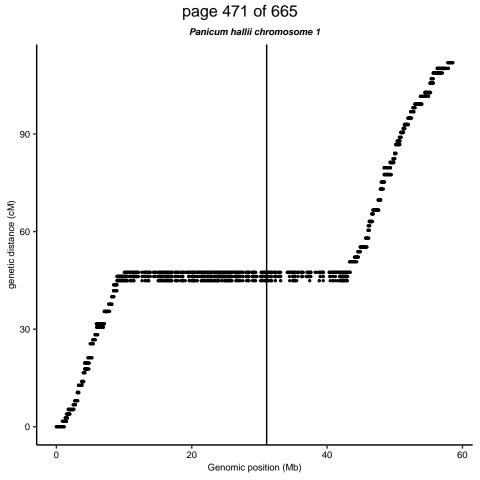


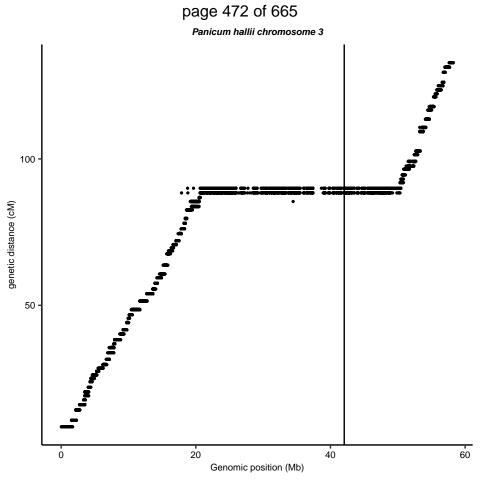






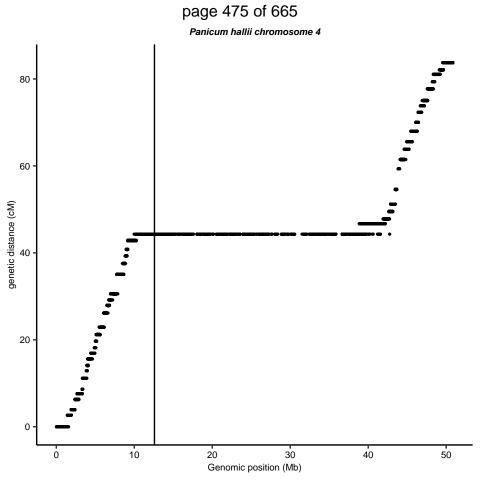


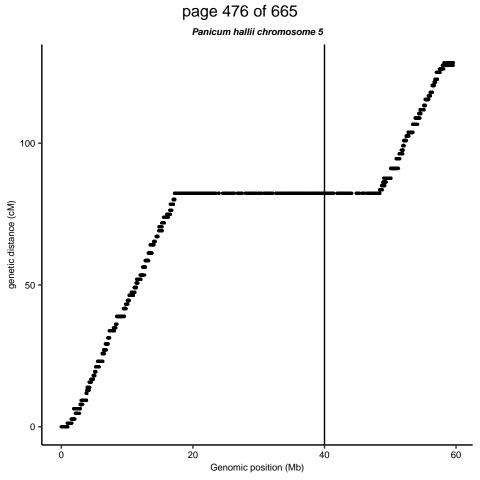


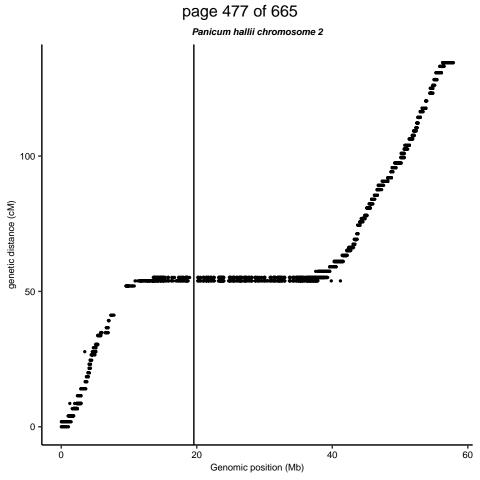


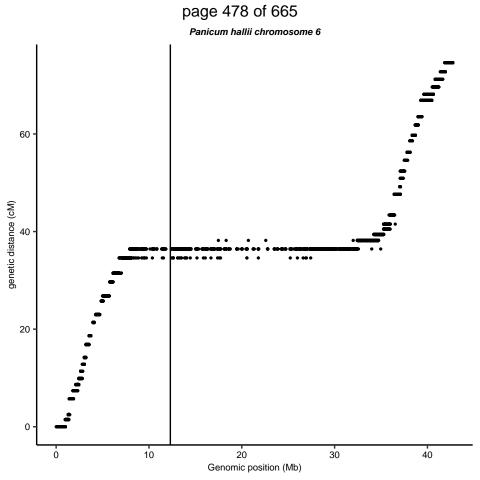
page 473 of 665 Panicum hallii chromosome 7 genetic distance (cM) Genomic position (Mb)

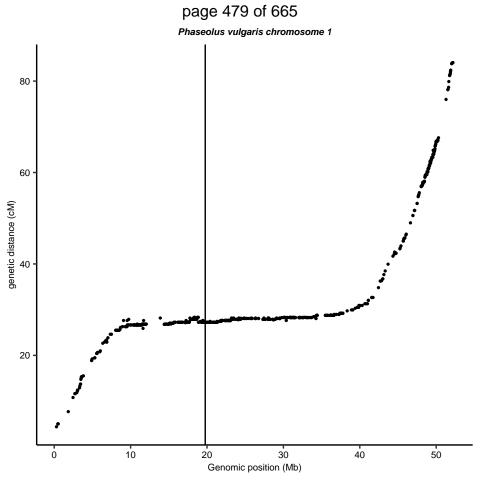
page 474 of 665 Panicum hallii chromosome 8 genetic distance (cM) ö Genomic position (Mb)

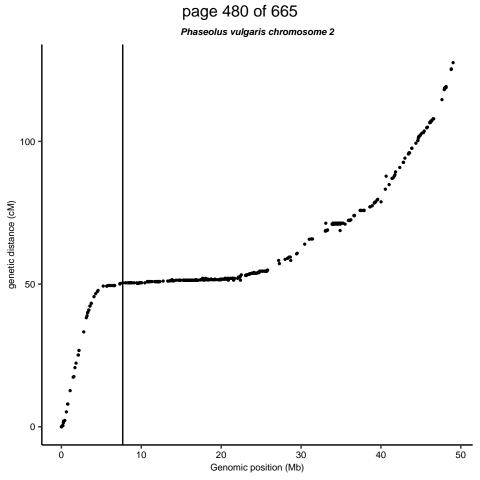


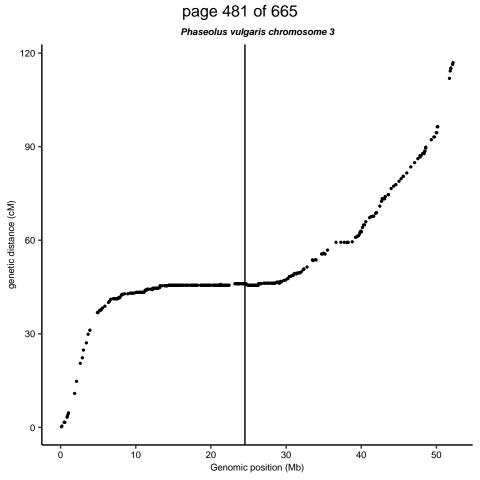


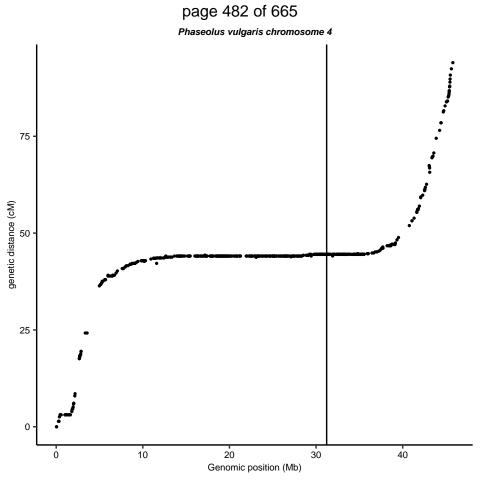


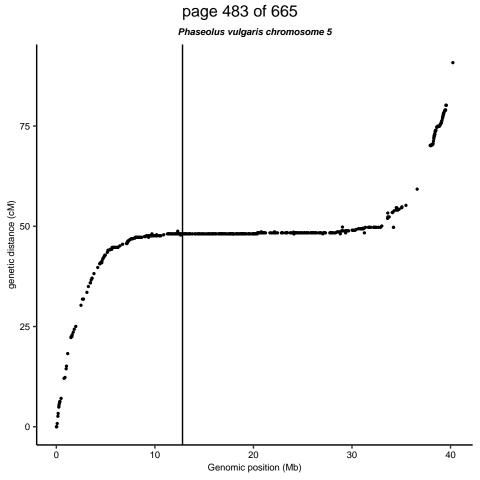


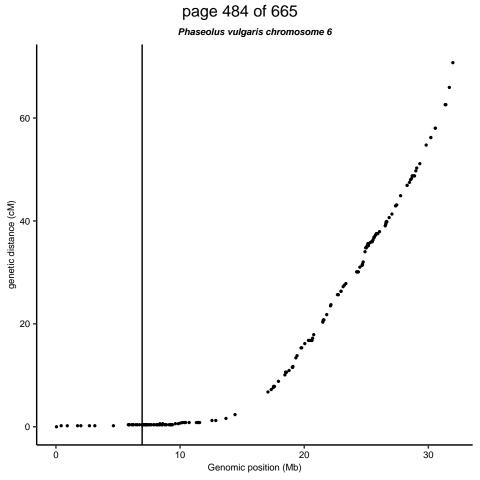




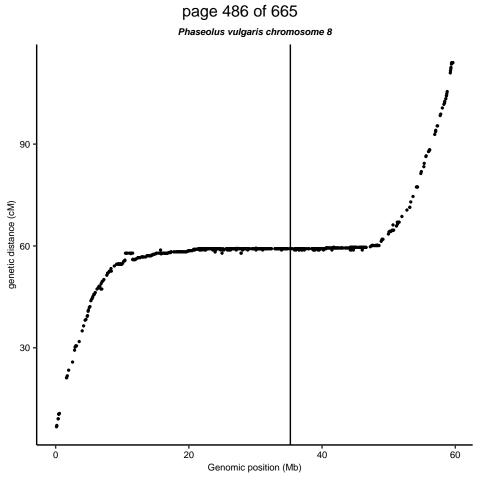


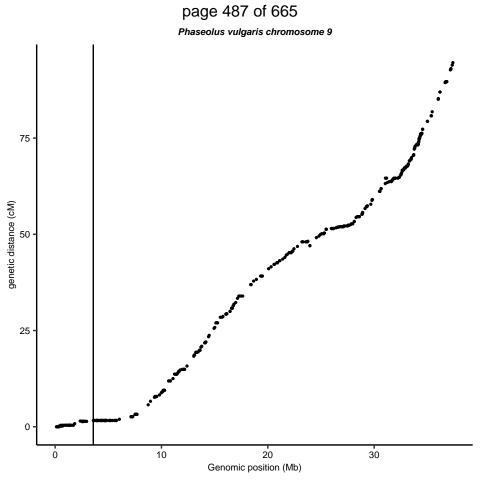




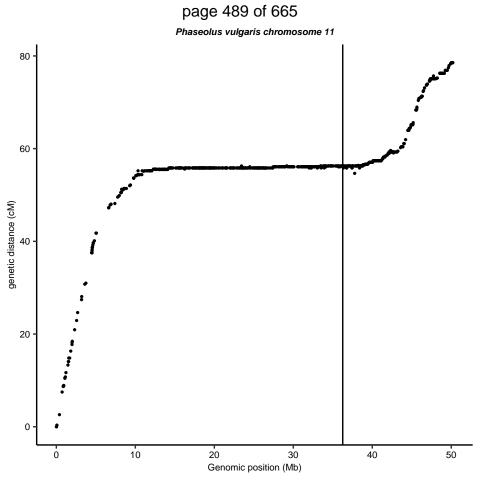


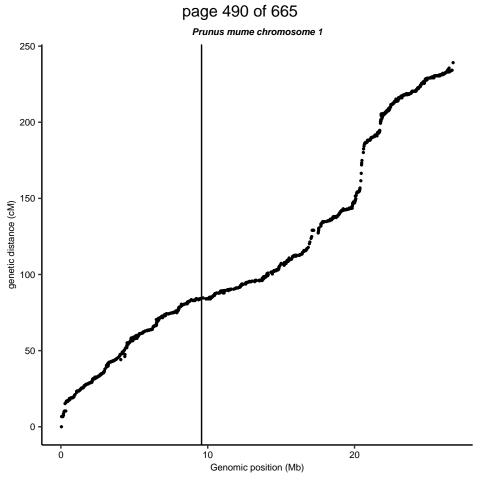
page 485 of 665 Phaseolus vulgaris chromosome 7 75 genetic distance (cM) Genomic position (Mb)

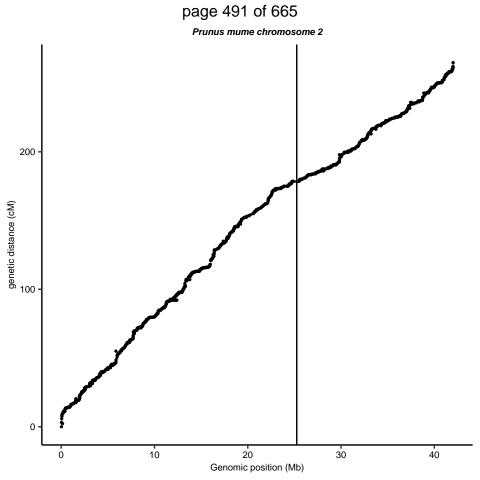


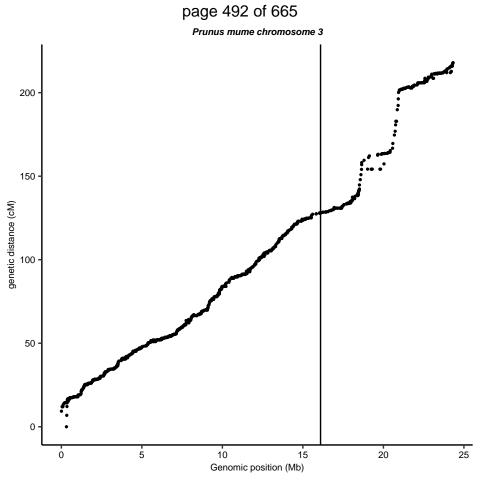


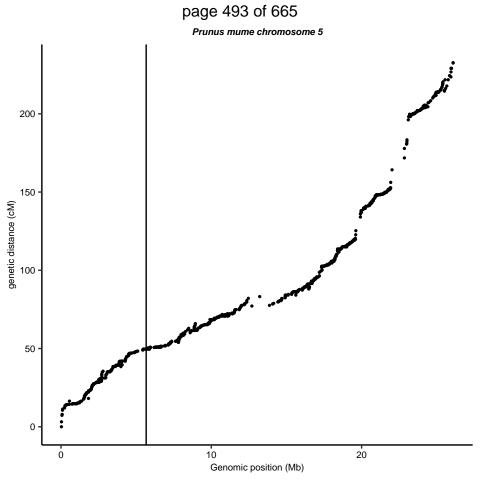
page 488 of 665 Phaseolus vulgaris chromosome 10 genetic distance (cM) Genomic position (Mb)

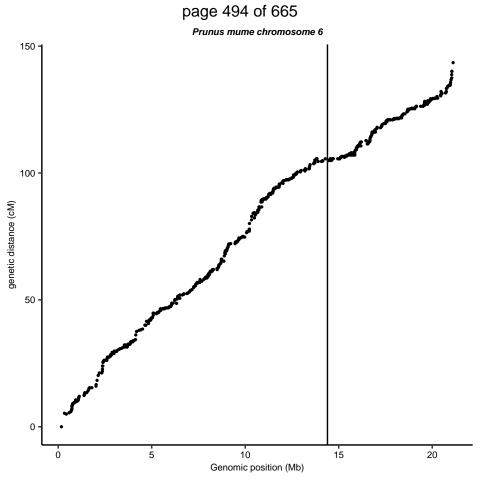


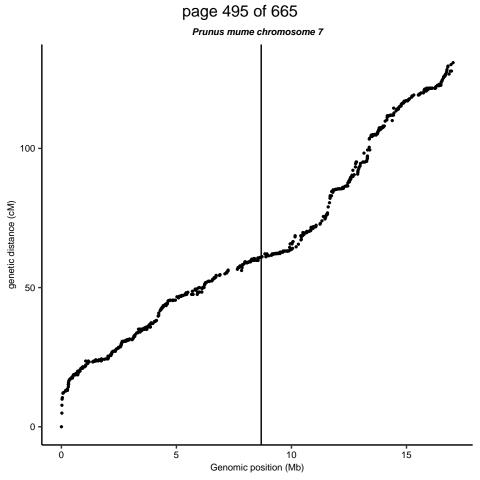


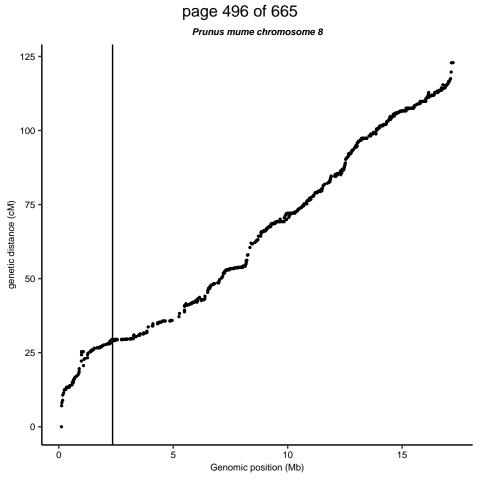


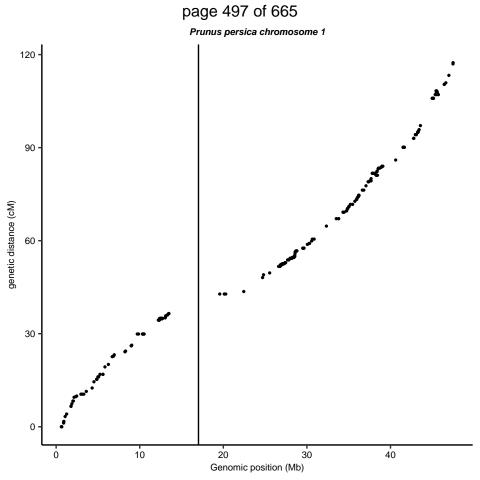


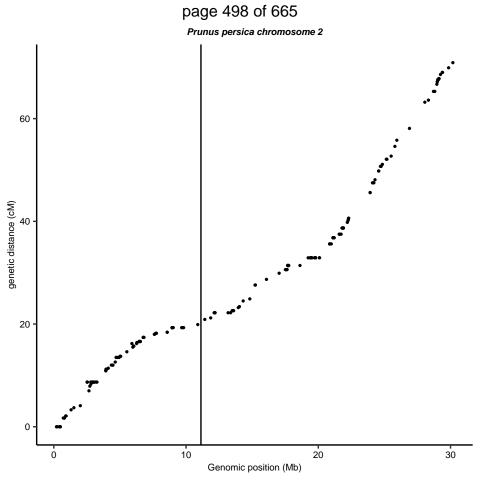


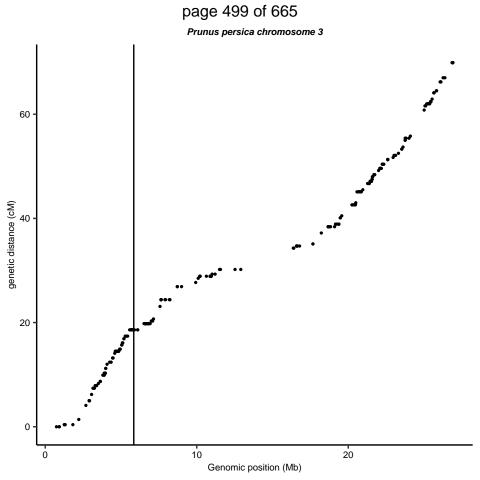


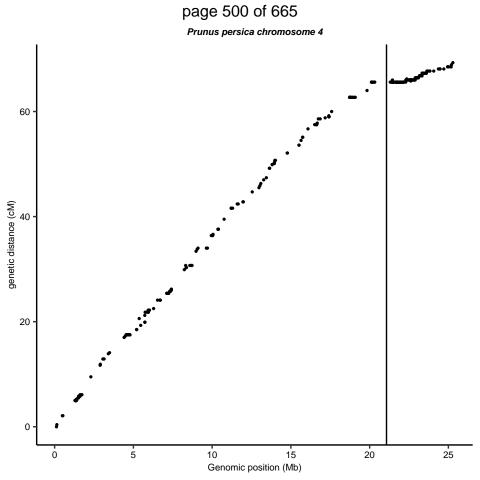


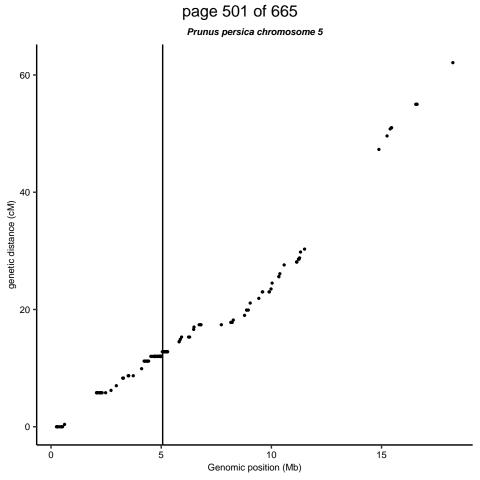


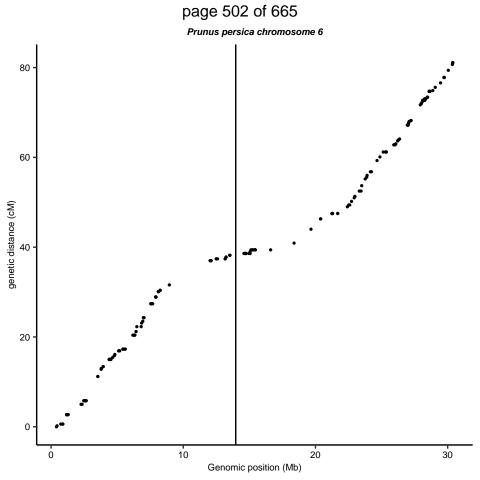


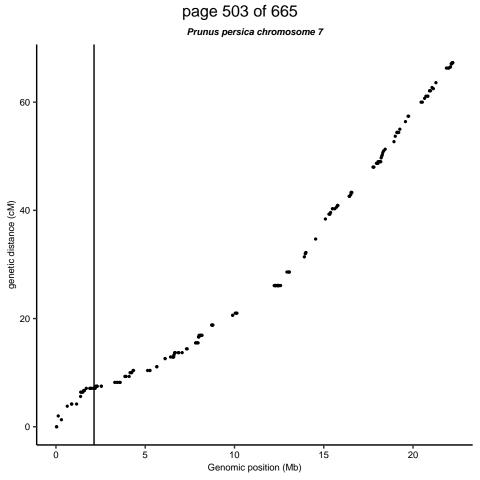


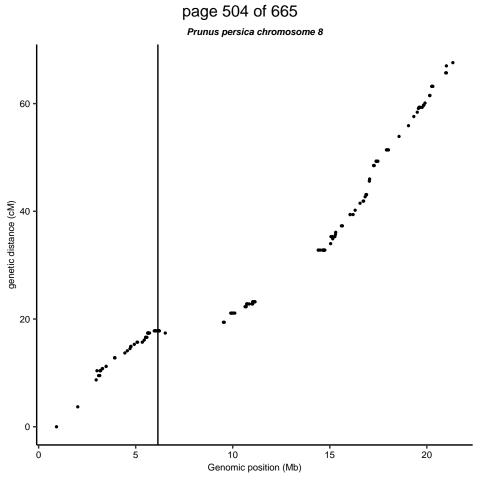


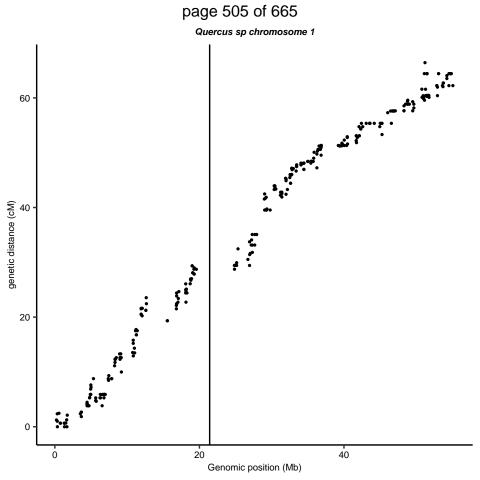


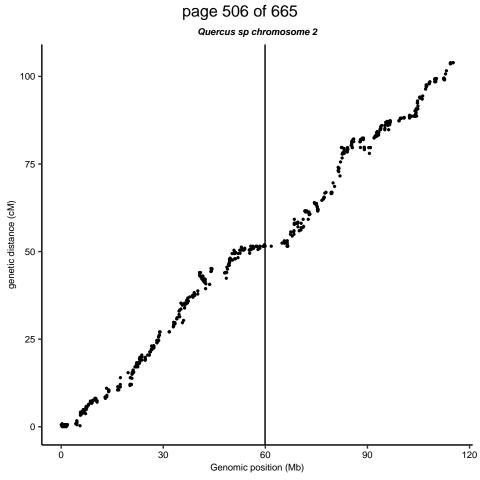


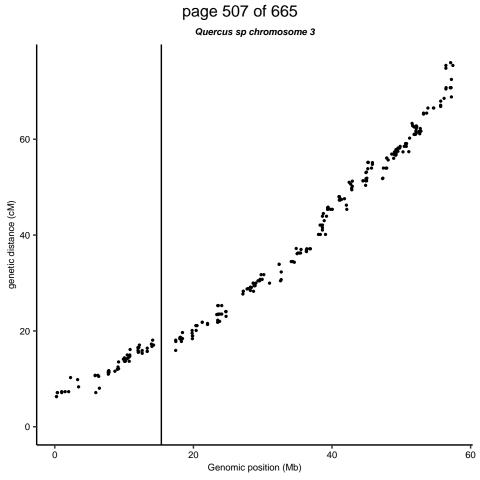


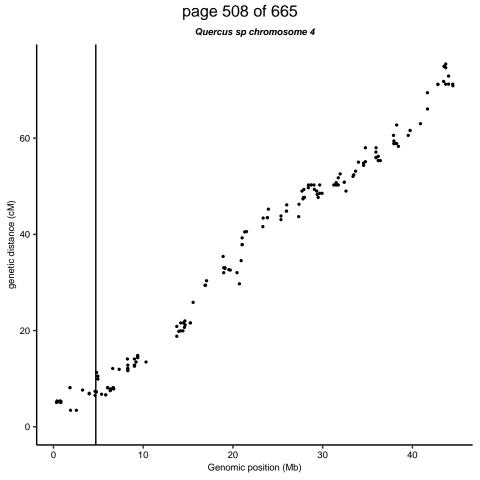


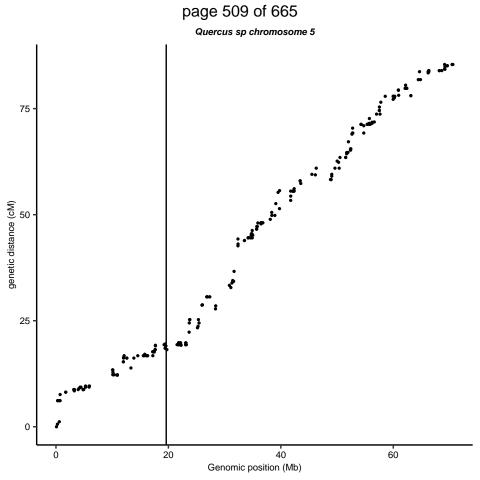


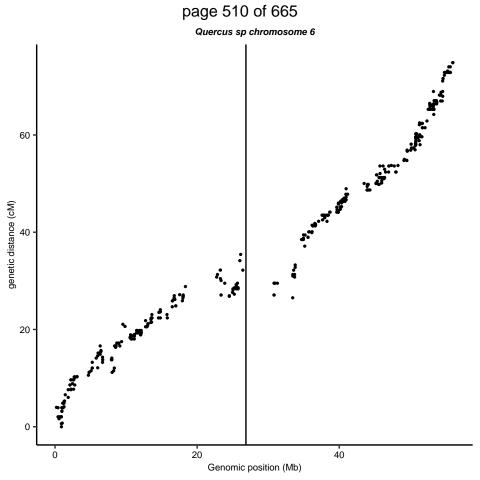


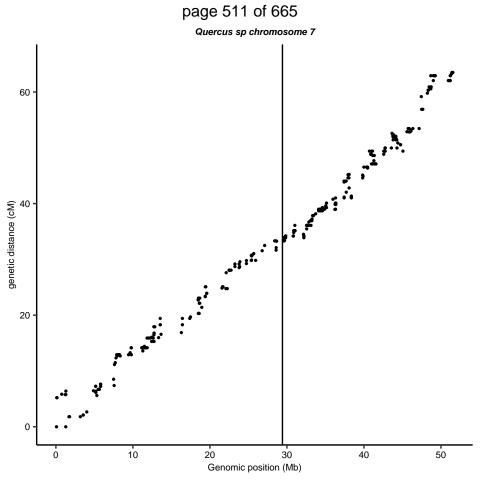


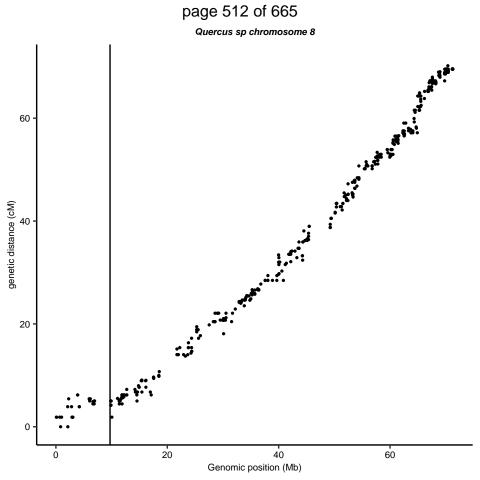


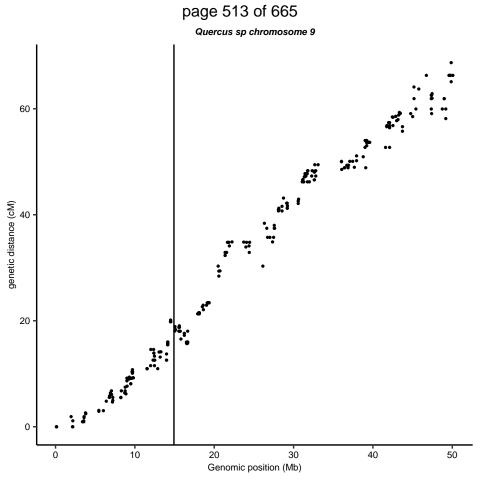


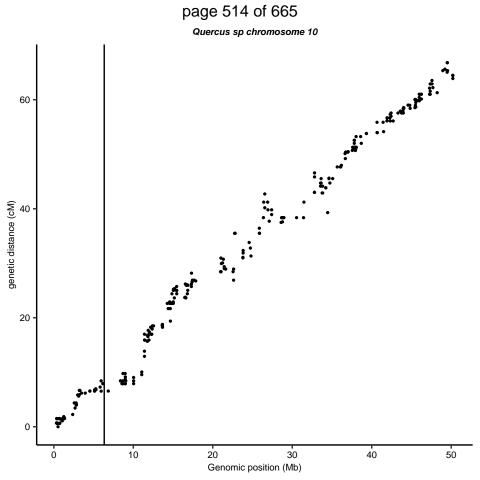


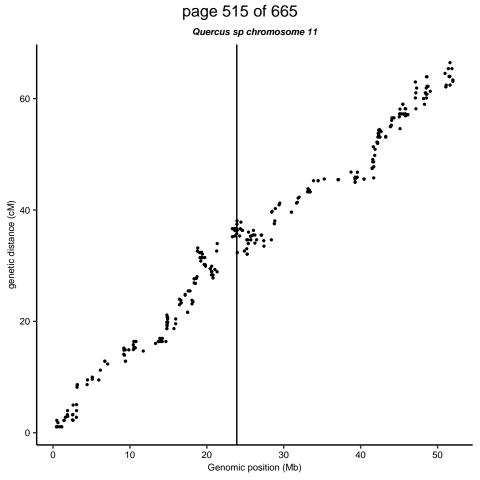


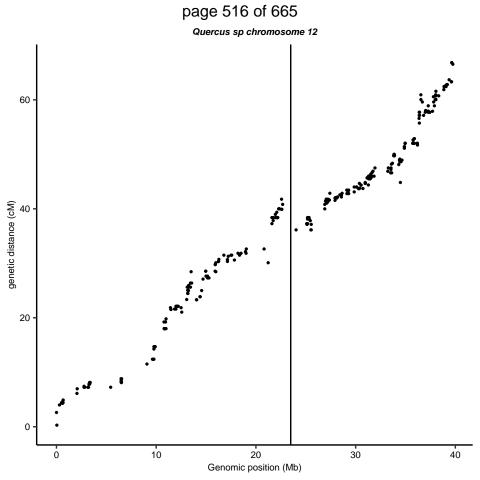


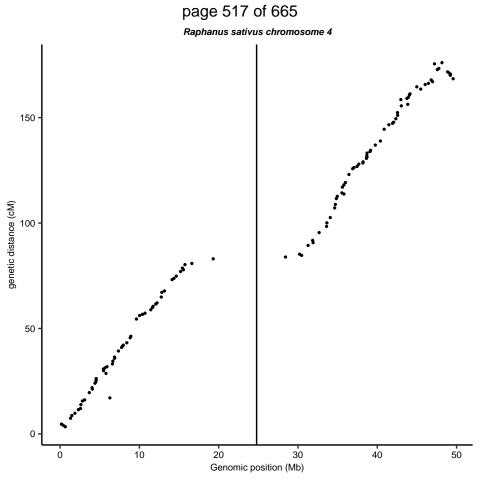


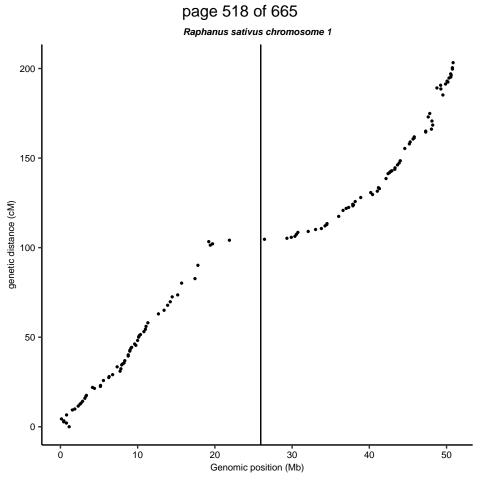


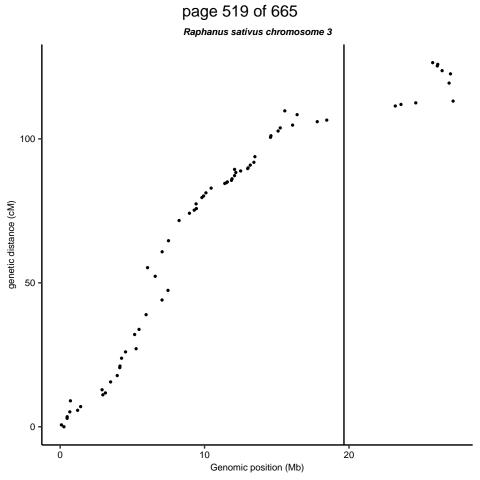


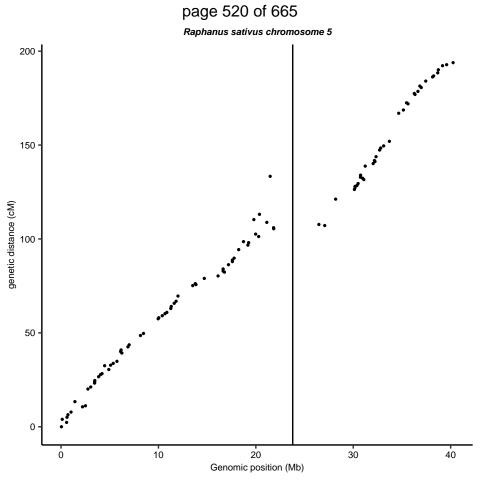


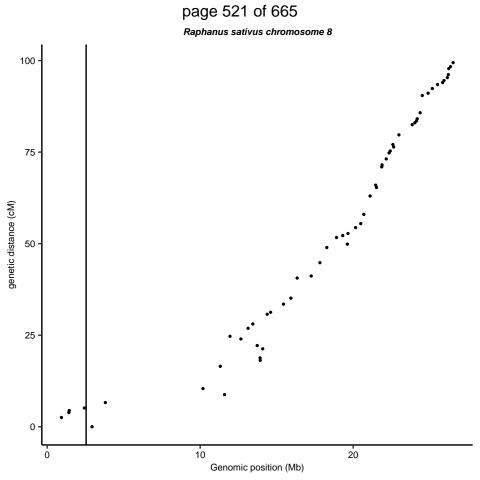


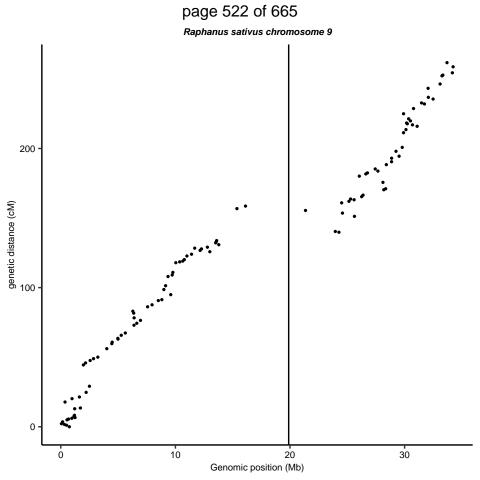


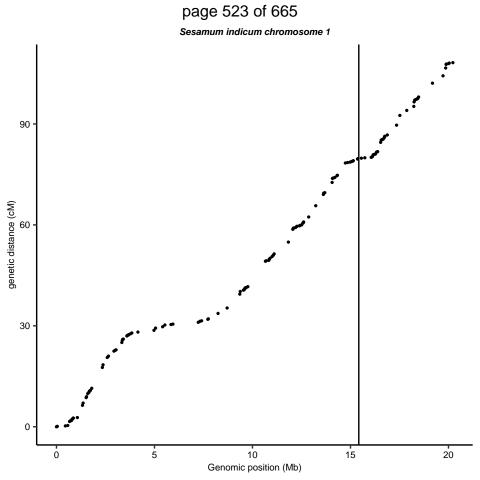


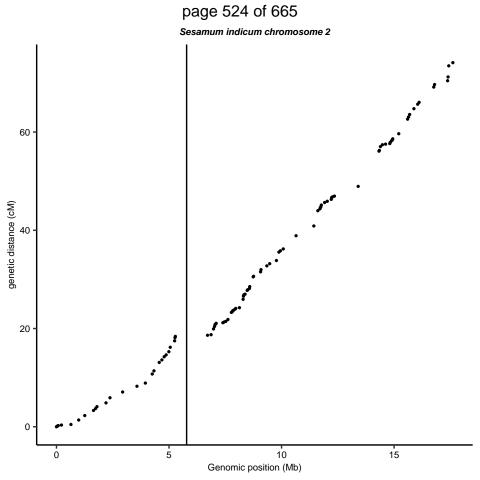


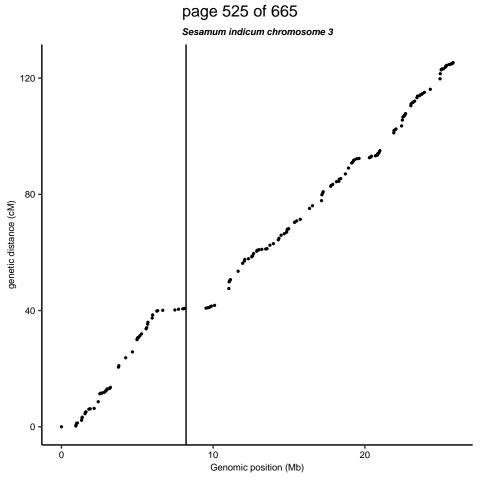




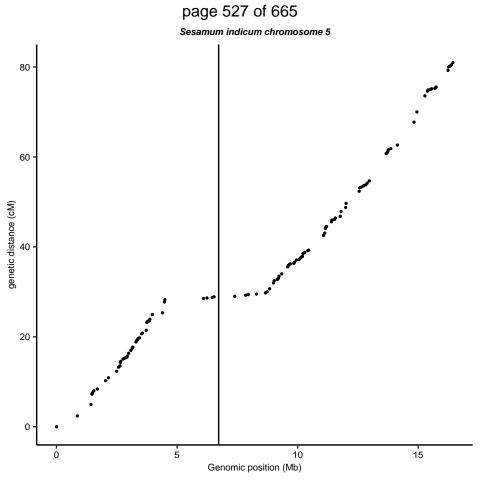


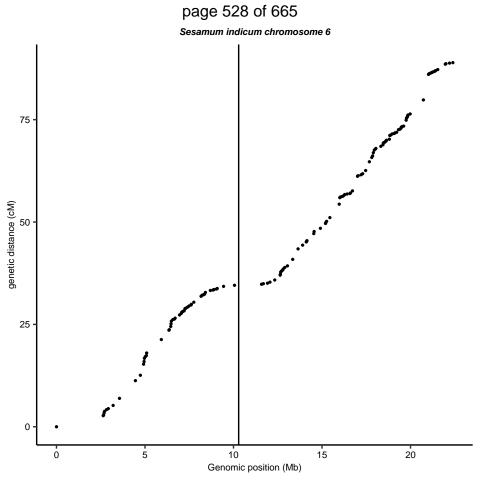


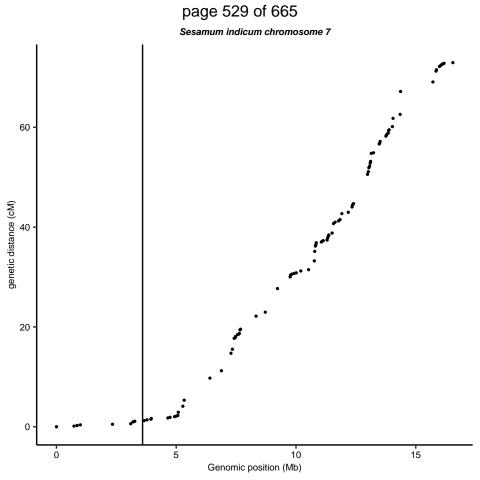


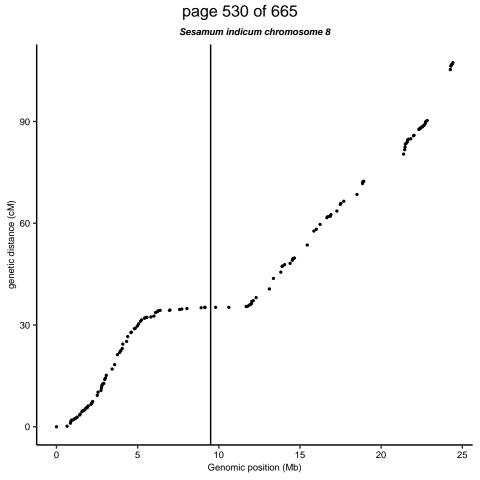


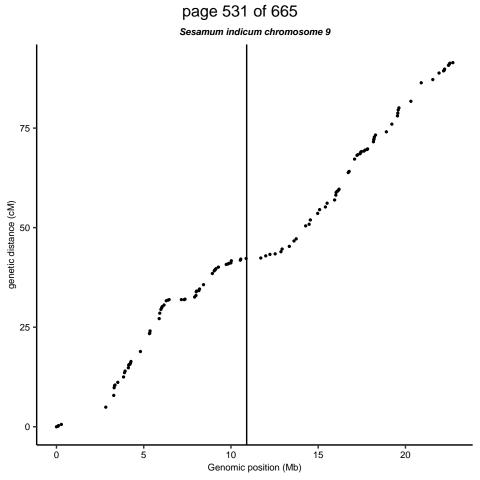
page 526 of 665 Sesamum indicum chromosome 4 80 -60 genetic distance (cM) 20 0 15 20 5 10 Genomic position (Mb)

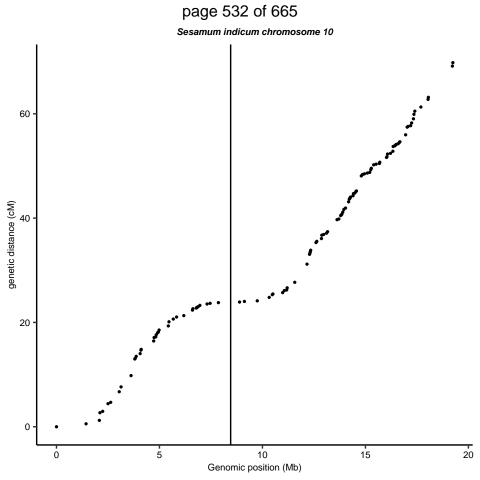


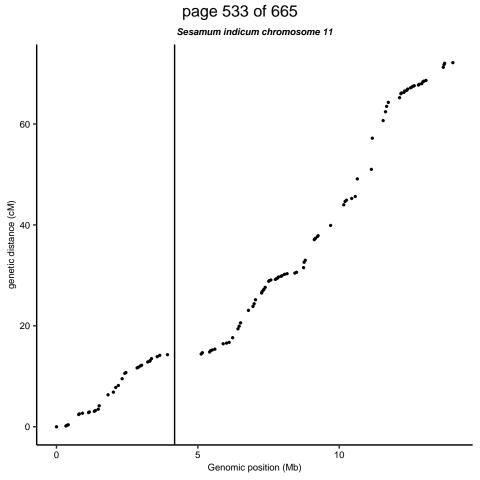


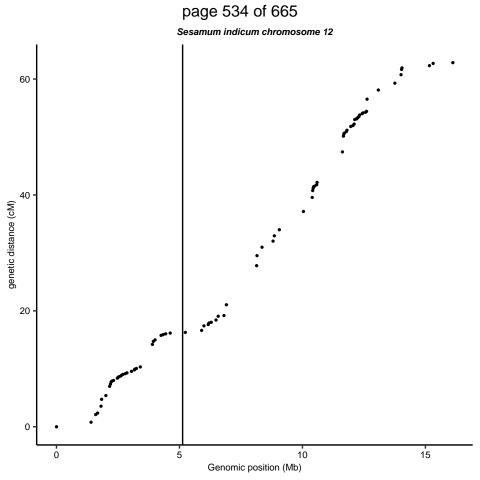


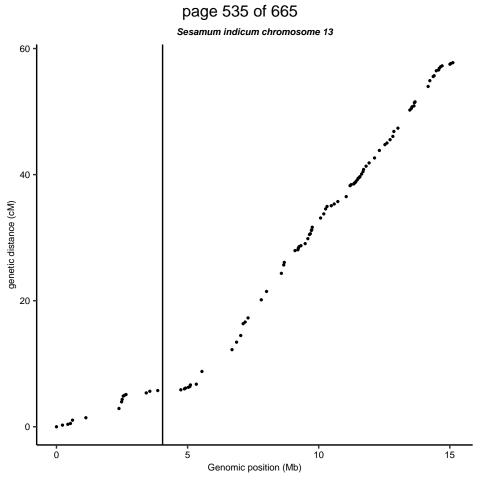


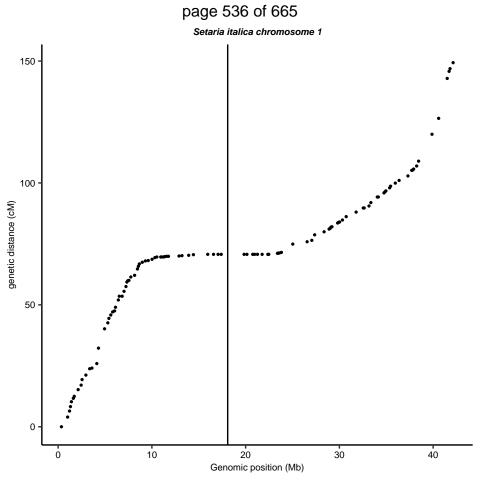




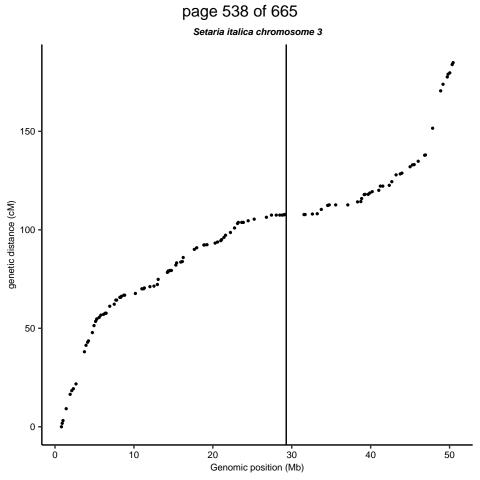


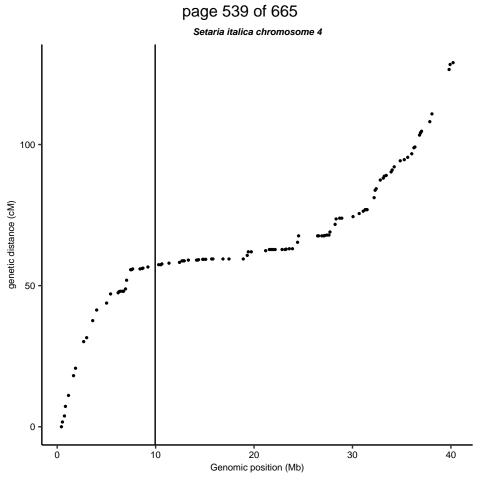


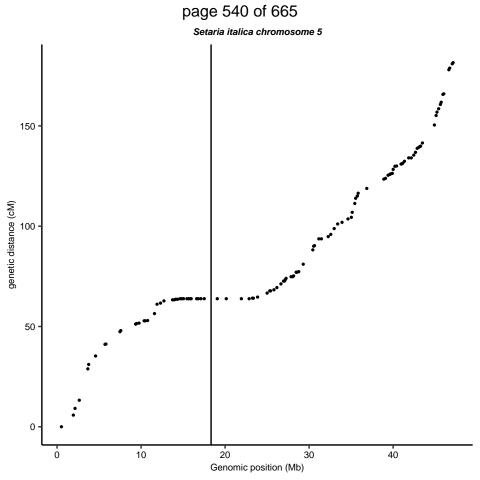


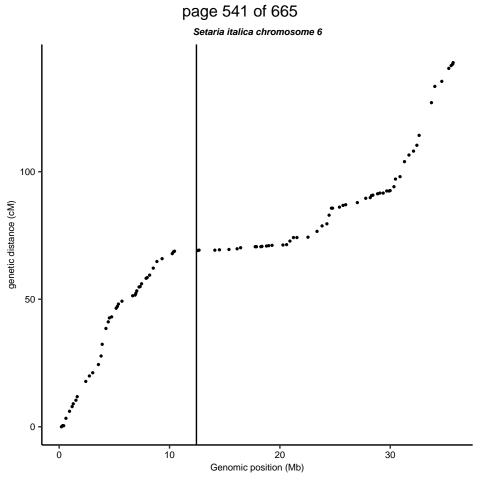


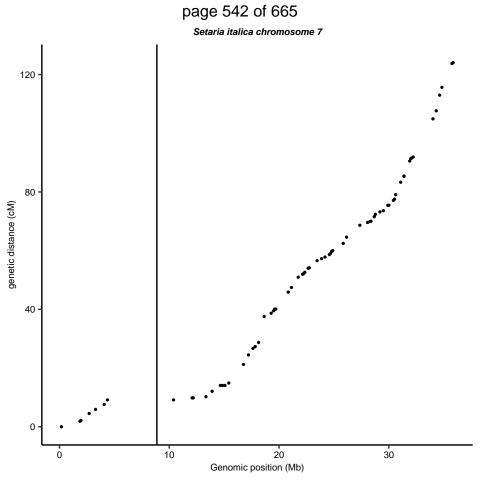
page 537 of 665 Setaria italica chromosome 2 genetic distance (cM) Genomic position (Mb)

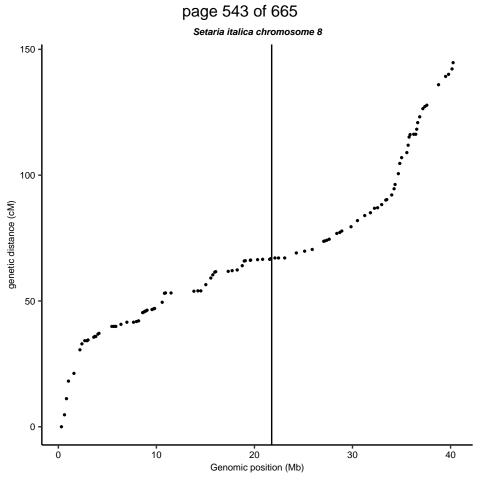


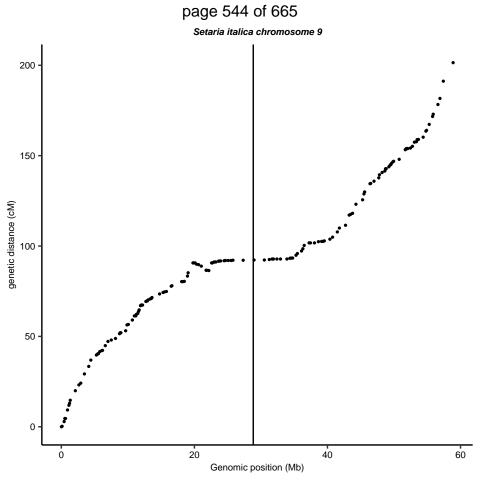






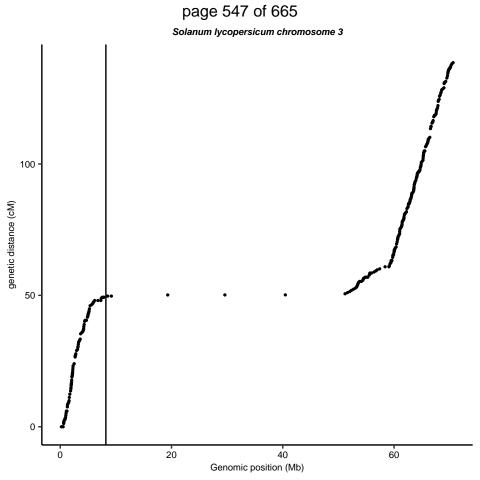






page 545 of 665 Solanum lycopersicum chromosome 1 150 genetic distance (cM) 50 0 -25 75 50 100 Genomic position (Mb)

page 546 of 665 Solanum lycopersicum chromosome 2 genetic distance (cM) Genomic position (Mb)

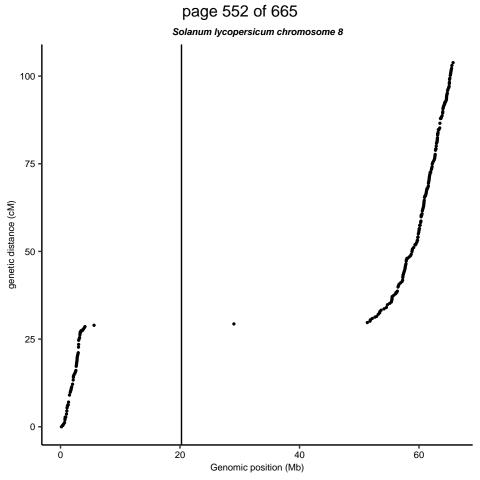


page 548 of 665 Solanum lycopersicum chromosome 4 genetic distance (cM) Genomic position (Mb)

page 549 of 665 Solanum lycopersicum chromosome 5 90 genetic distance (cM) 30 0 -20 40 60 ò Genomic position (Mb)

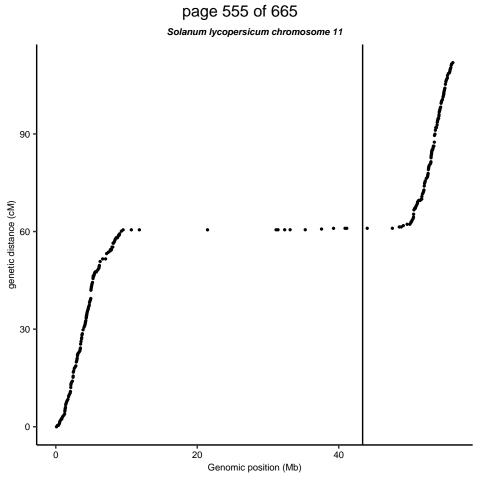
page 550 of 665 Solanum lycopersicum chromosome 6 60 genetic distance (cM) 20 0 -10 20 40 50 30 Genomic position (Mb)

page 551 of 665 Solanum lycopersicum chromosome 7 100 75 genetic distance (cM) 50 25 0 20 60 40 Genomic position (Mb)



page 553 of 665 Solanum lycopersicum chromosome 9 125 100 genetic distance (cM) 75 -50 25 -0 -20 40 60 Genomic position (Mb)

page 554 of 665 Solanum lycopersicum chromosome 10 100 75 genetic distance (cM) 50 25 -0 20 40 60 Genomic position (Mb)



page 556 of 665 Solanum lycopersicum chromosome 12 120 -90 genetic distance (cM) 60 30

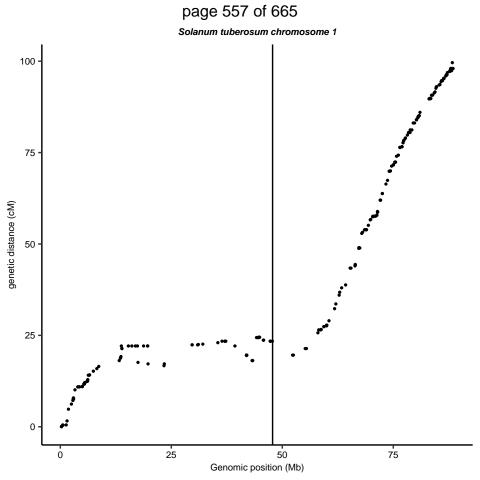
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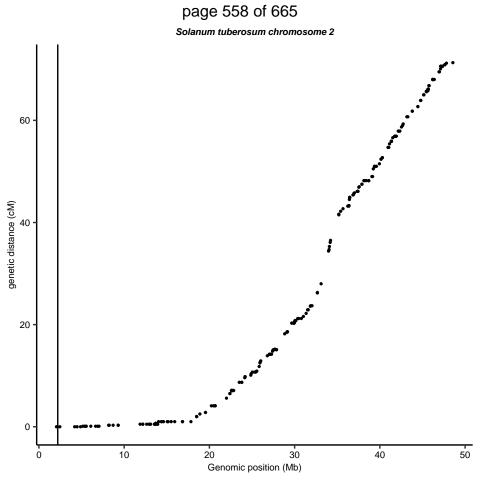
Genomic position (Mb)

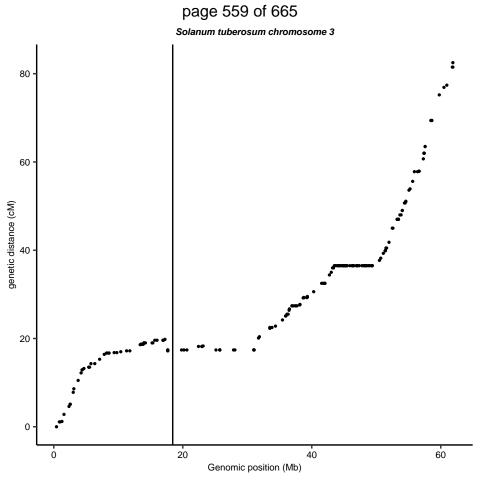
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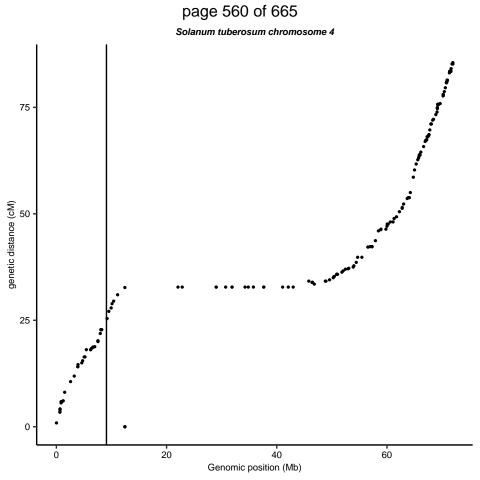
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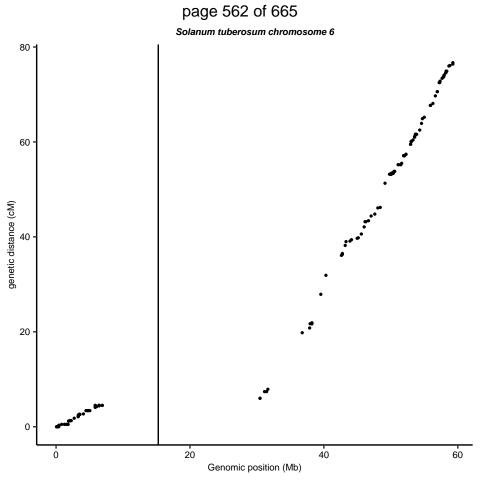


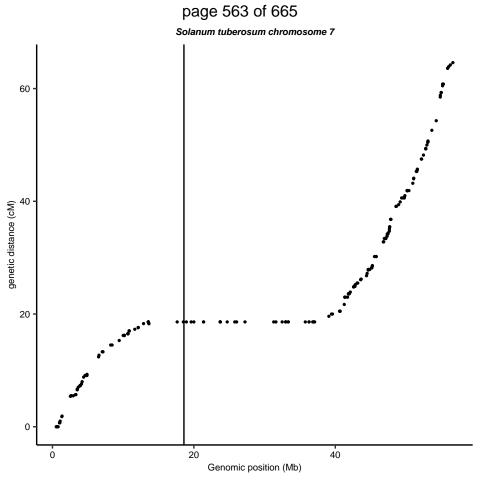


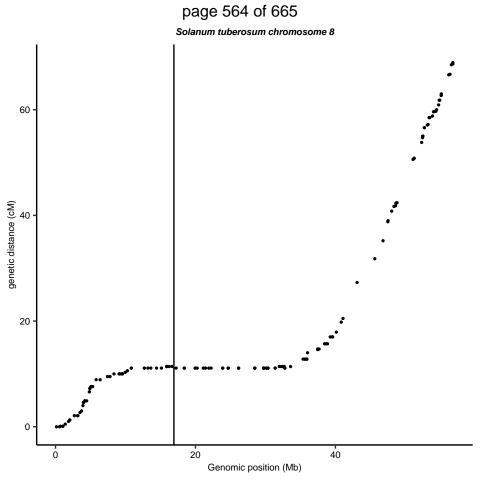


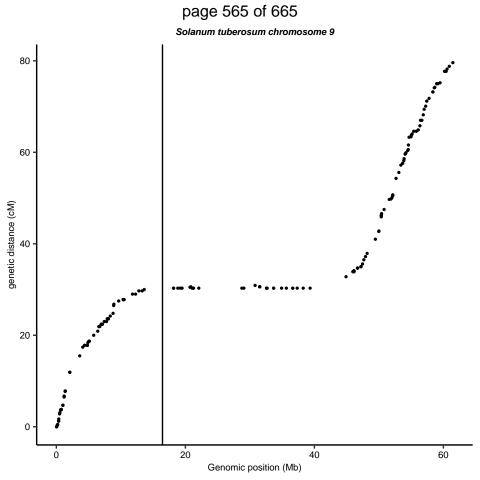


page 561 of 665 Solanum tuberosum chromosome 5 60 genetic distance (cM) 20 0 -10 20 50 40 Ö 30 Genomic position (Mb)





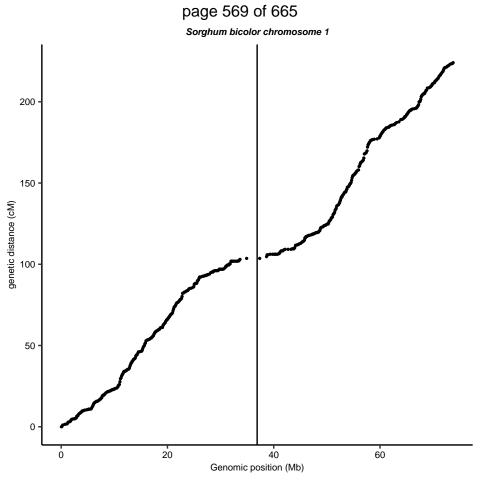


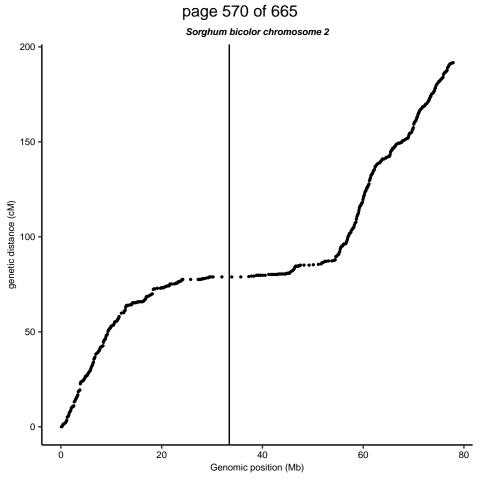


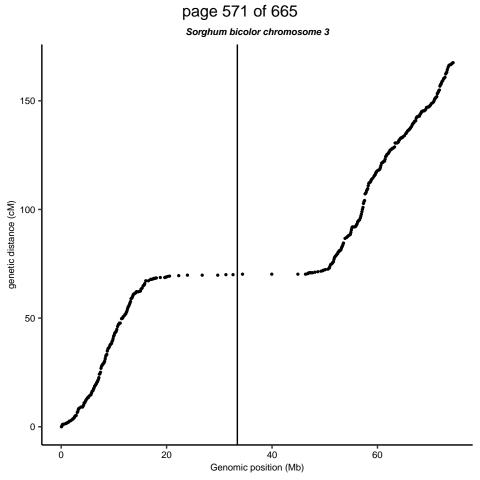
page 566 of 665 Solanum tuberosum chromosome 10 genetic distance (cM) Genomic position (Mb)

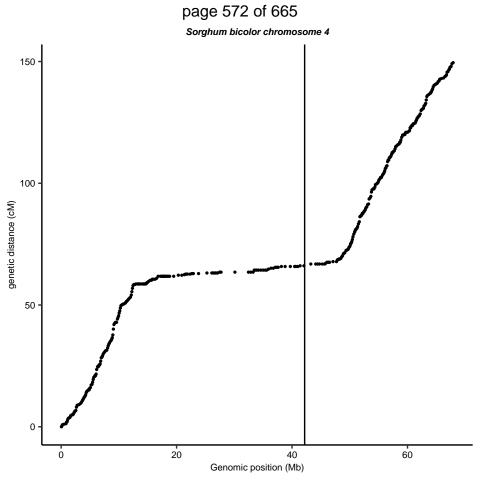
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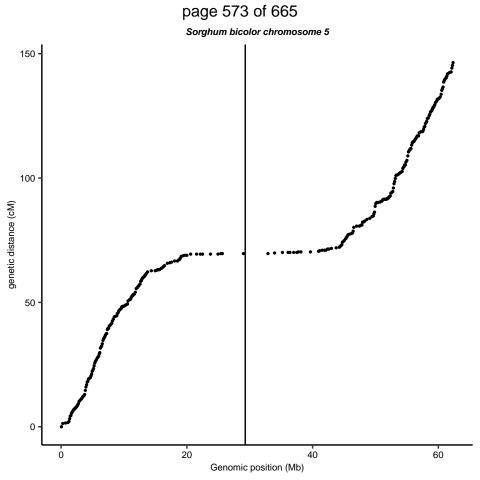
page 568 of 665 Solanum tuberosum chromosome 12 60 genetic distance (cM) 20 0 20 60 40 Genomic position (Mb)

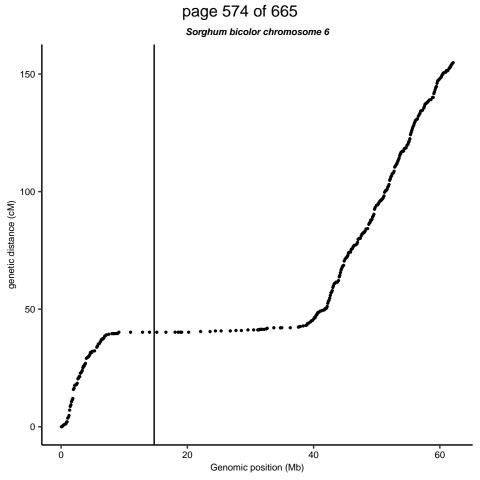


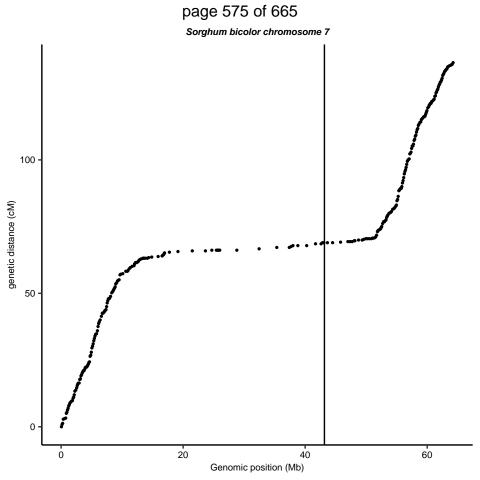


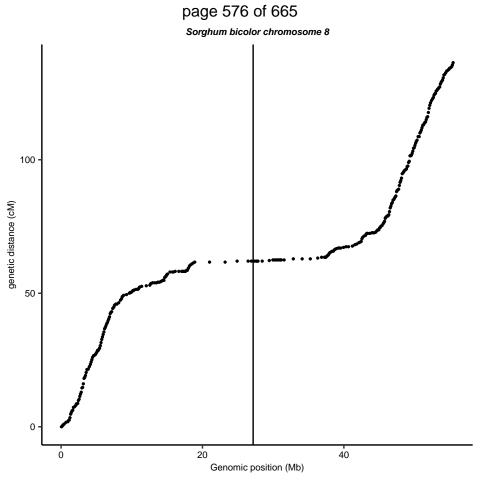


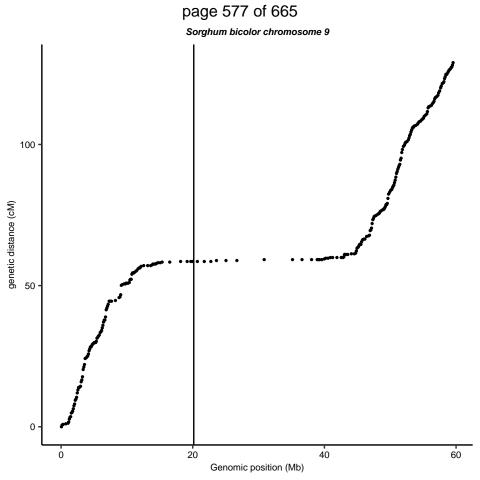


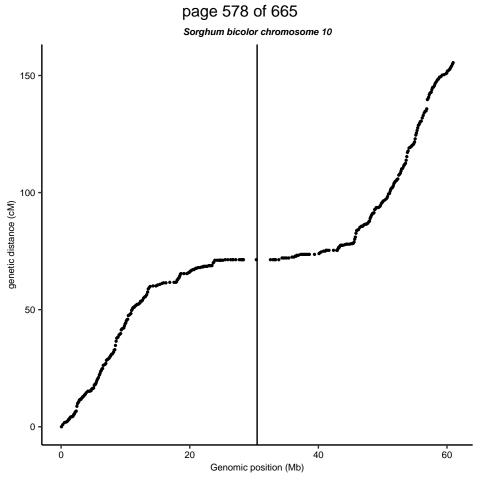


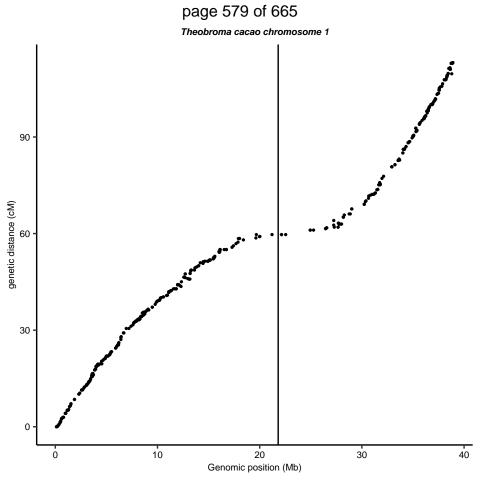


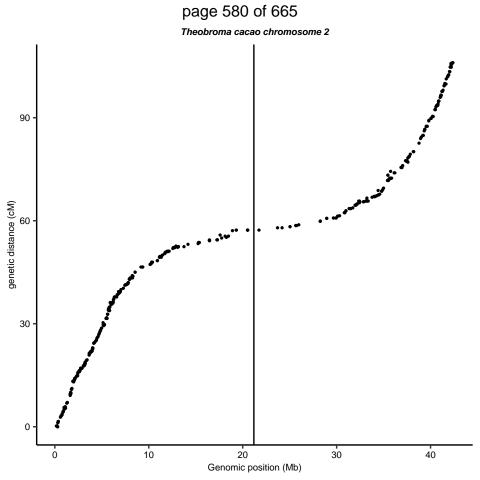


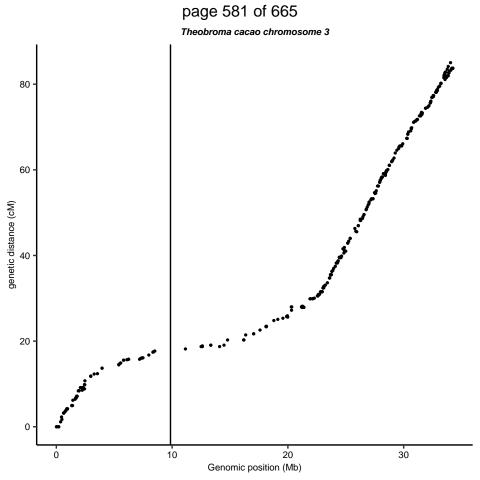


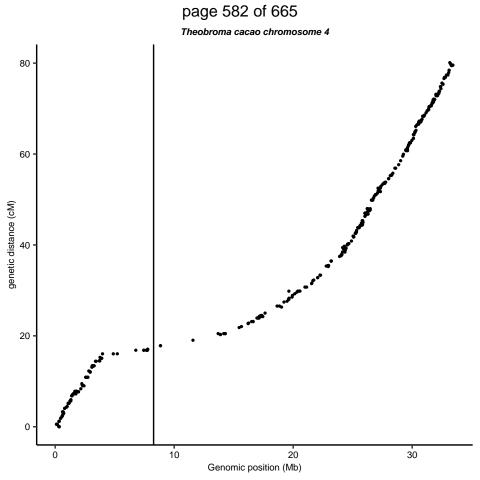


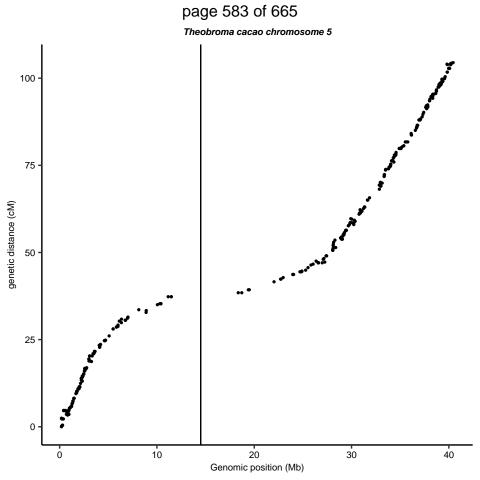


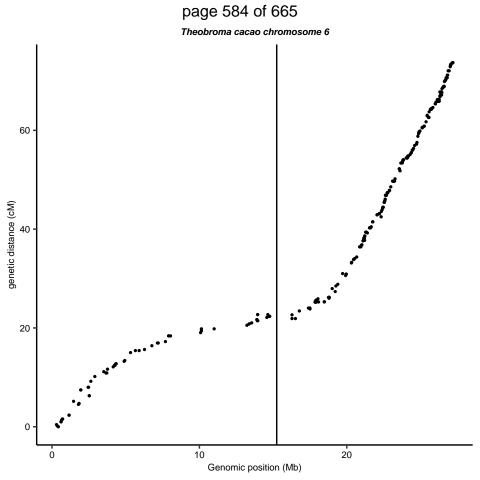


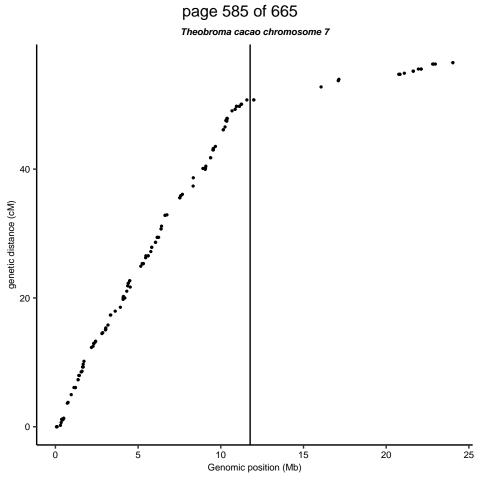


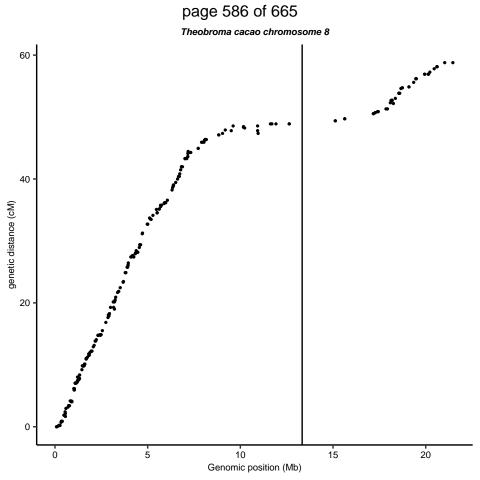


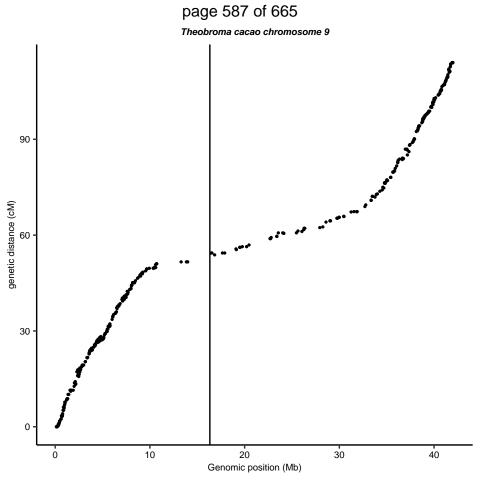




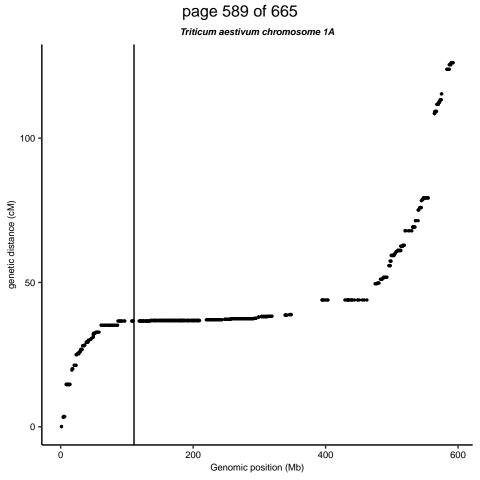


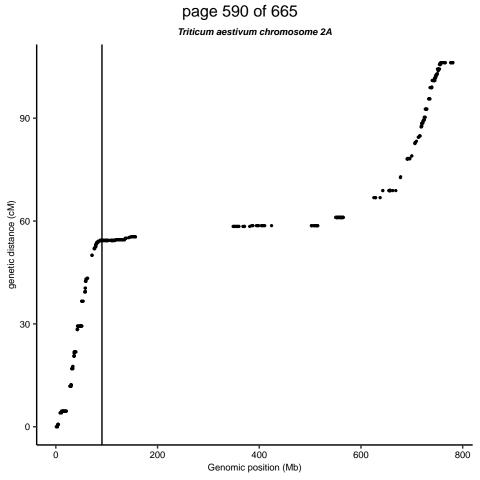


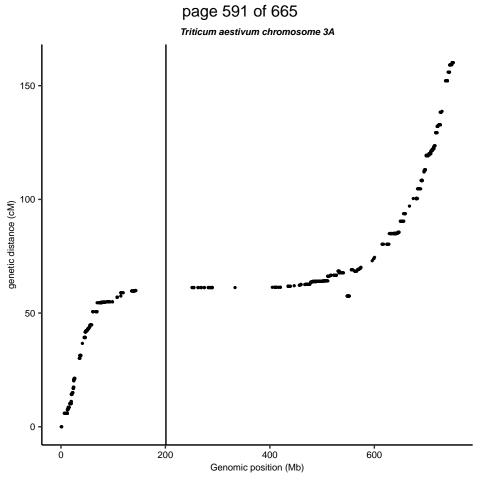




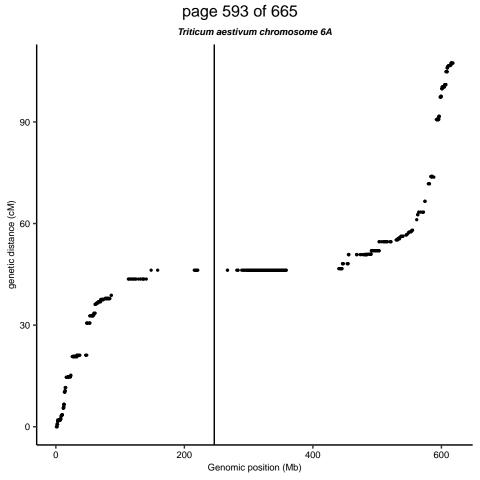
page 588 of 665 Theobroma cacao chromosome 10 genetic distance (cM) Genomic position (Mb)

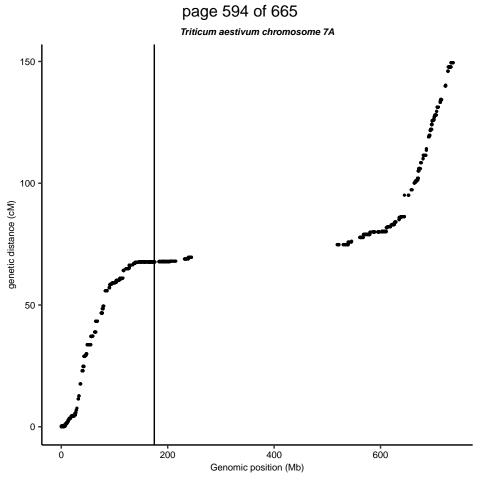


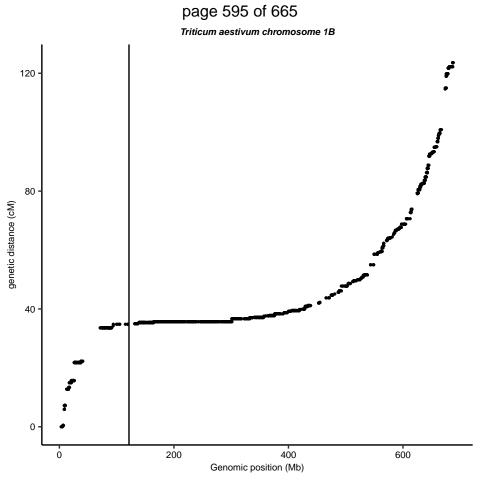


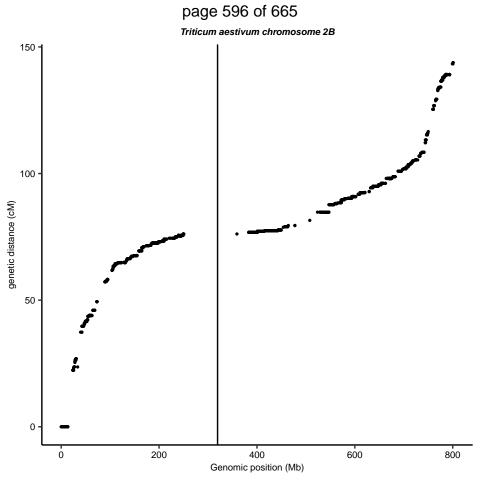


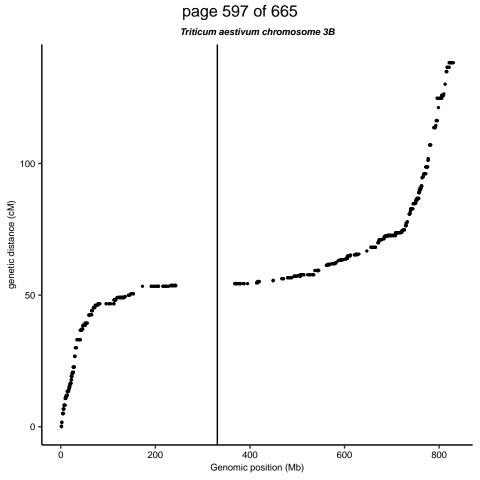
page 592 of 665 Triticum aestivum chromosome 4A 100 genetic distance (cM) 50 200 600 400 Genomic position (Mb)

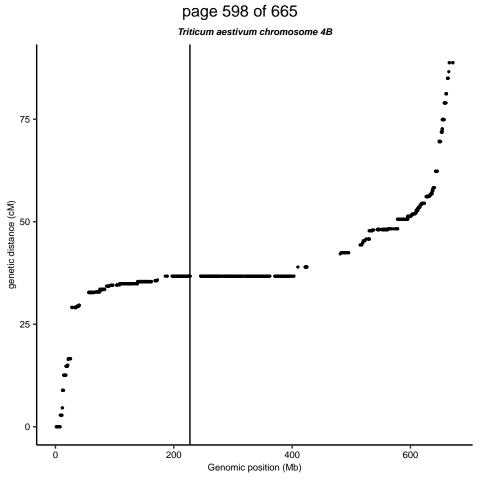


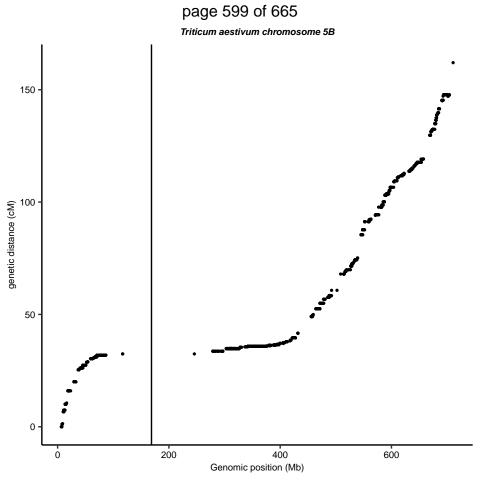


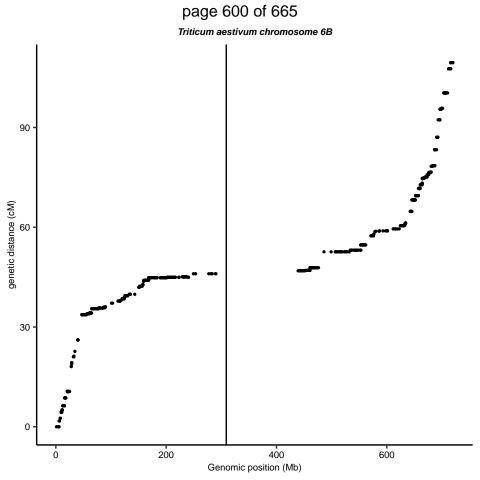


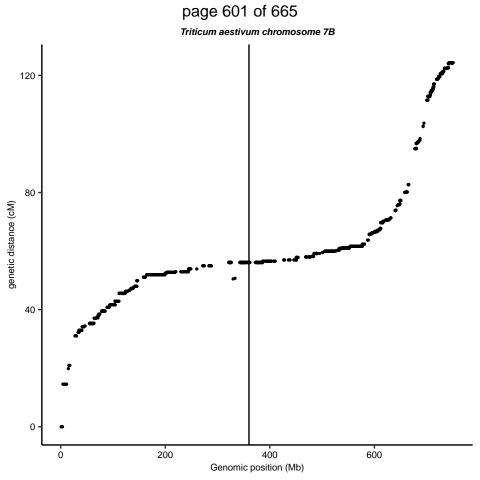


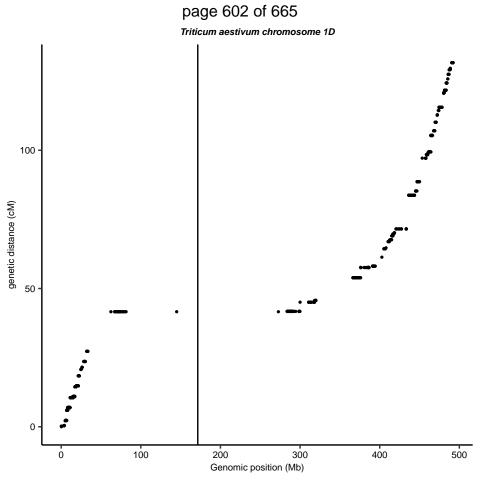


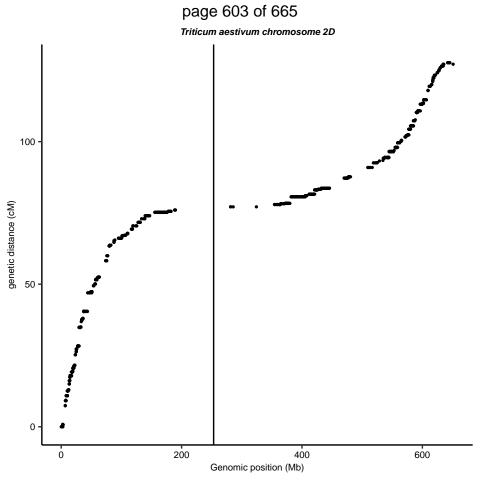


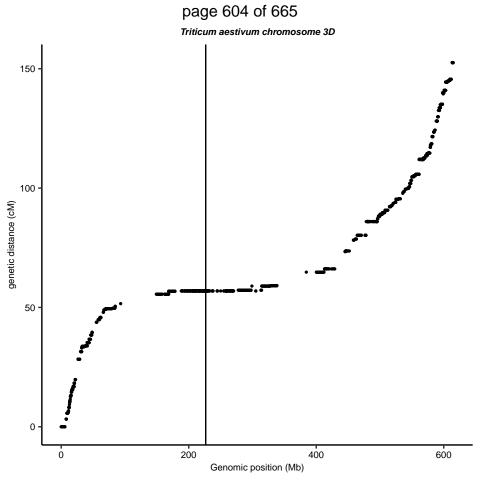


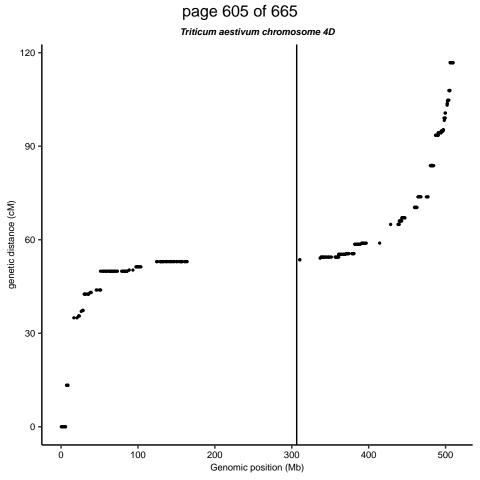


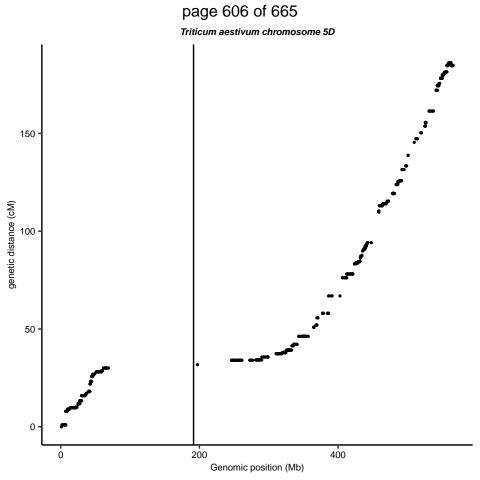


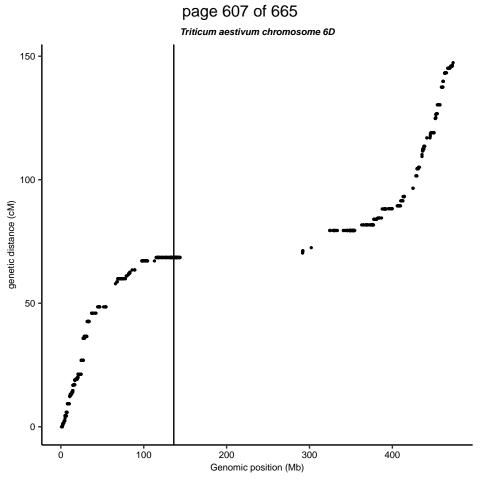


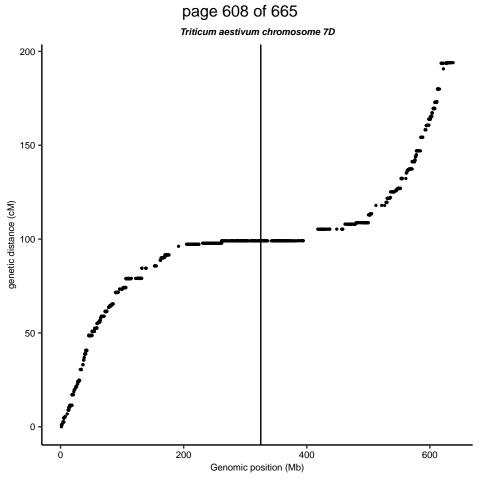


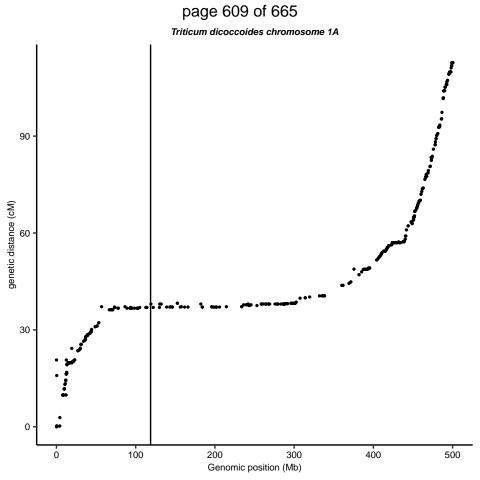


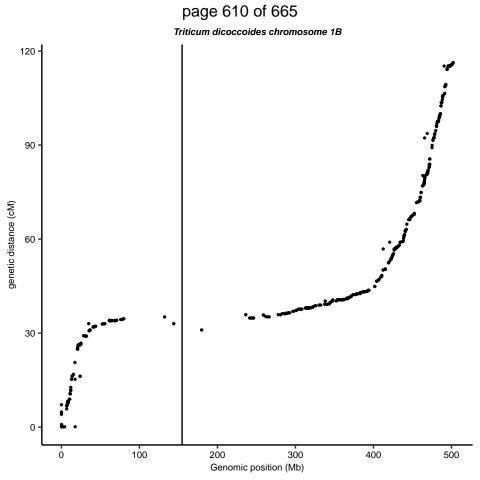


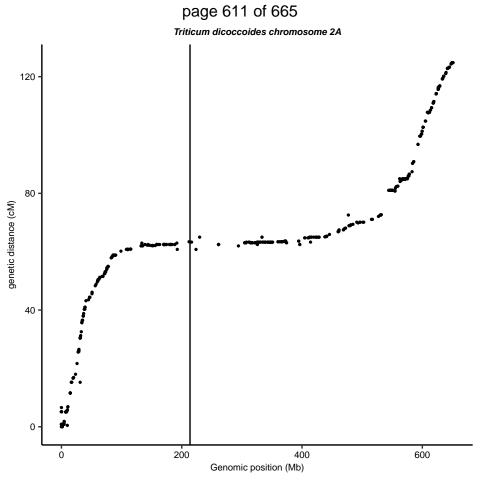


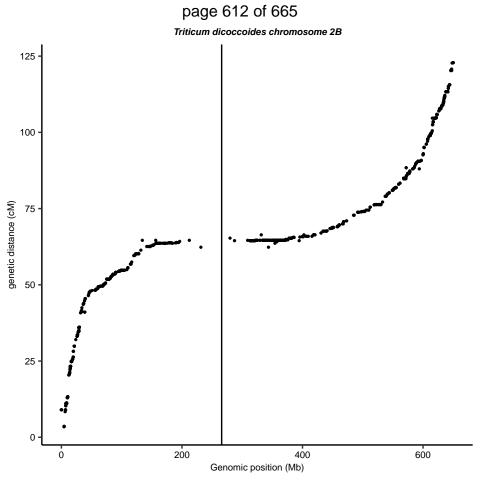


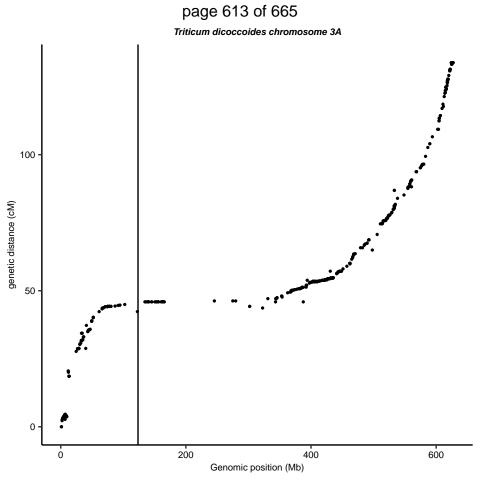




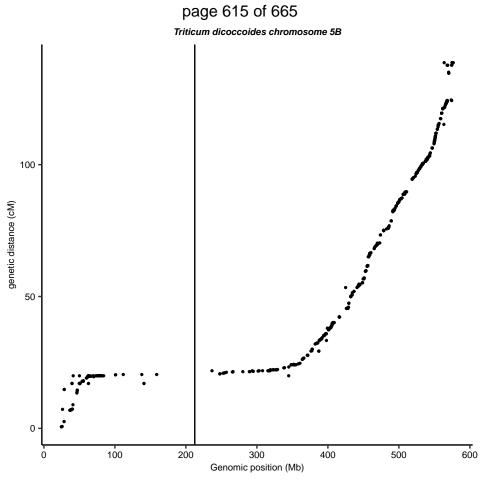








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page 616 of 665 Triticum dicoccoides chromosome 6A 75 genetic distance (cM) Genomic position (Mb)

