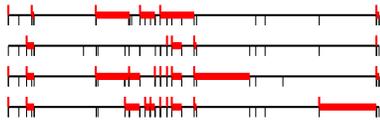


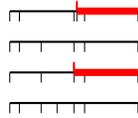
Haplotype blocks identified in Gabriel et al. (2002) data

Schematic of the haplotype blocks identified in fifty genomic regions (data of Gabriel et al. 2002), using the block definition of Gabriel et al. (2002), with a slight modification (described in Wall and Pritchard 2003, AJHG, submitted). The four lines for each region represent data from four population samples: European-Americans, African-Americans, East Asians and sub-Saharan Africans, respectively. The horizontal red lines denote haplotype blocks; the left-hand end of each block is marked by an upper red tick mark. The lower tick marks indicate the locations of the markers typed in each population. This is an expanded version of Figure 3 in Wall and Pritchard (2003, Nature Reviews Genetics, submitted).

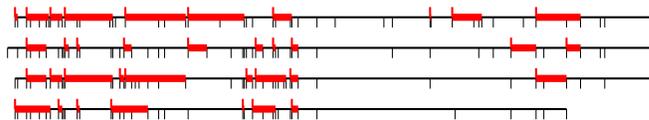
Region 1a
(3.35 cMMb)



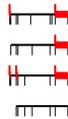
Region 1b
(3.35 cMMb)



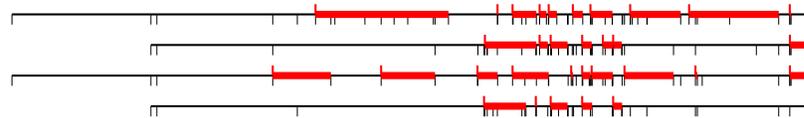
Region 2a
(2.99 cMMb)



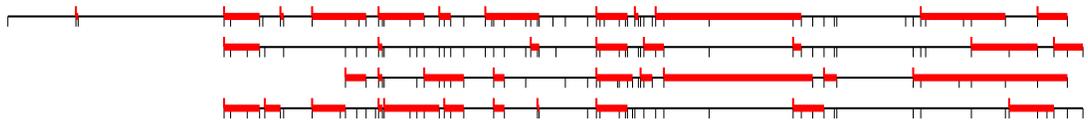
Region 2b
(2.99 cMMb)



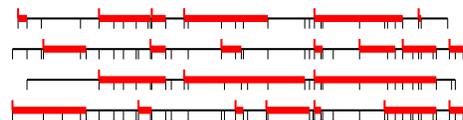
Region 3a
(1.07 cMMb)



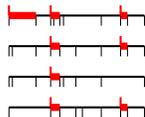
Region 4a
(1.17 cMMb)



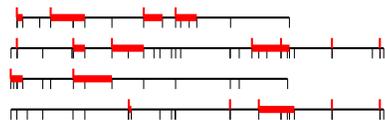
Region 7a
(0.82 cMMb)



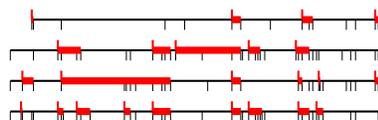
Region 7b
(1.00 cMMb)



Region 8a
(2.12 cMMb)

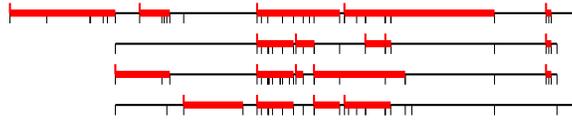


Region 9a
(1.37 cMMb)

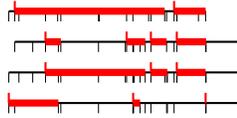


100 Kb

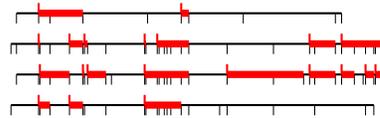
Region 10a
(0.49 cMMb)



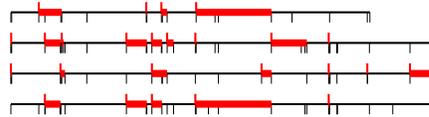
Region 11a
(0.85 cMMb)



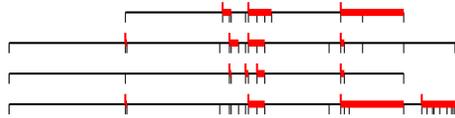
Region 13a
(0.92 cMMb)



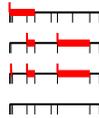
Region 14a
(1.50 cMMb)



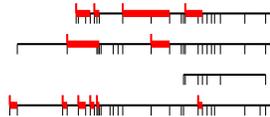
Region 15a
(1.28 cMMb)



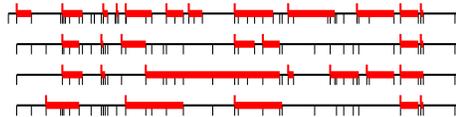
Region 16a
(1.36 cMMb)



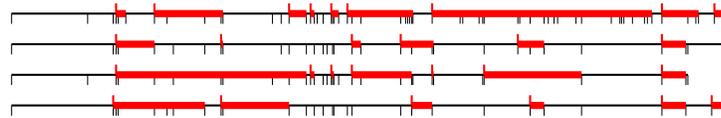
Region 16b
(1.36 cMMb)



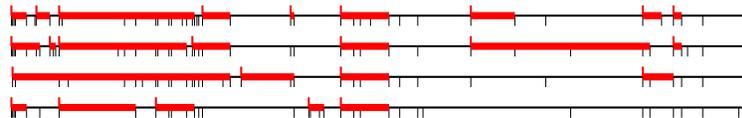
Region 17a
(0.58 cMMb)



Region 18a
(1.19 cMMb)

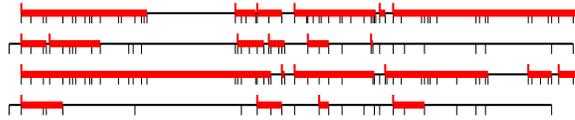


Region 19a
(0.49 cMMb)

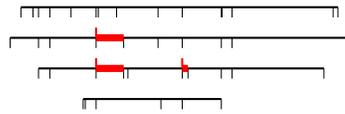


100 Kb

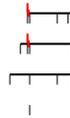
Region 20a
(0.80 cMMb)



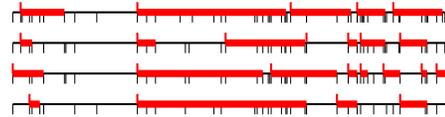
Region 21a
(2.55 cMMb)



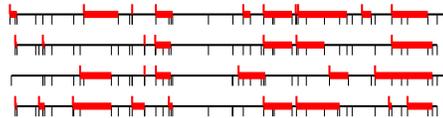
Region 21b
(2.55 cMMb)



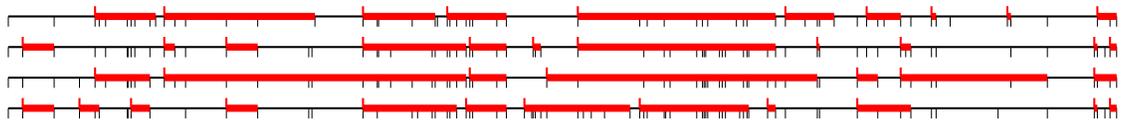
Region 22a
(0.85 cMMb)



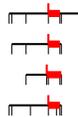
Region 23a
(1.05 cMMb)



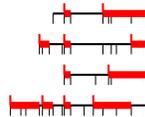
Region 24a
(1.33 cMMb)



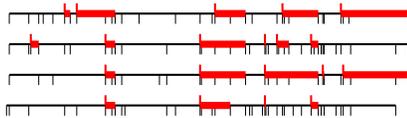
Region 25a
(0.57 cMMb)



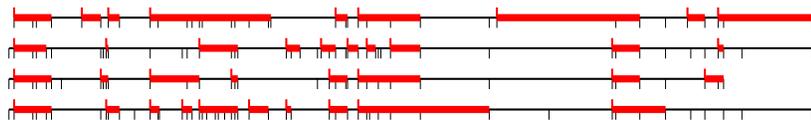
Region 25b
(0.57 cMMb)



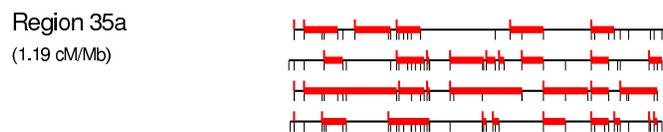
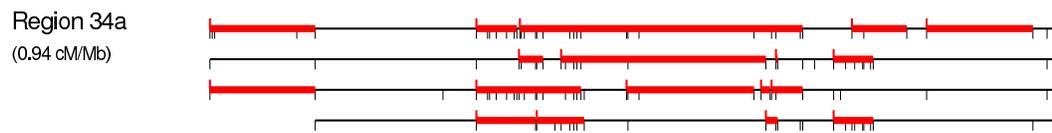
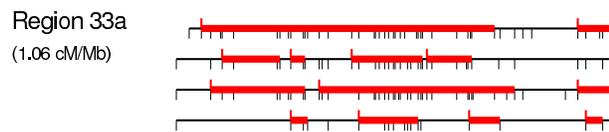
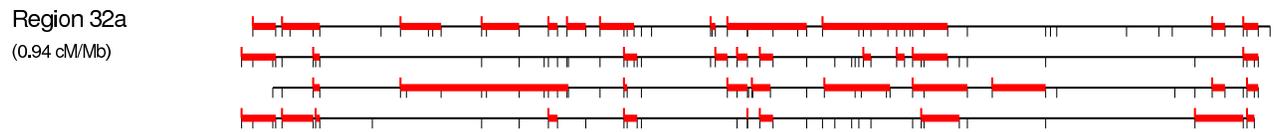
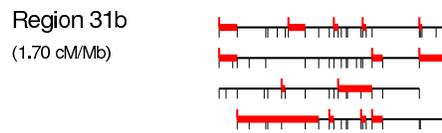
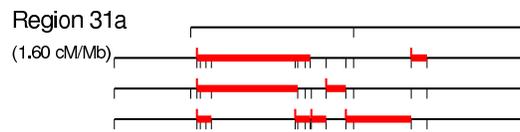
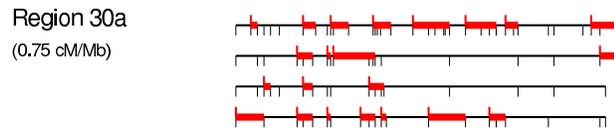
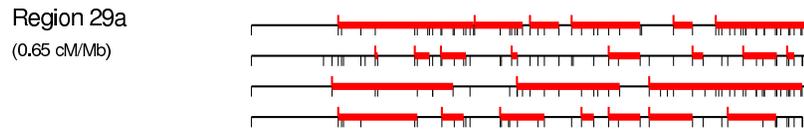
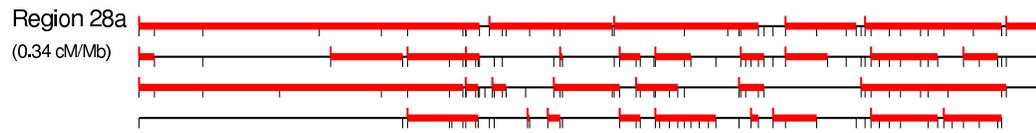
Region 26a
(1.57 cMMb)



Region 27a
(1.04 cMMb)

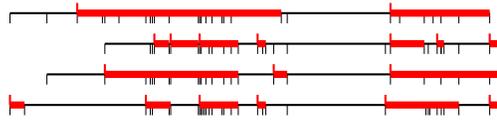


100 Kb

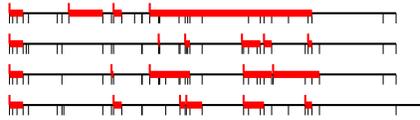


100 Kb

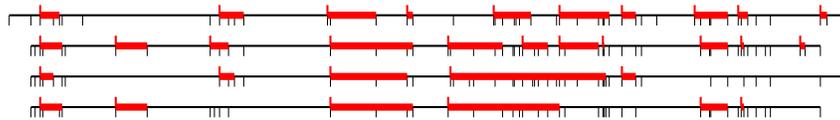
Region 36a
(0.61 cMMb)



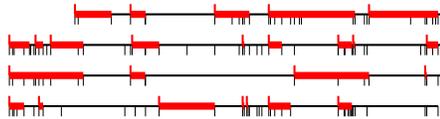
Region 37a
(1.03 cMMb)



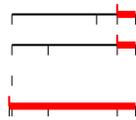
Region 38a
(1.41 cMMb)



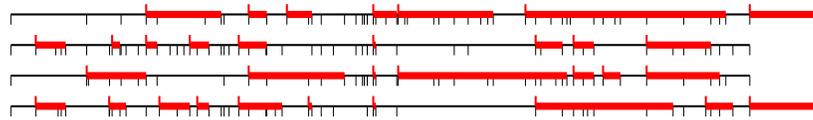
Region 39a
(1.19 cMMb)



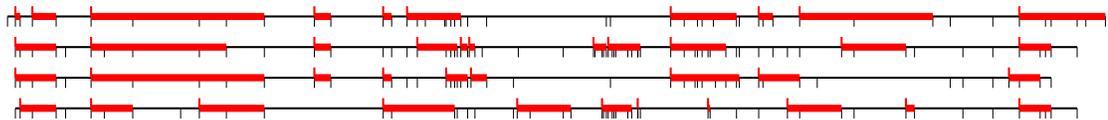
Region 39b
(1.19 cMMb)



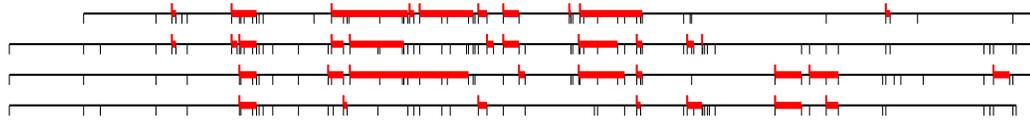
Region 40a
(1.44 cMMb)



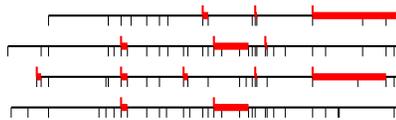
Region 41a
(0.65 cMMb)



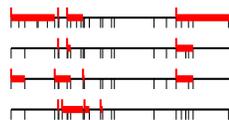
Region 42a
(1.84 cMMb)



Region 43a
(3.02 cMMb)

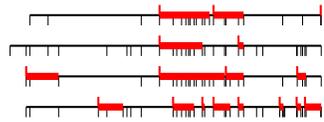


Region 44a
(1.62 cMMb)

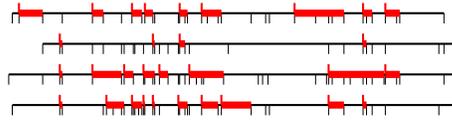


100 Kb

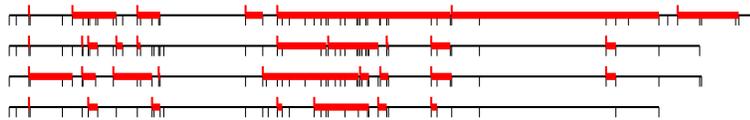
Region 44b
(1.62 cM/Mb)



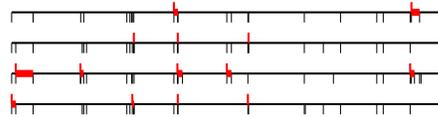
Region 45a
(2.22 cM/Mb)



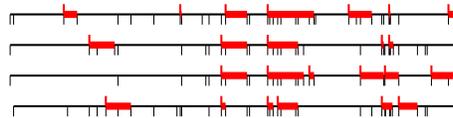
Region 46a
(1.37 cM/Mb)



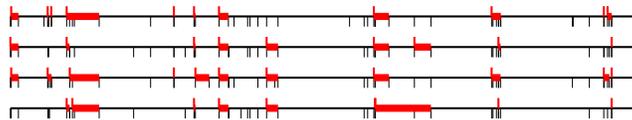
Region 47a
(2.22 cM/Mb)



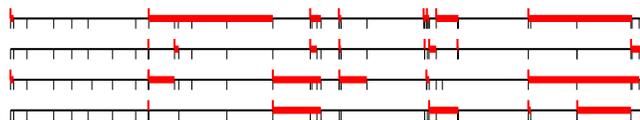
Region 48a
(1.57 cM/Mb)



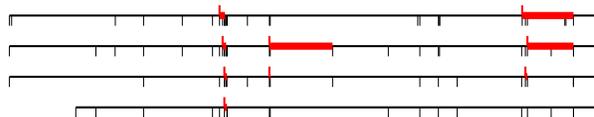
Region 50a
(3.51 cM/Mb)



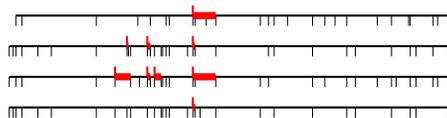
Region 51a
(1.49 cM/Mb)



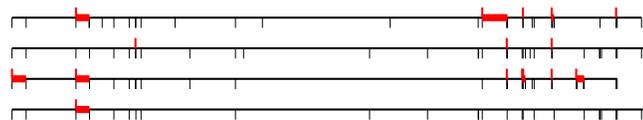
Region 52a
(2.02 cM/Mb)



Region 53a
(2.51 cM/Mb)



Region 54a
(2.51 cM/Mb)



100 Kb