

Region Plots for 139 novel signals of association with lung function

The region plots are from the meta-analysis of UK Biobank and SpiroMeta, from which the effect sizes and standard errors were used to calculate the 99% credible sets. The variants in the 99% credible set are shown as filled circles. The green bar indicates the extent of the credible set. The red line is $P = 5 \times 10^{-9}$, used as our discovery threshold in UK Biobank and the meta-analysis.

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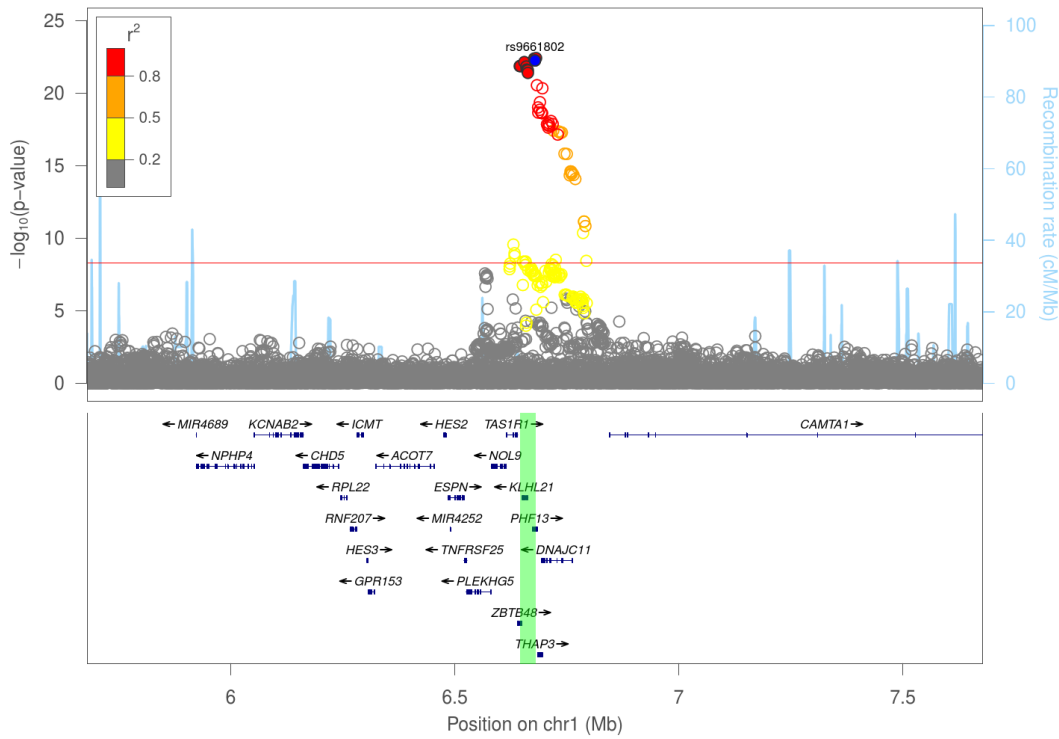
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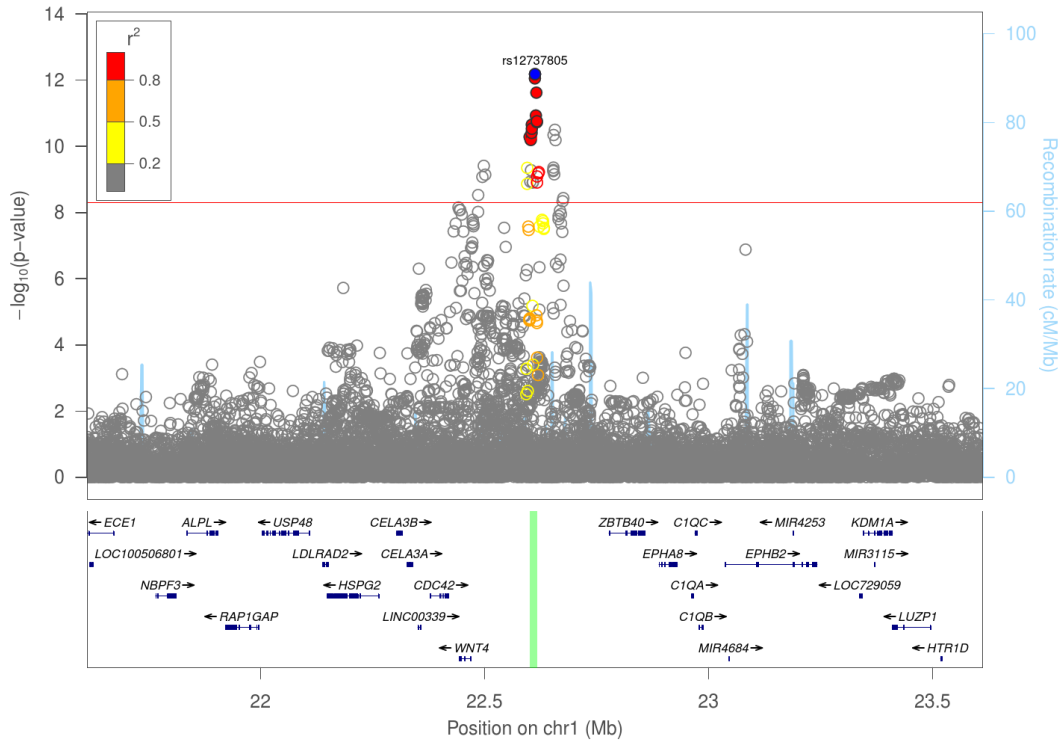
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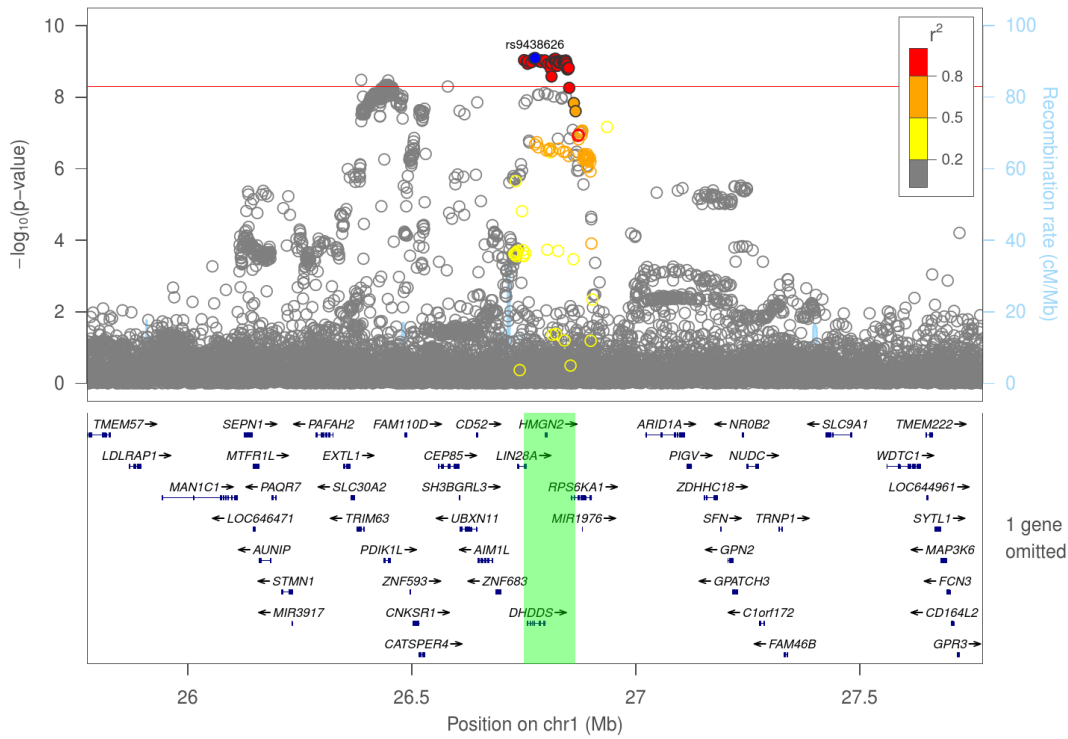
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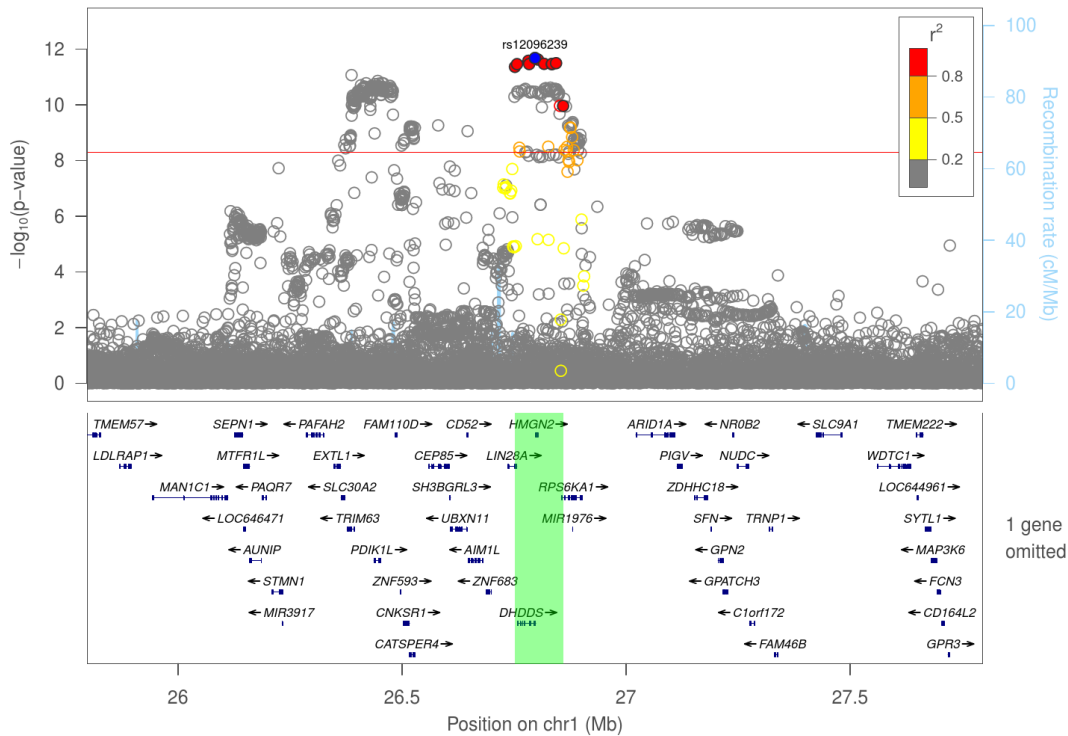
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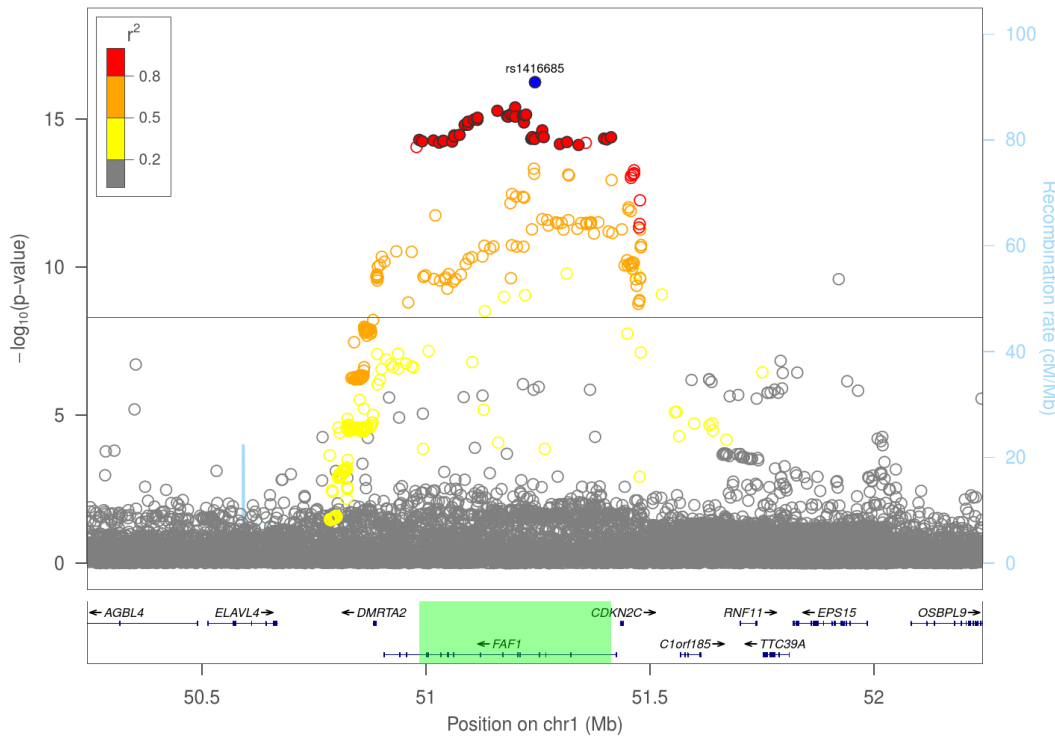
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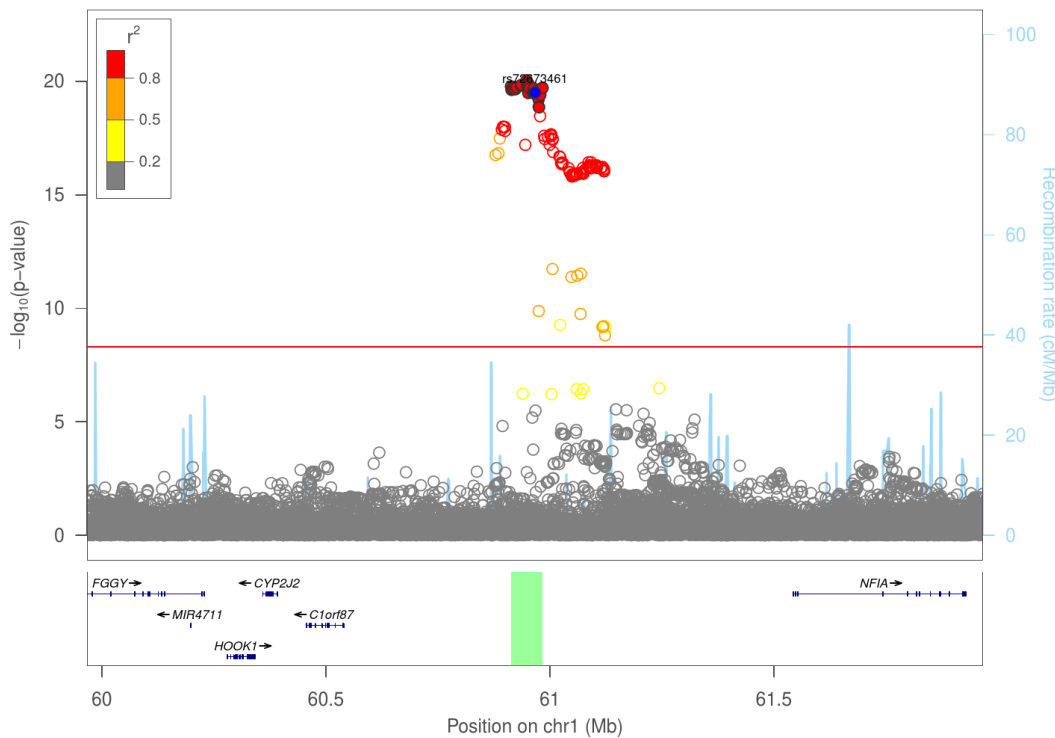
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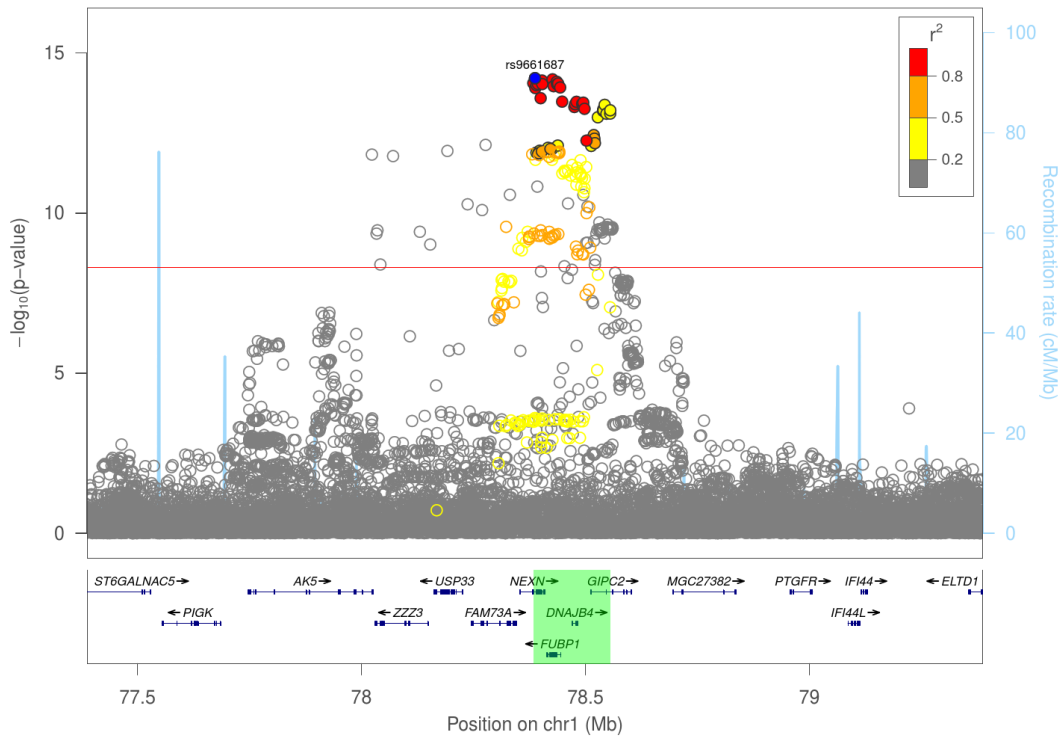
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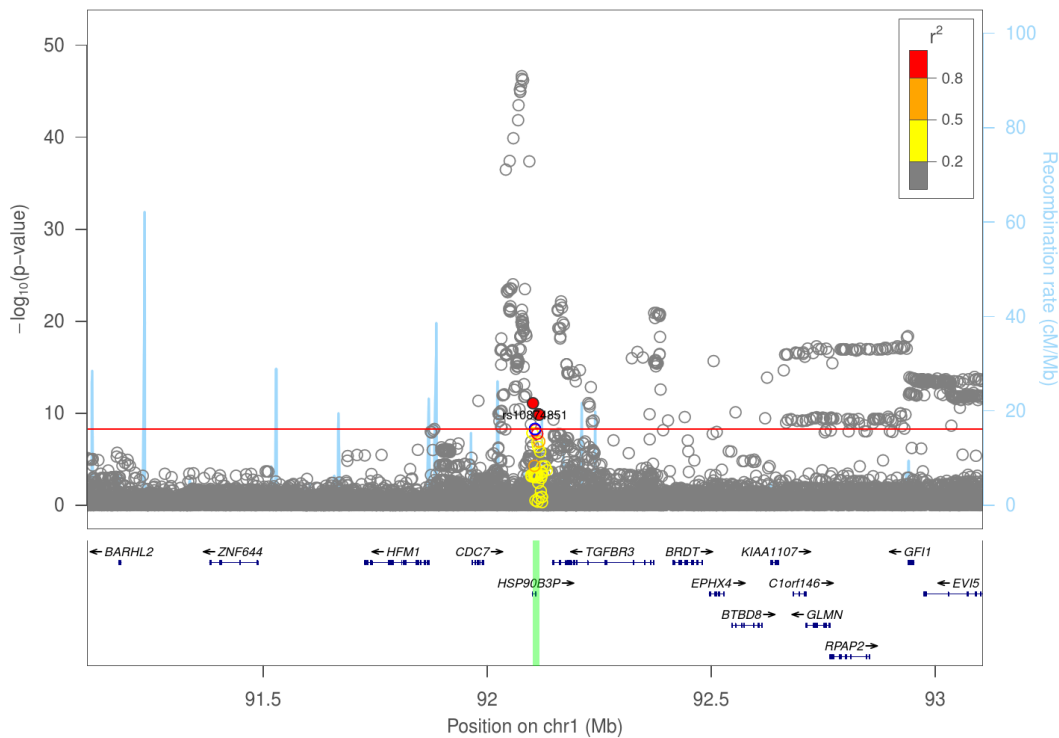
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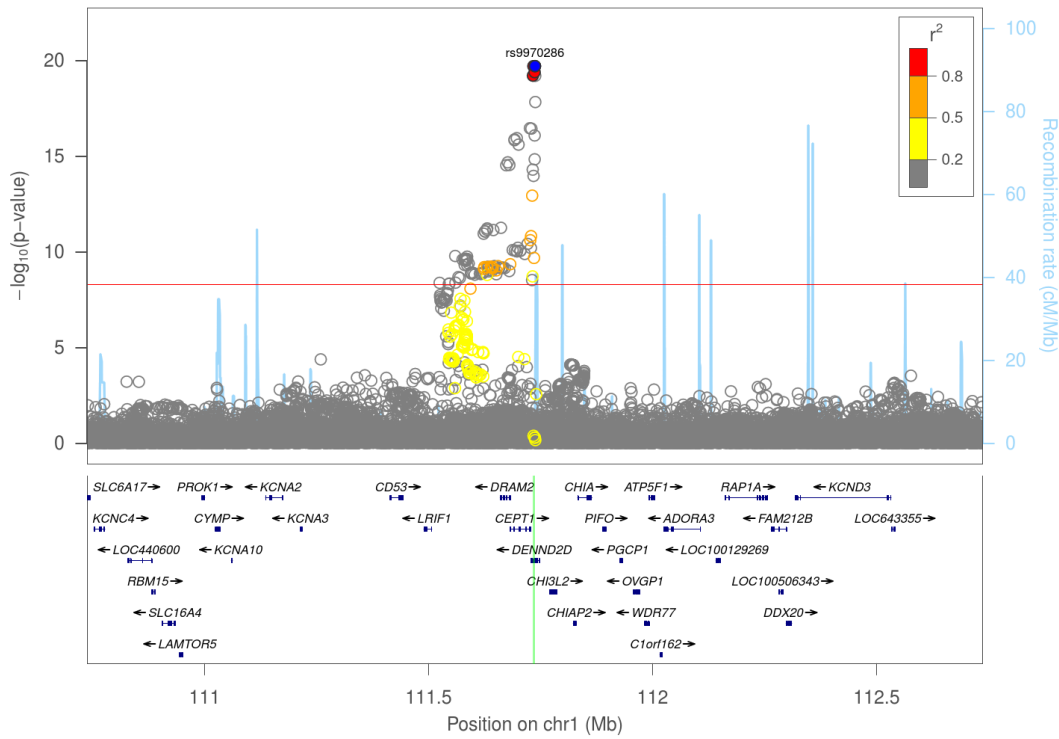
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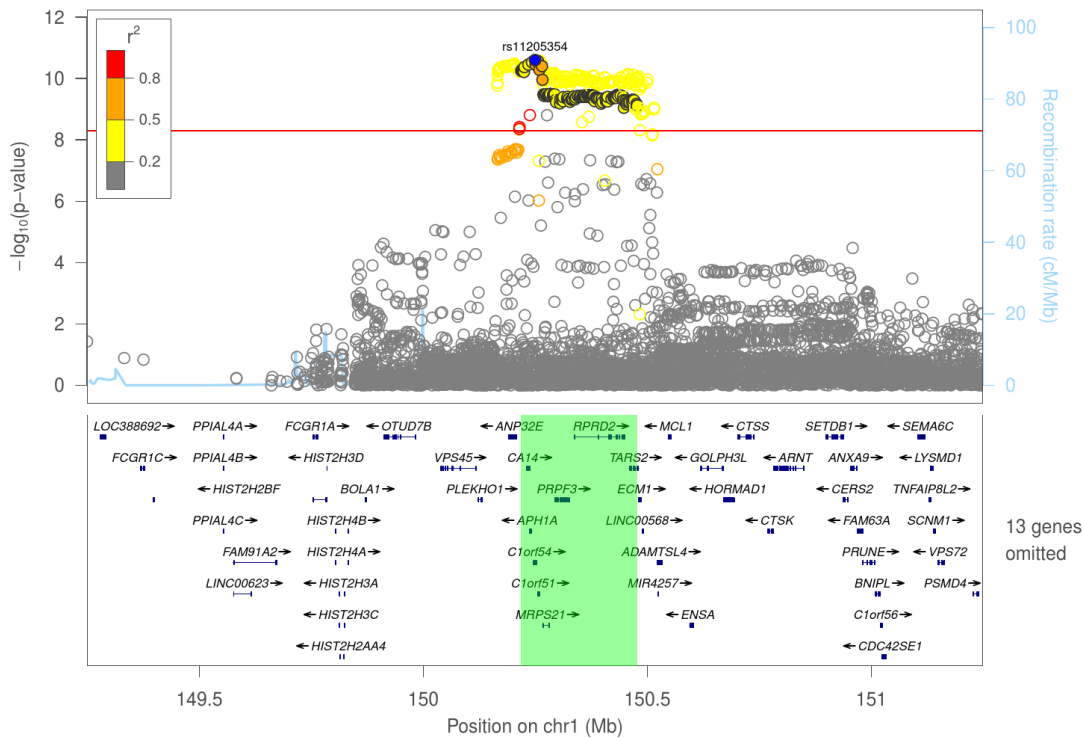
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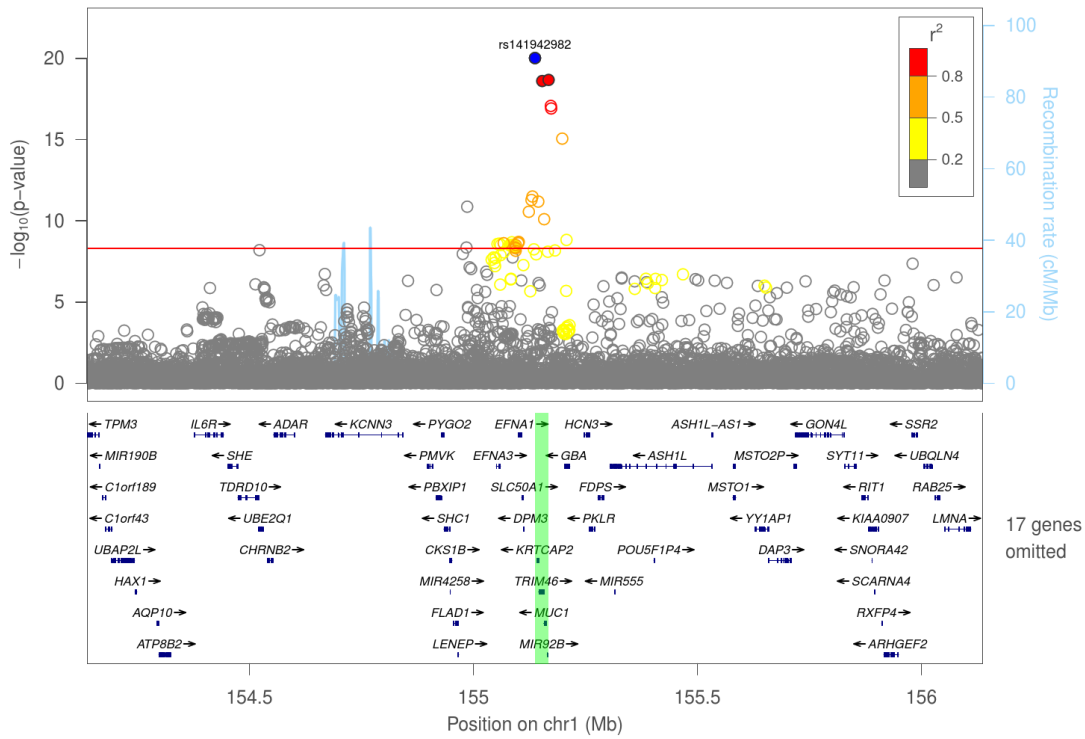
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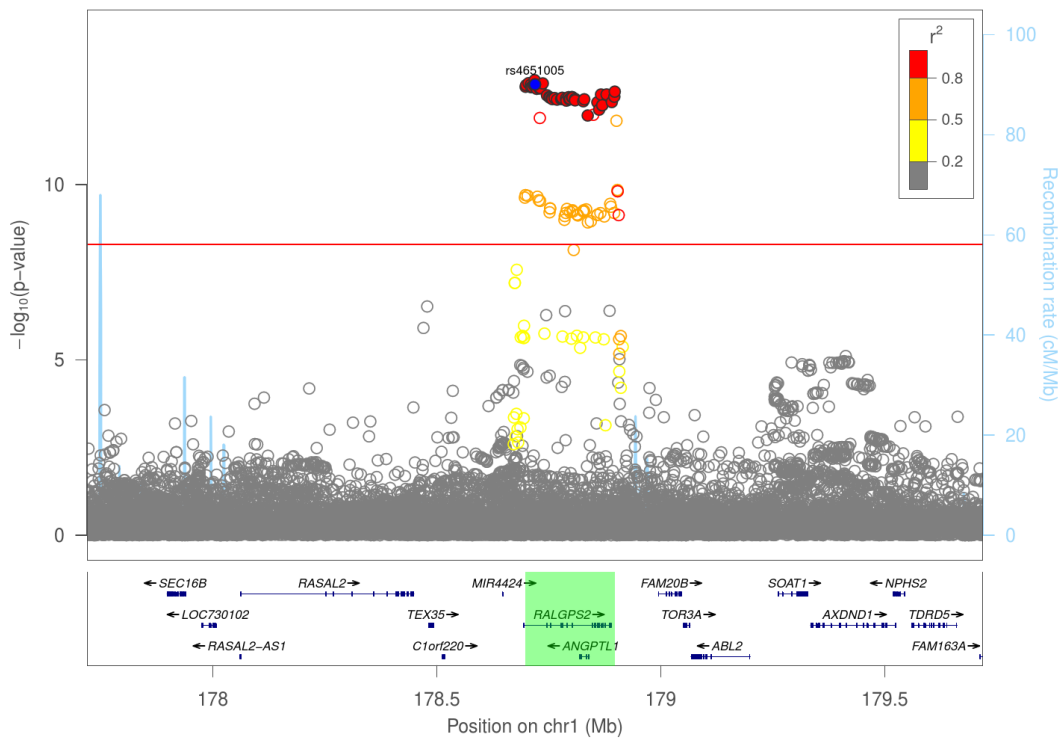
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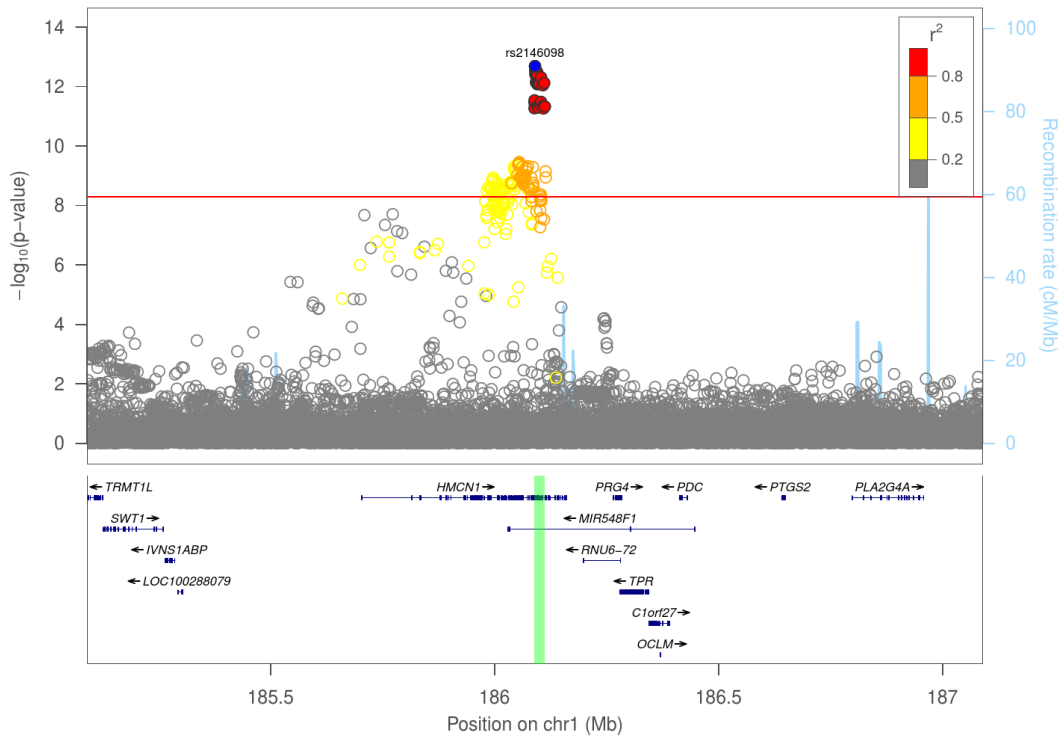
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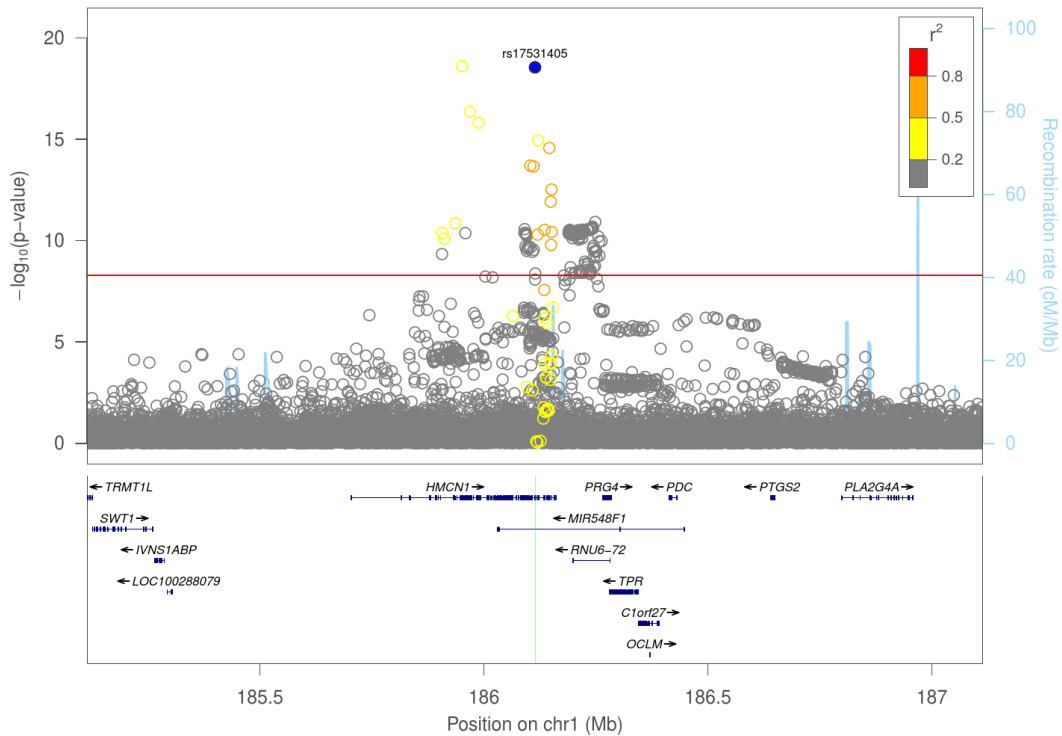
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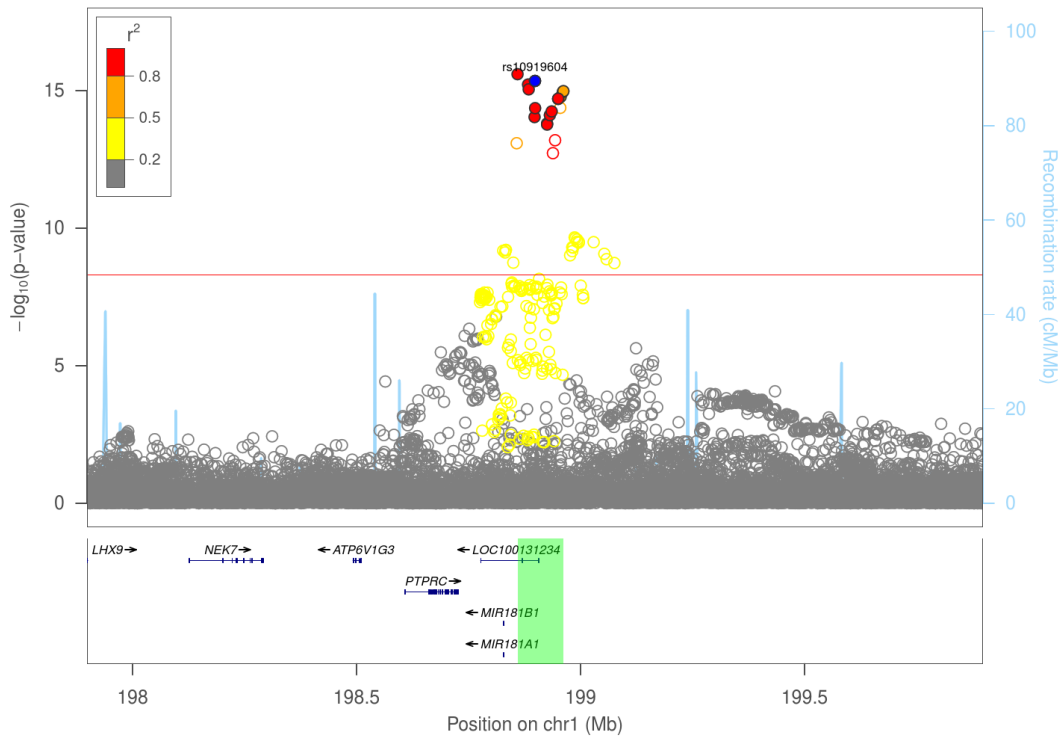
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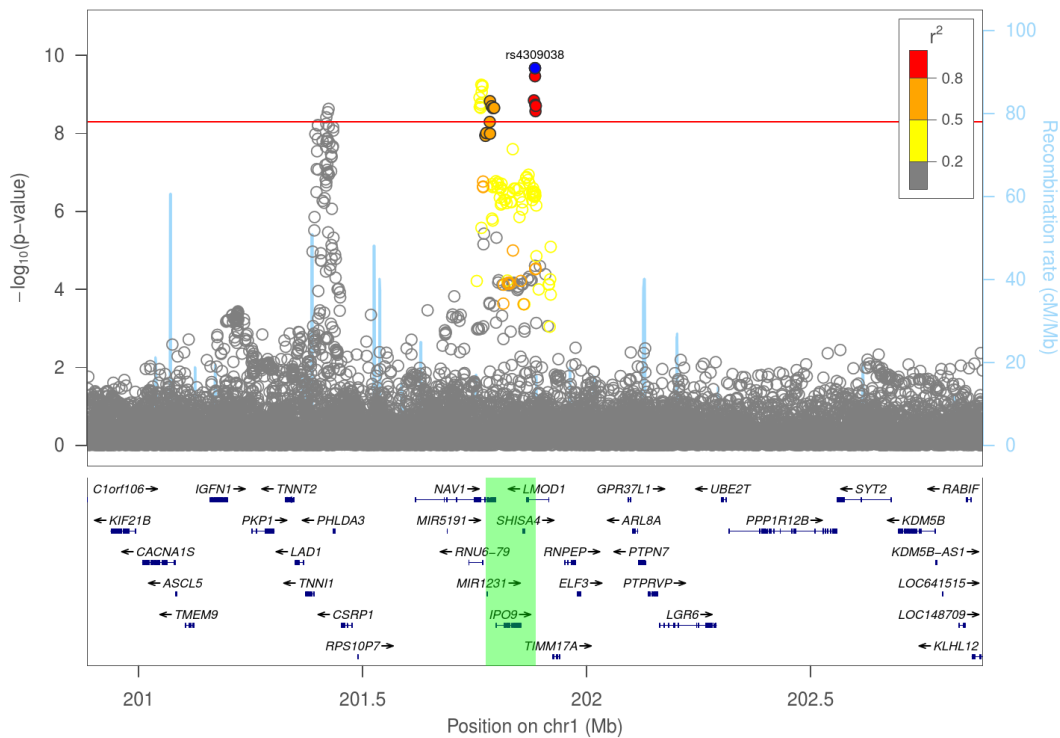
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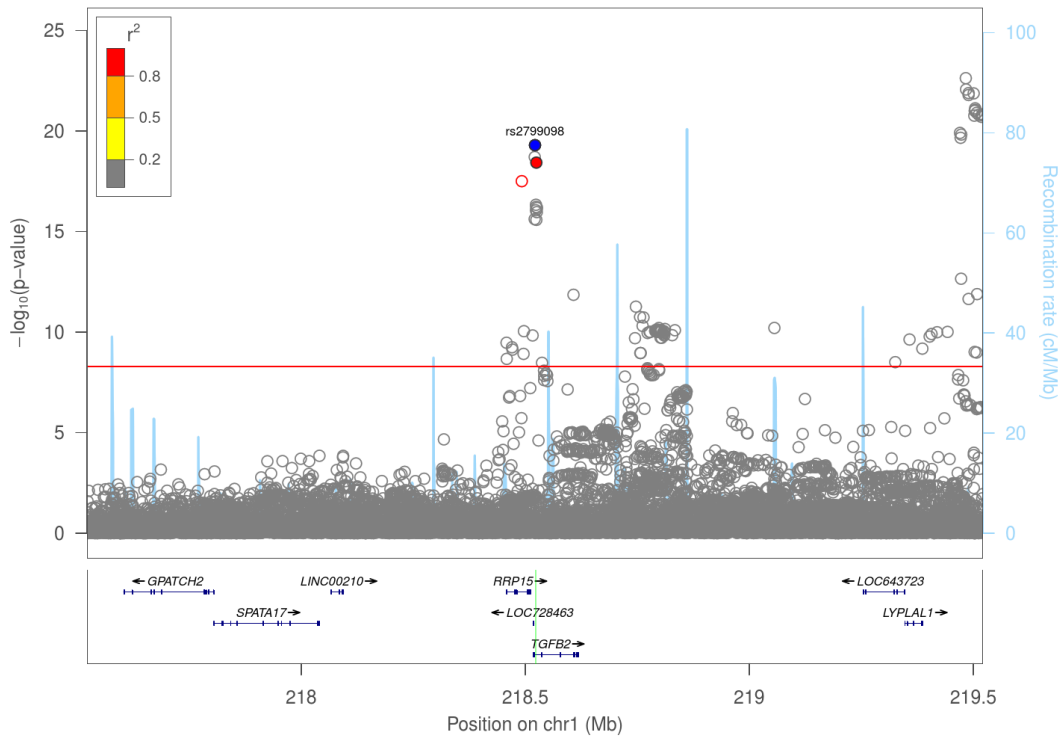
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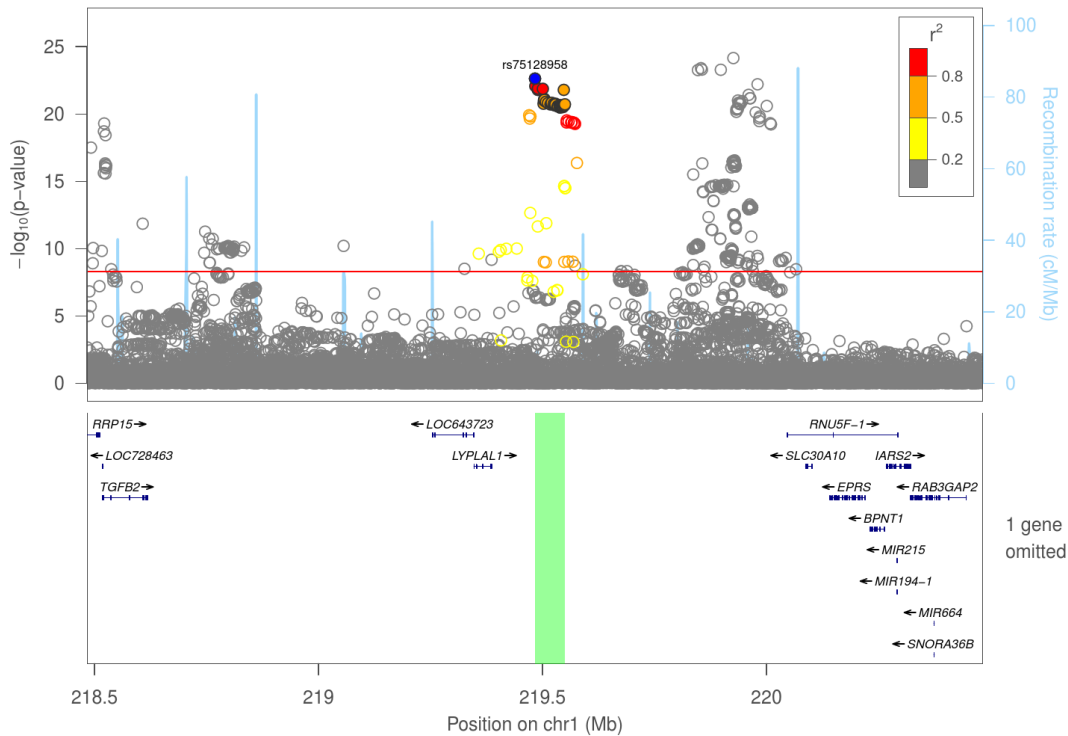
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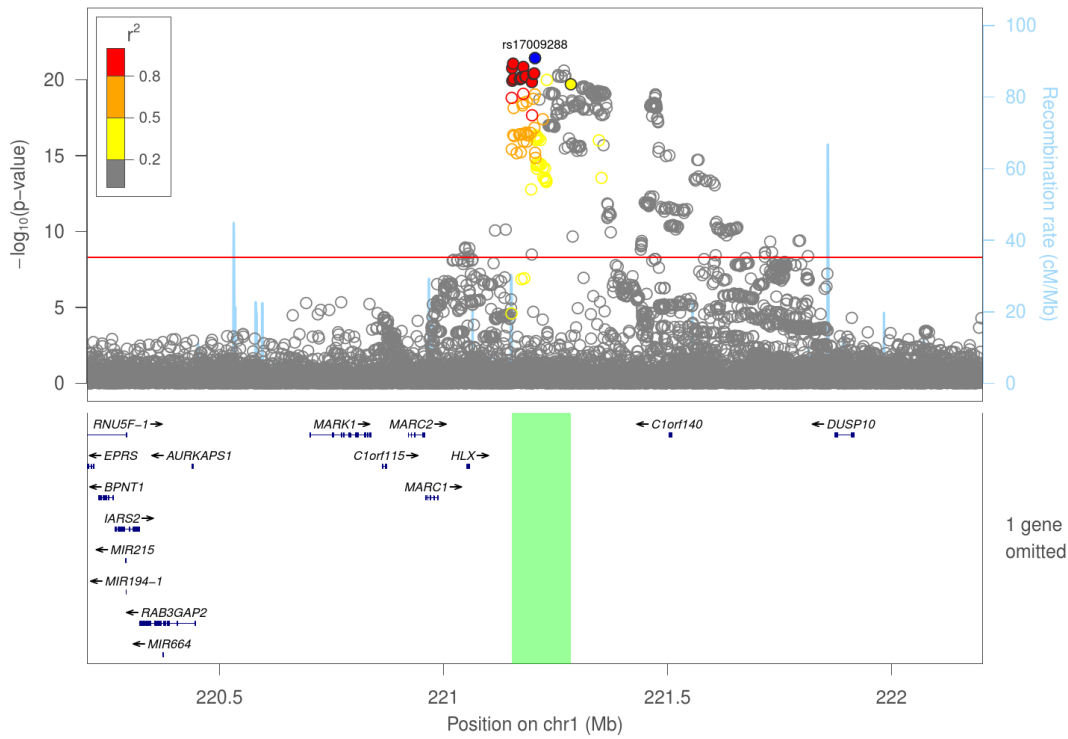
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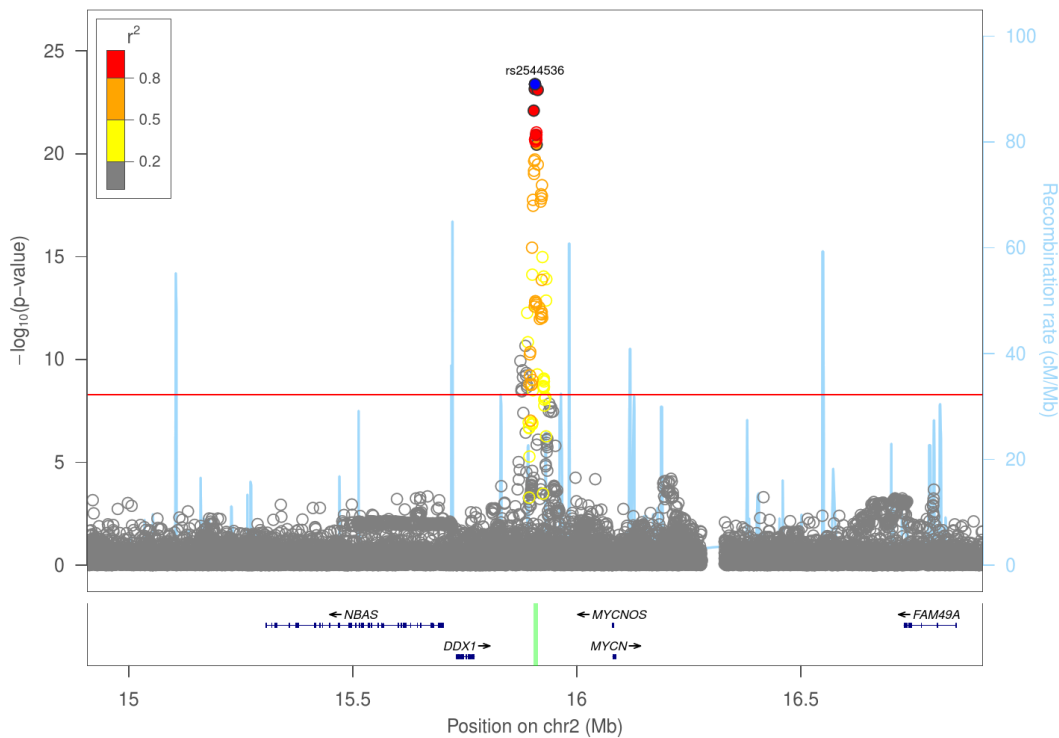
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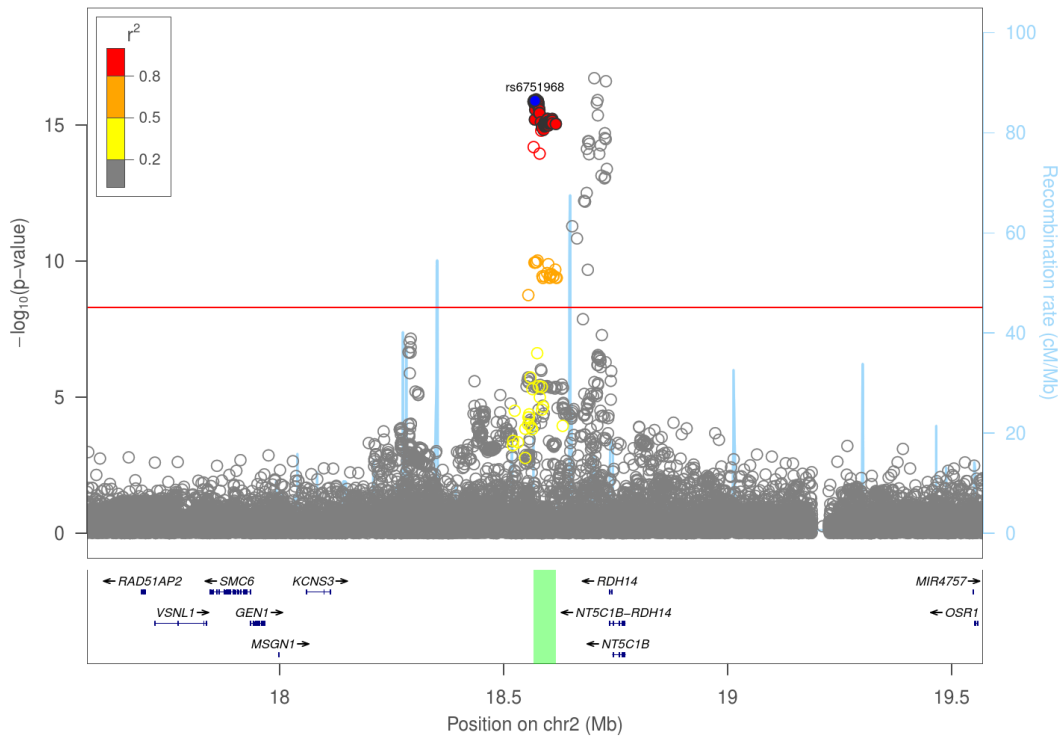
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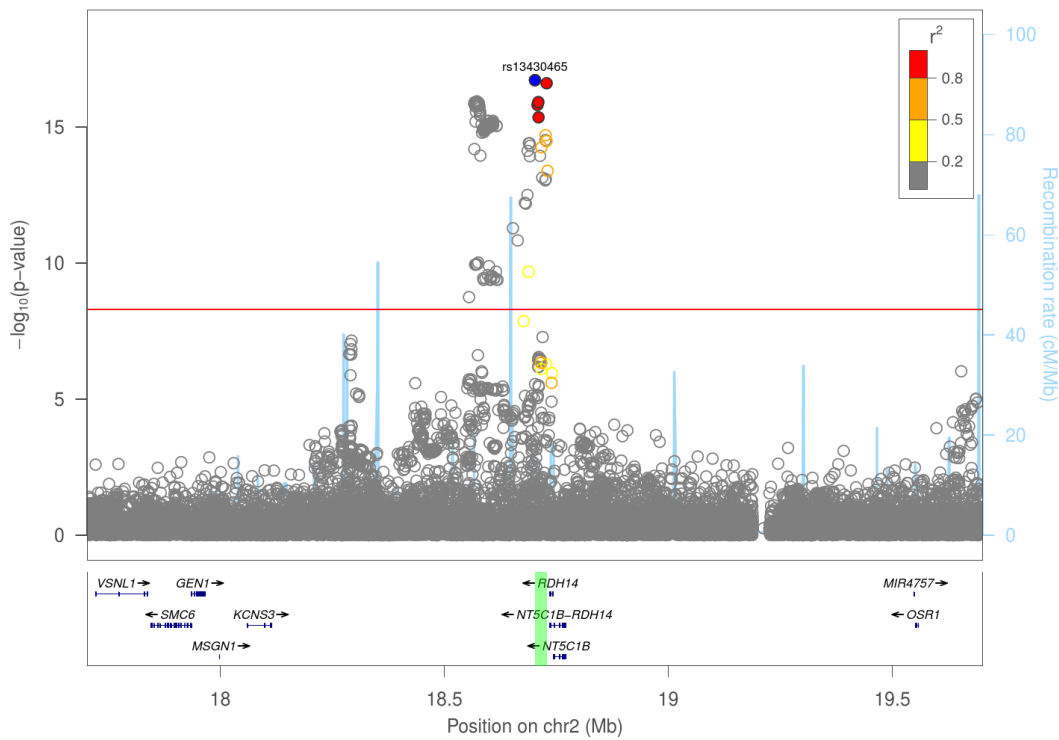
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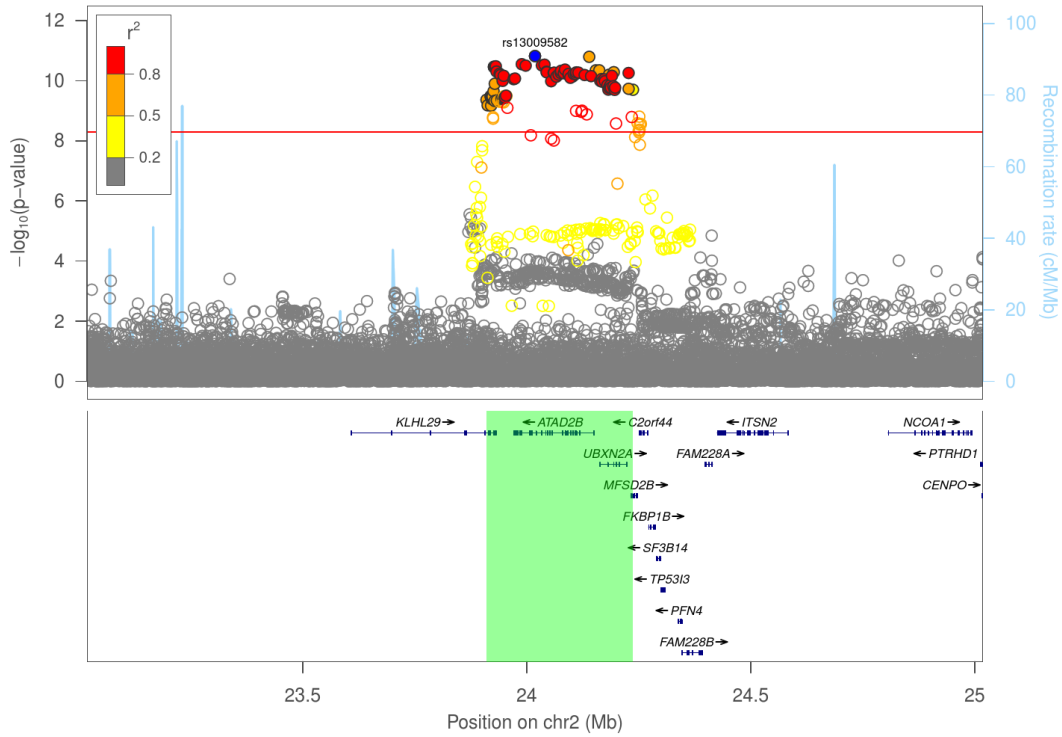
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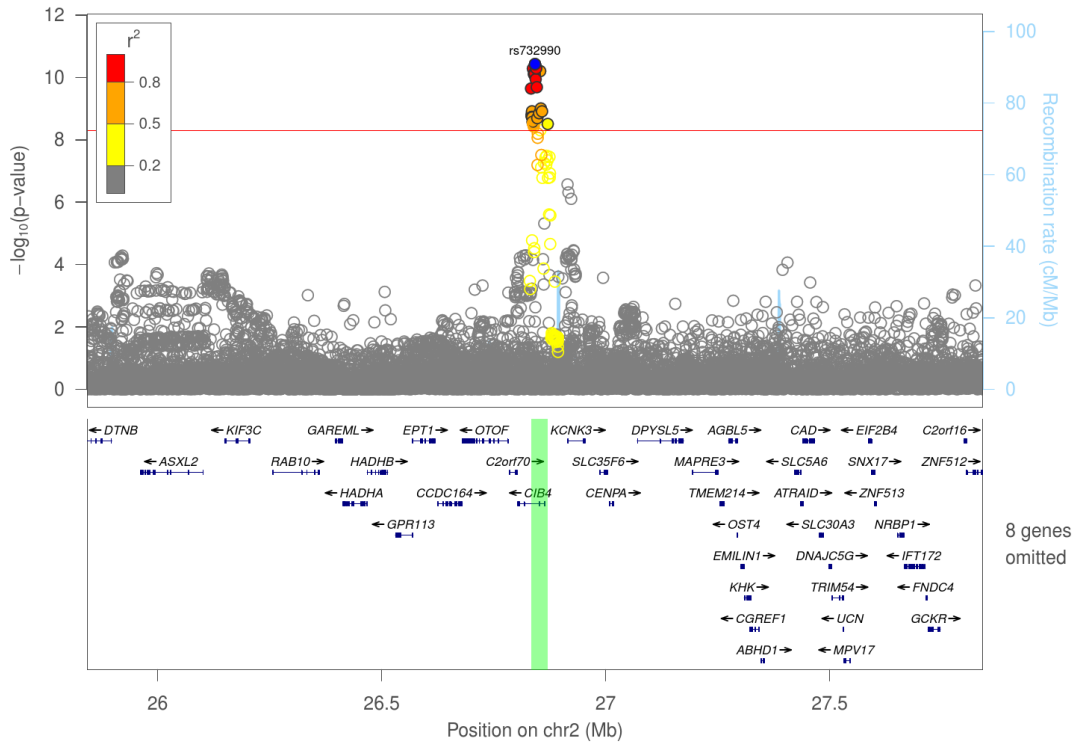
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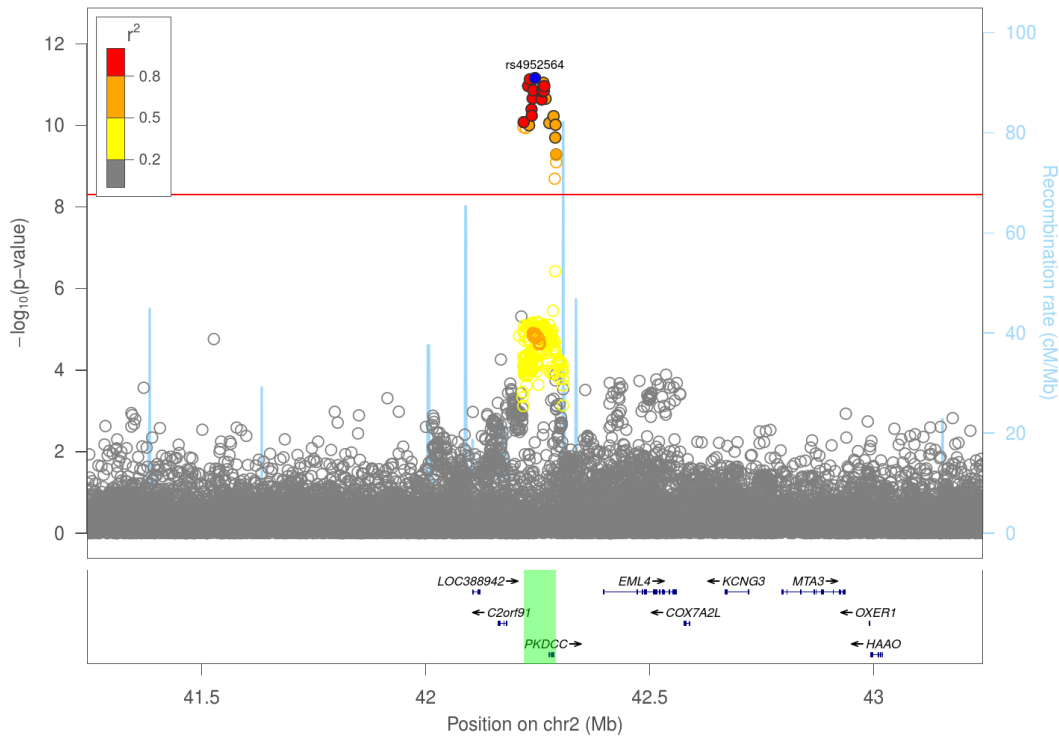
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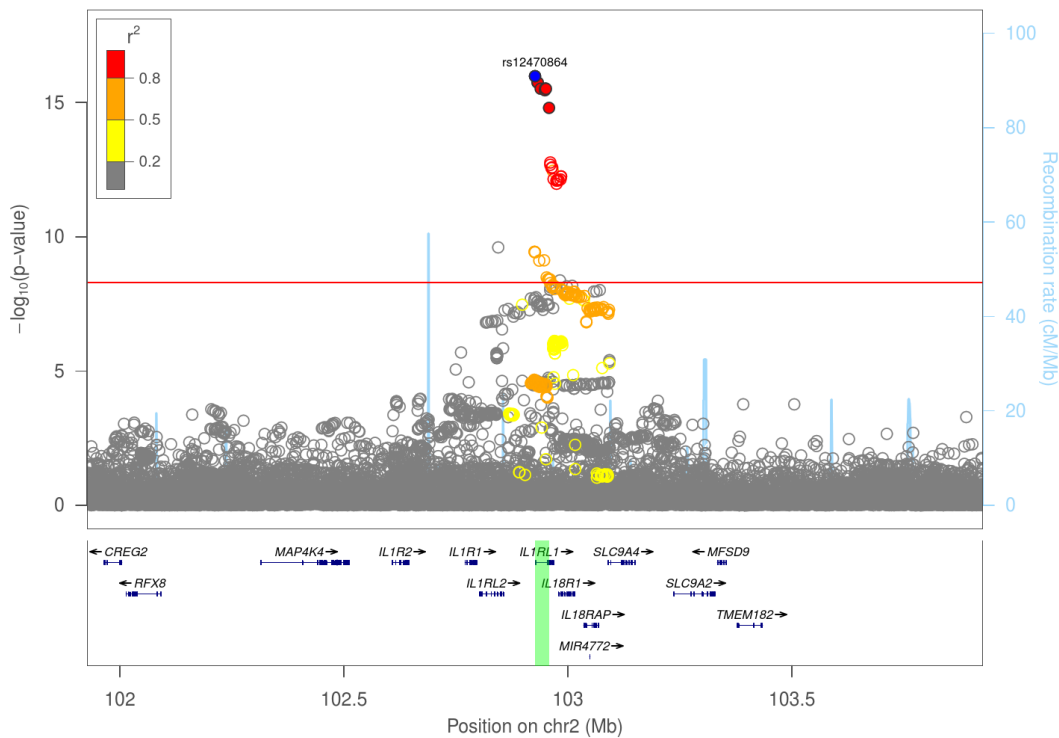
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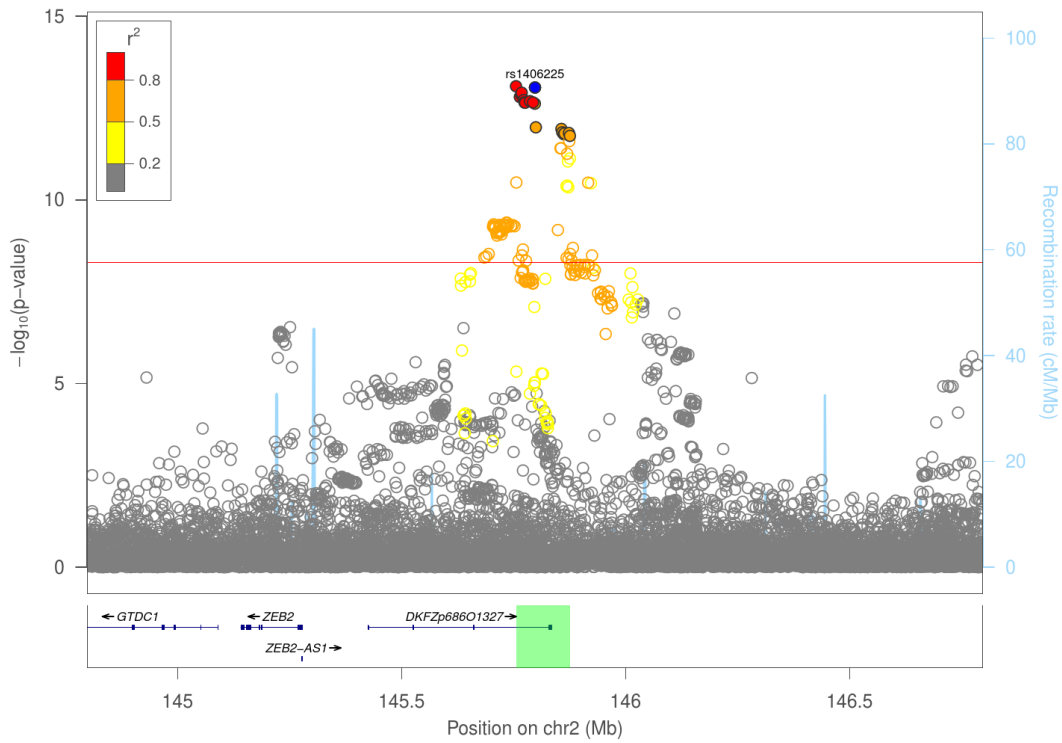
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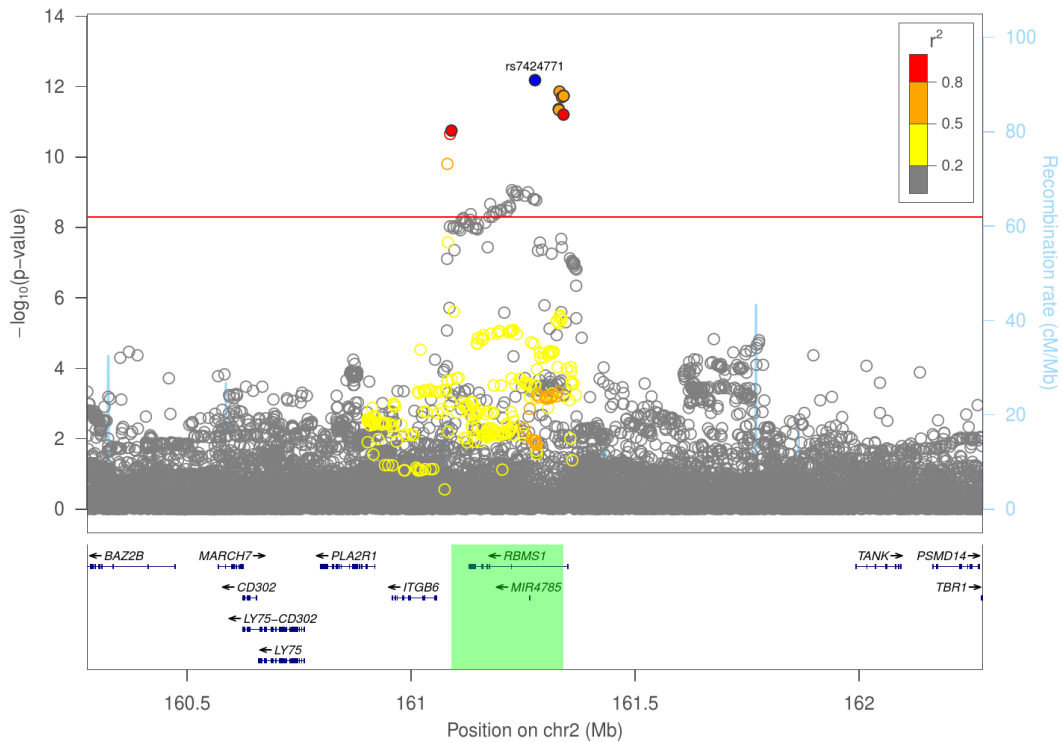
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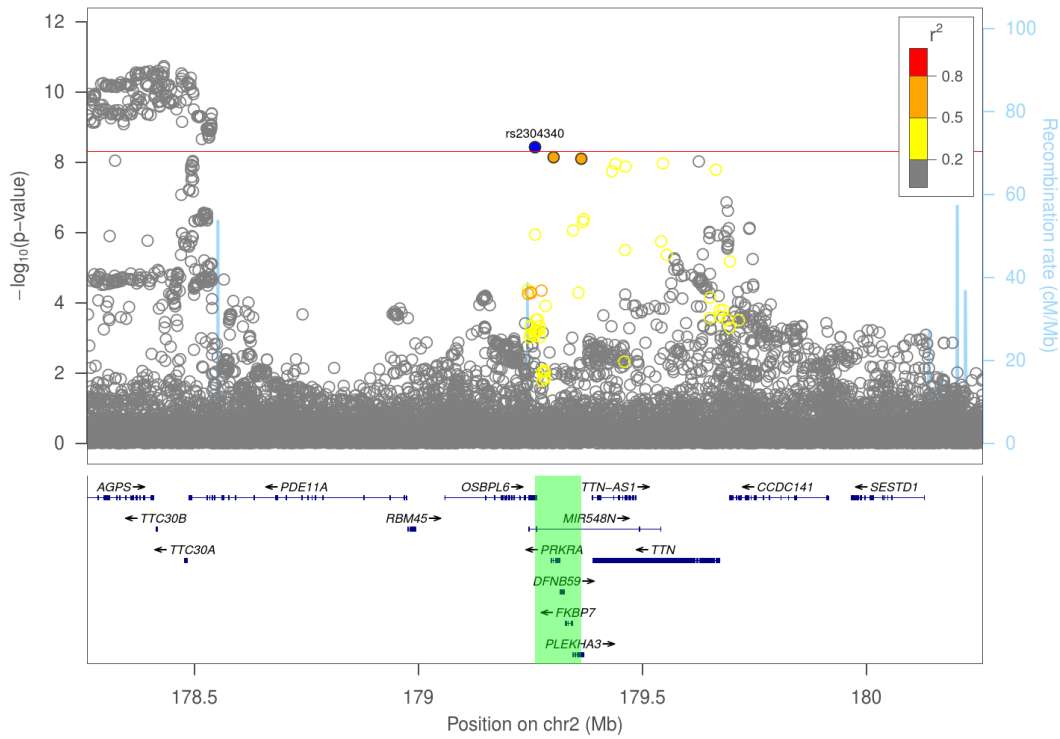
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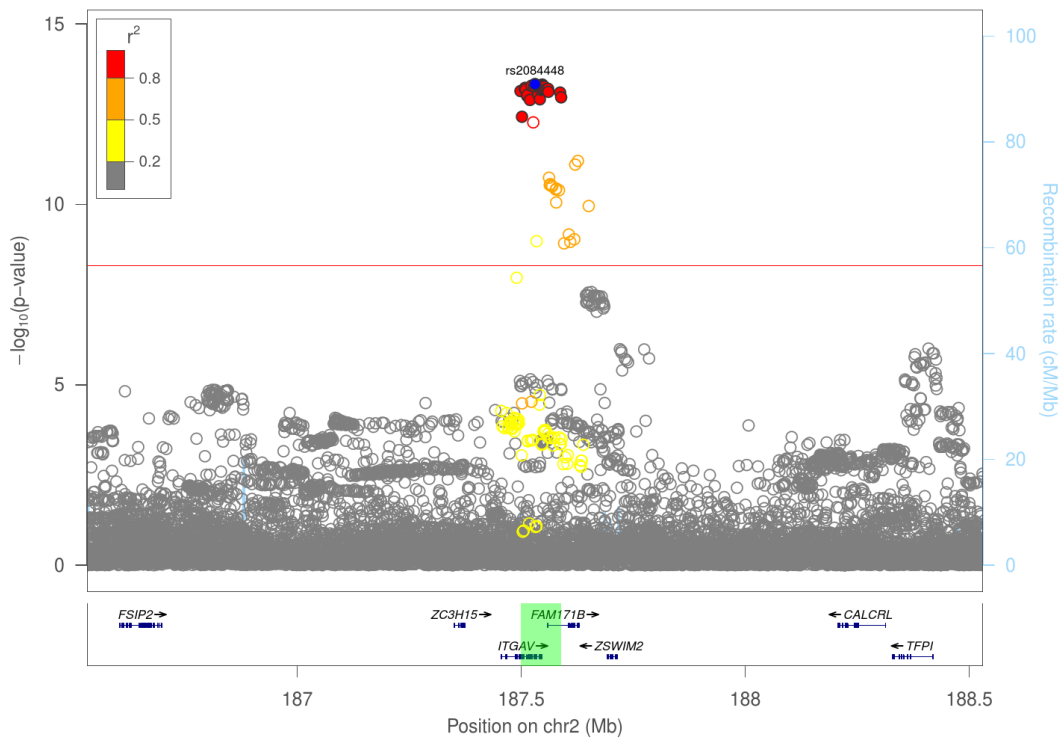
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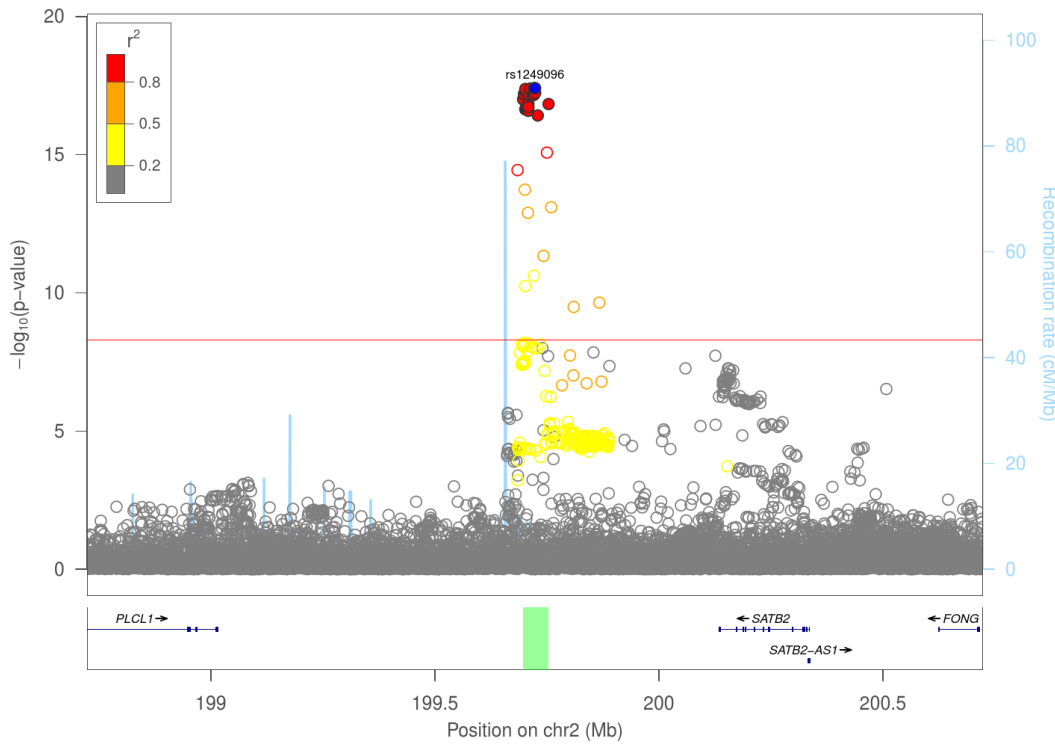
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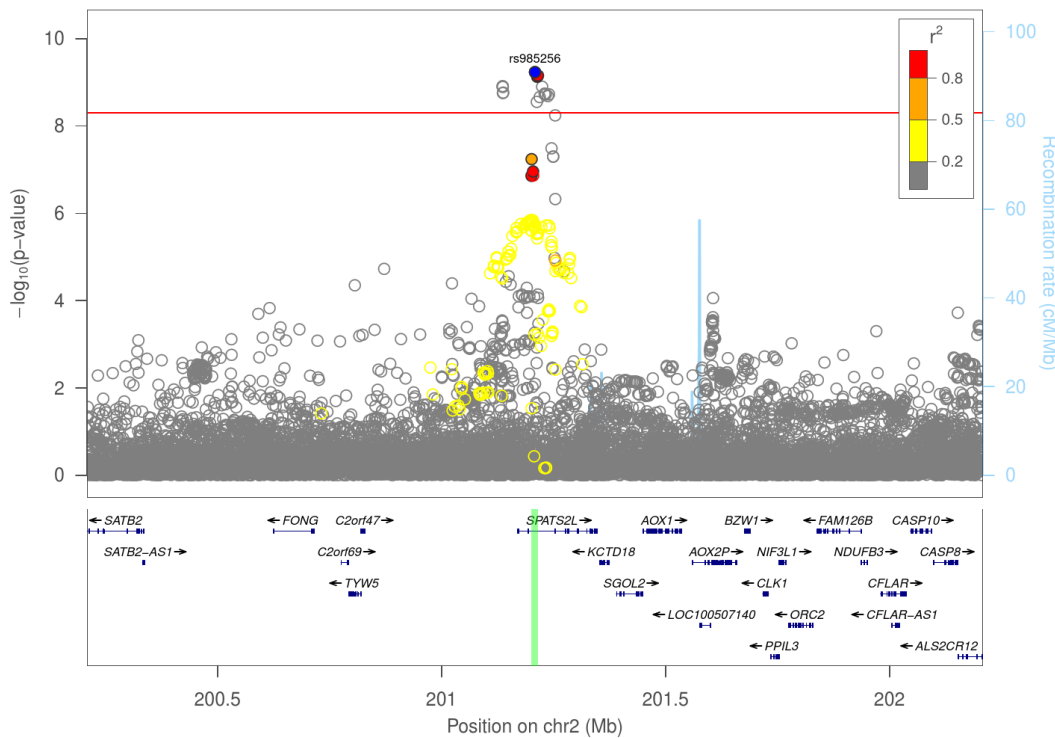
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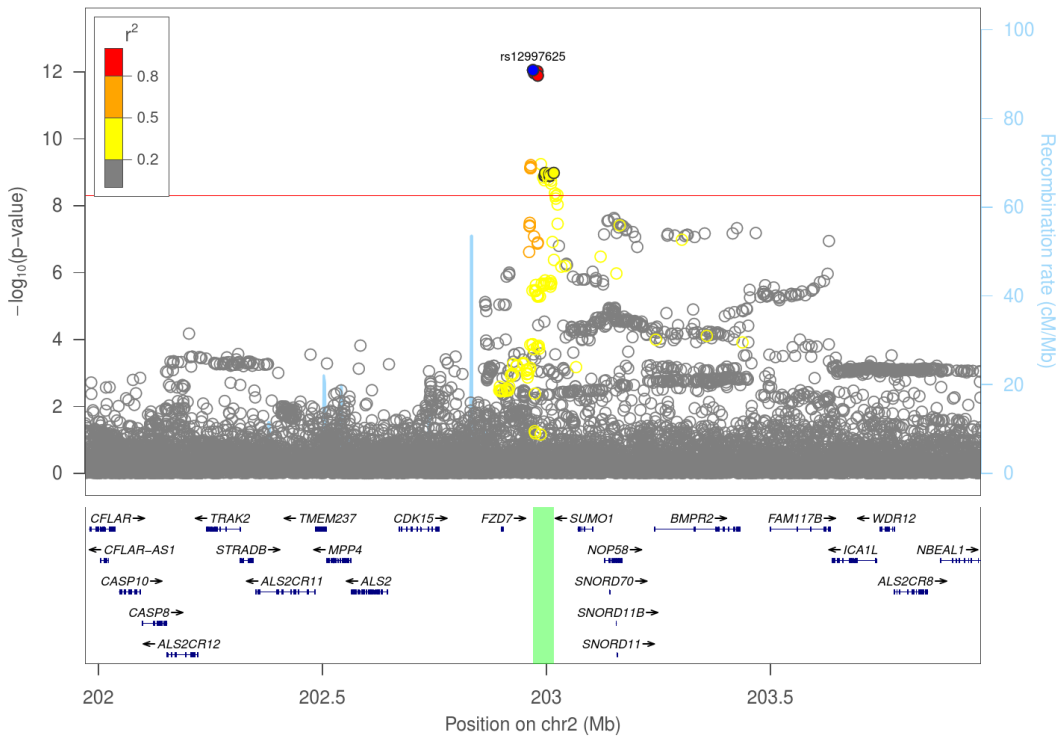
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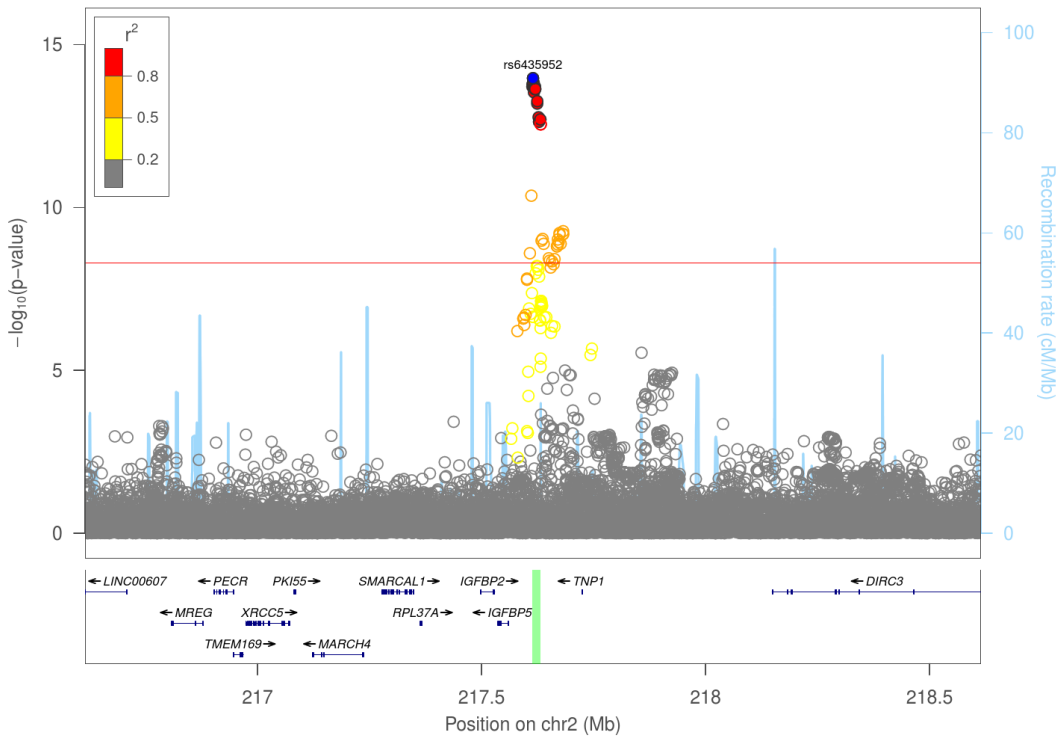
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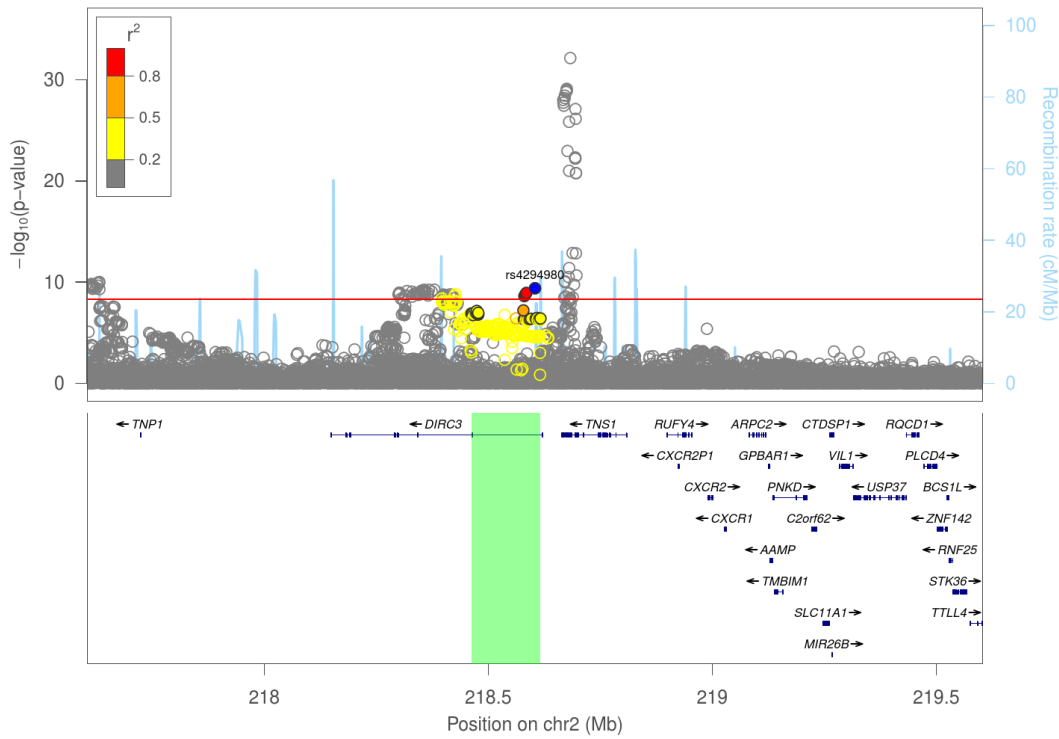
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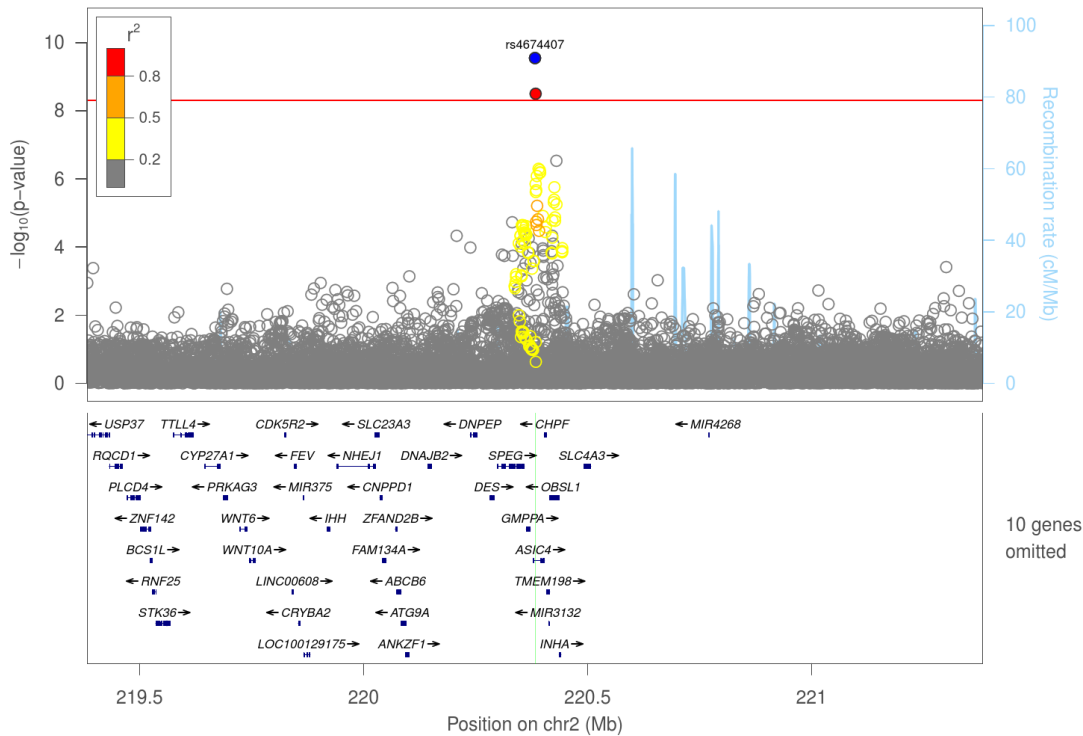
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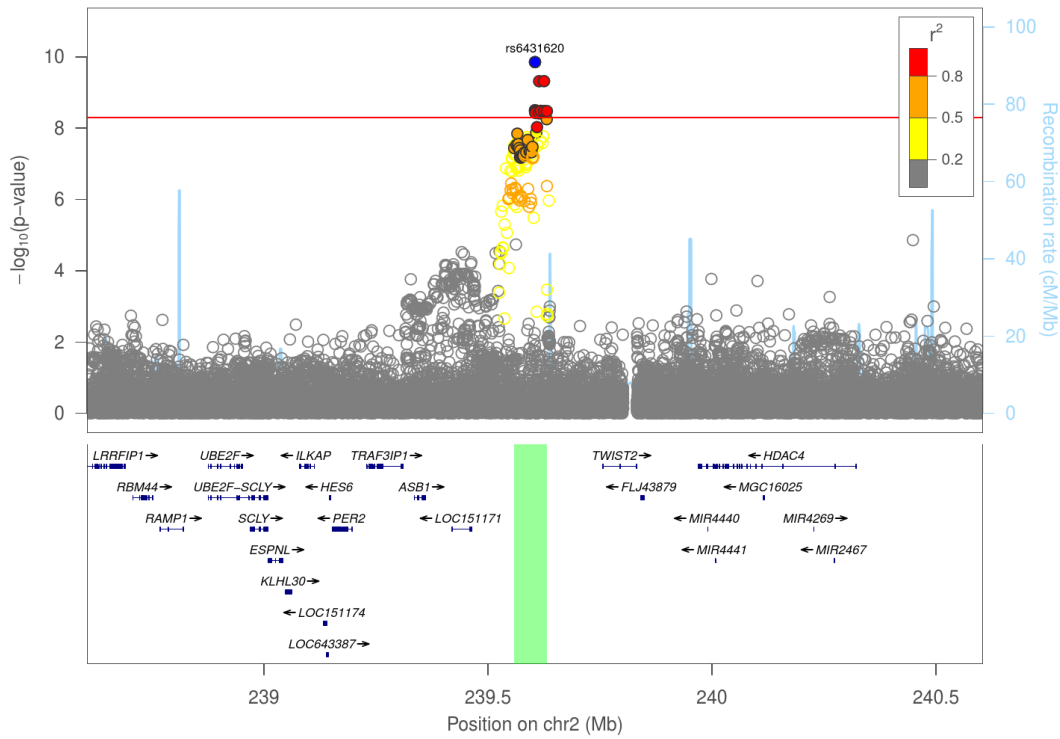
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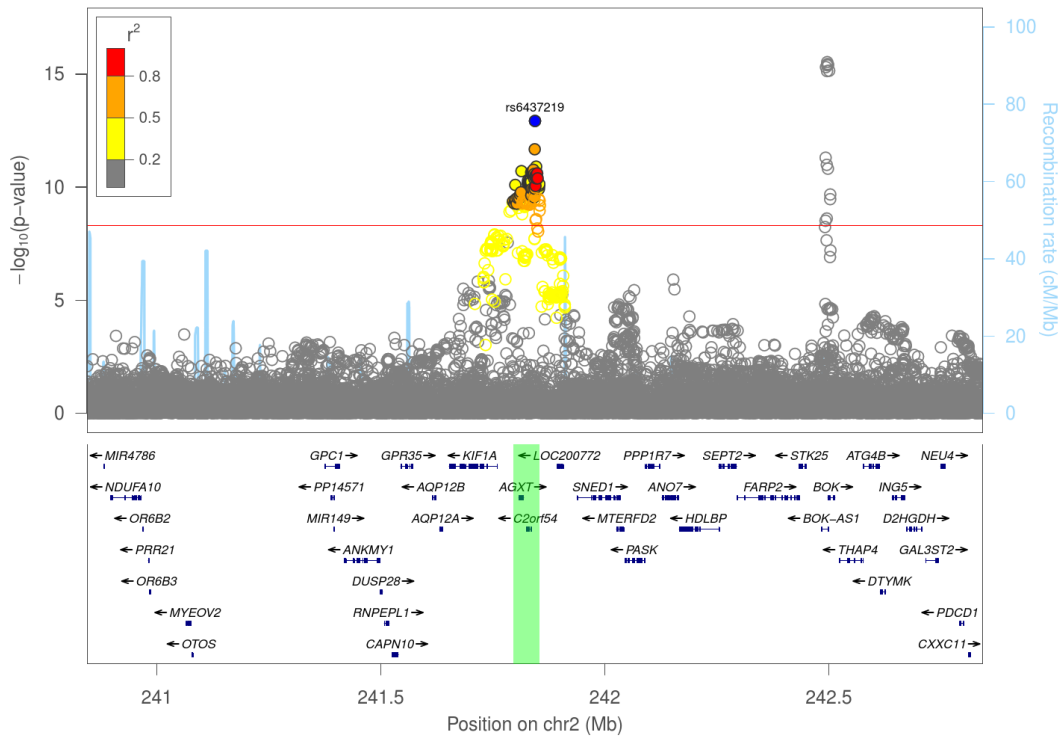
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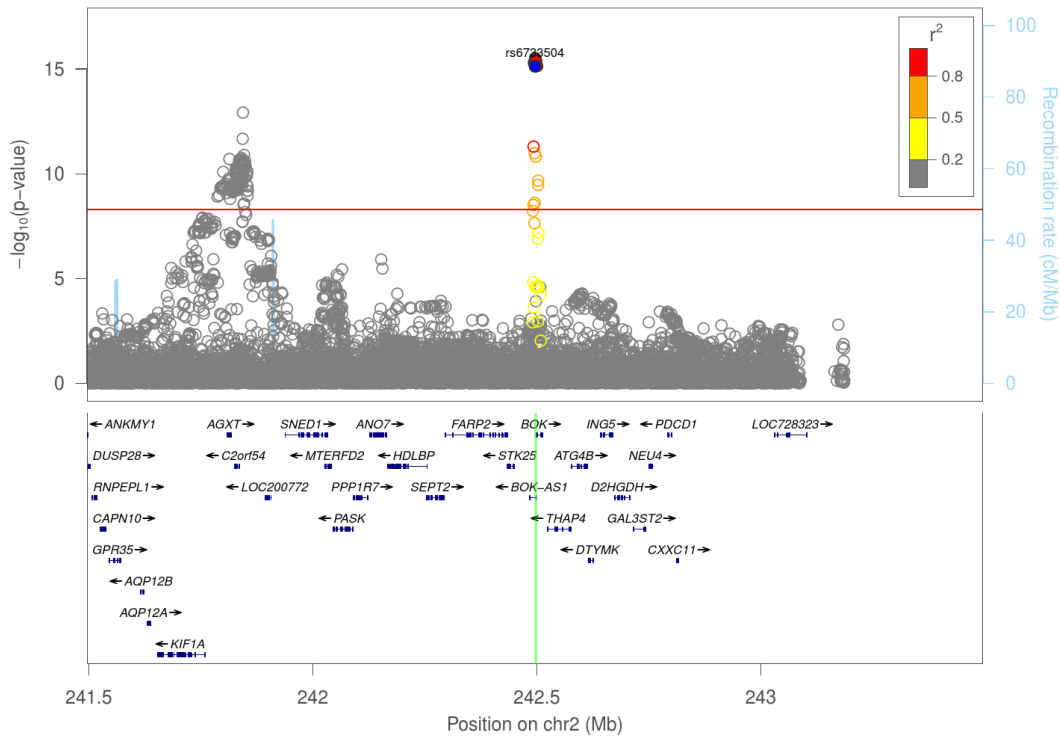
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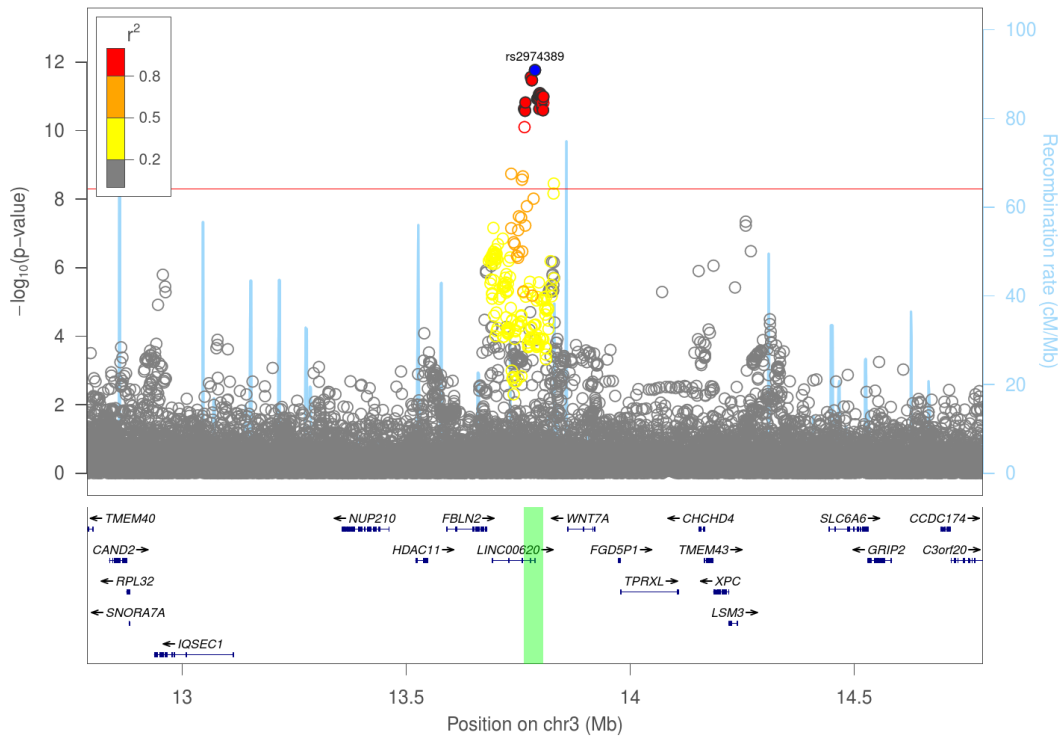
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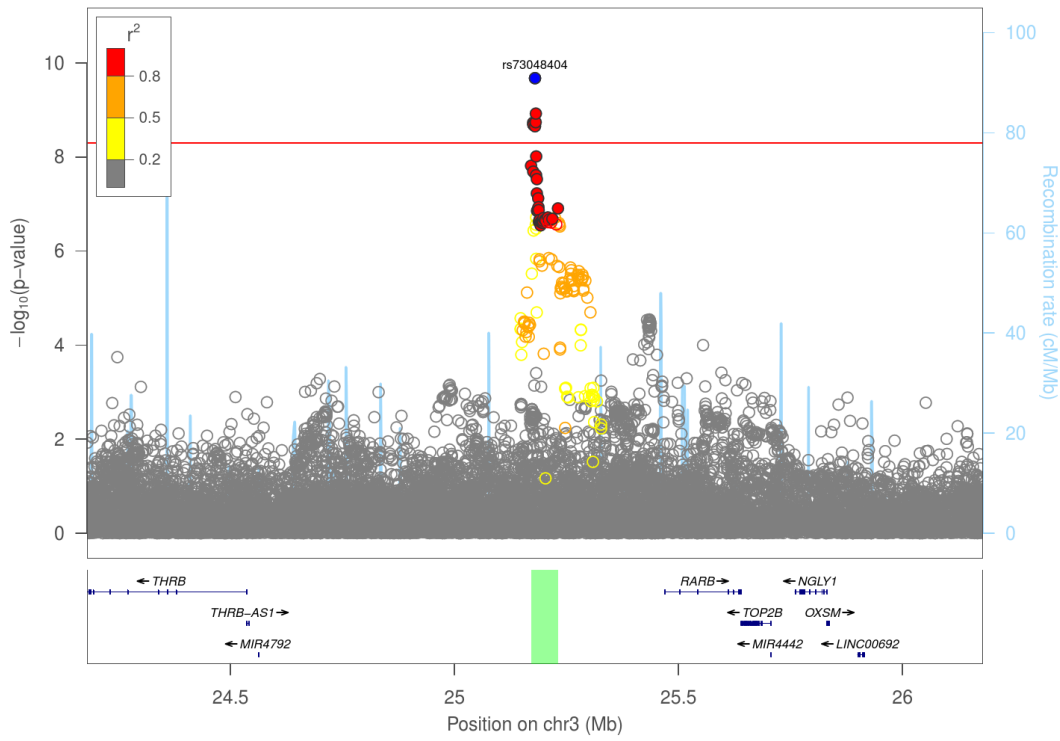
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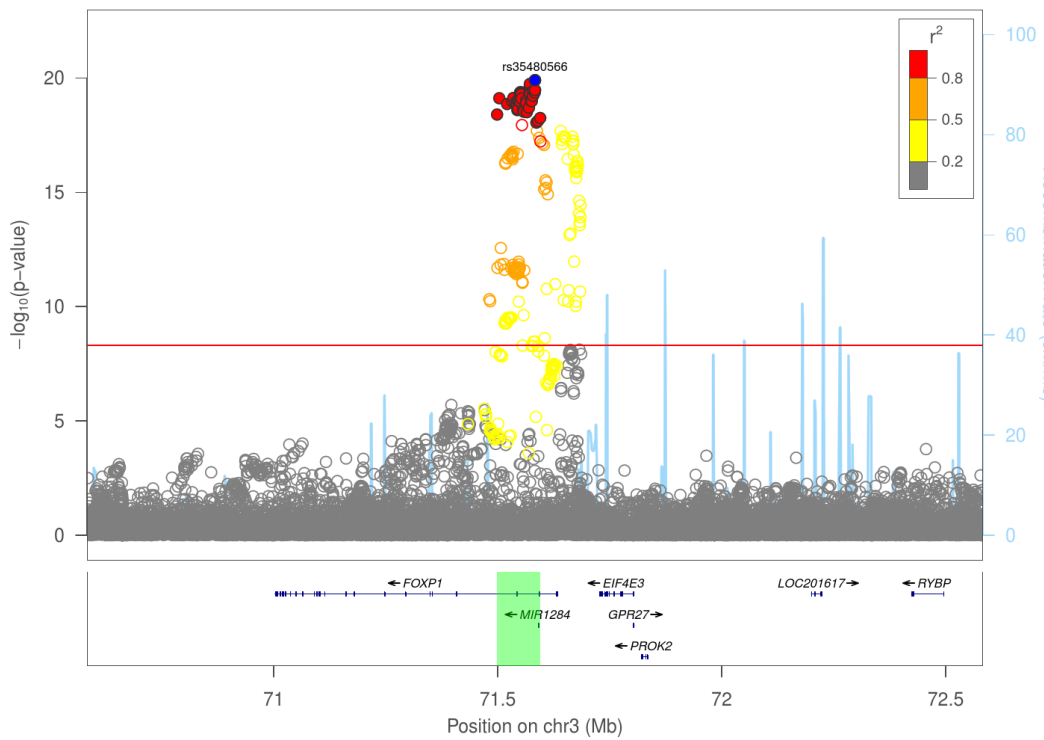
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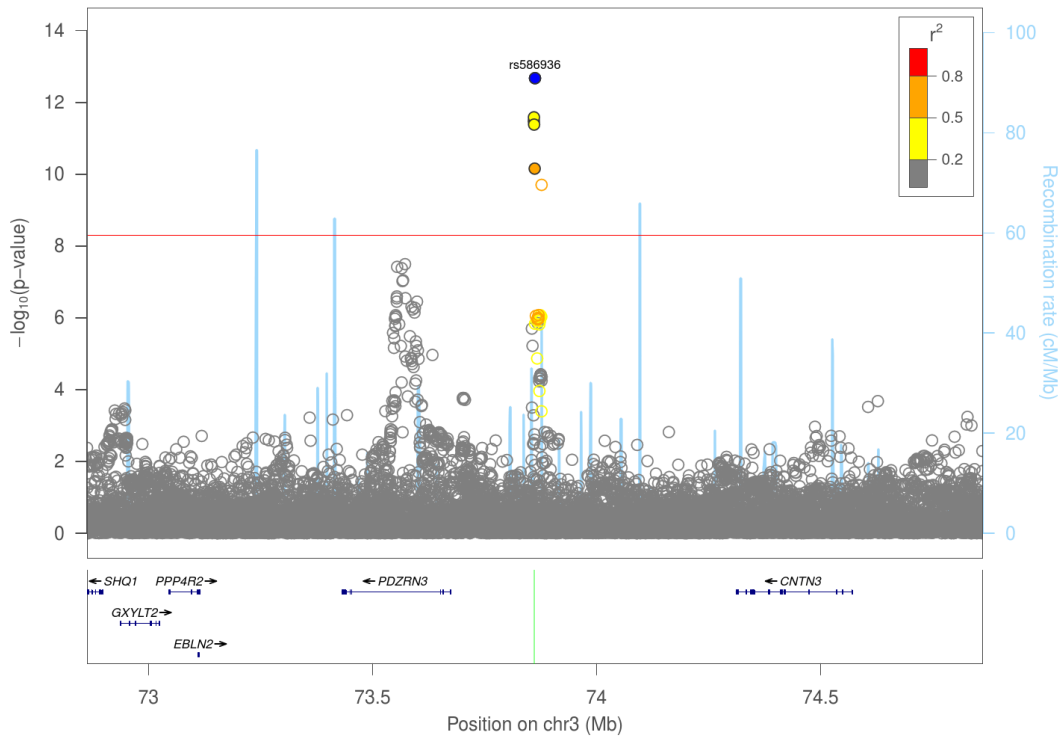
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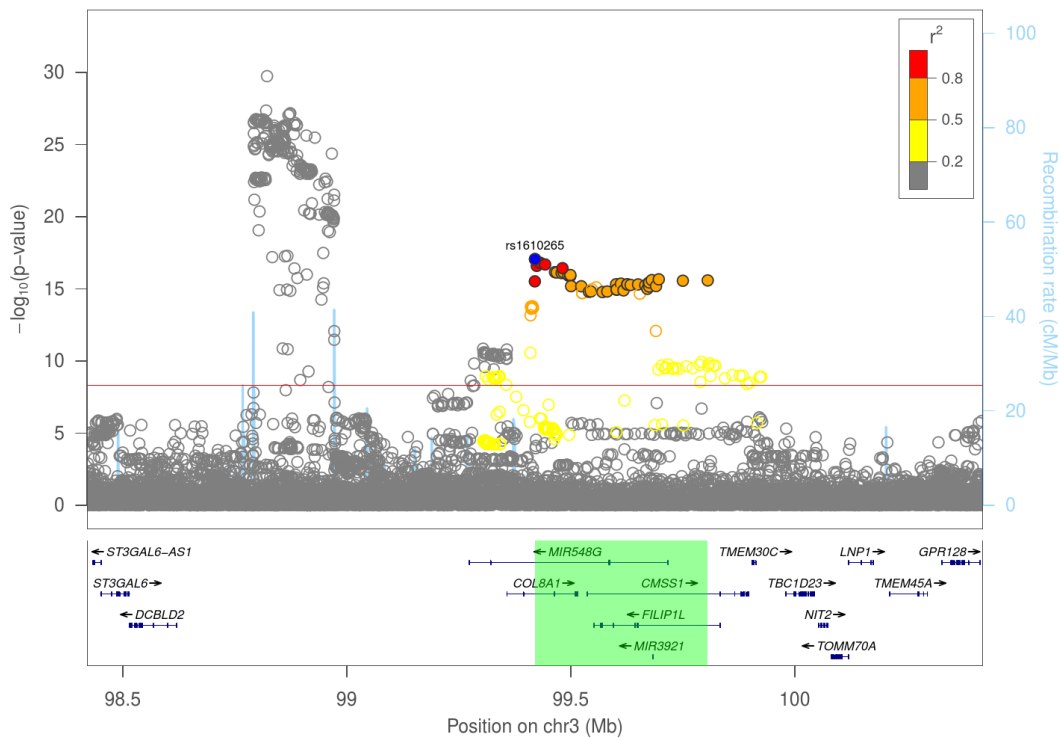
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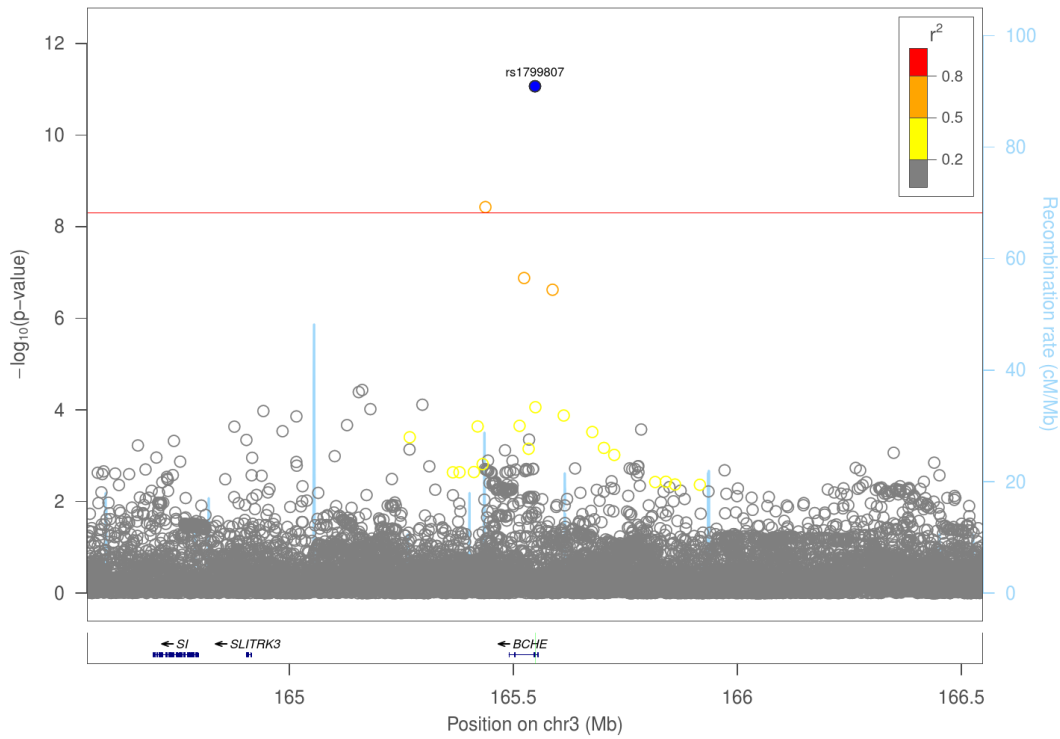
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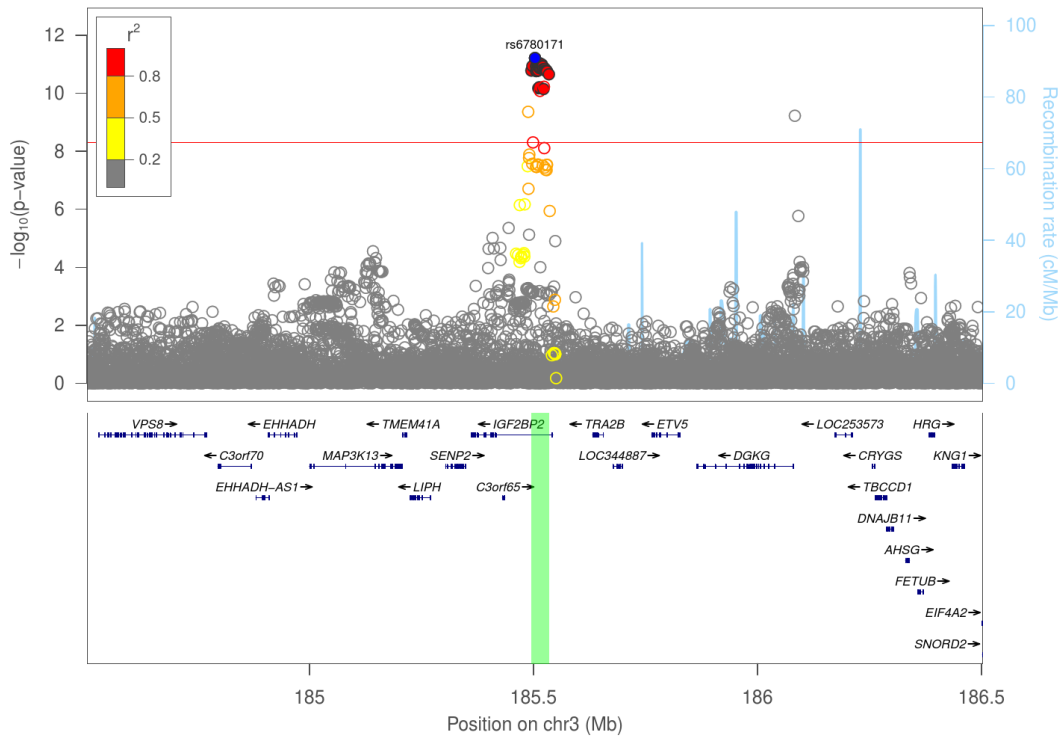
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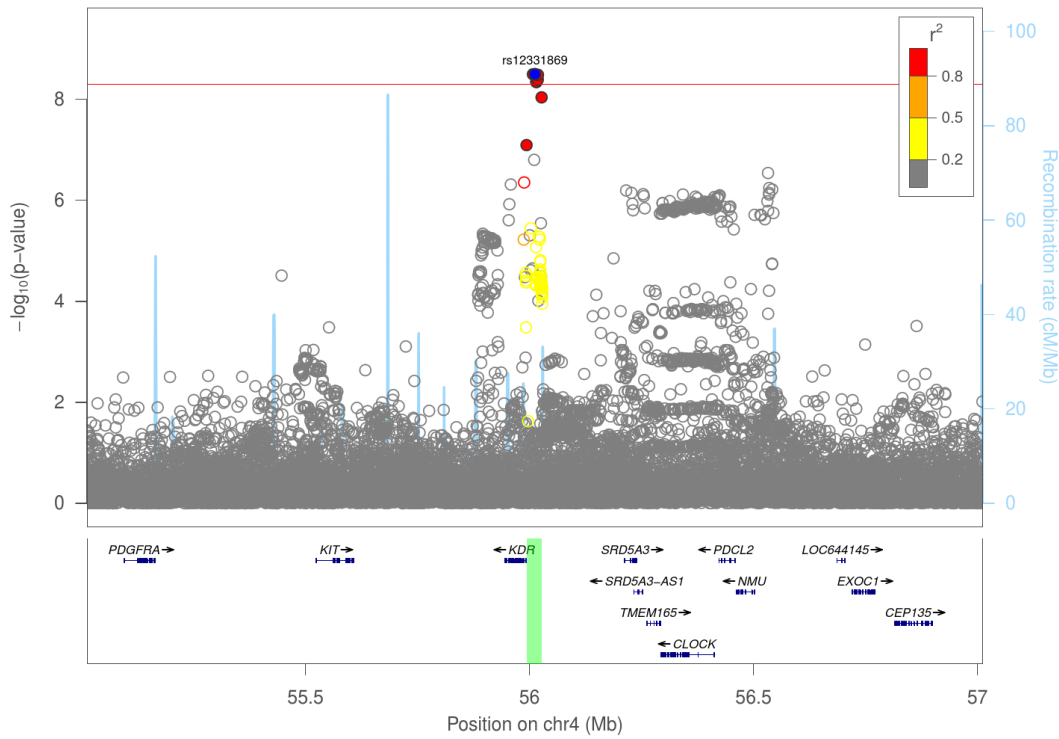
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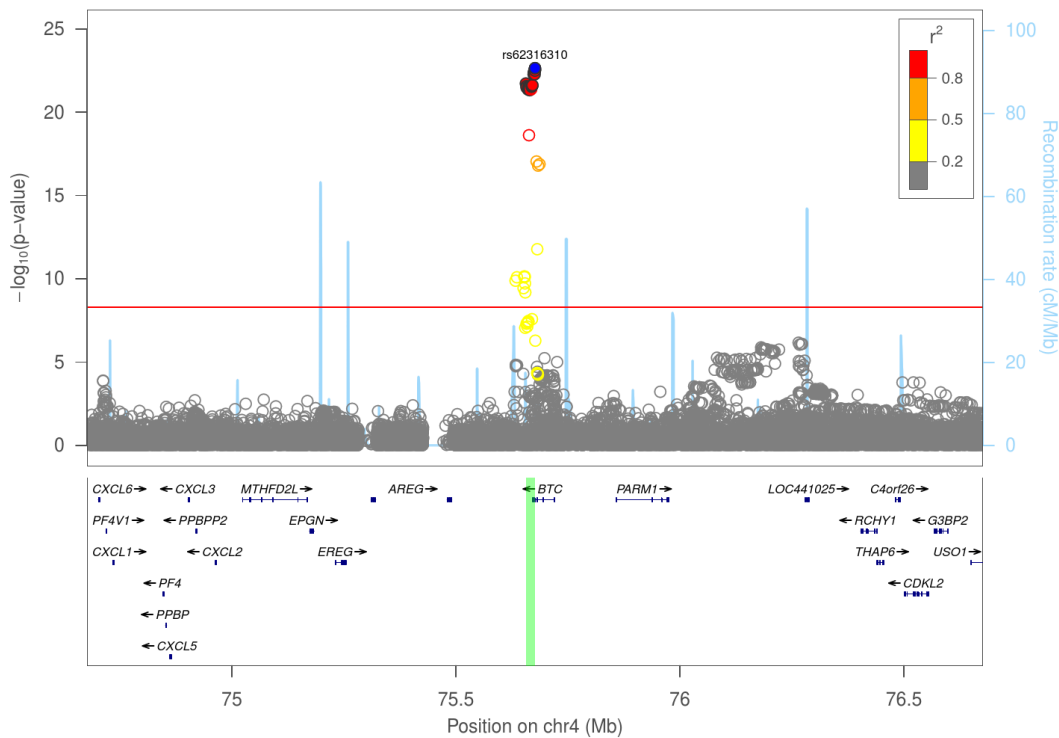
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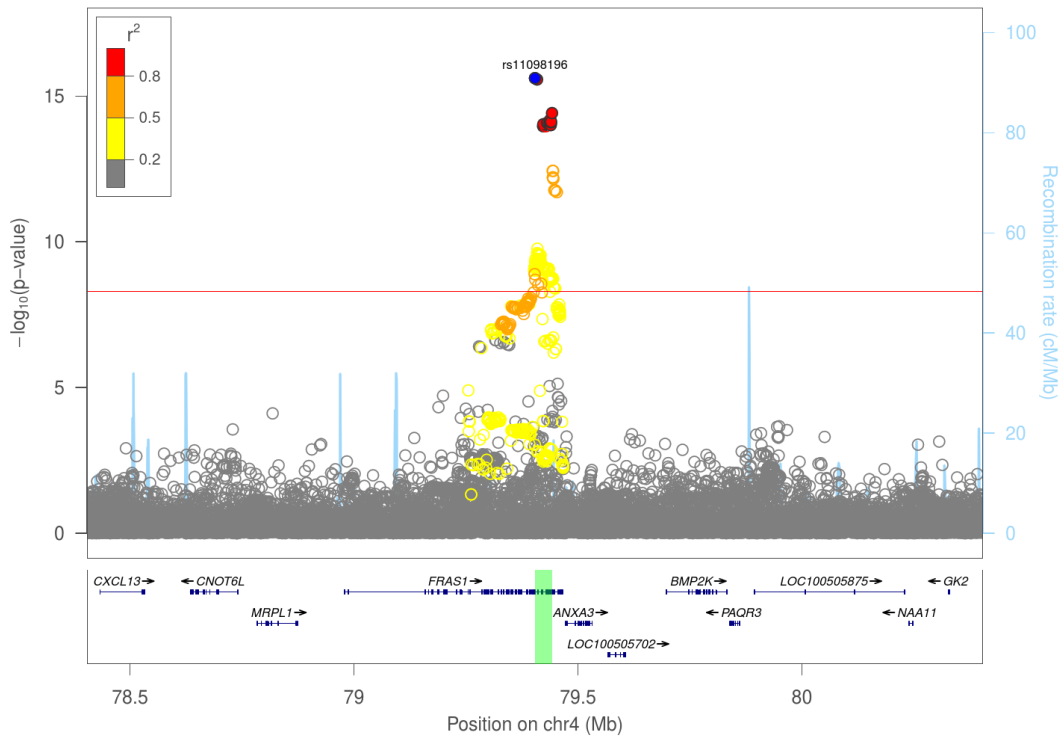
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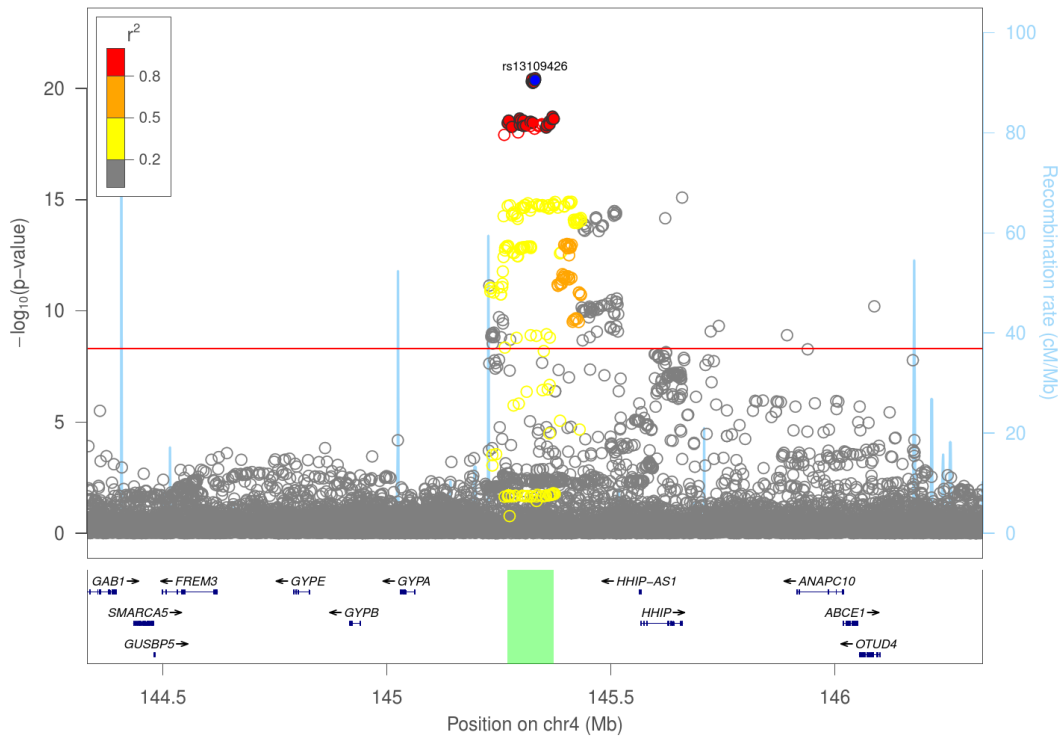
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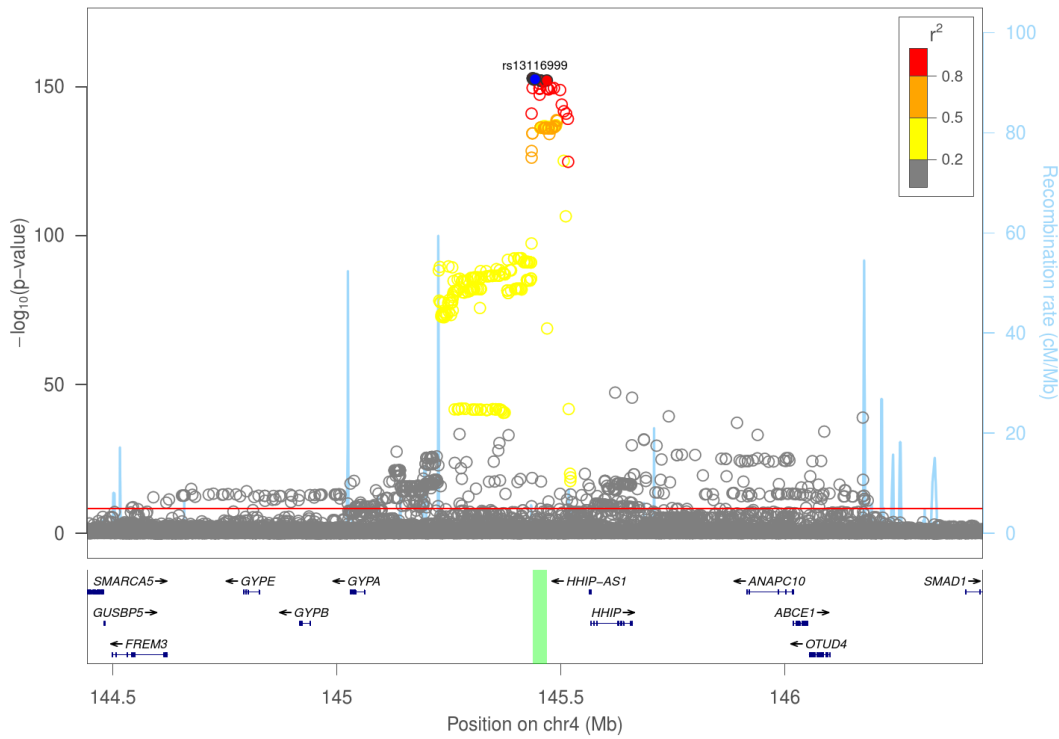
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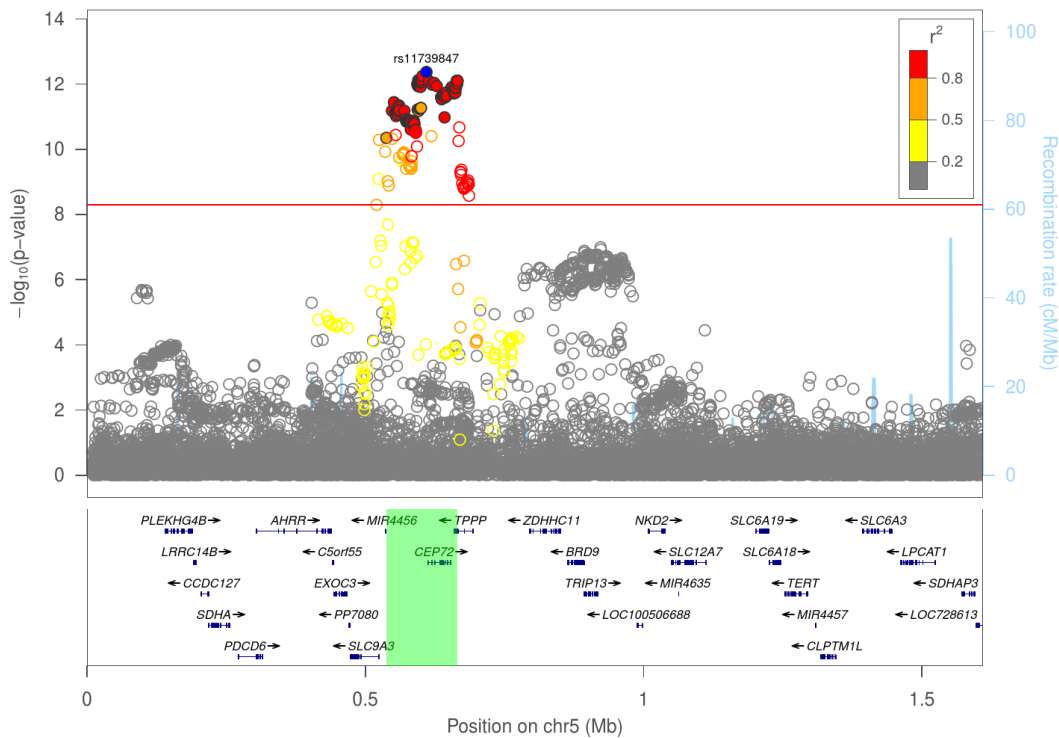
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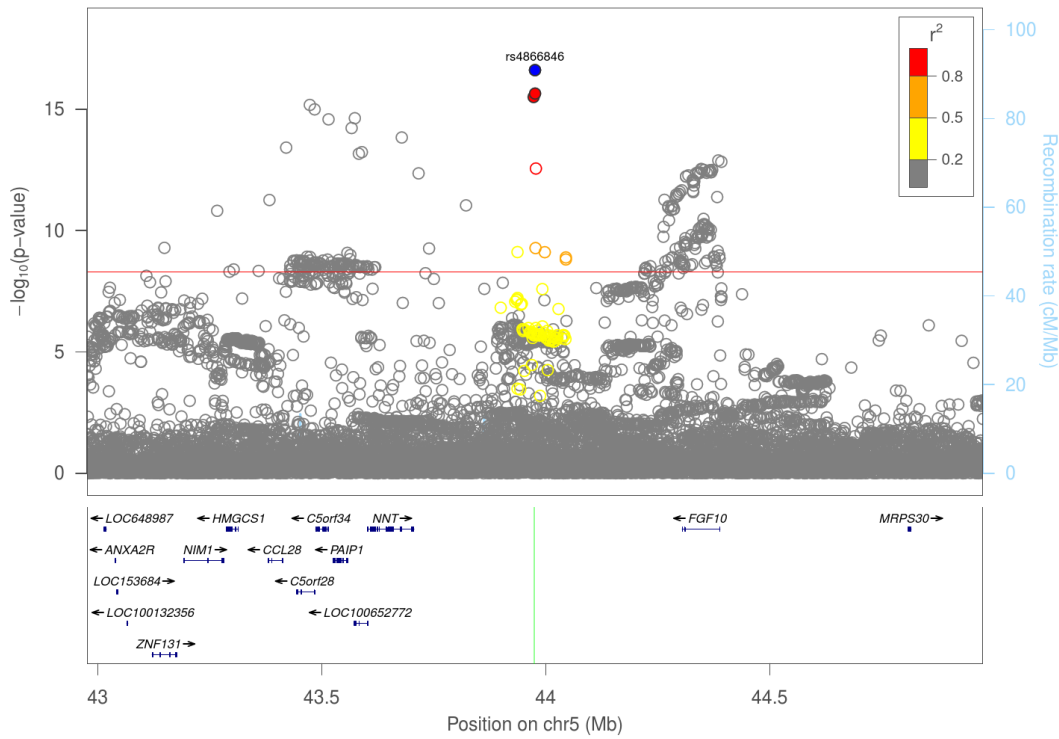
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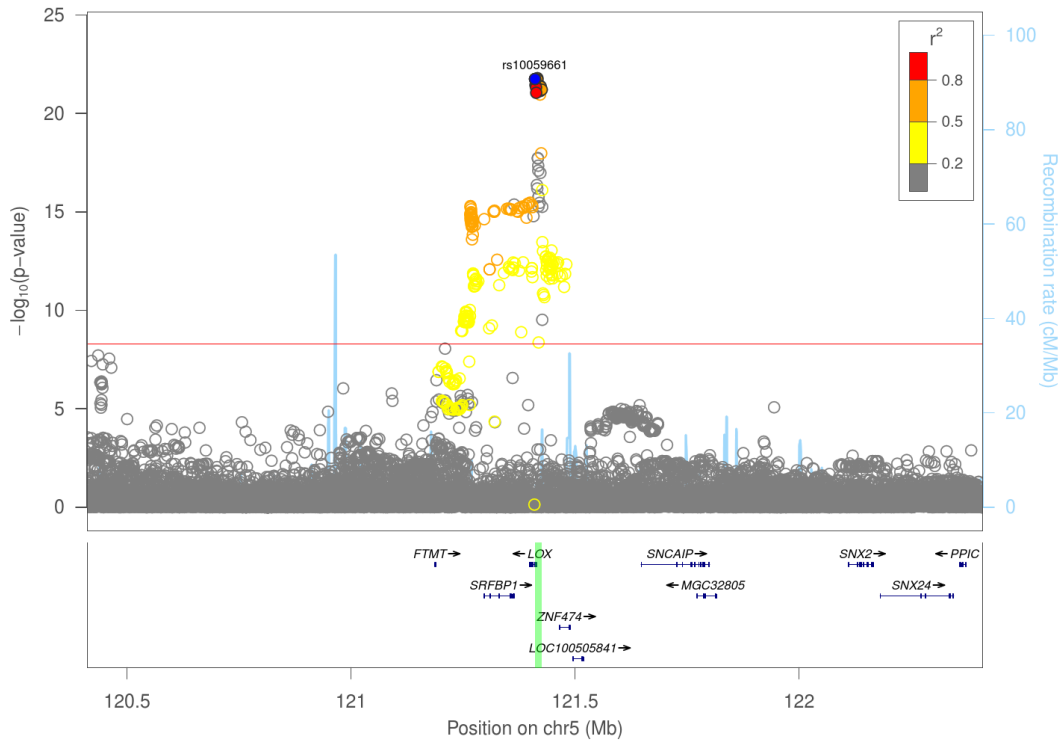
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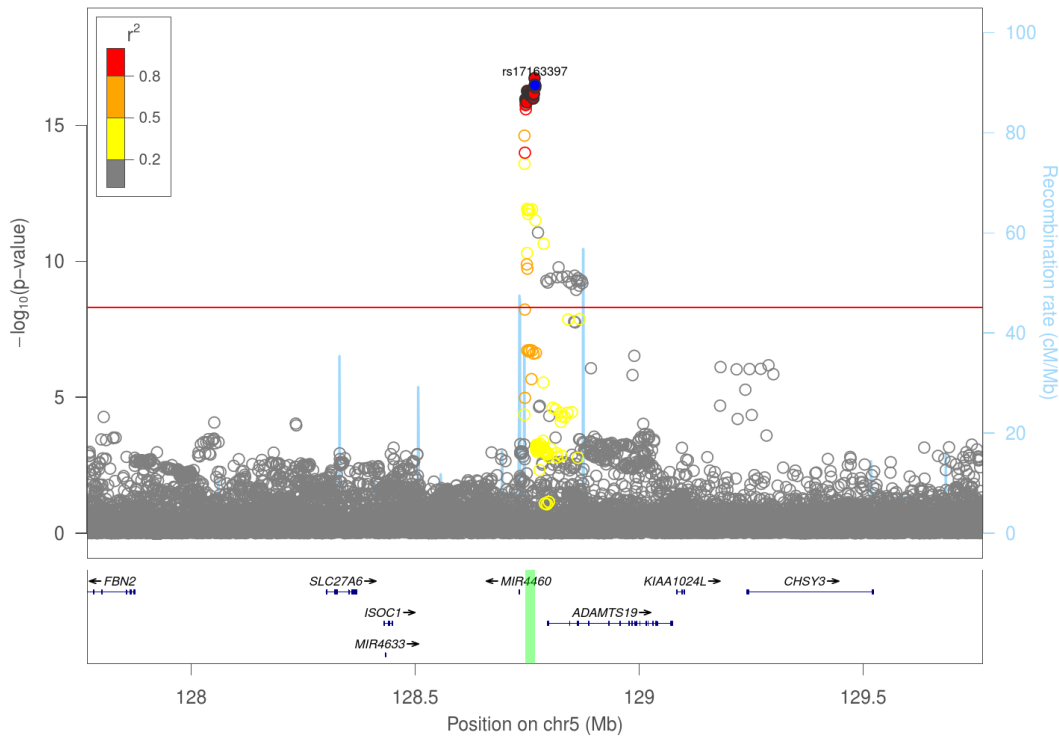
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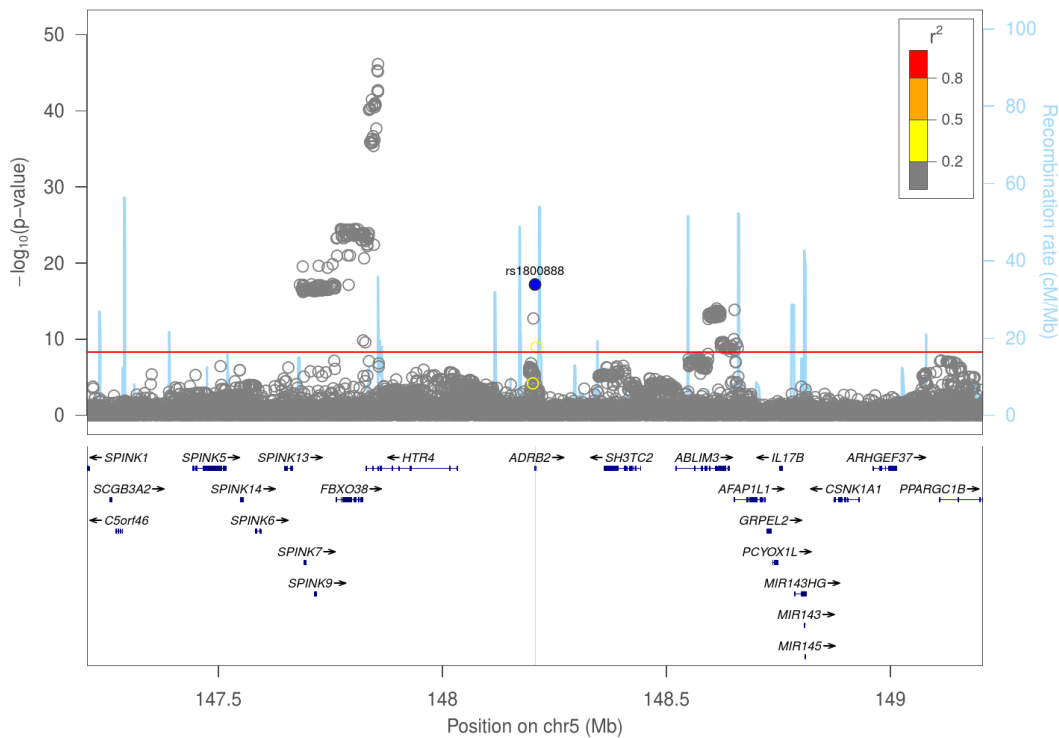
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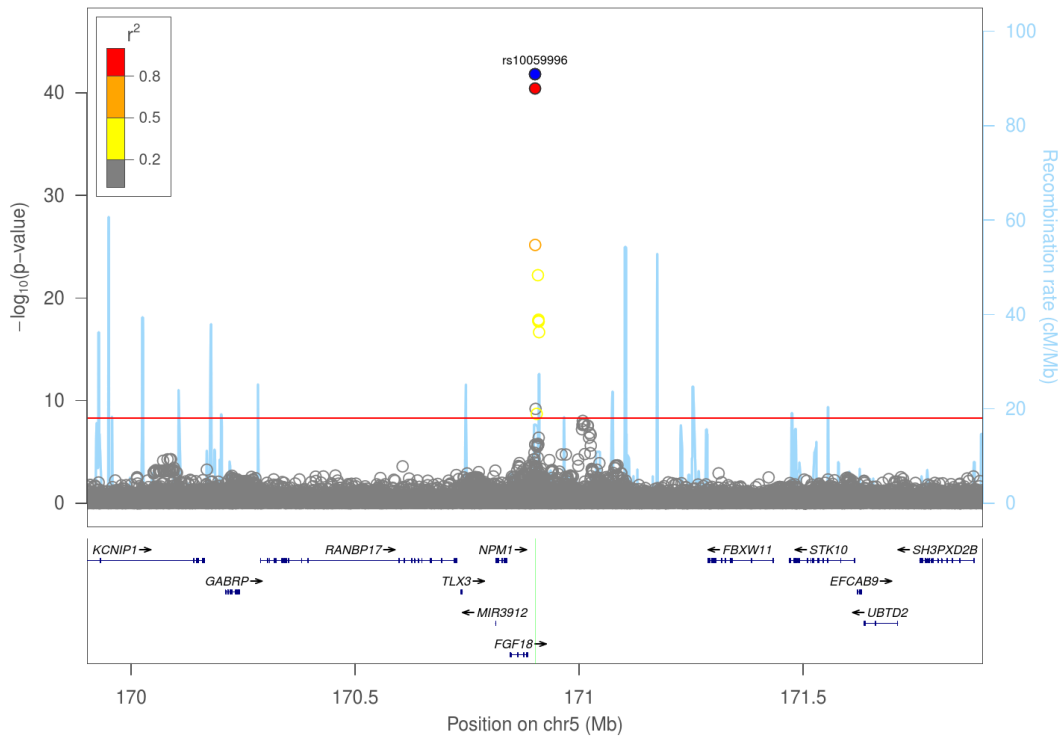
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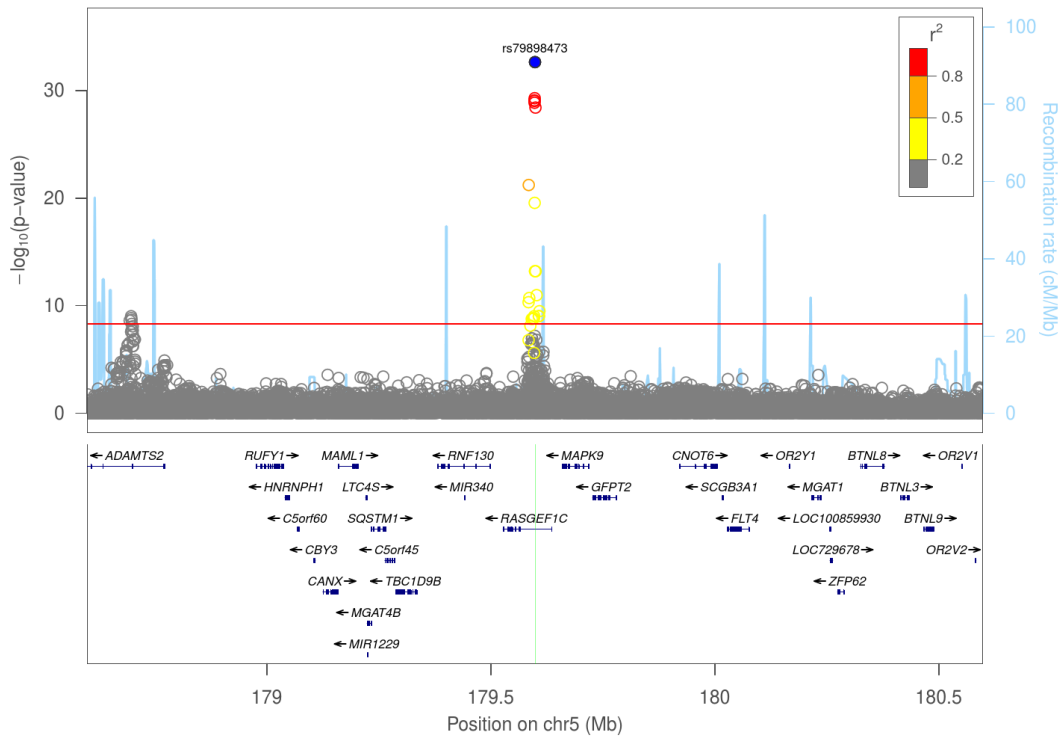
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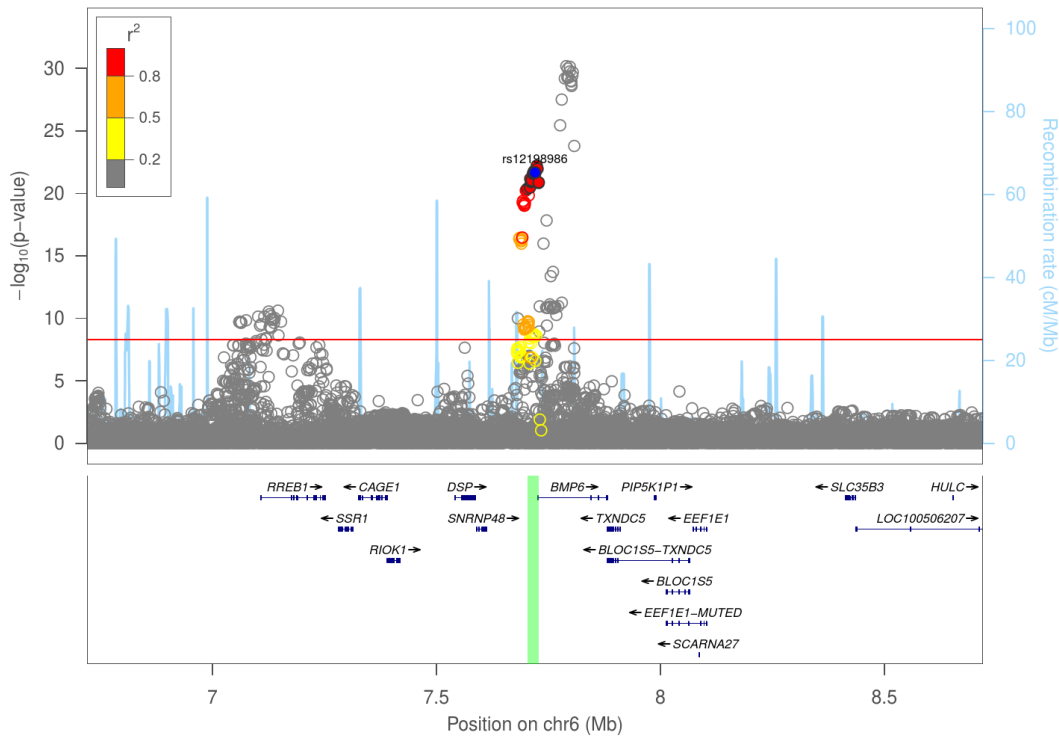
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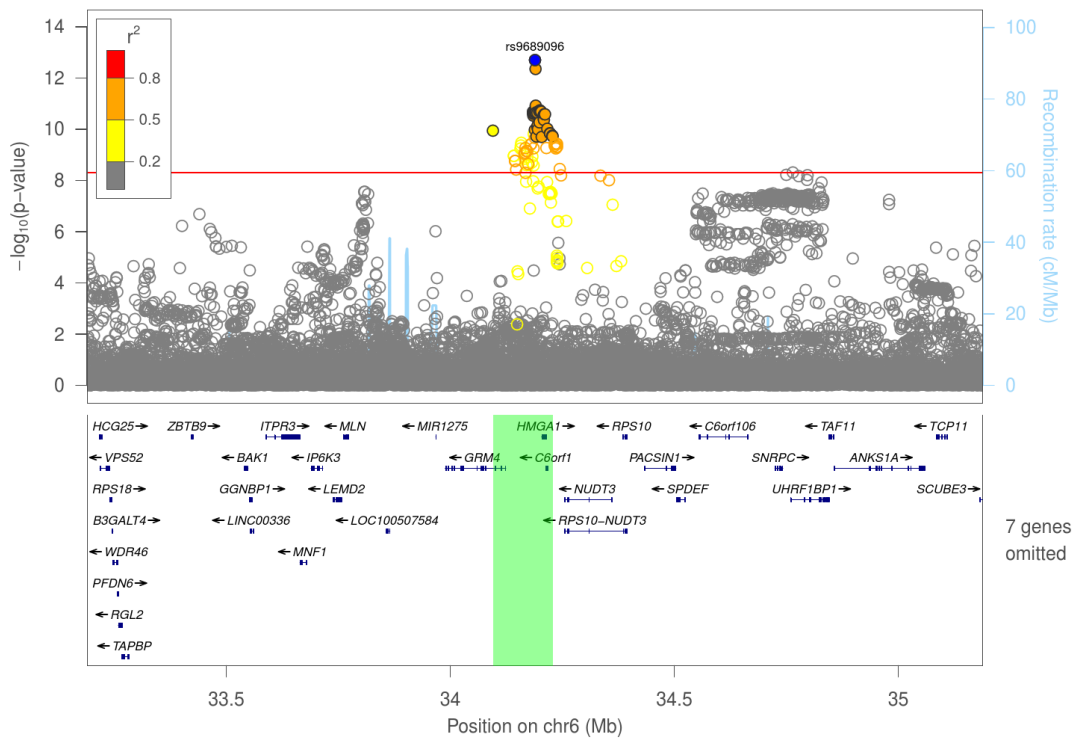
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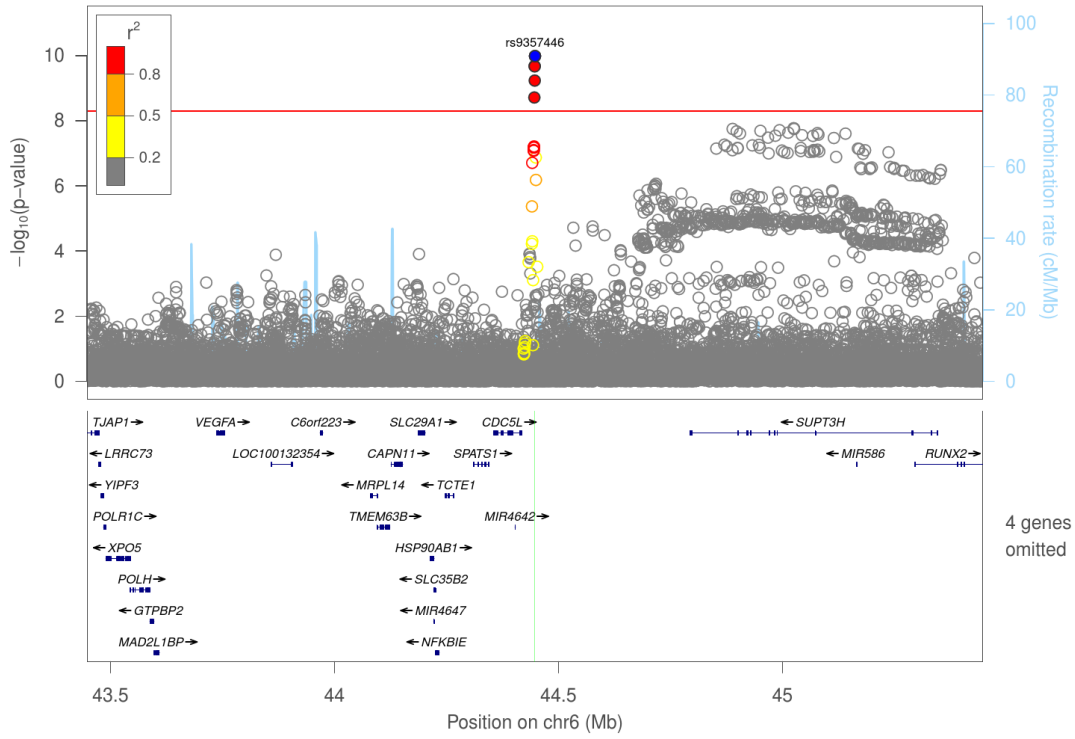
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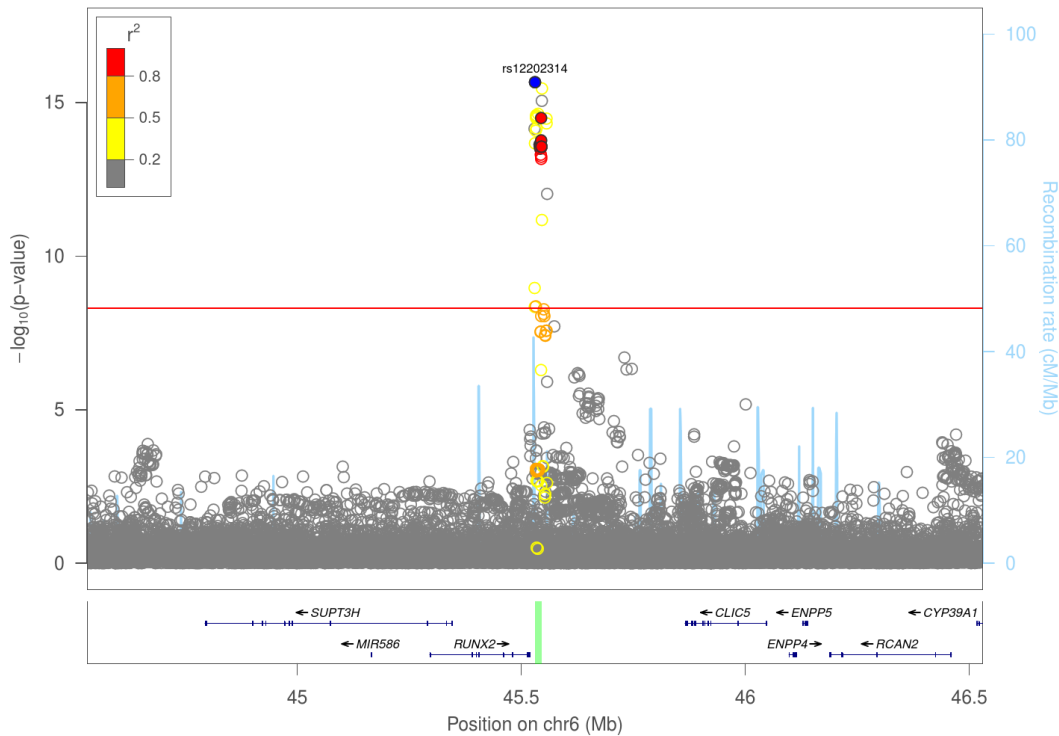
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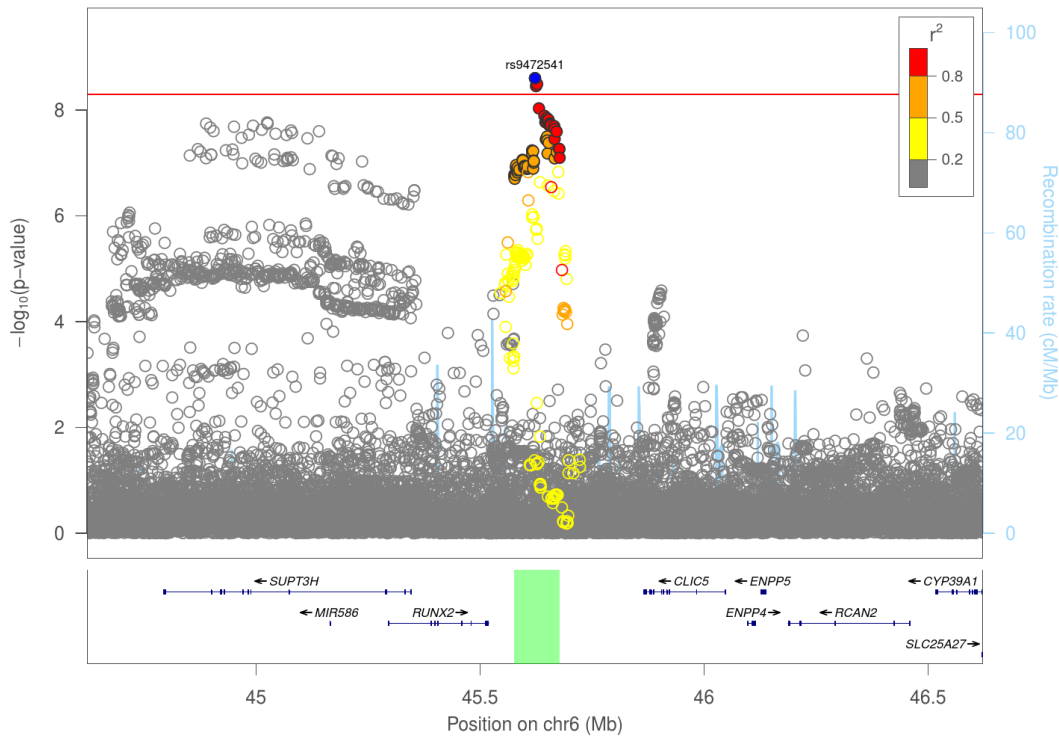
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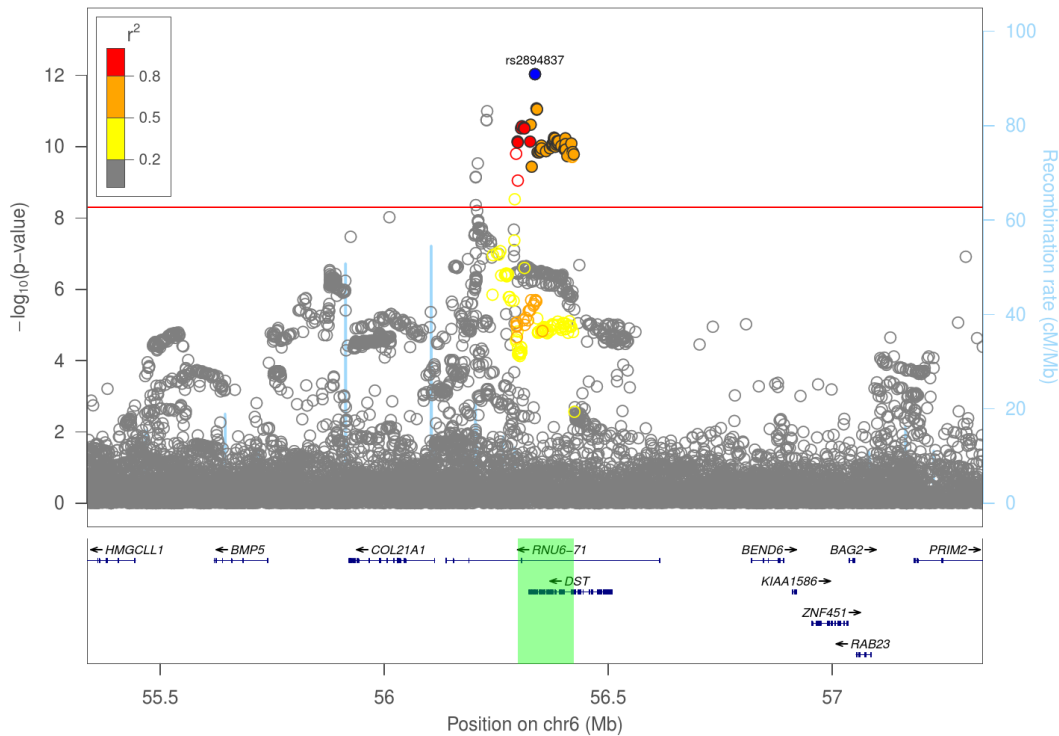
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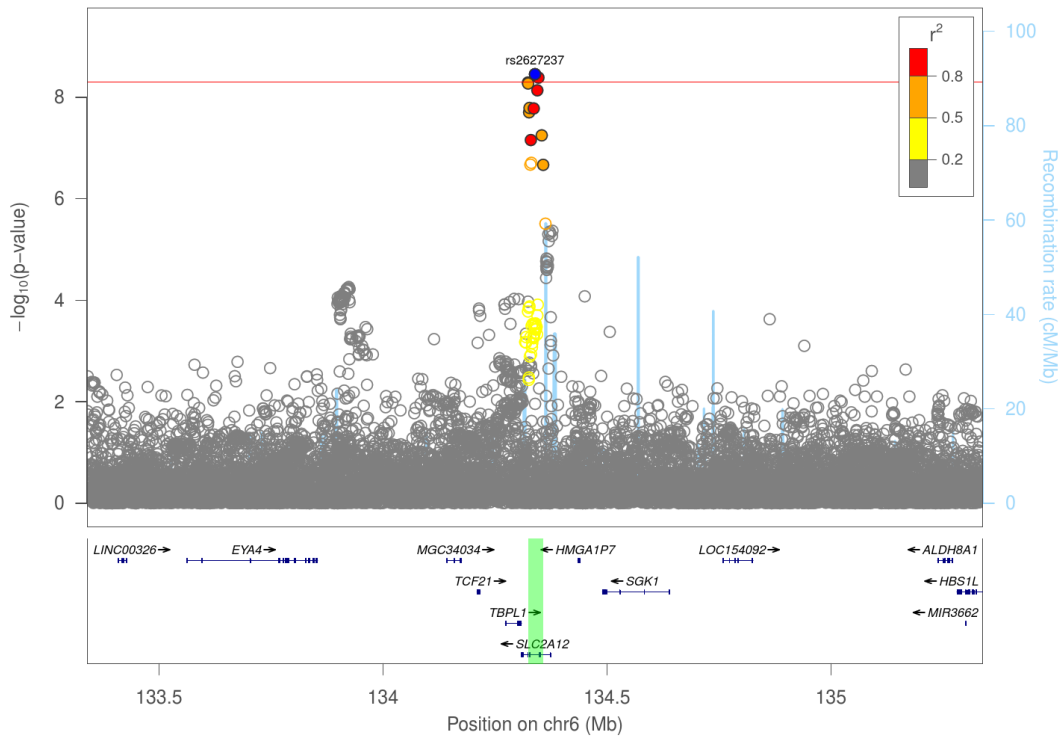
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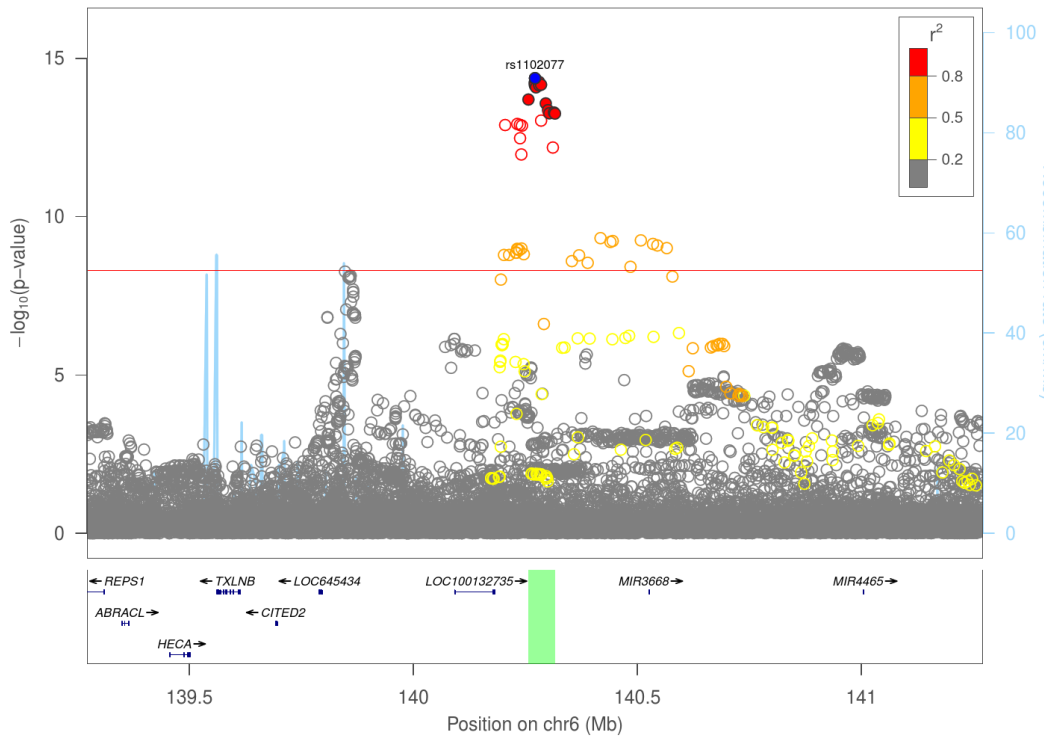
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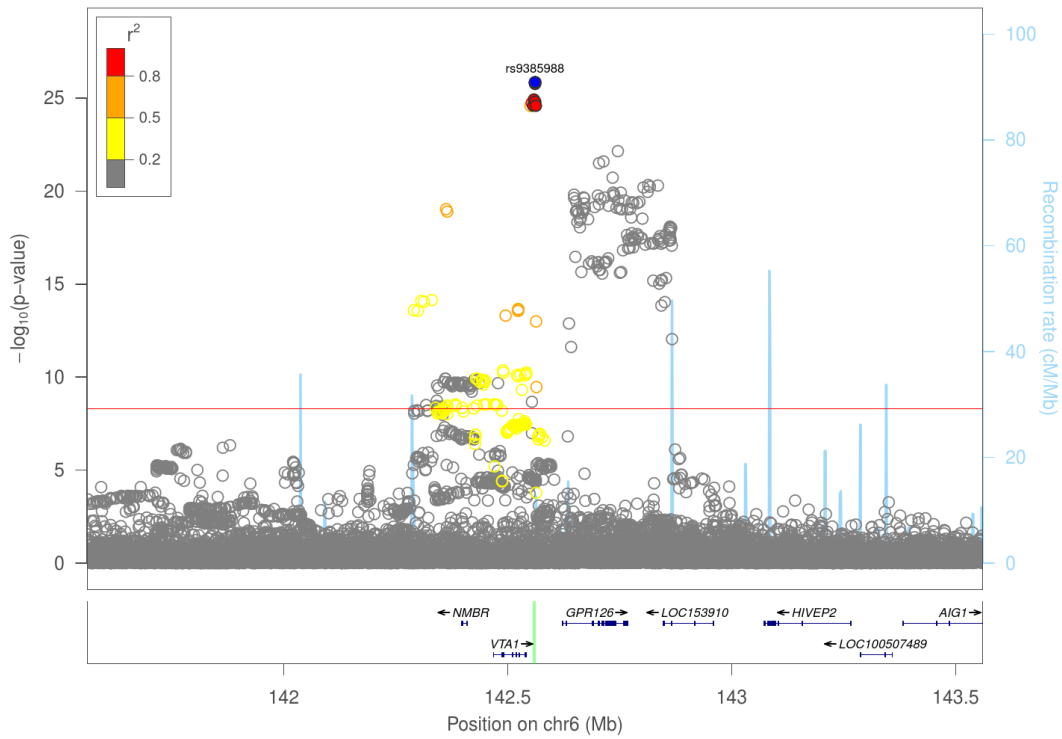
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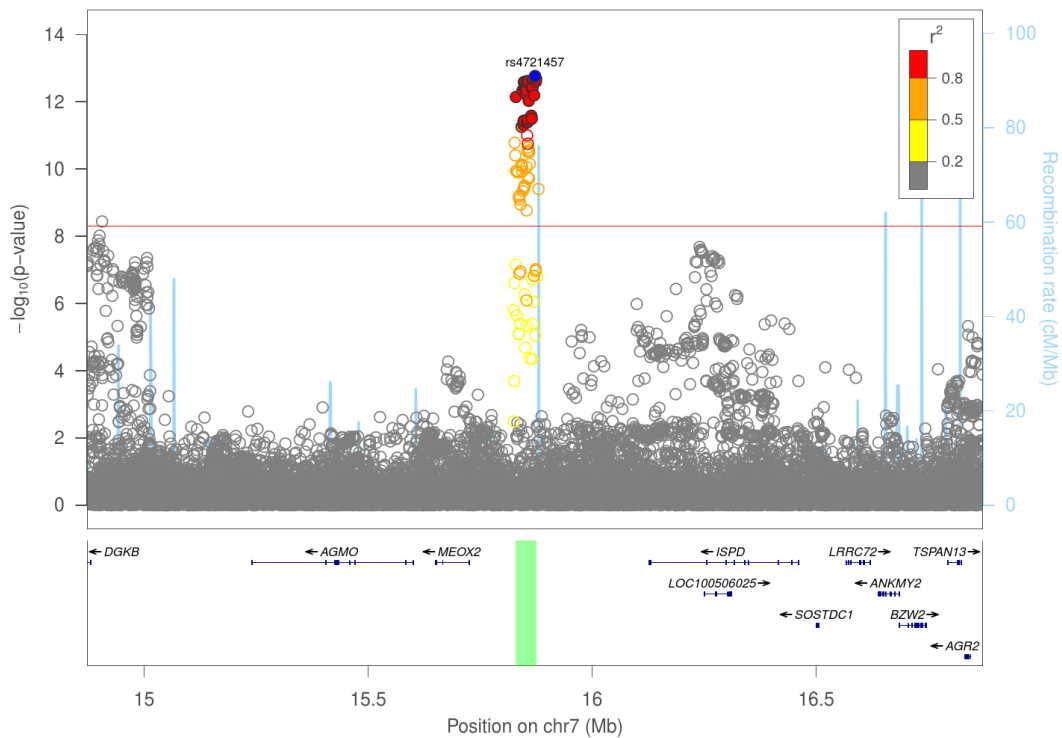
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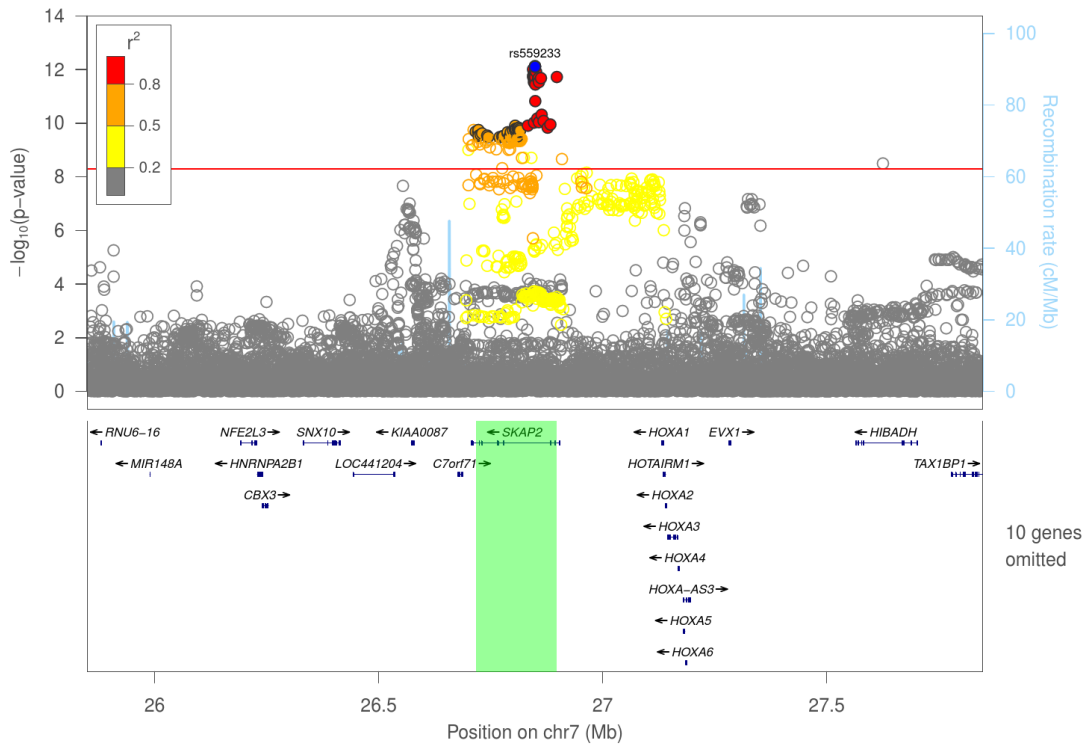
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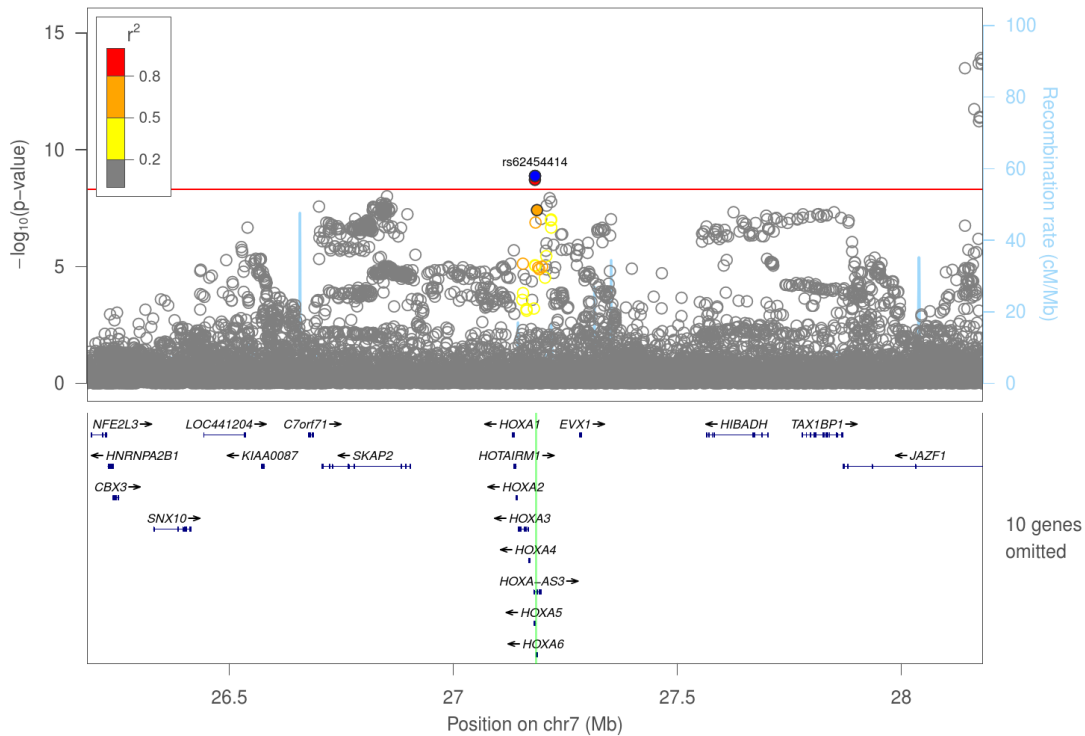
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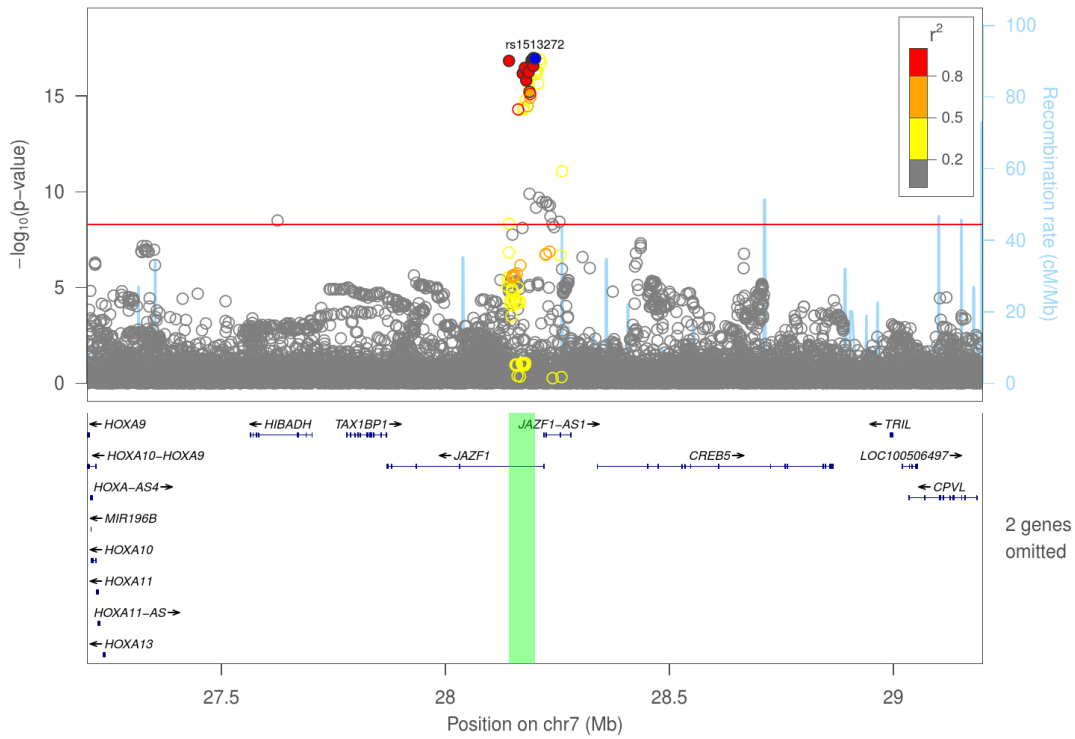
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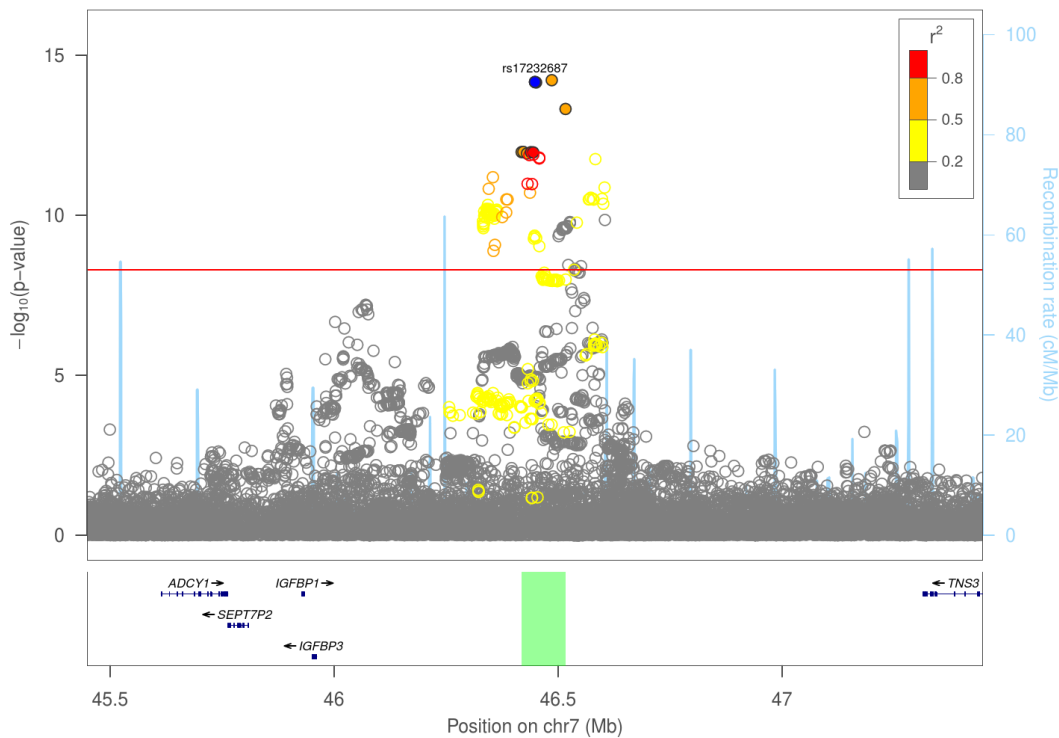
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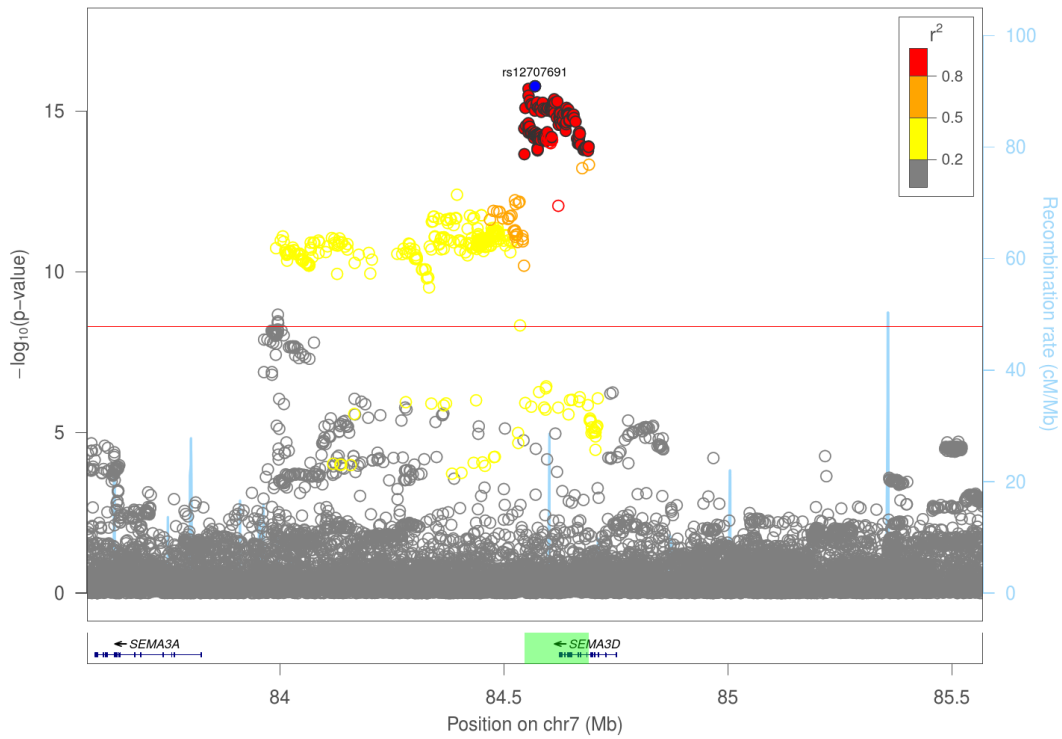
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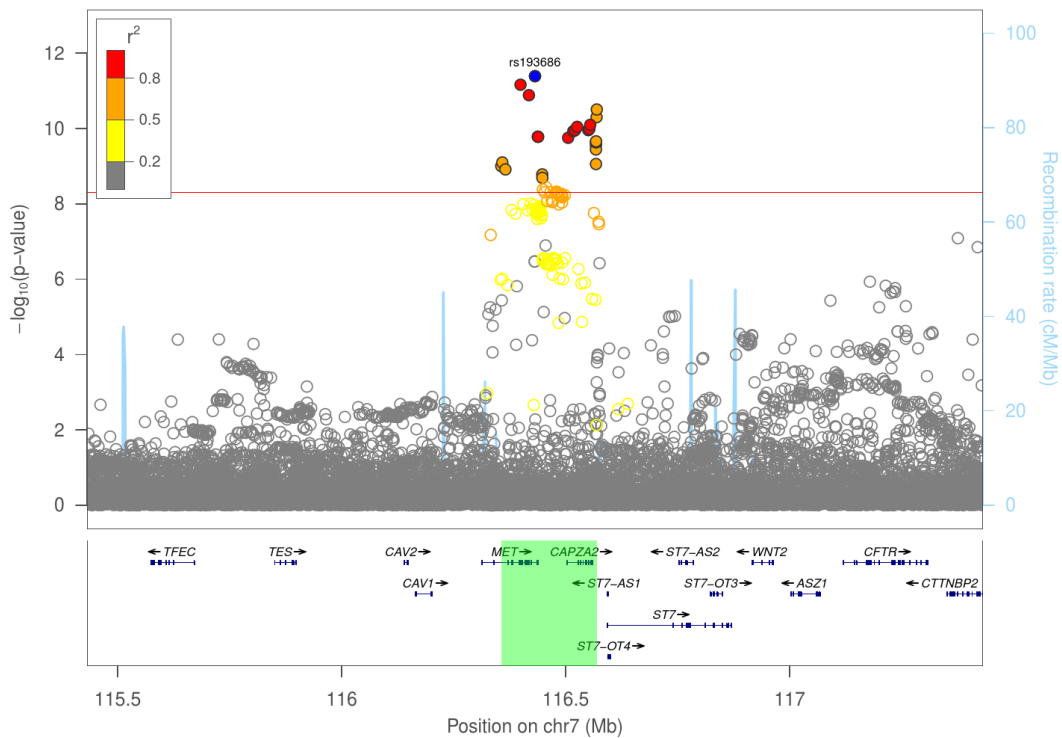
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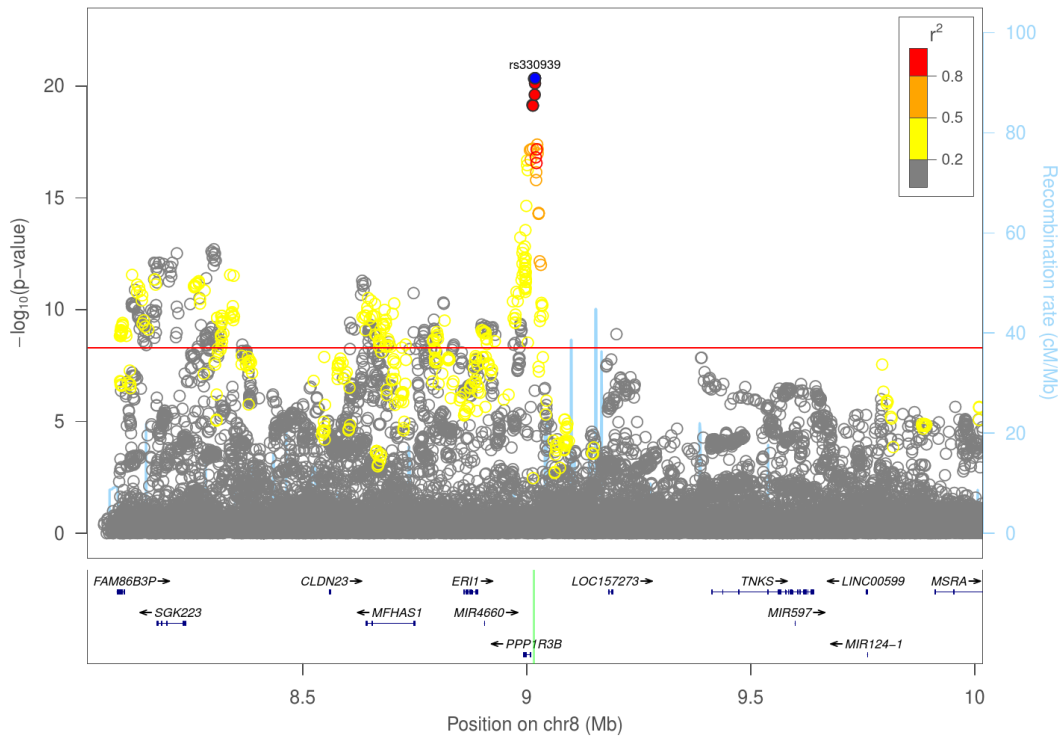
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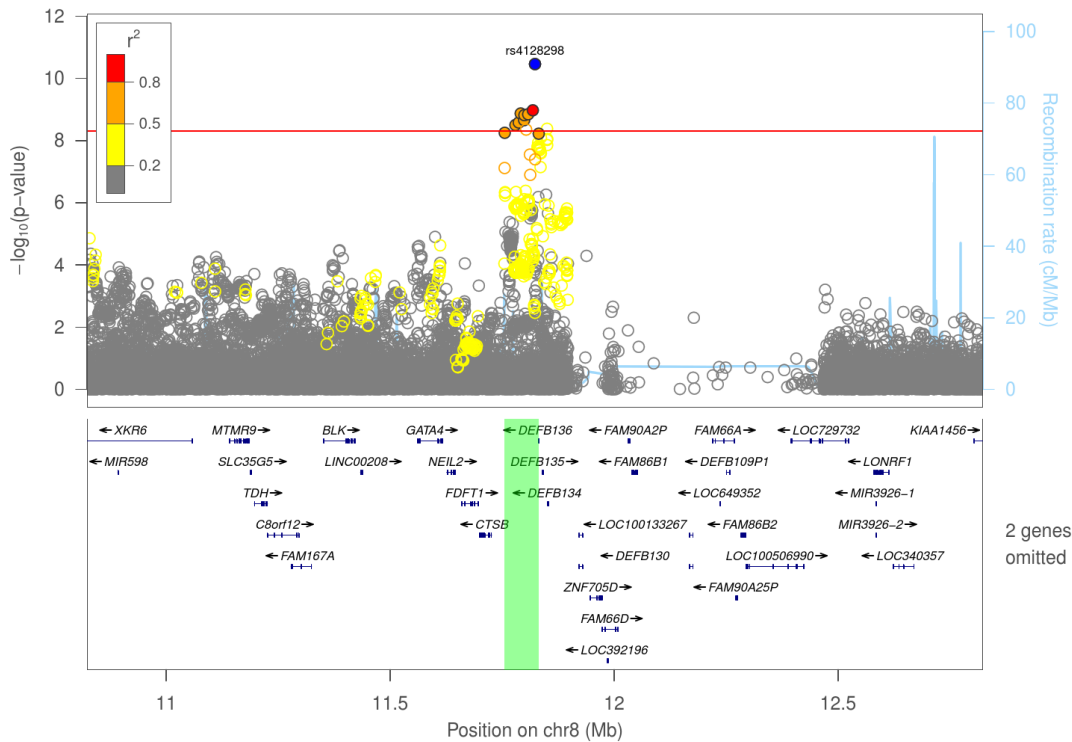
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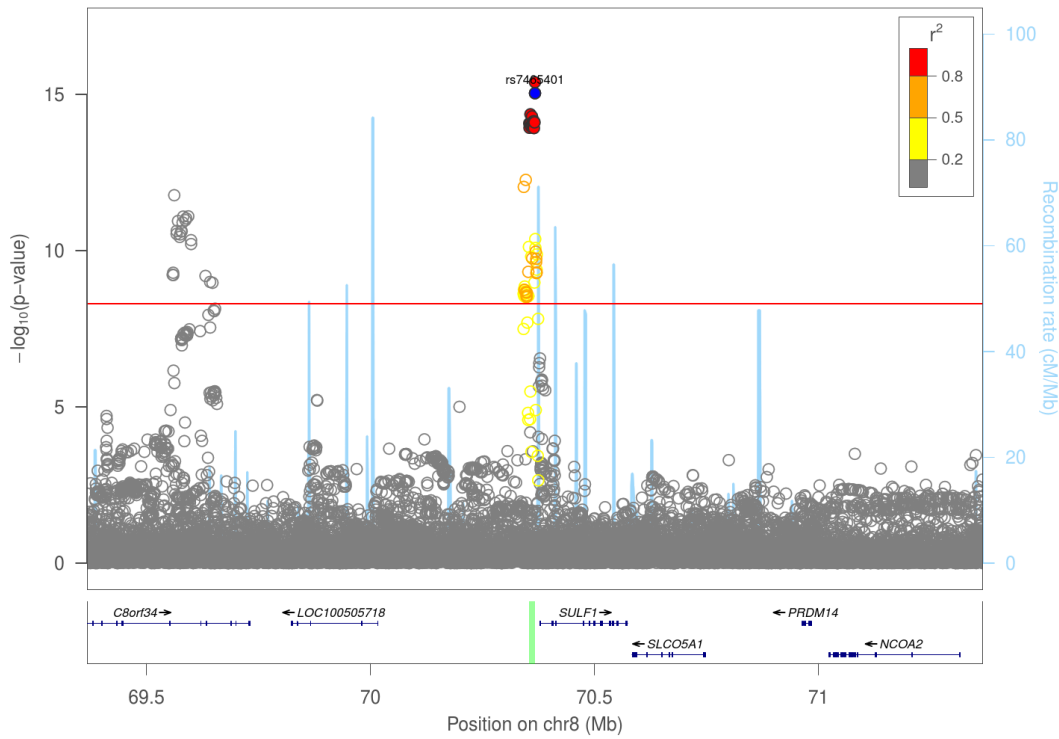
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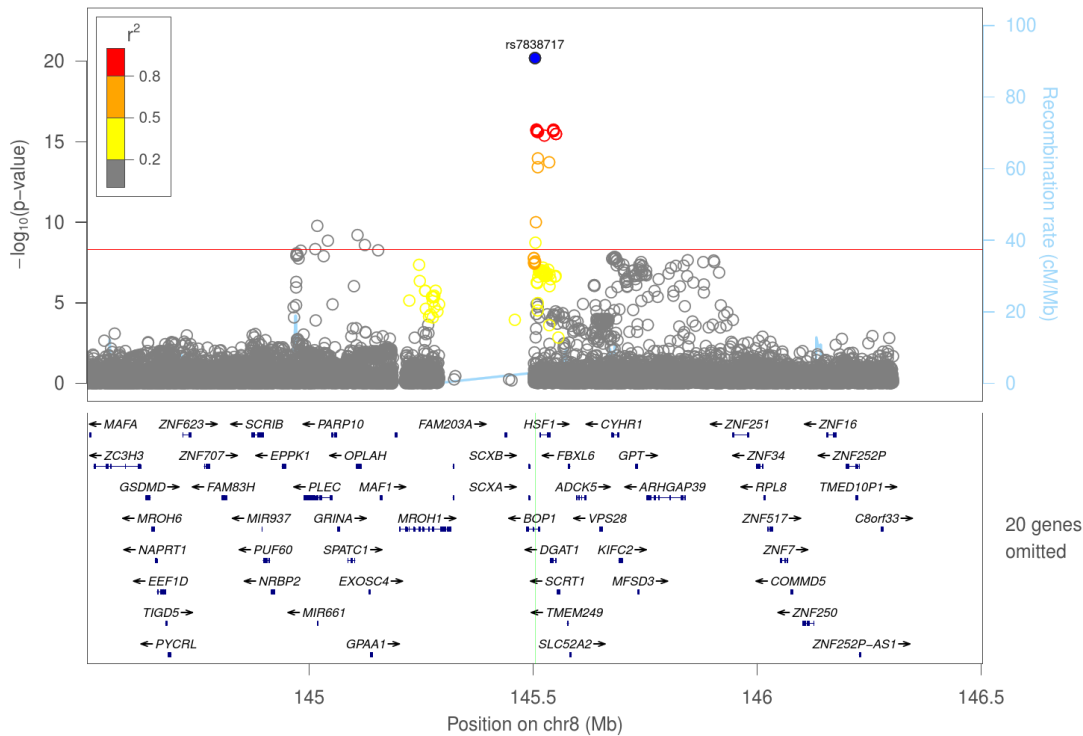
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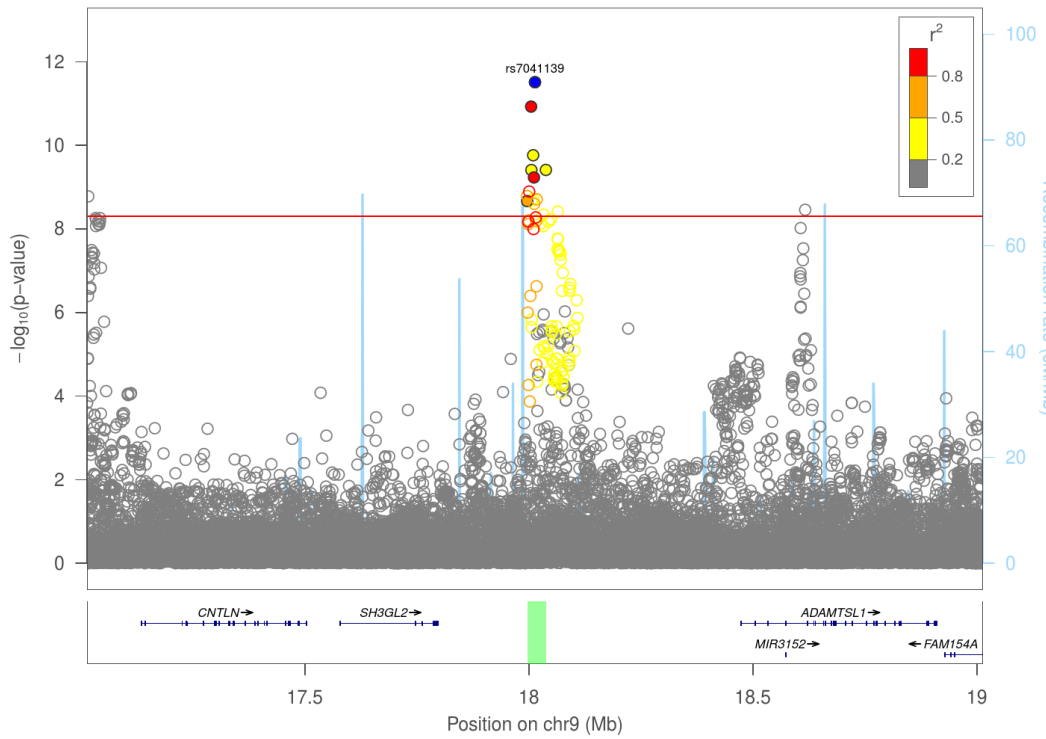
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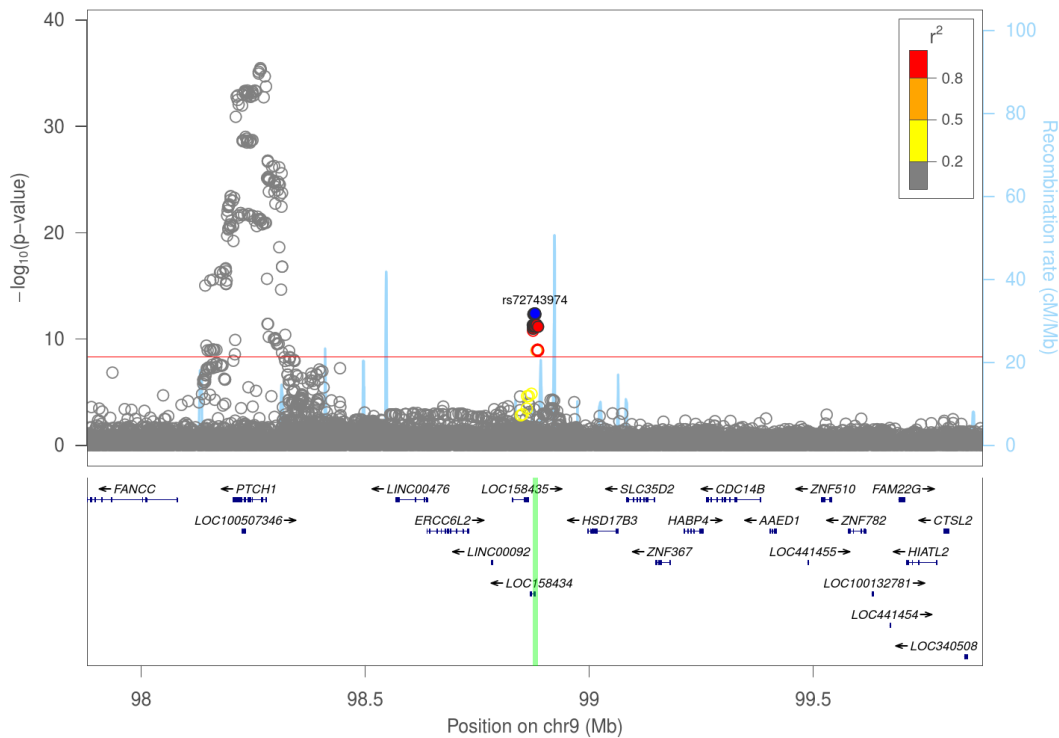
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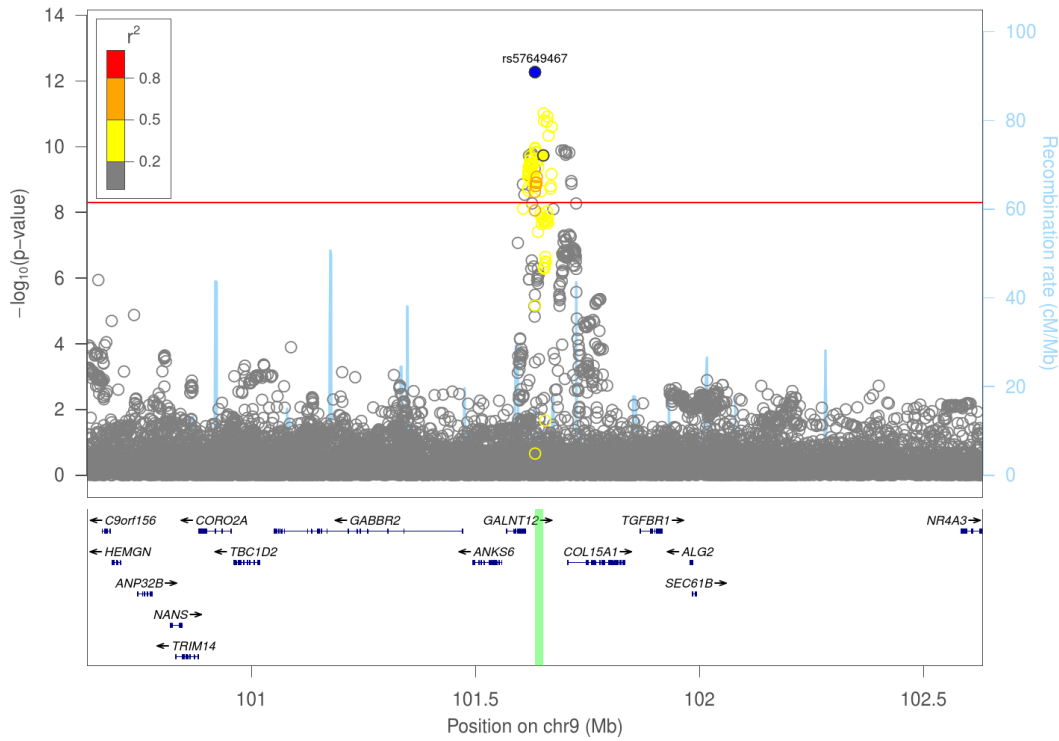
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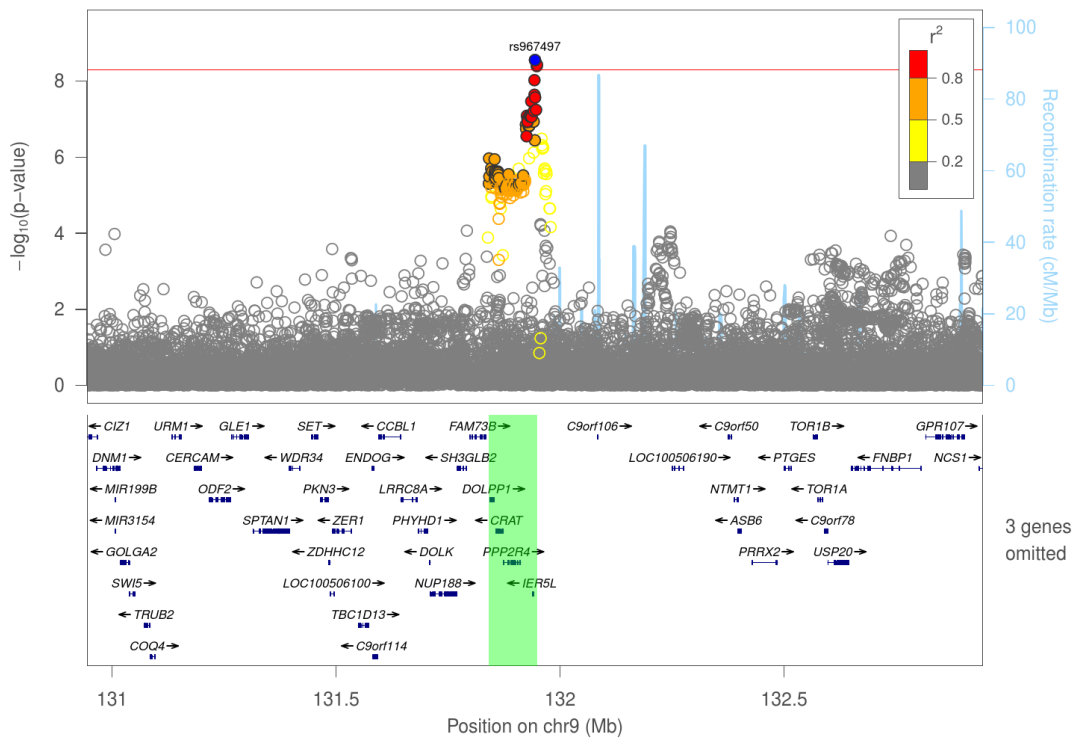
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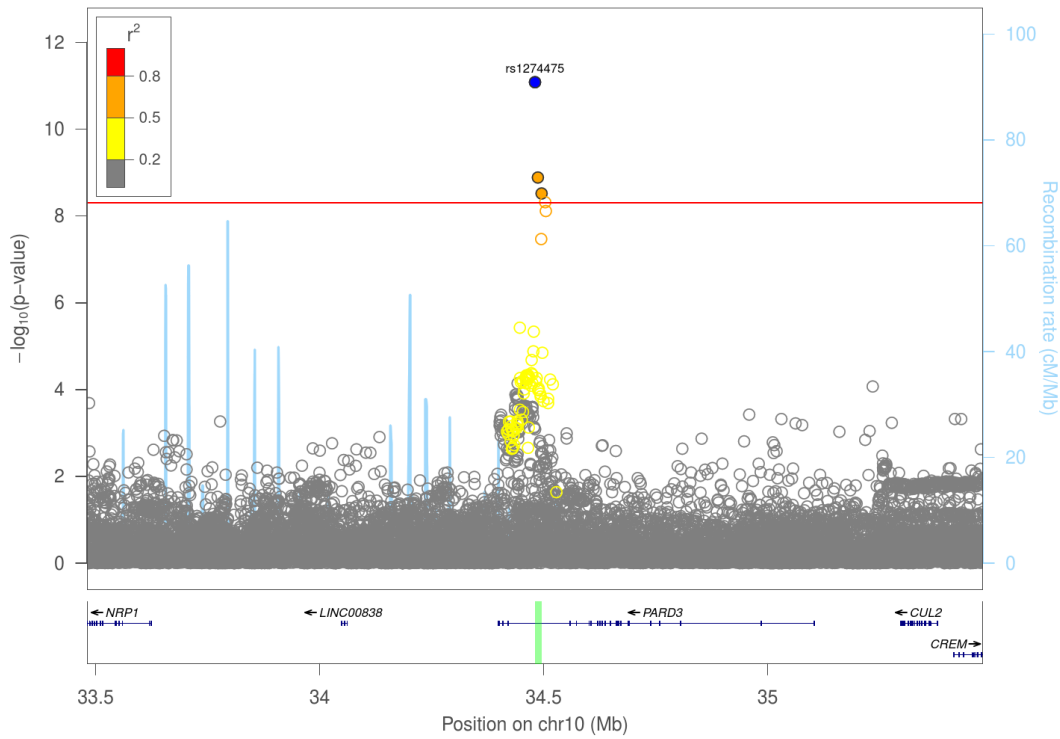
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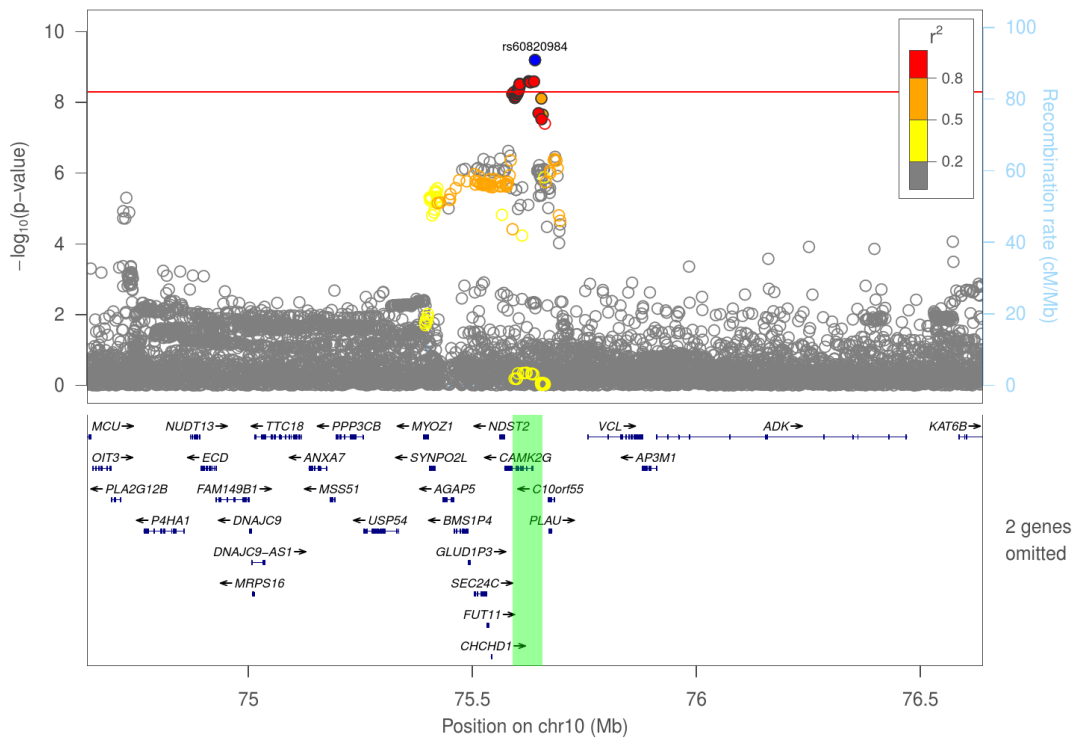
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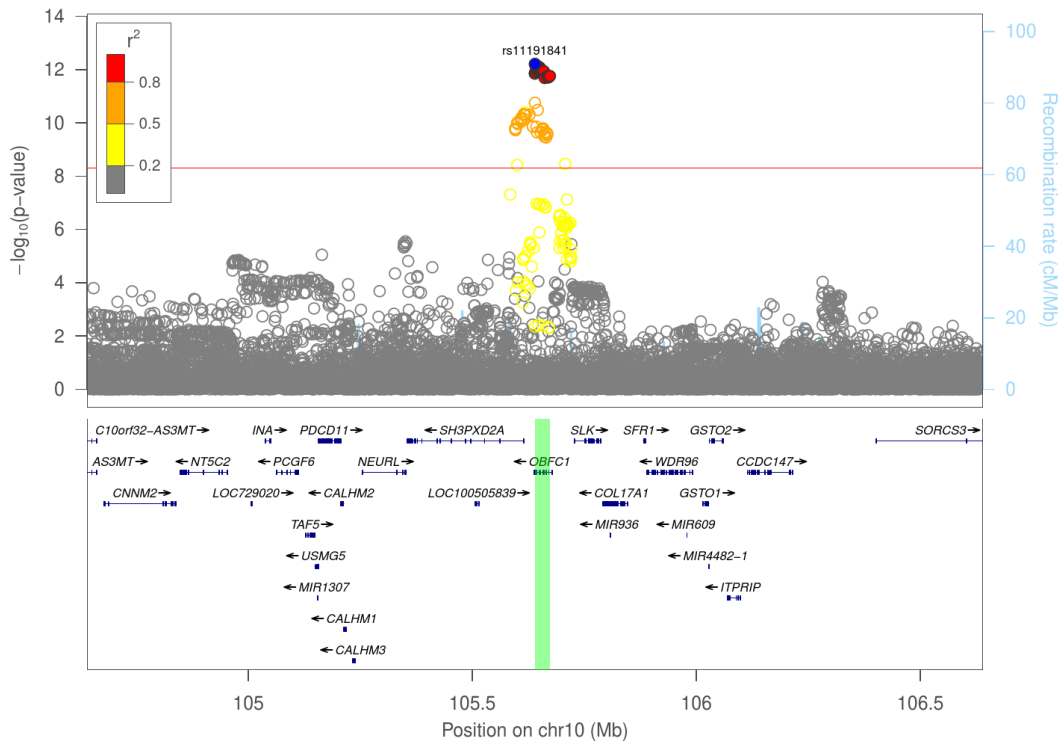
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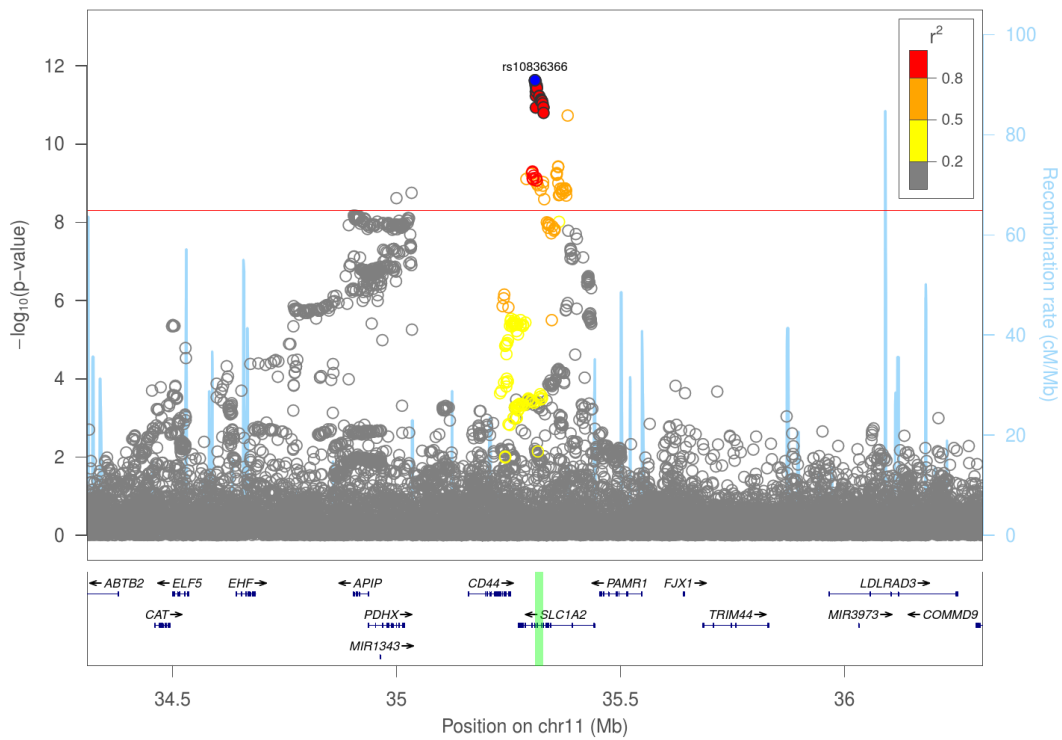
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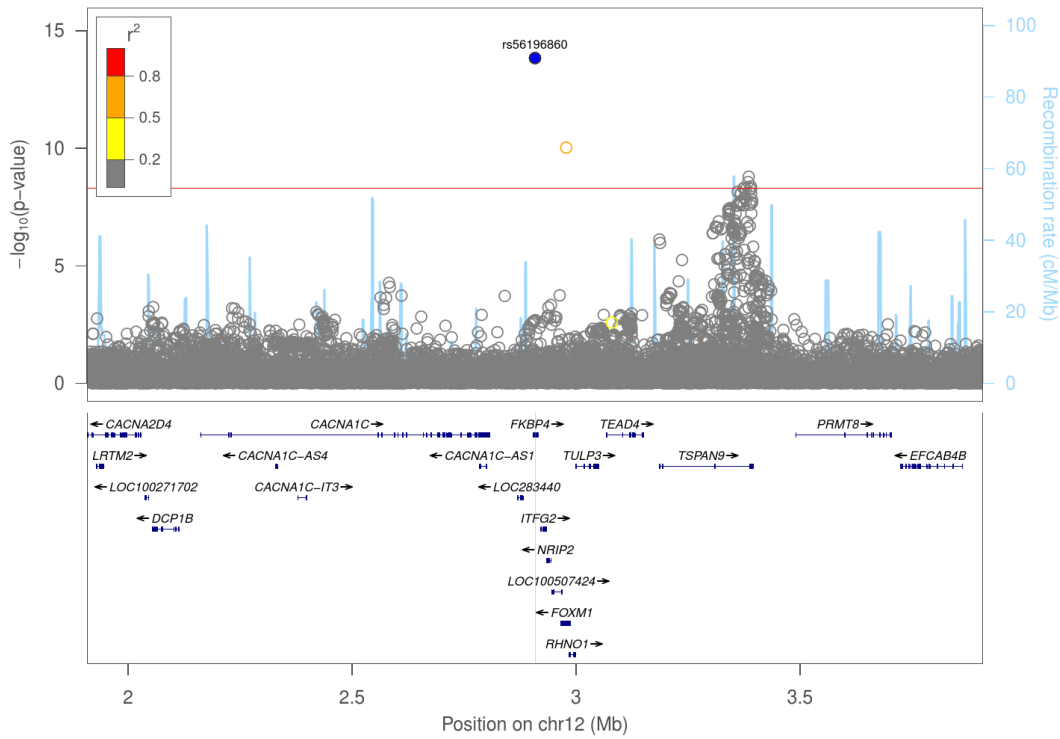
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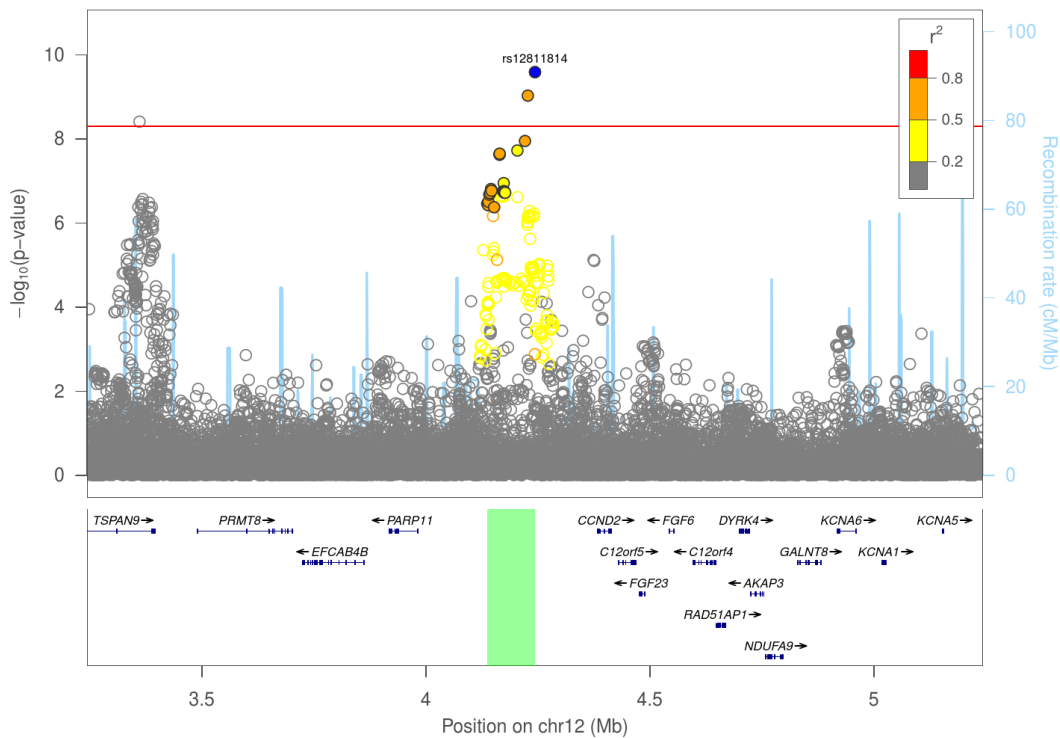
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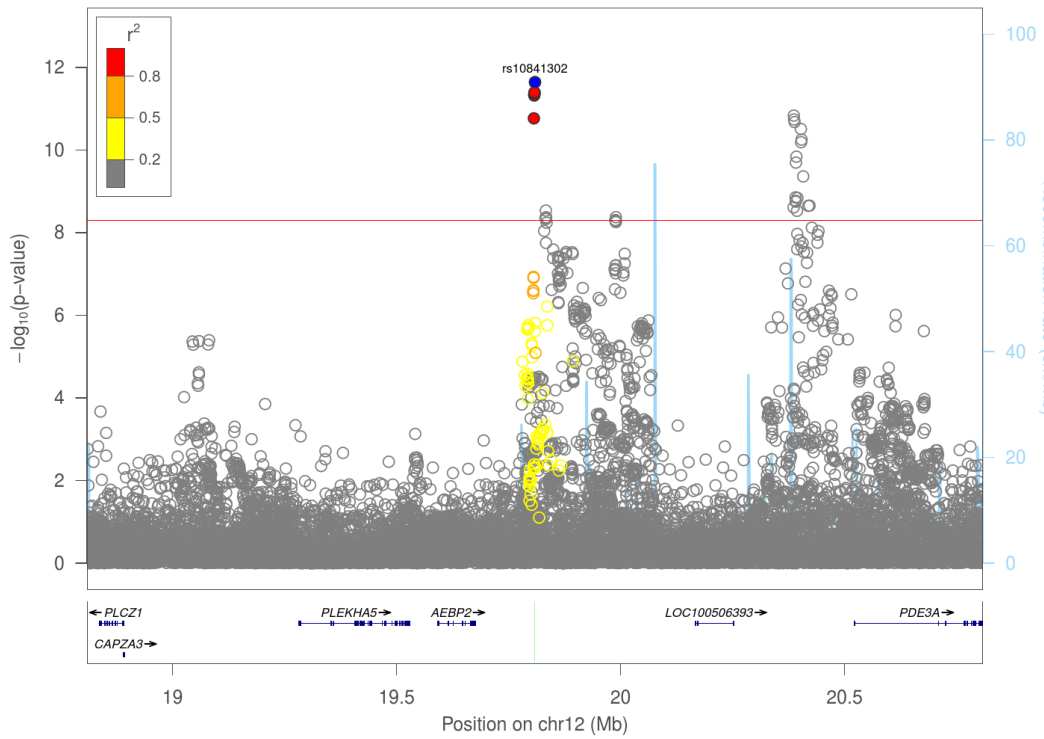
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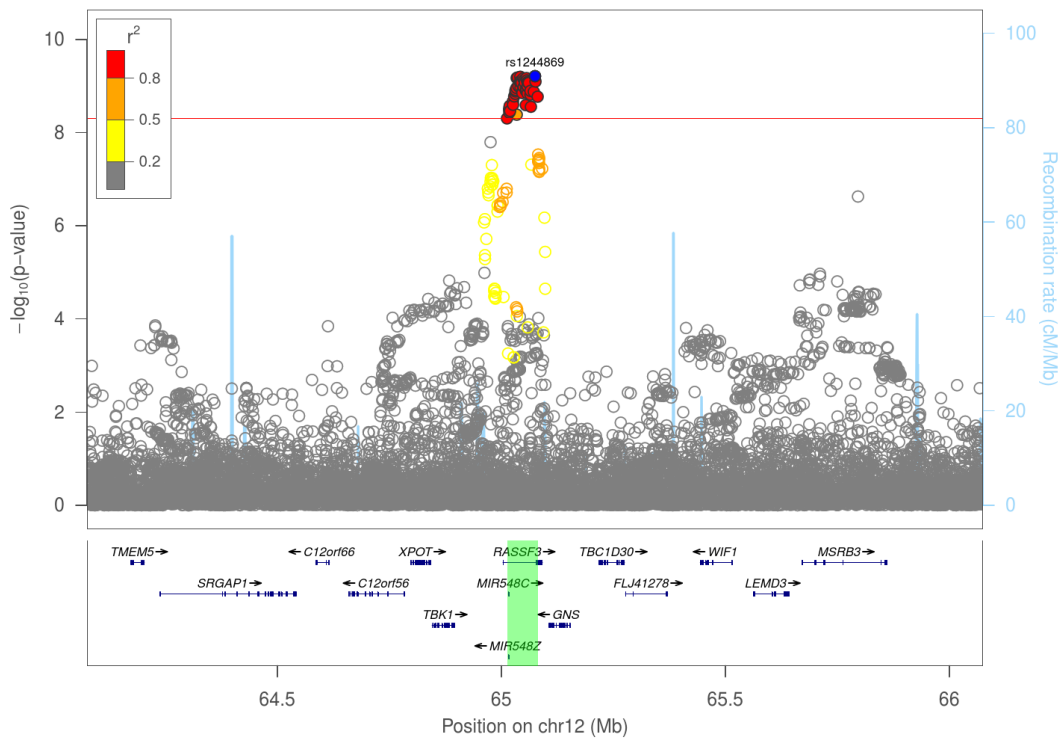
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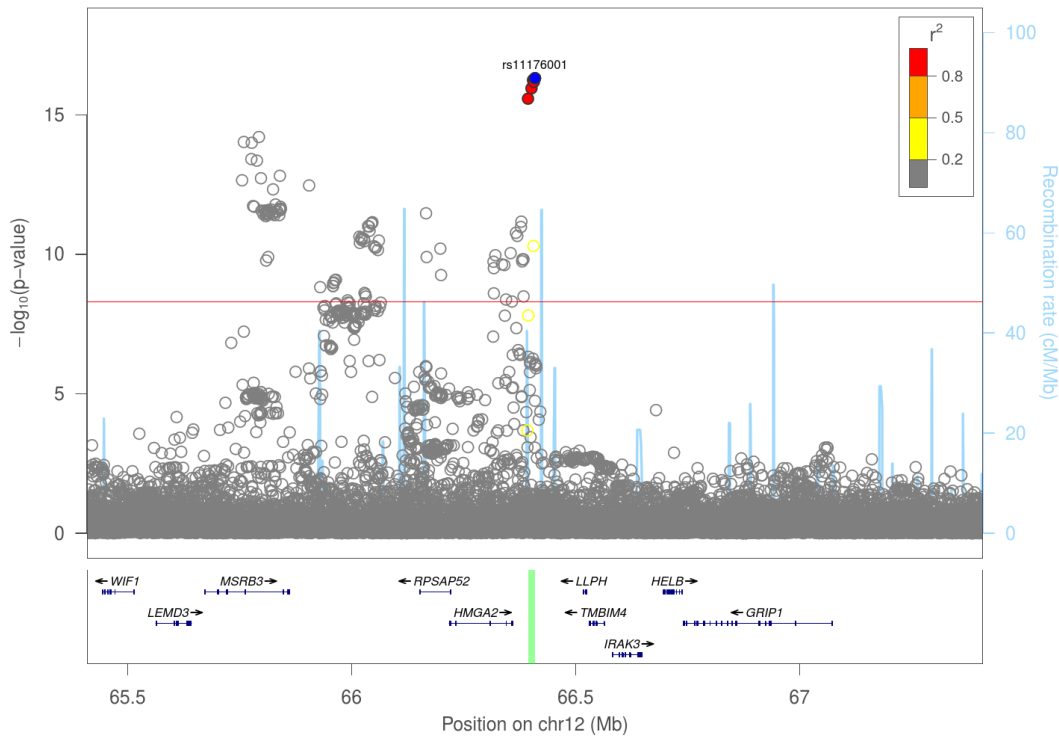
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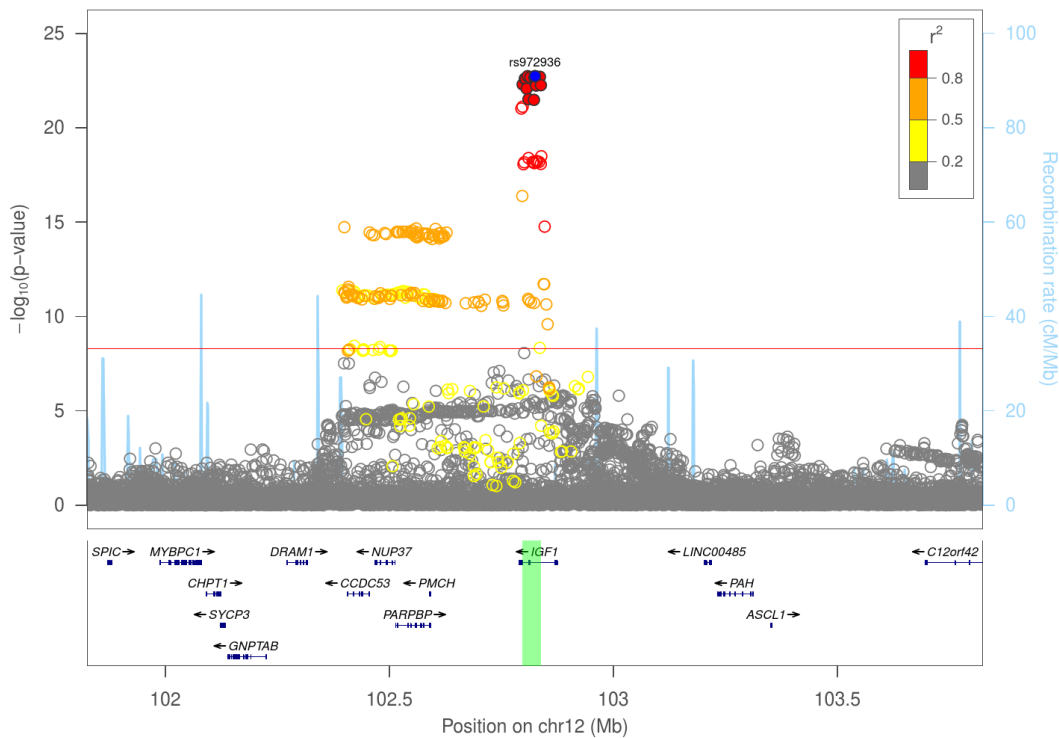
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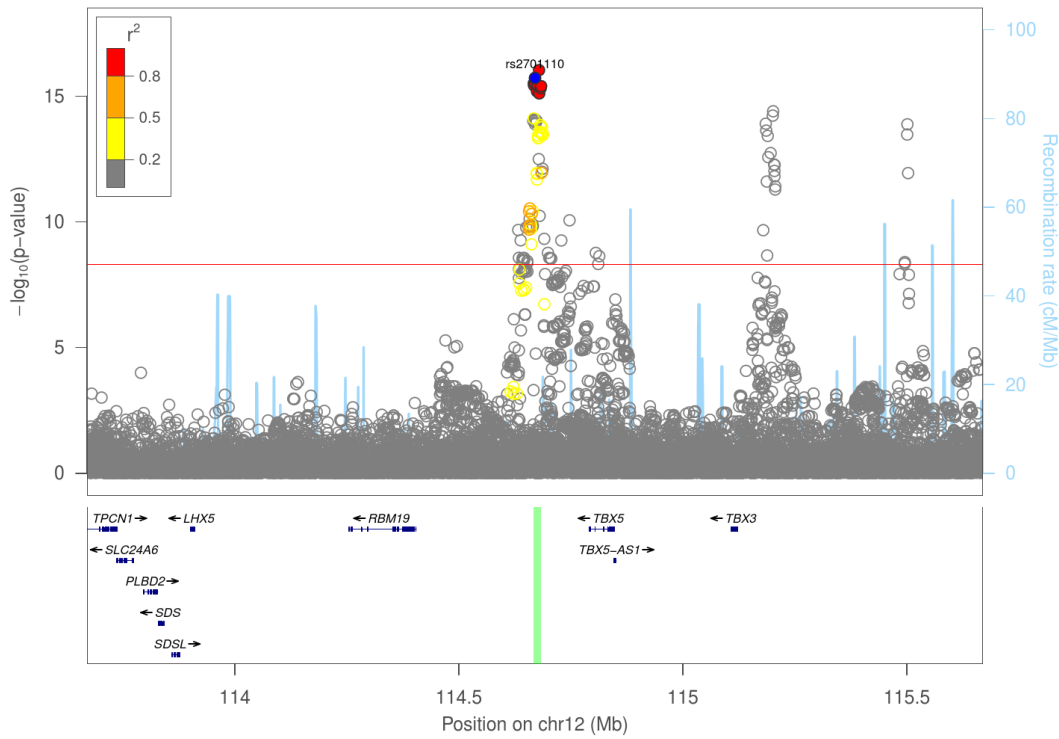
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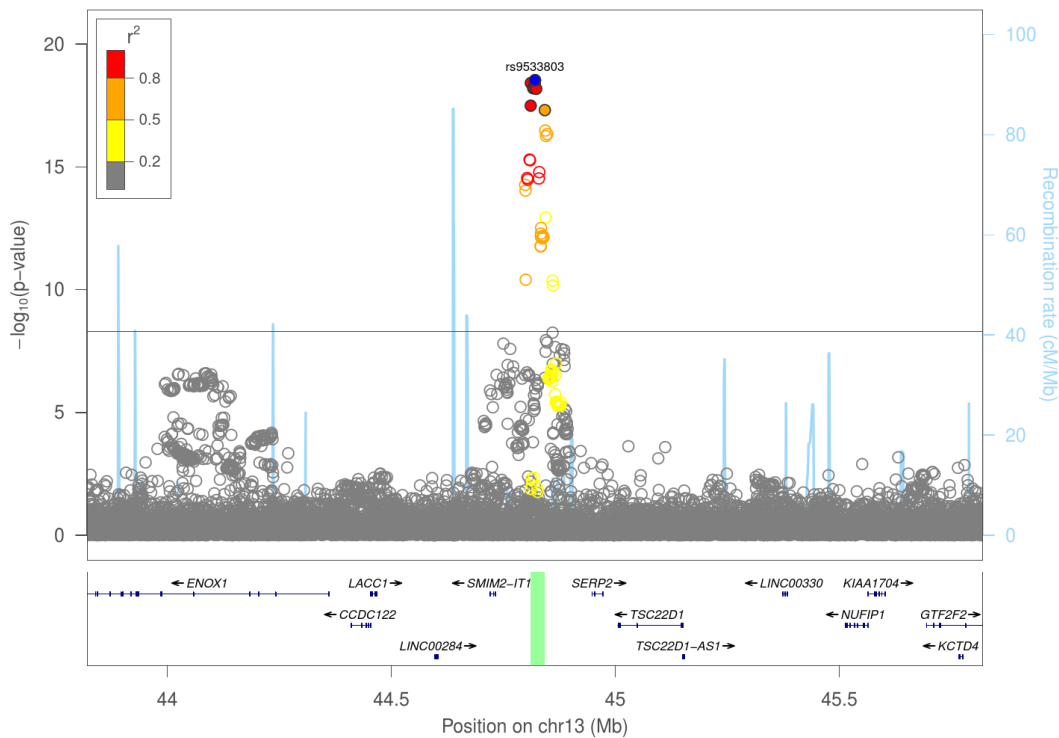
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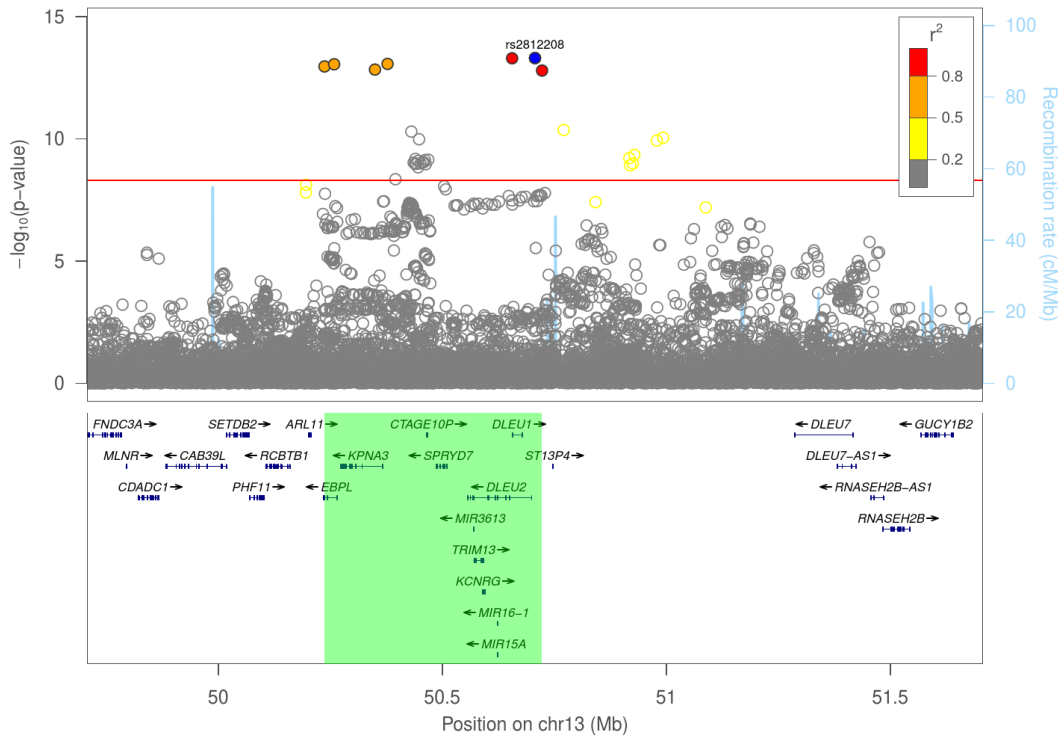
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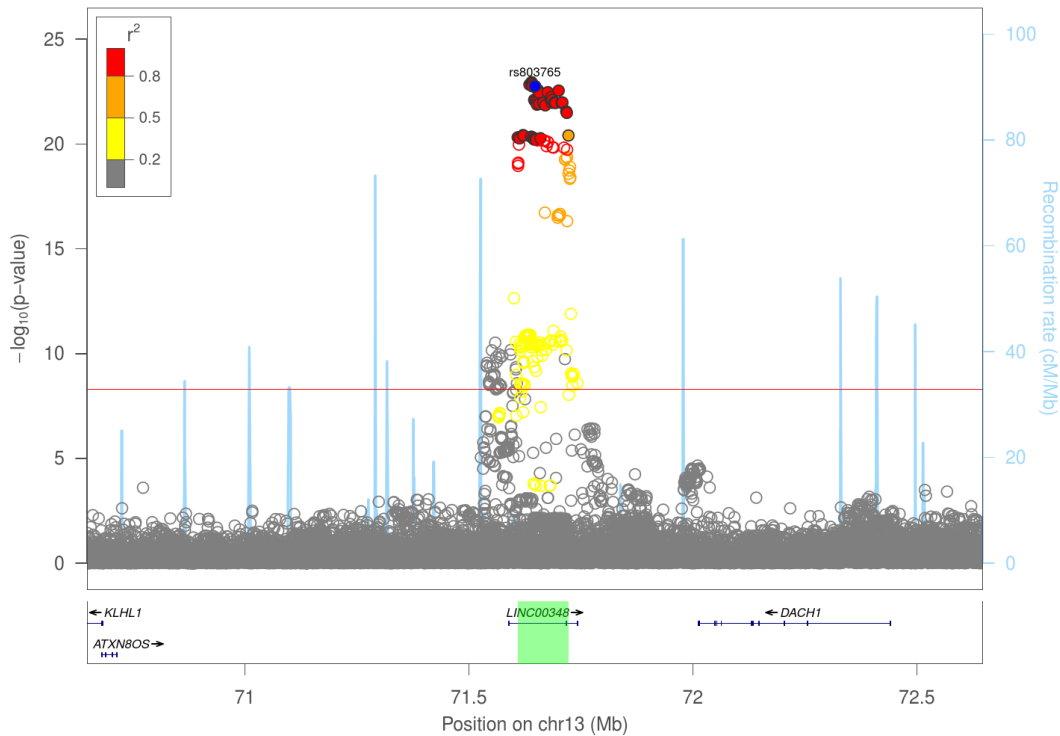
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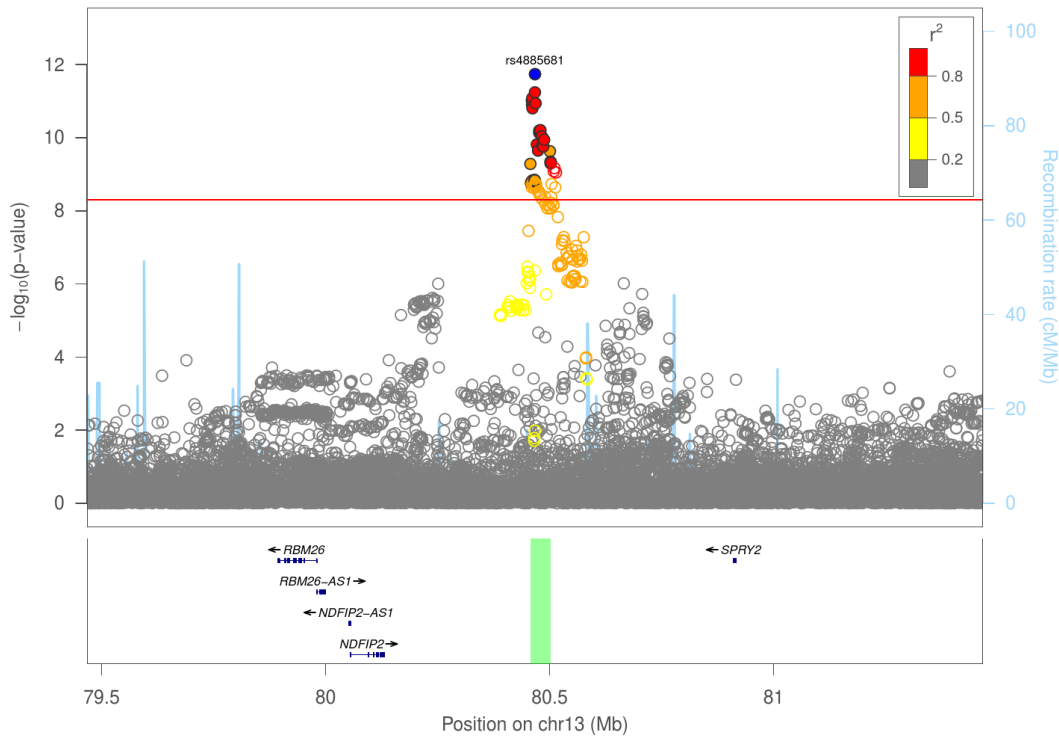
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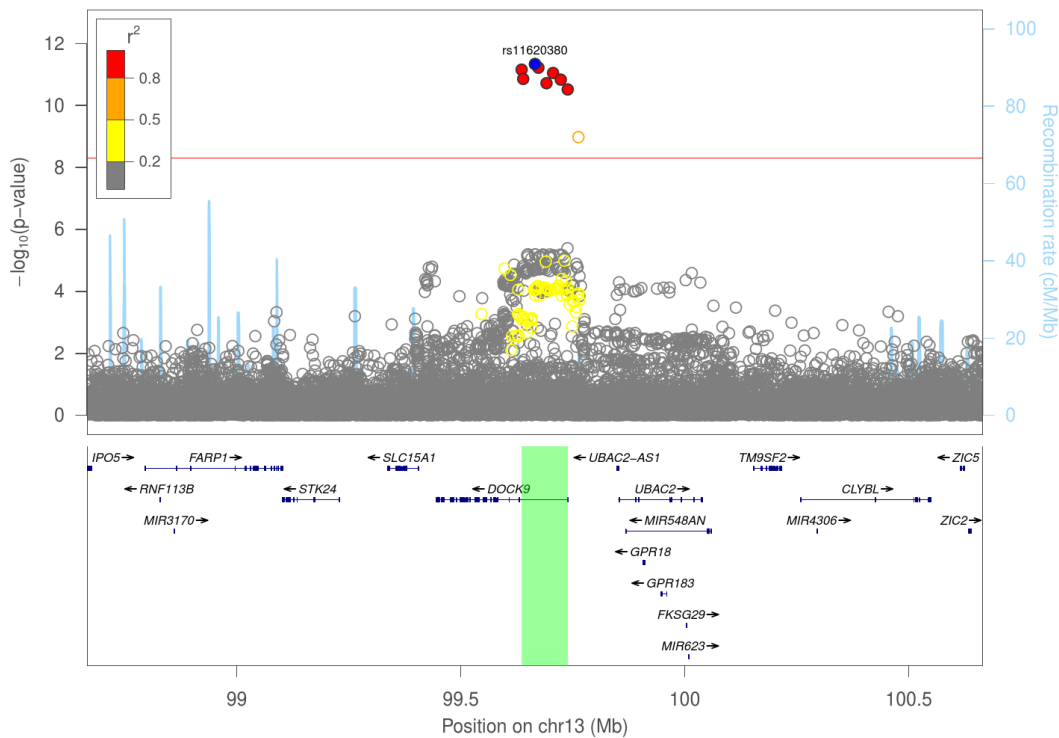
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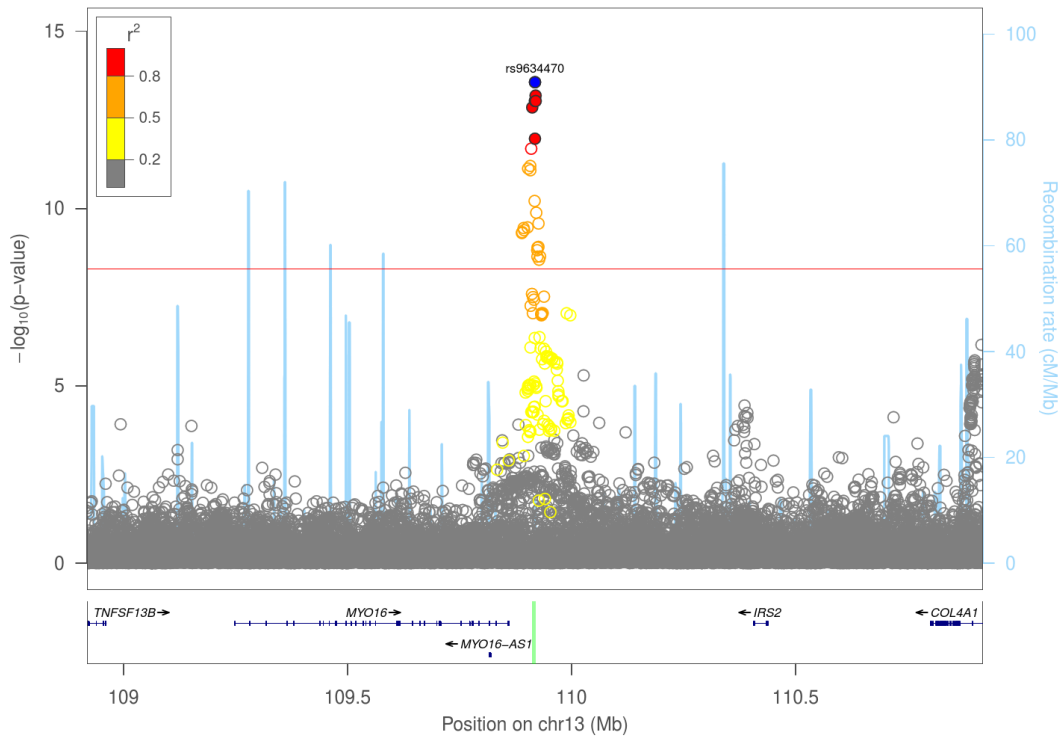
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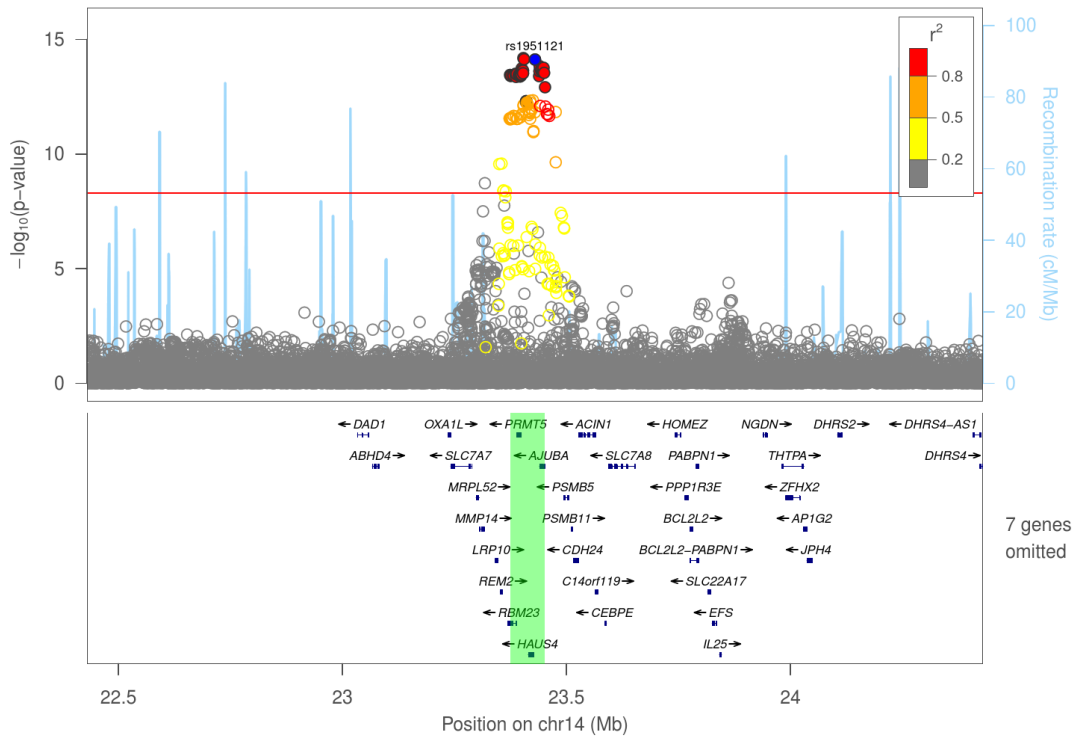
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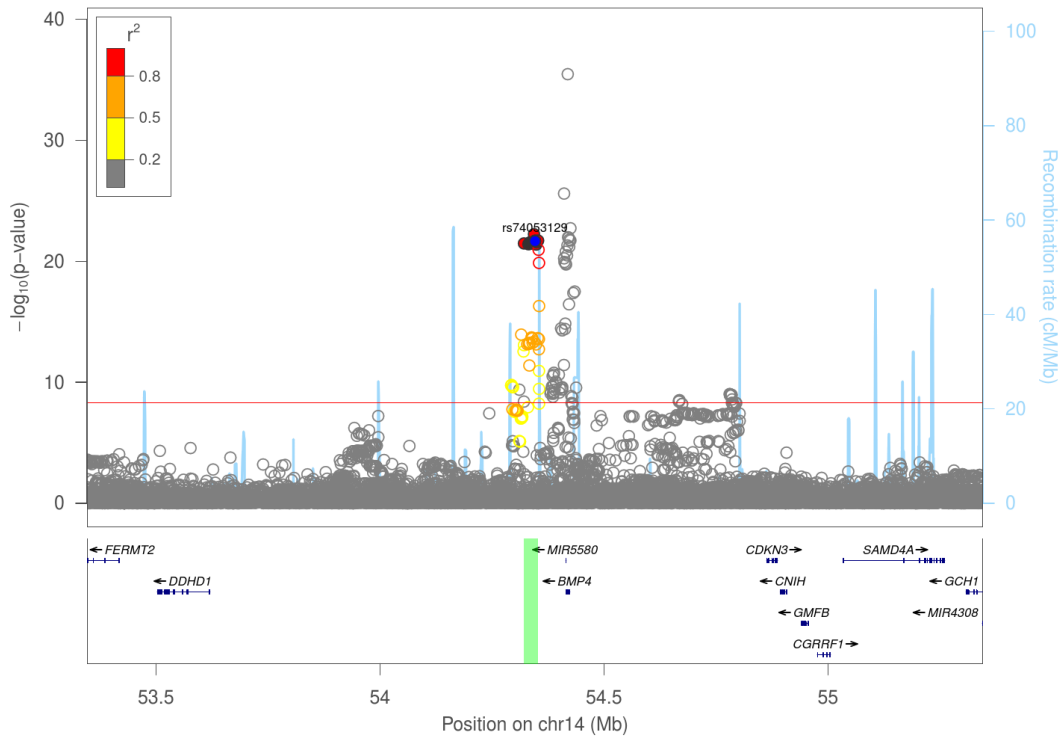
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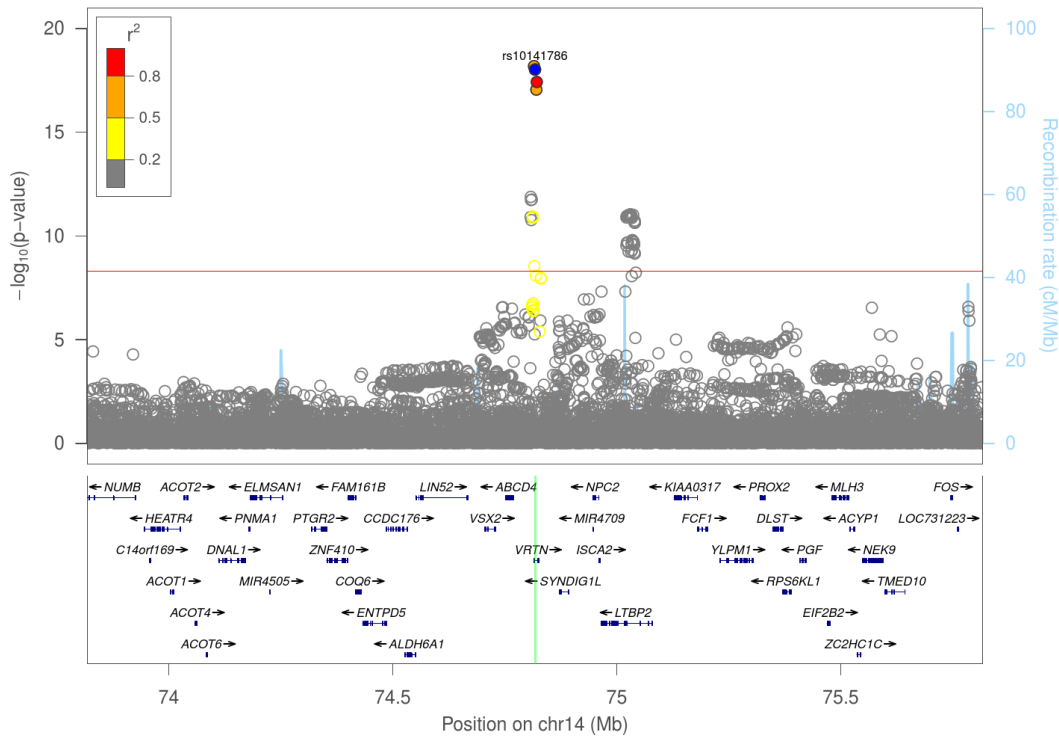
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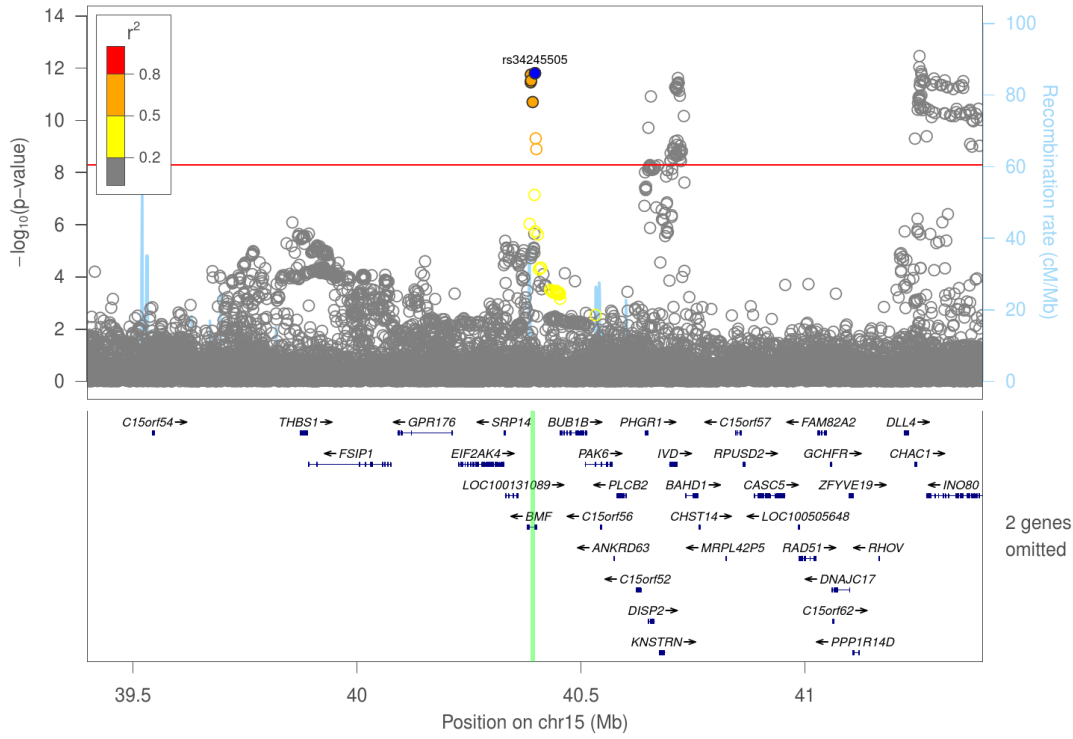
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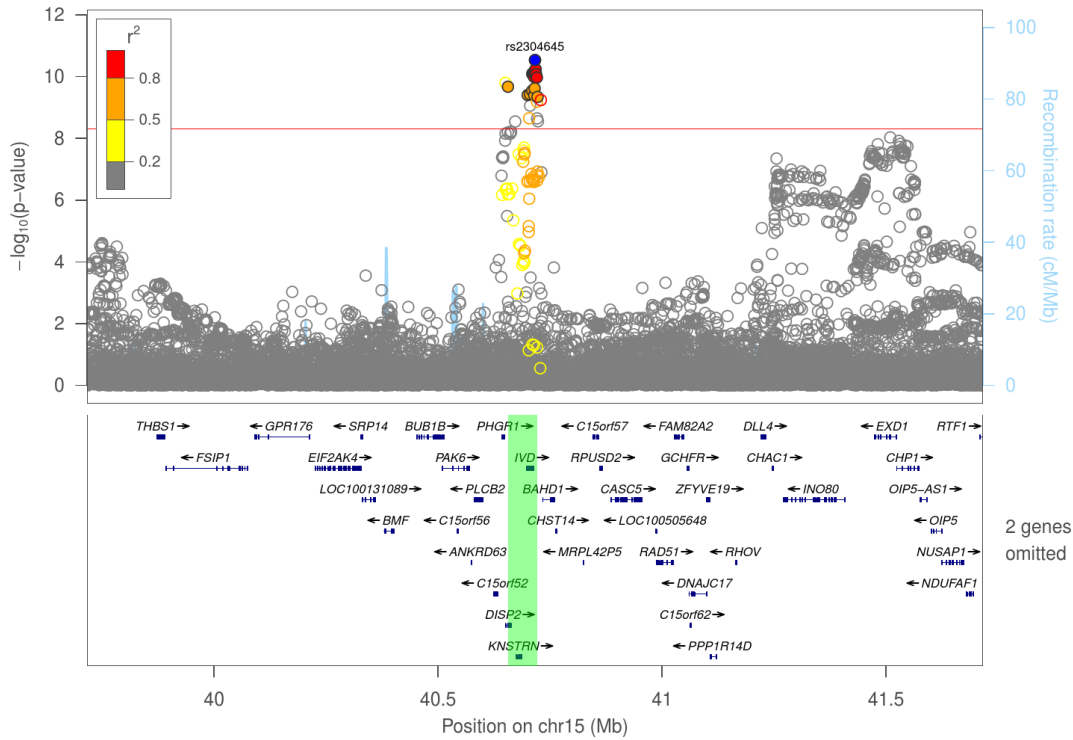
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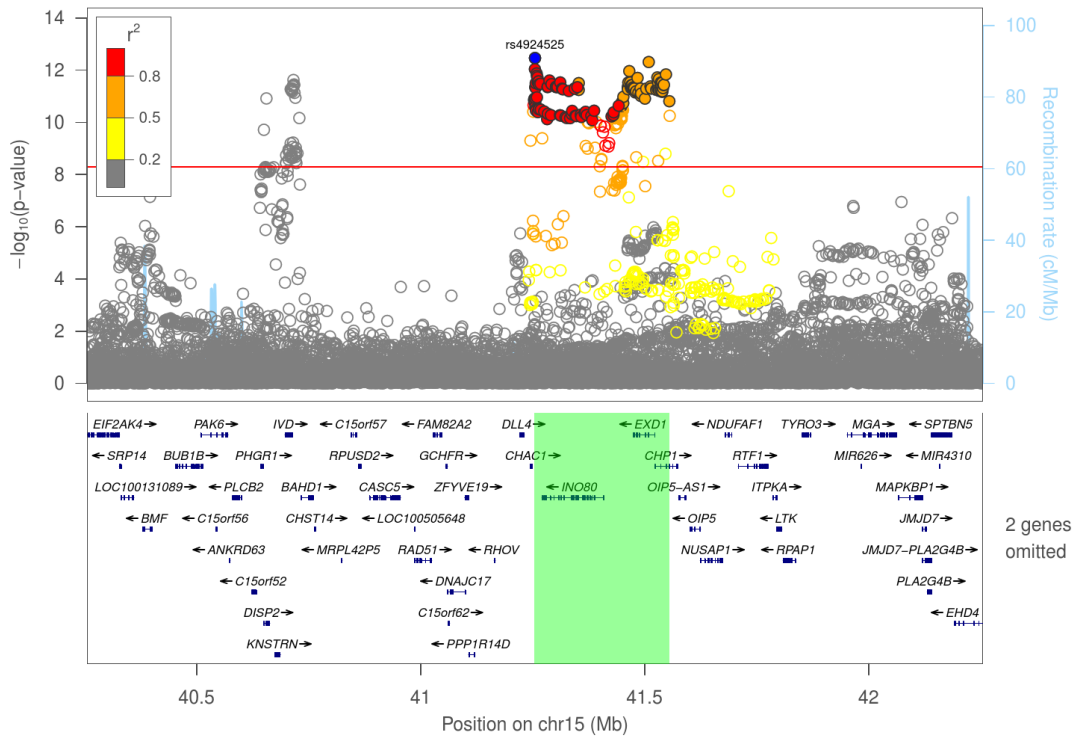
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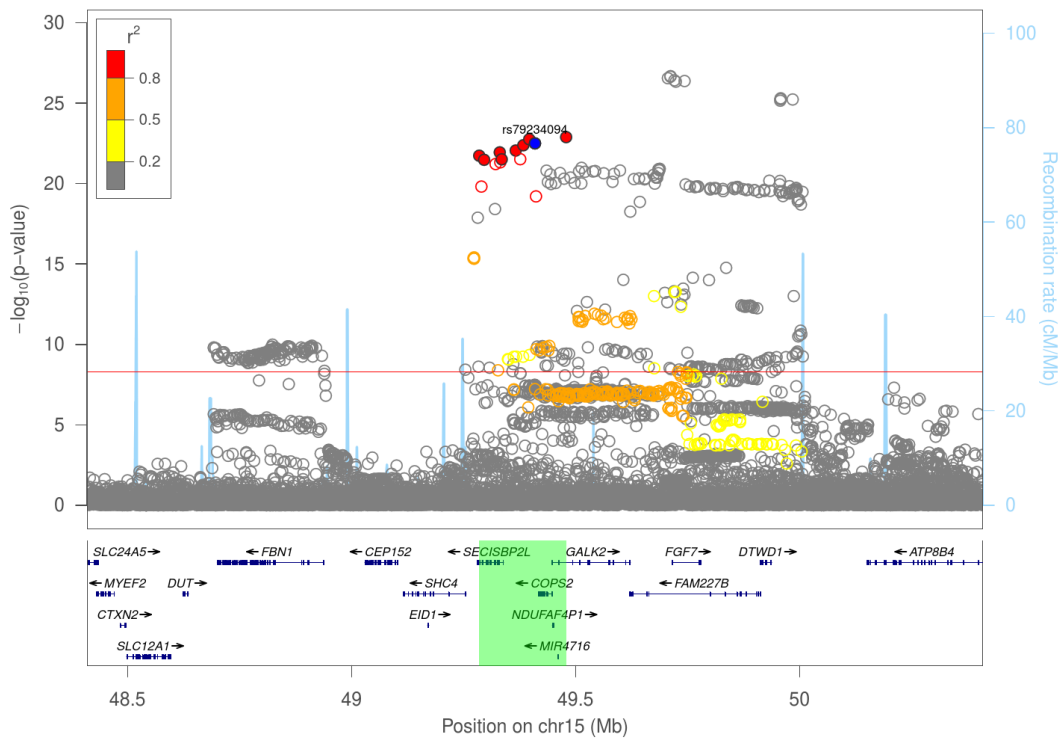
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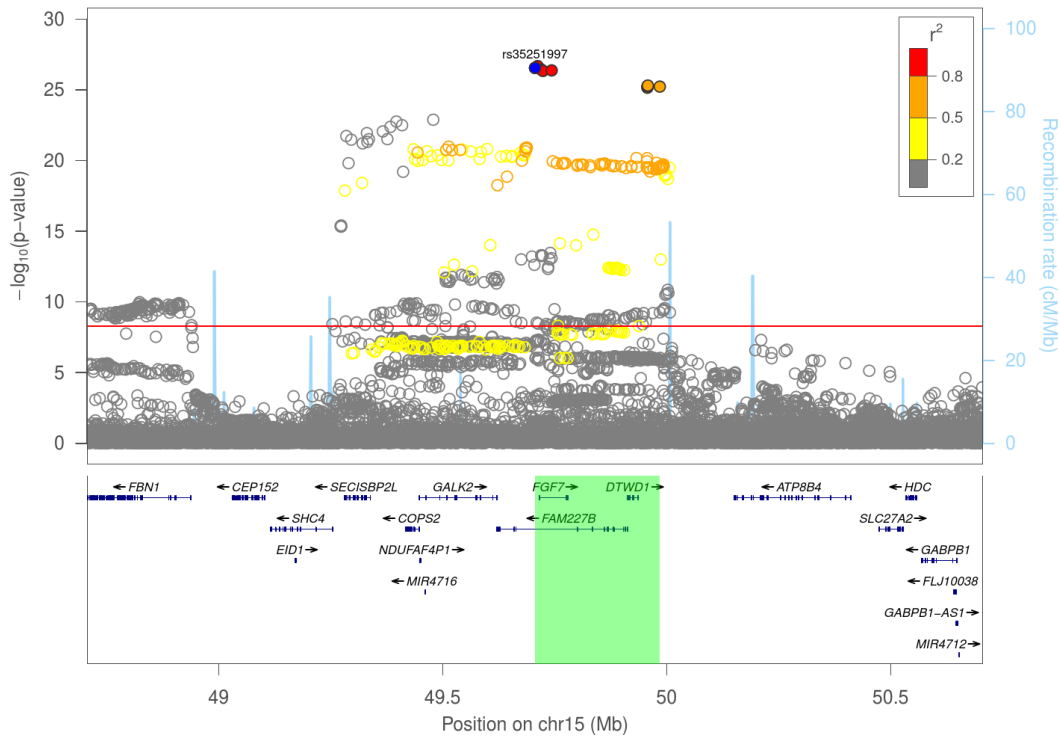
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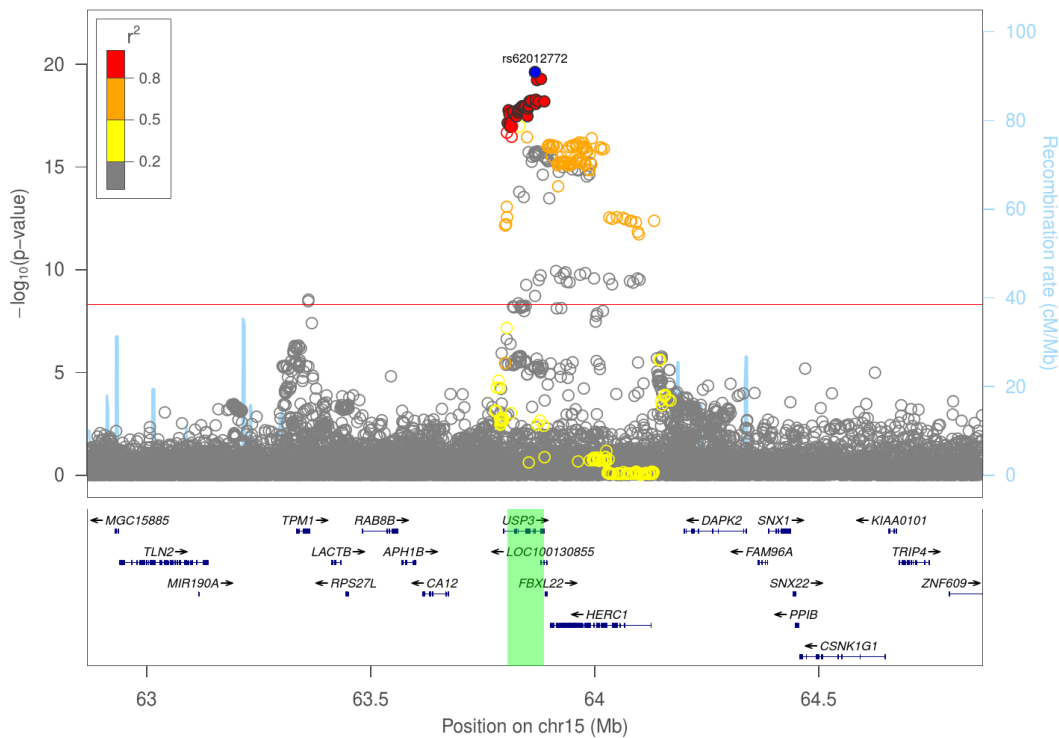
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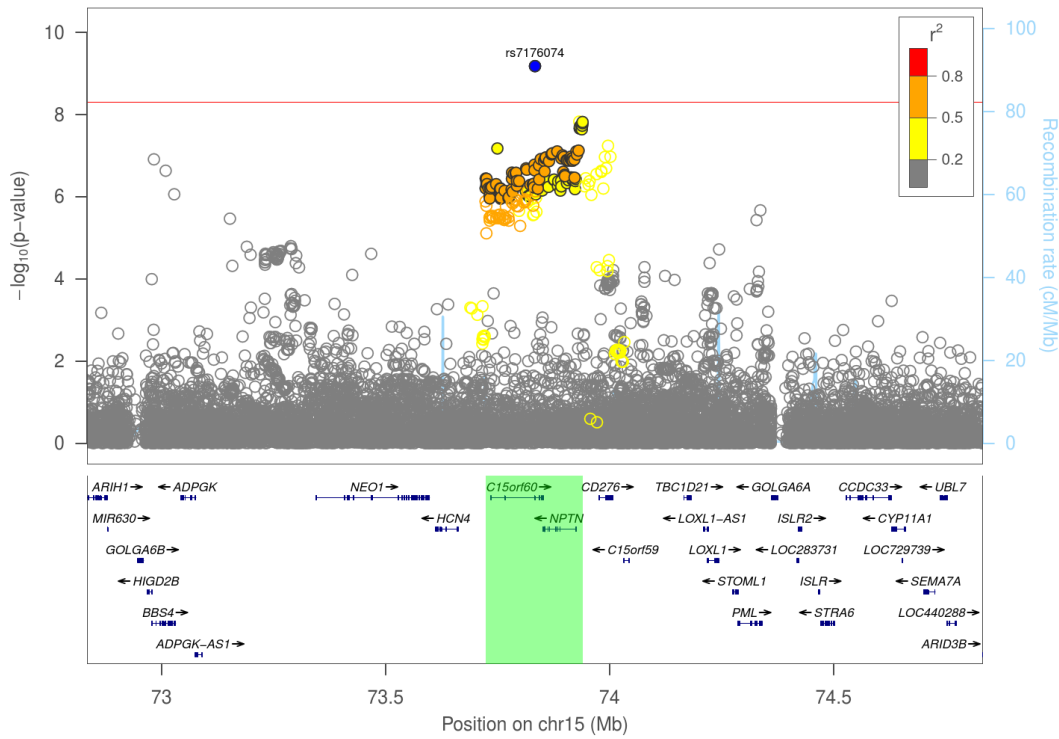
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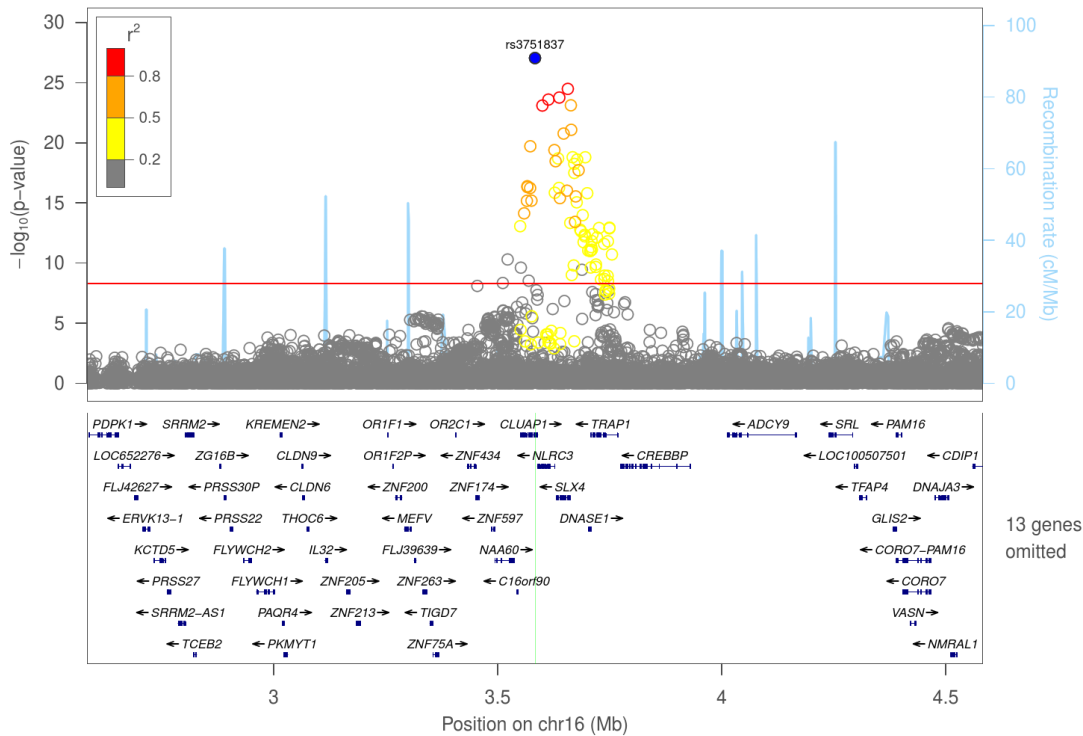
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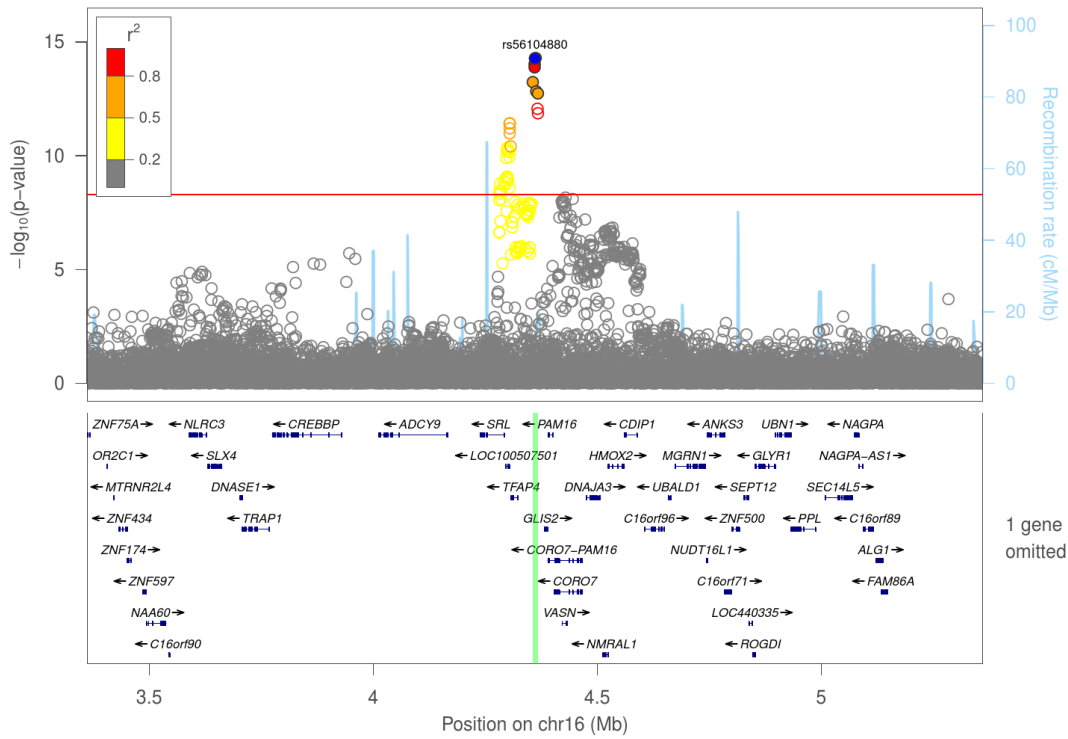
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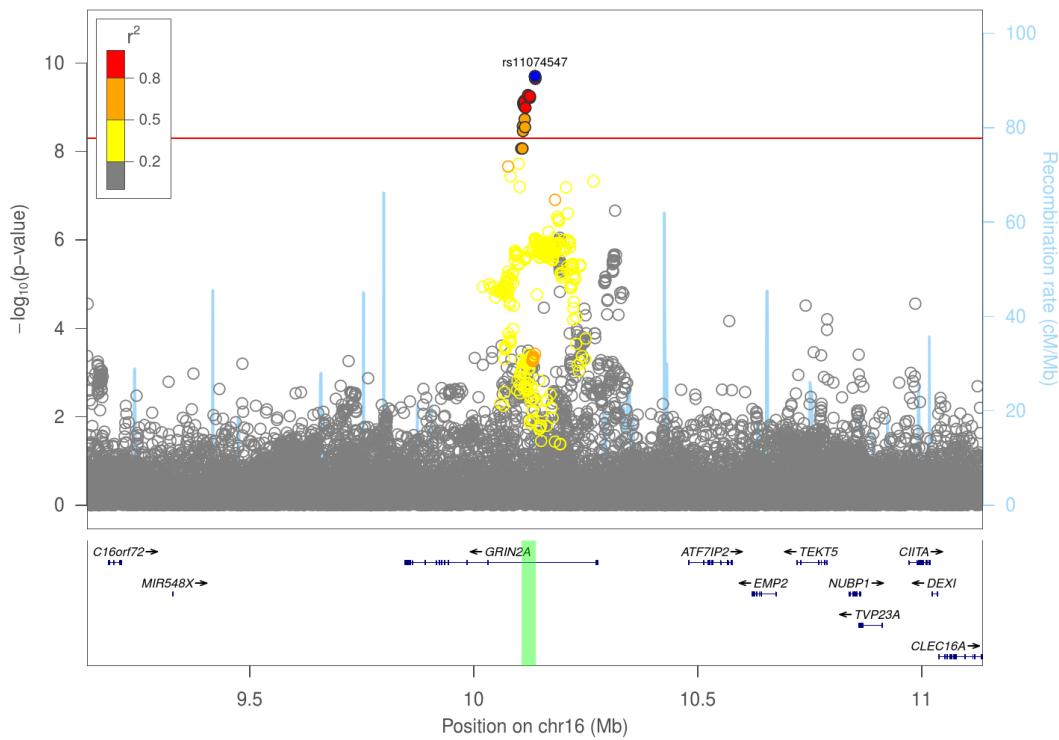
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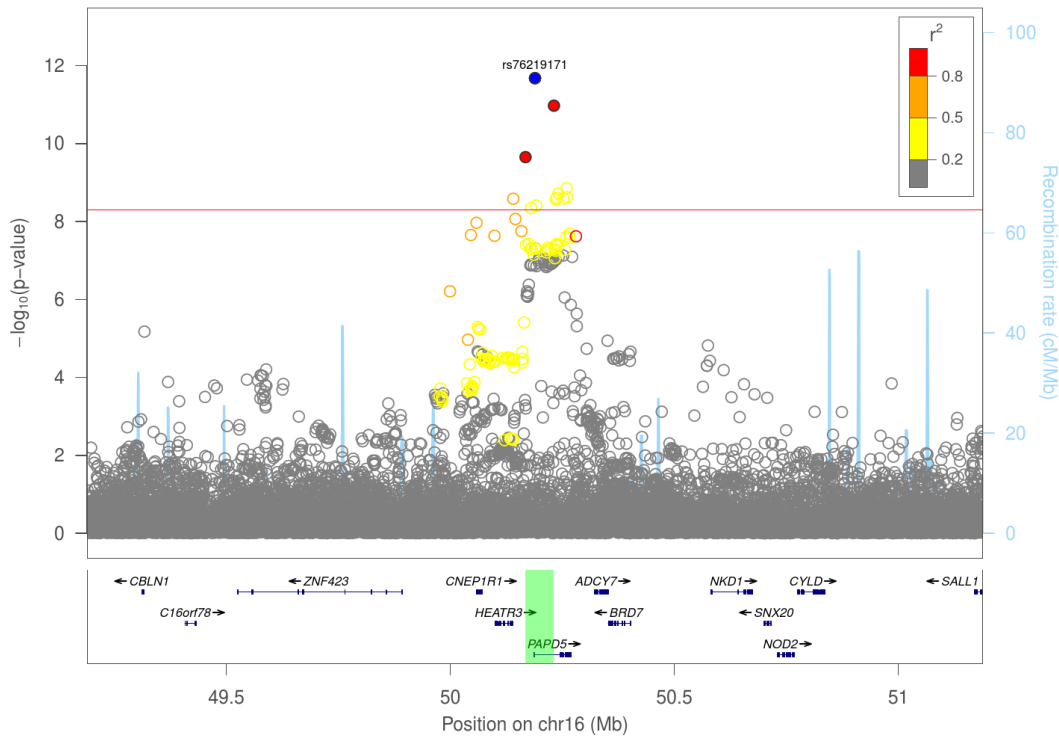
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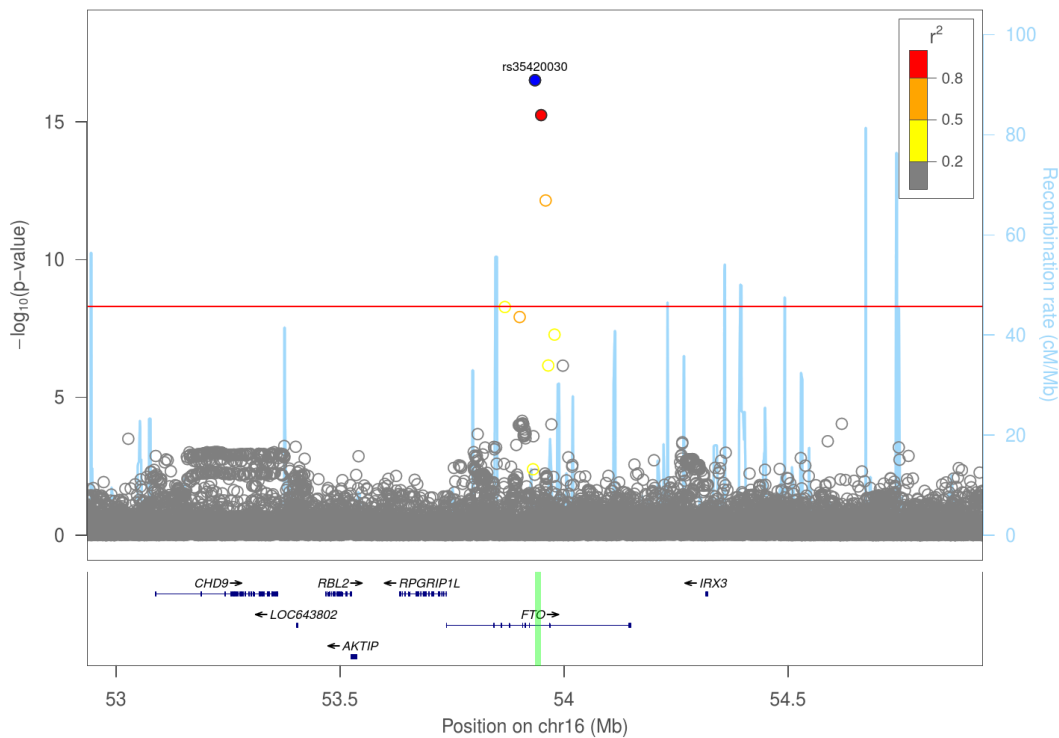
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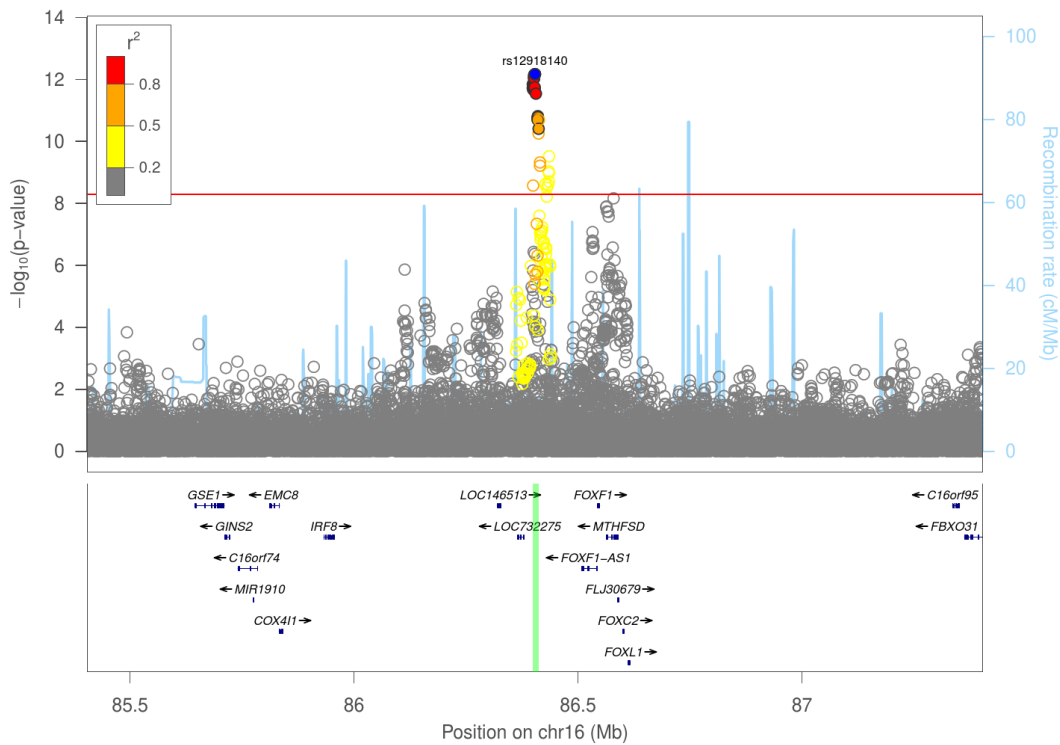
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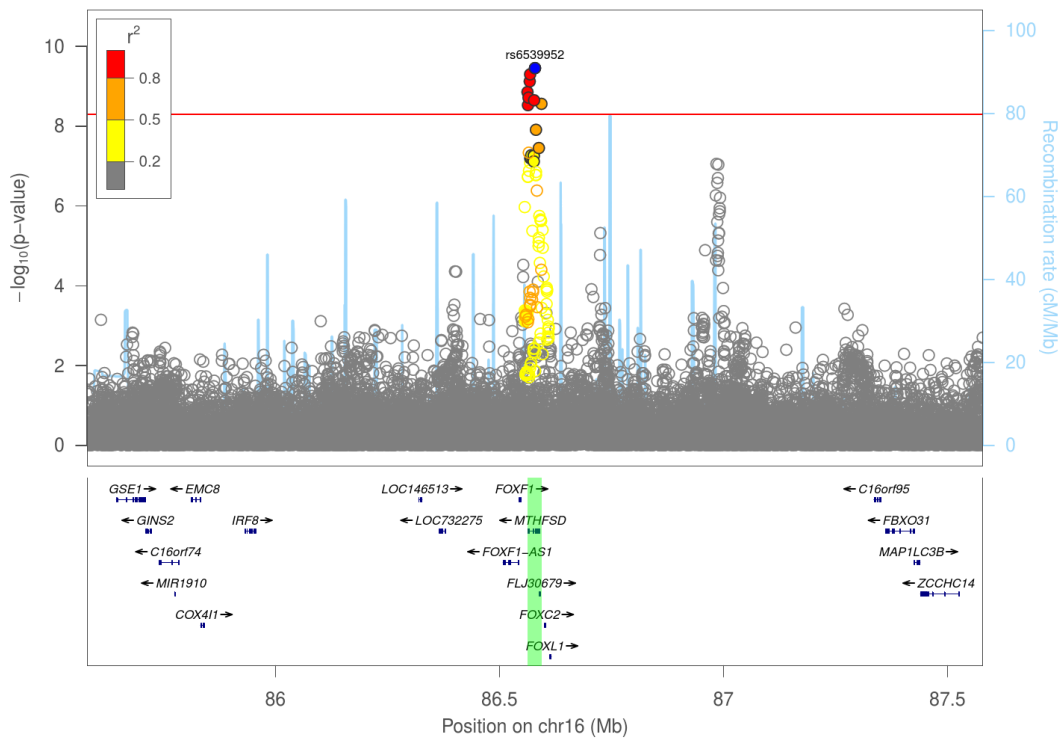
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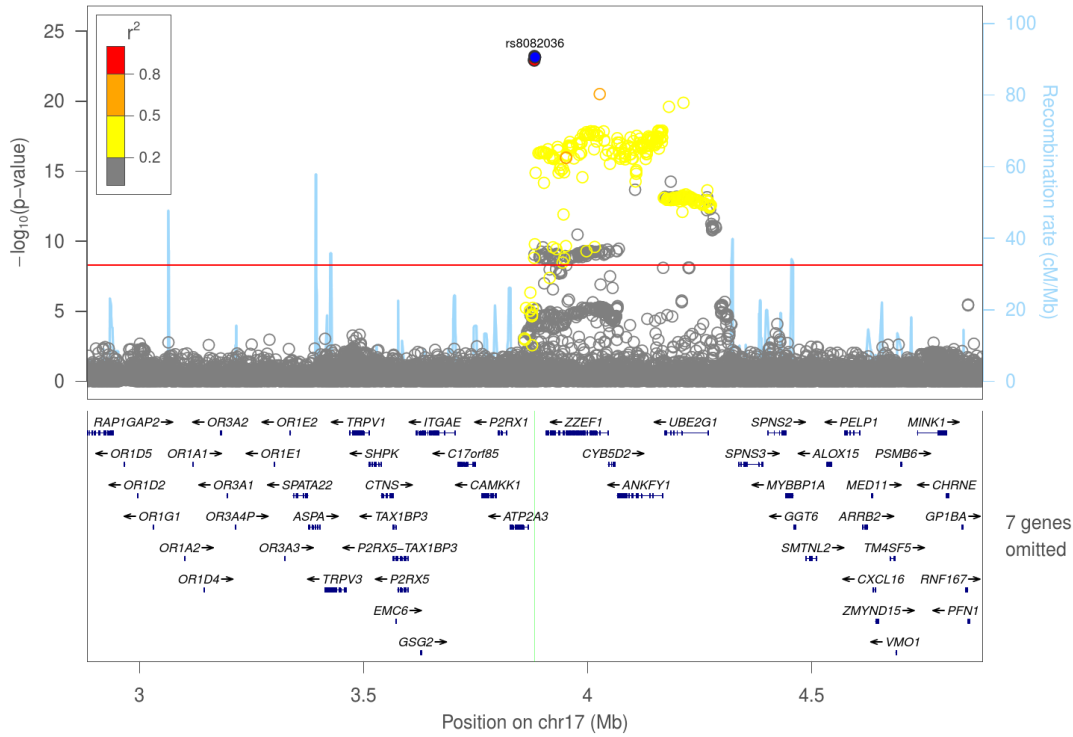
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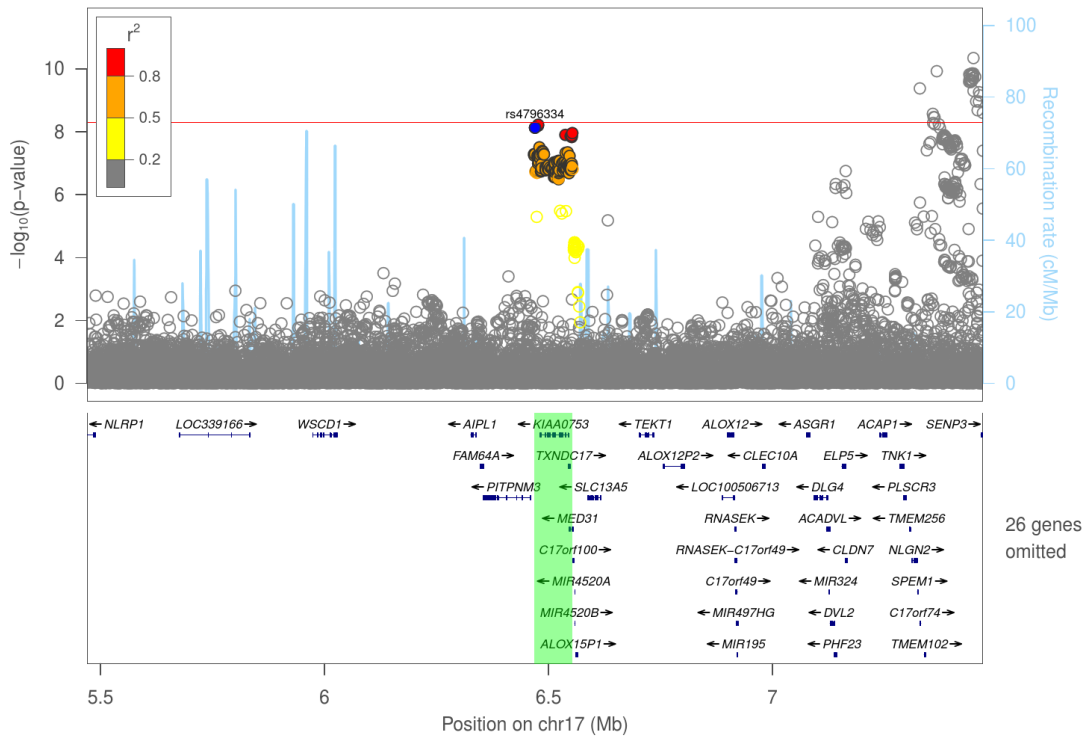
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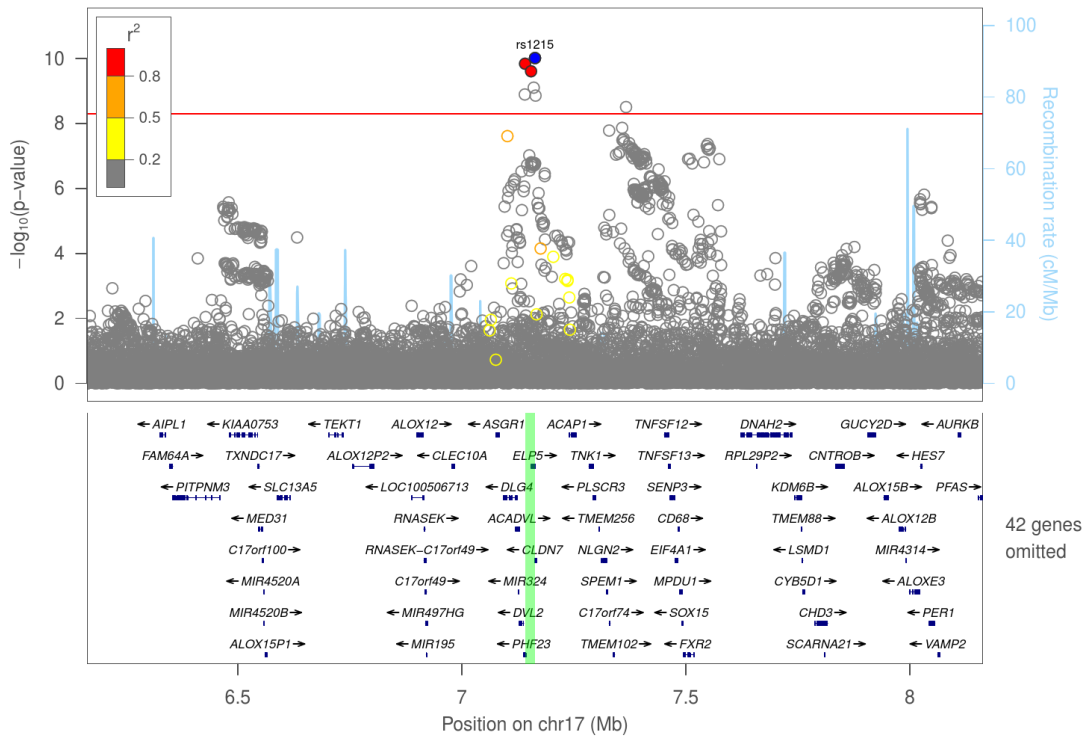
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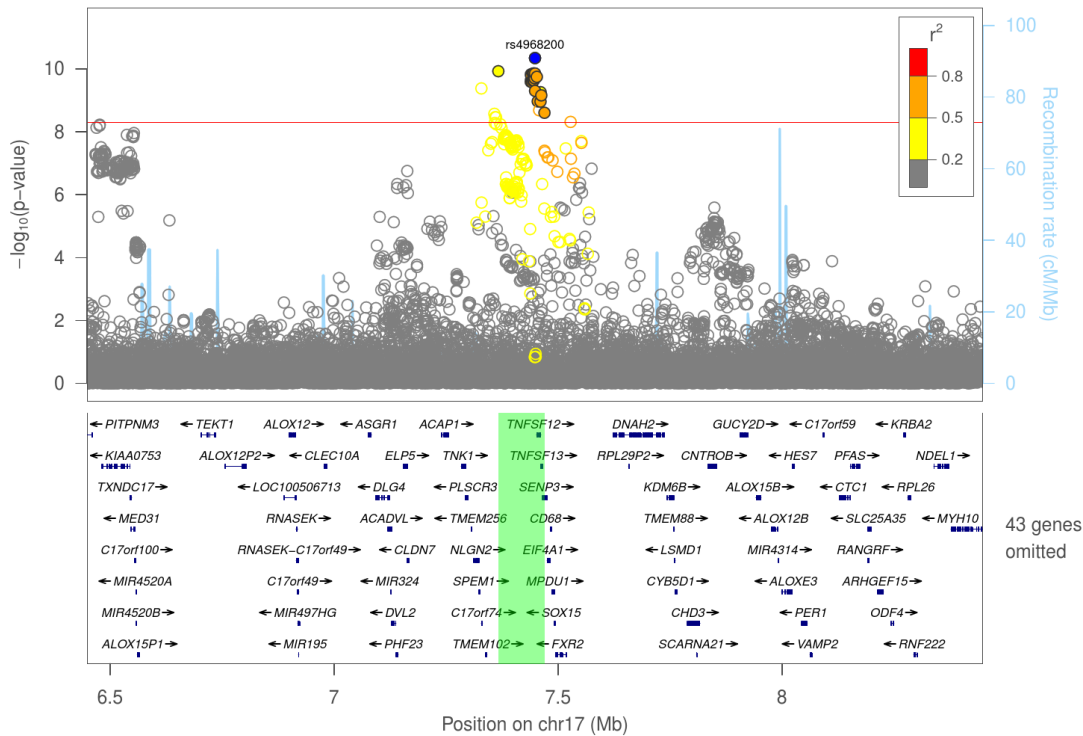
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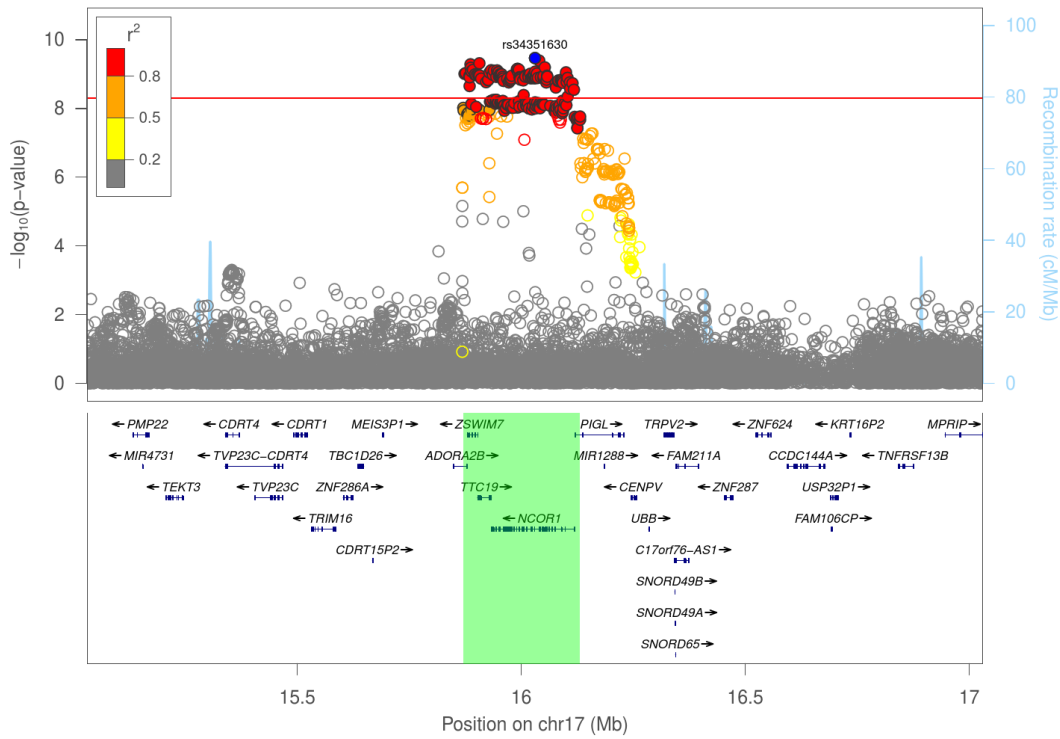
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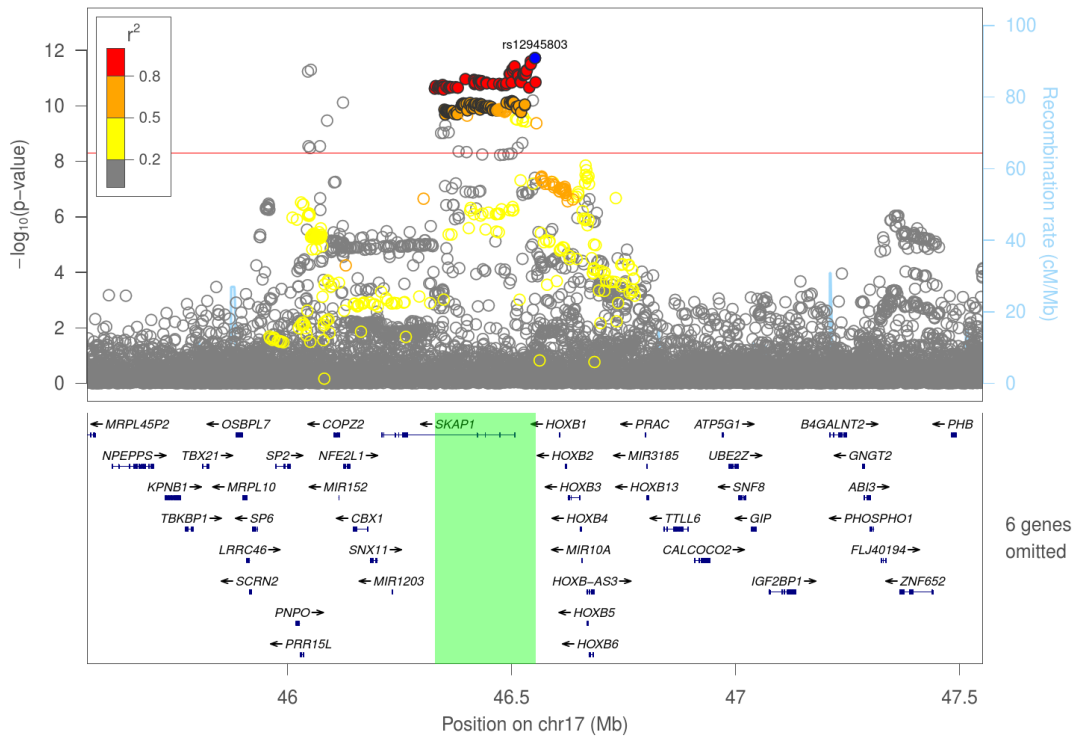
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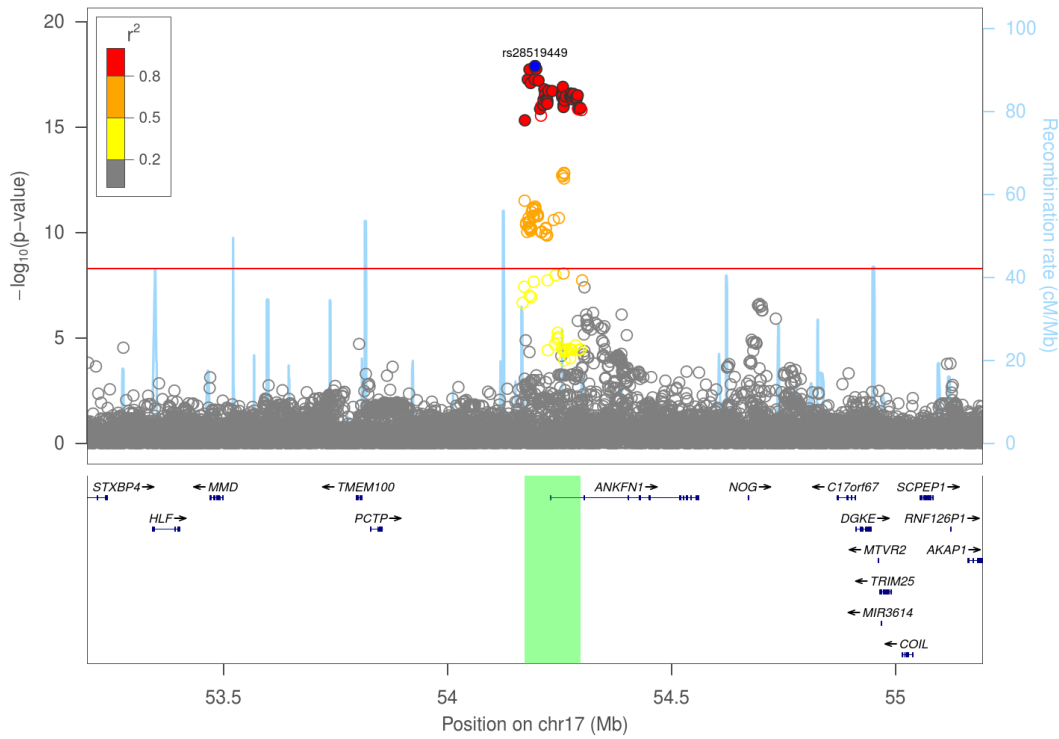
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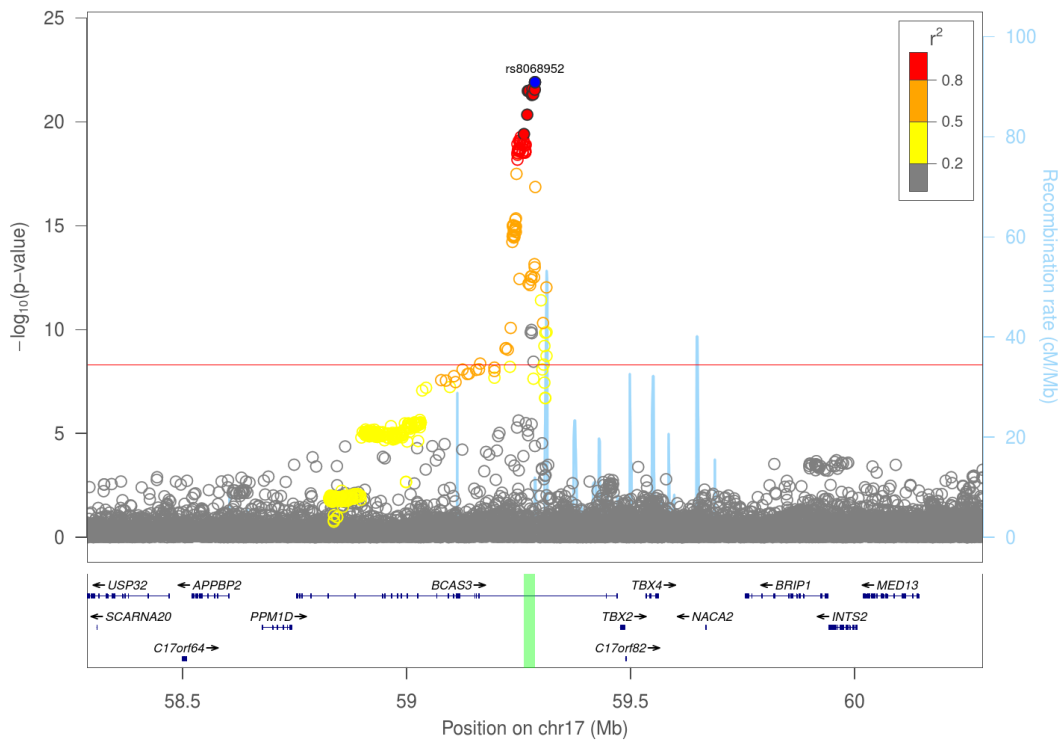
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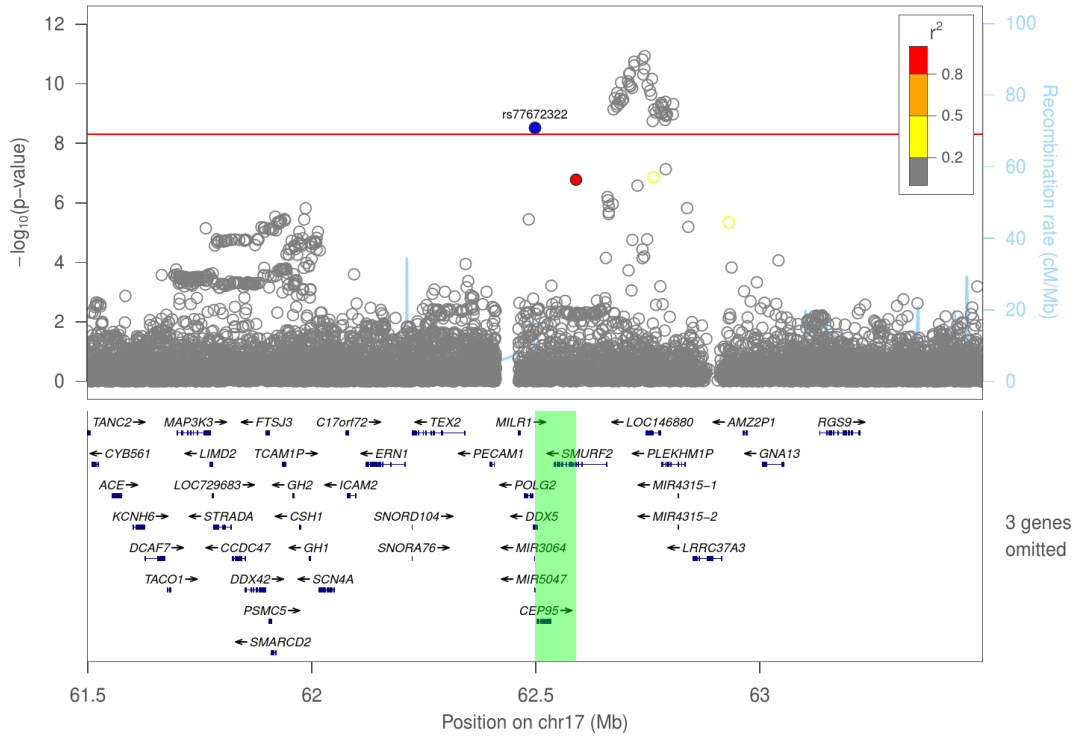
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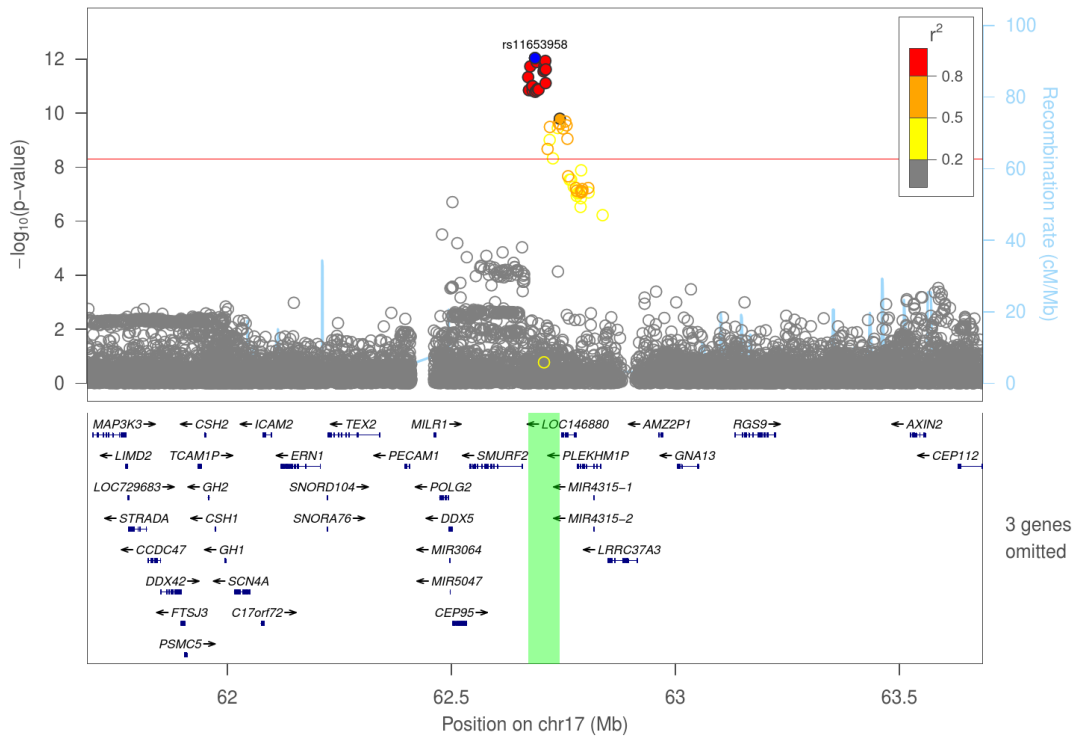
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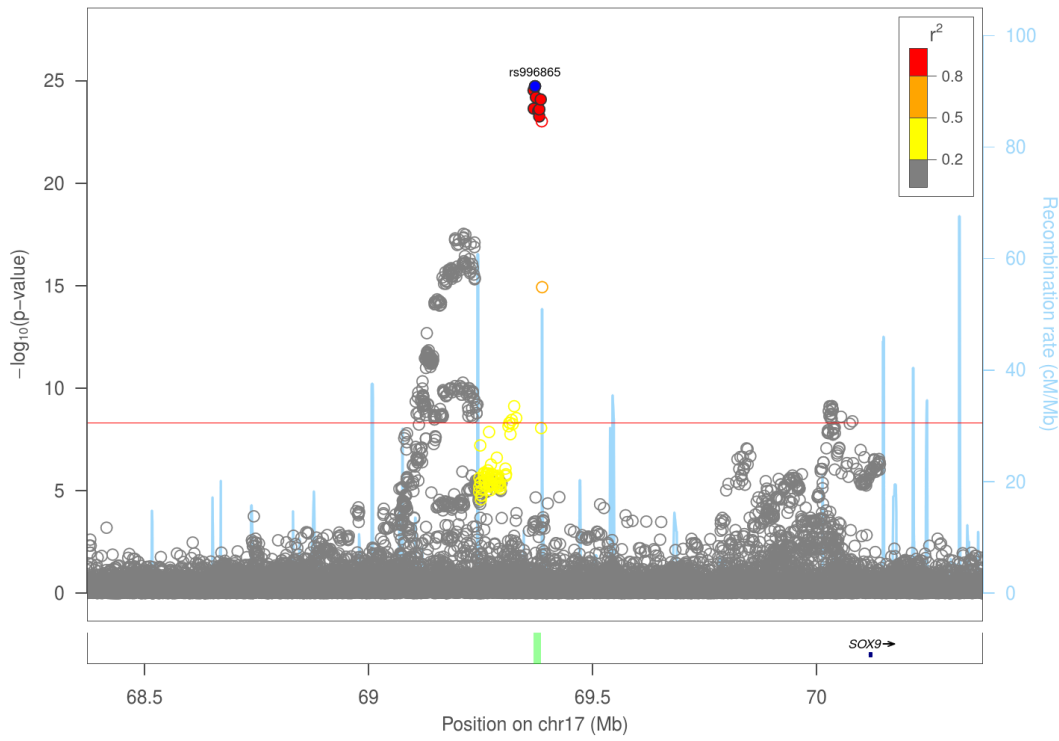
DDX5, FVC tier 2 rs77672322



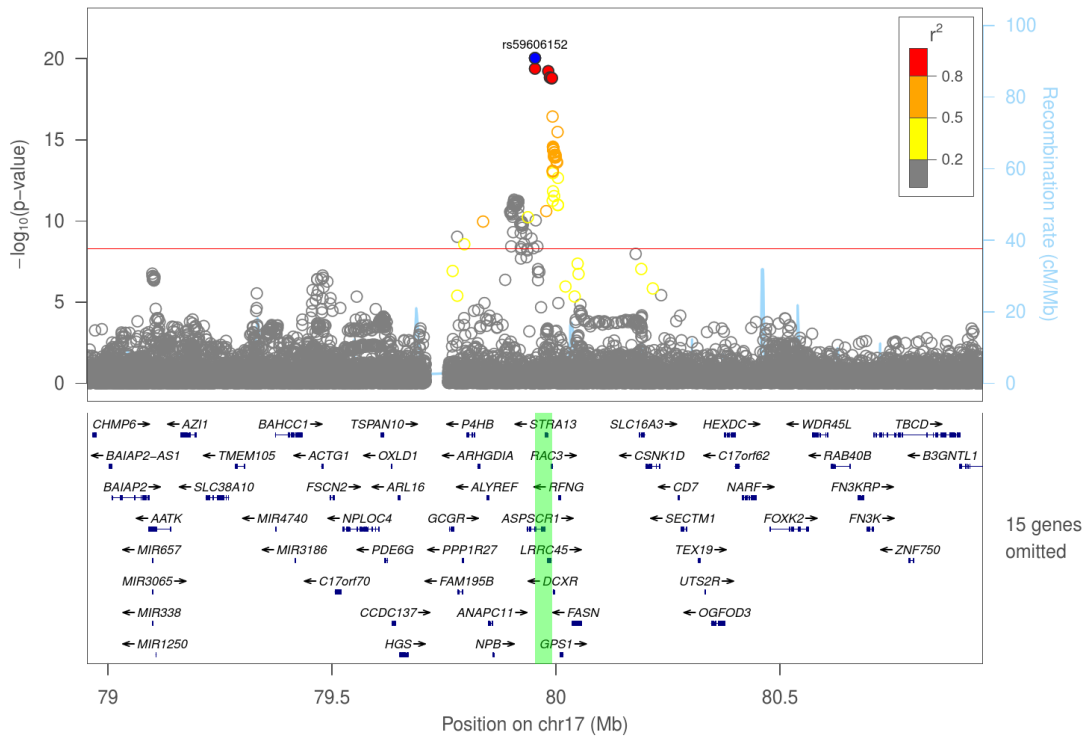
SMURF2, FEV₁/FVC tier 1 rs11653958



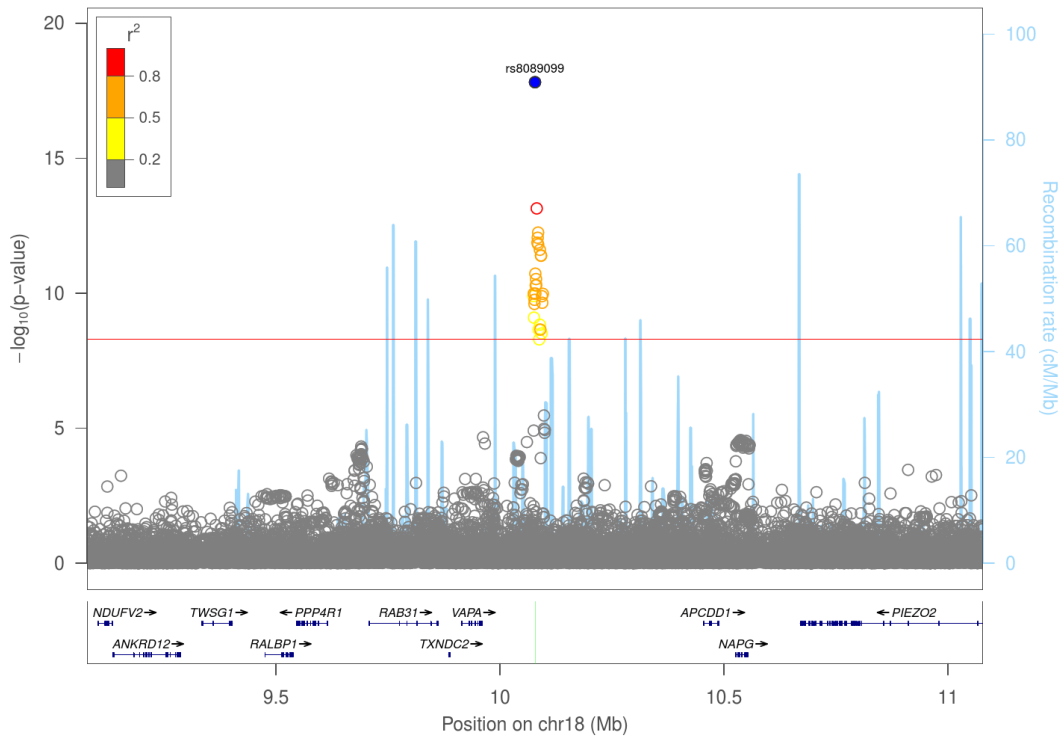
CASC17, FEV₁/FVC tier 1 rs996865



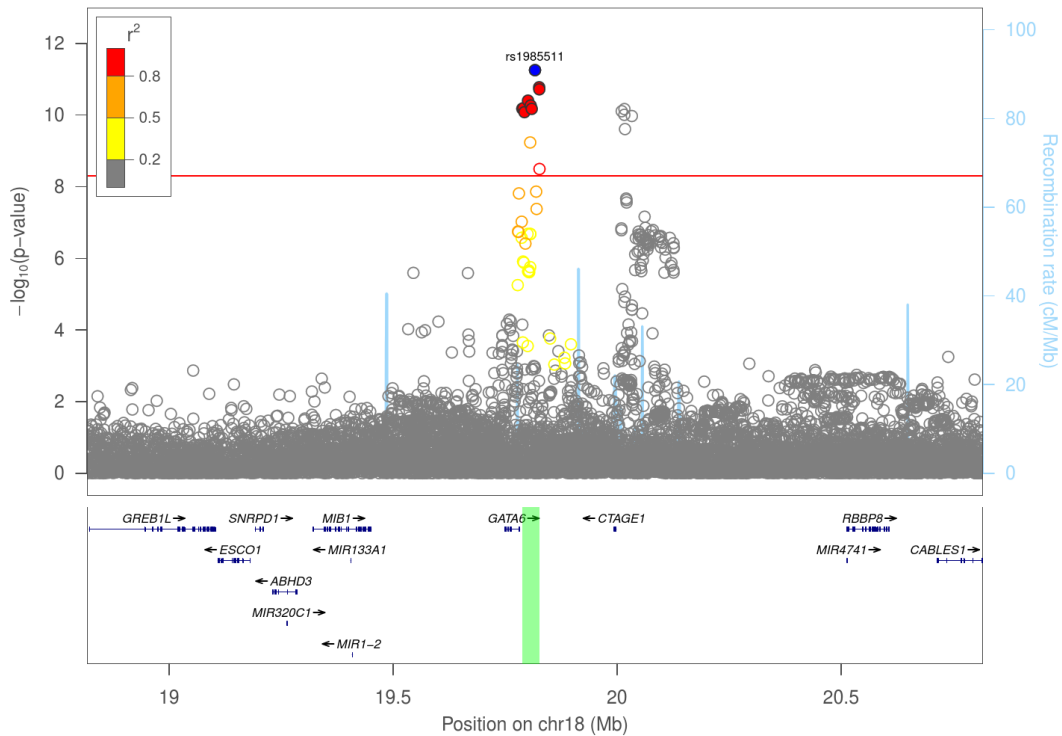
ASPCR1, FVC tier 1 rs59606152



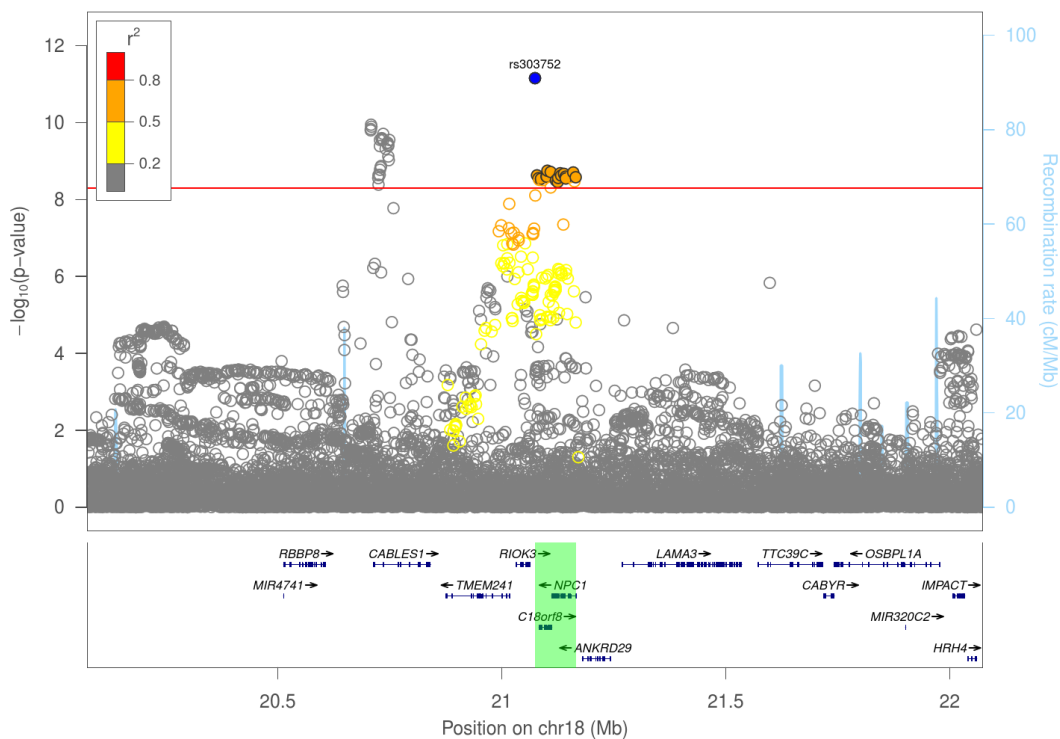
VAPA, FEV₁/FVC tier 1 rs8089099



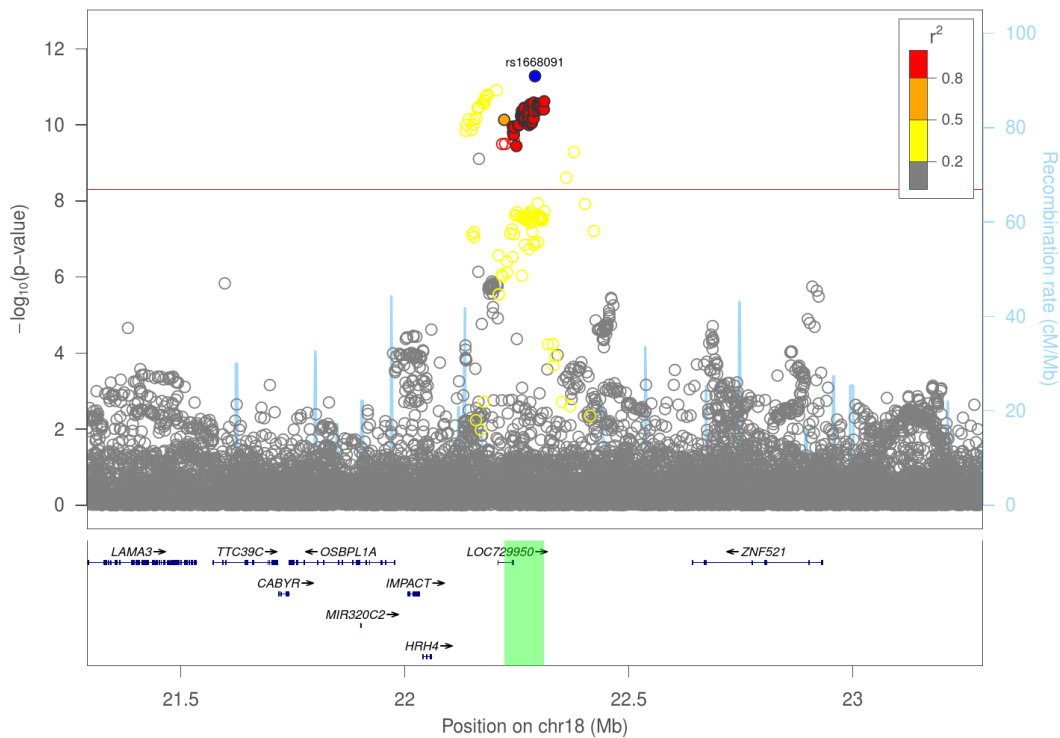
GATA6, FEV₁/FVC tier 2 rs1985511



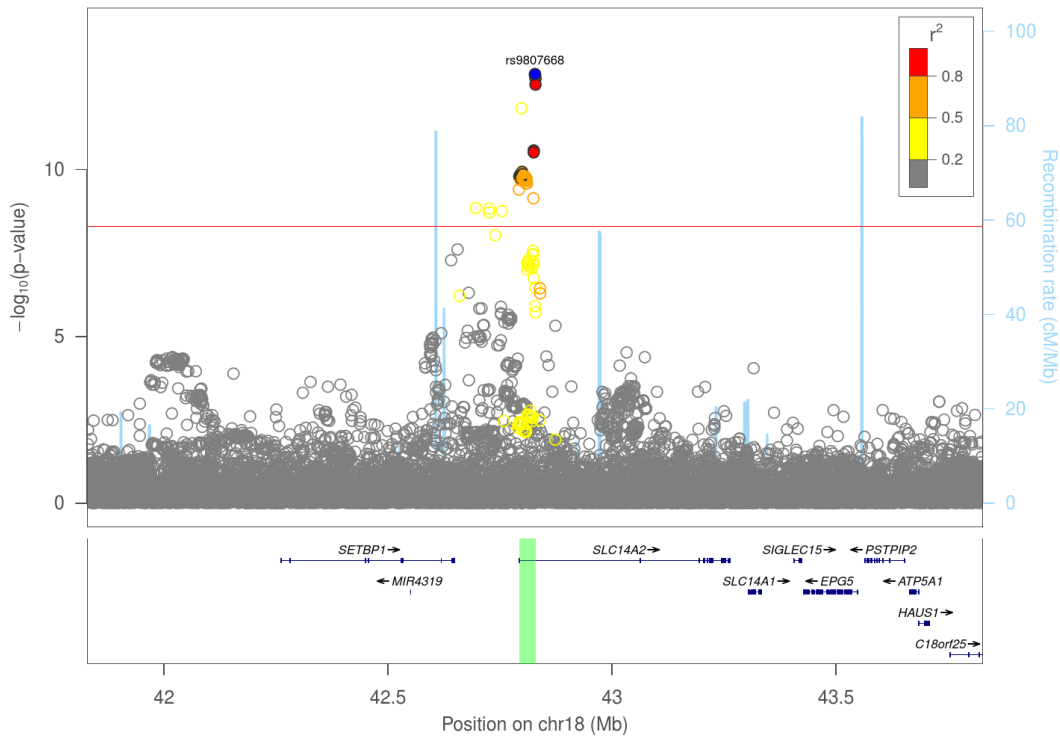
C18orf8, FVC tier 1 rs303752



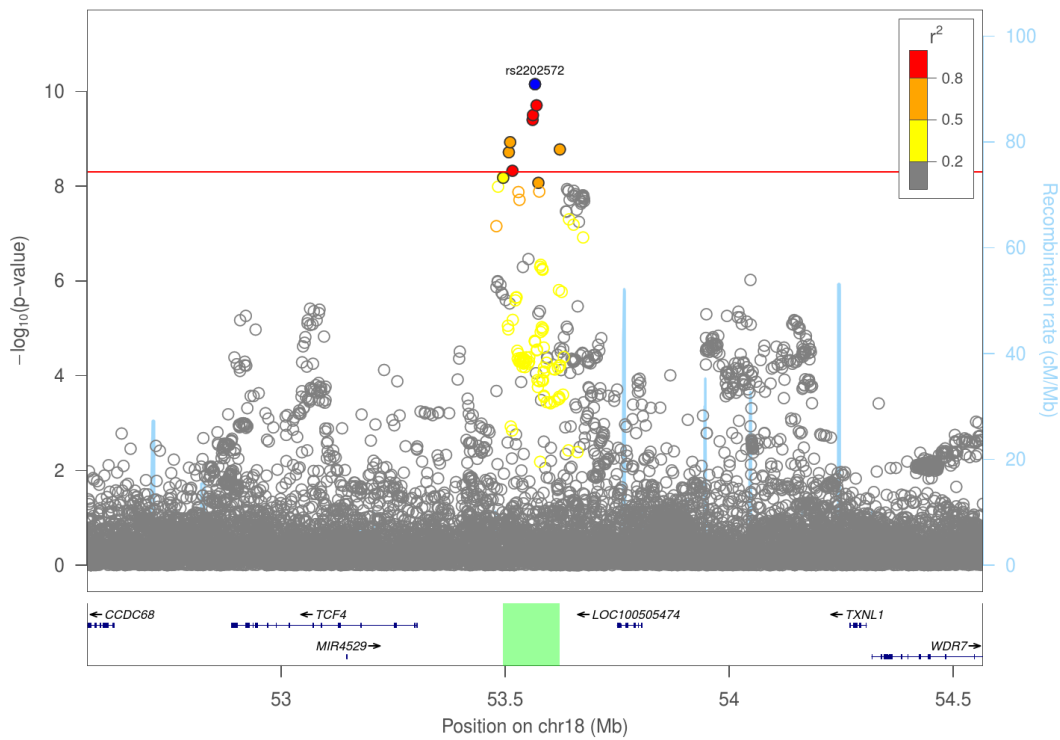
LOC729950, FVC tier 1 rs1668091



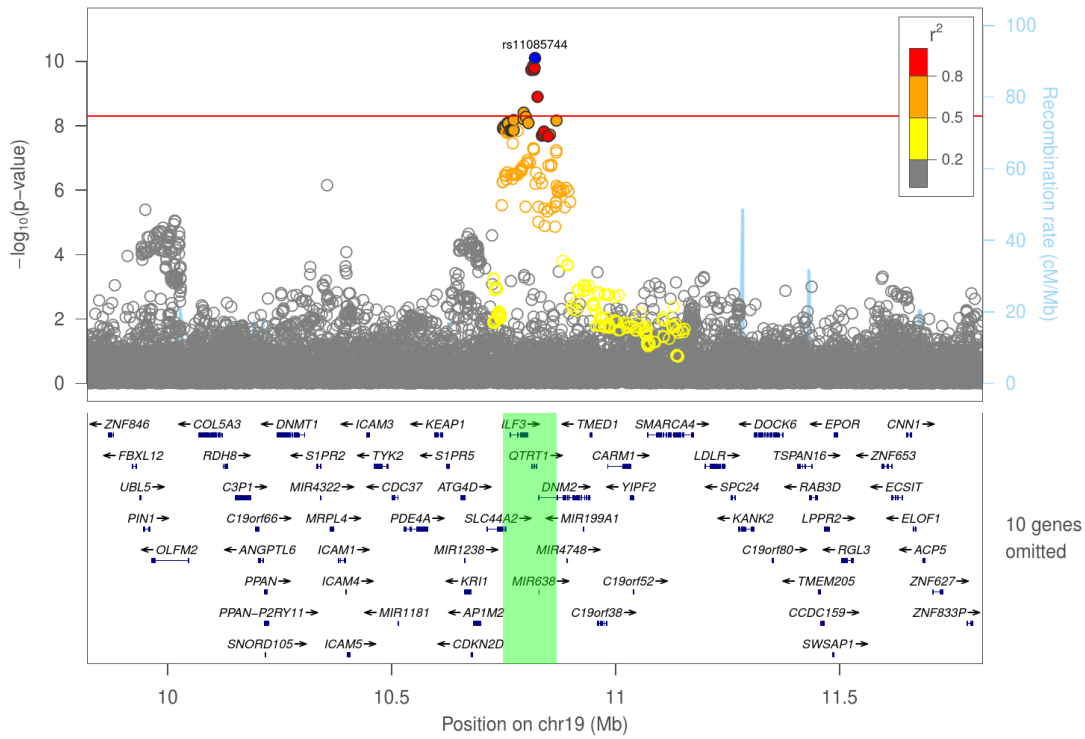
SLC14A2, FEV₁ tier 1 rs9807668



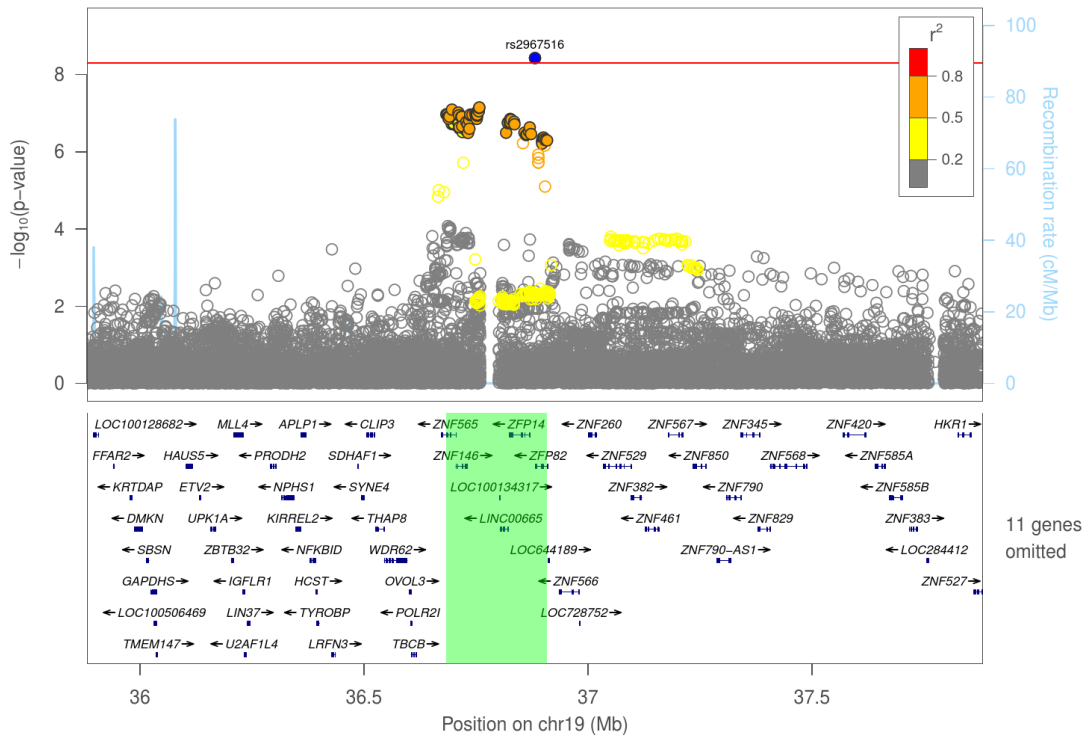
LOC101927273, FVC tier 2 rs2202572



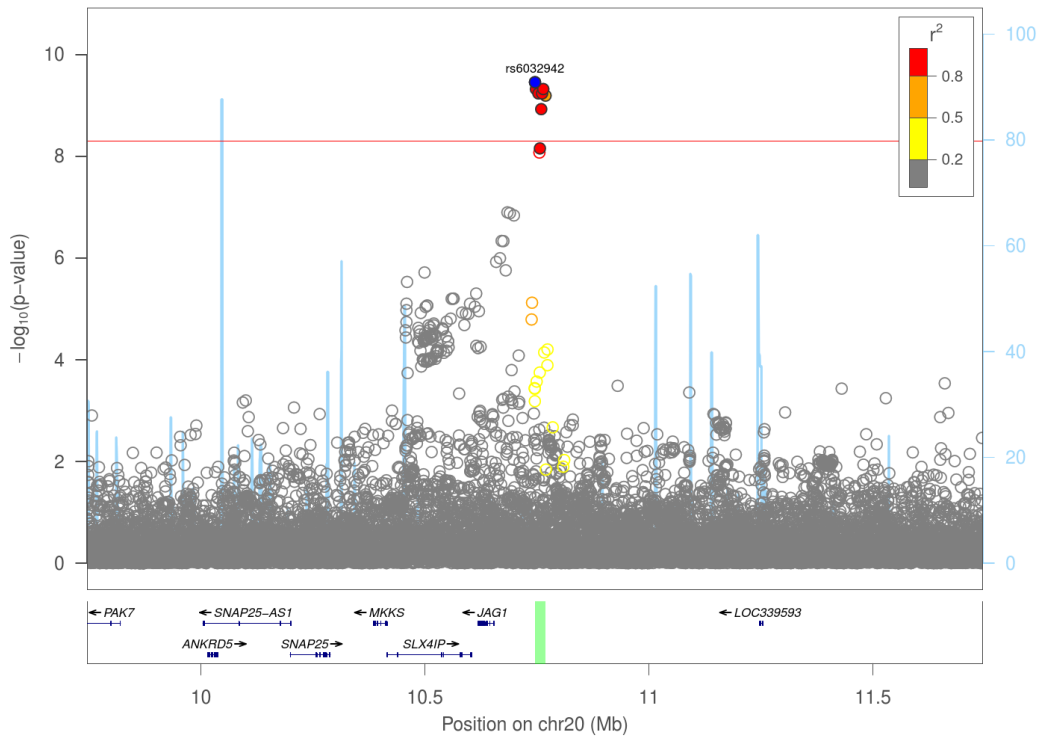
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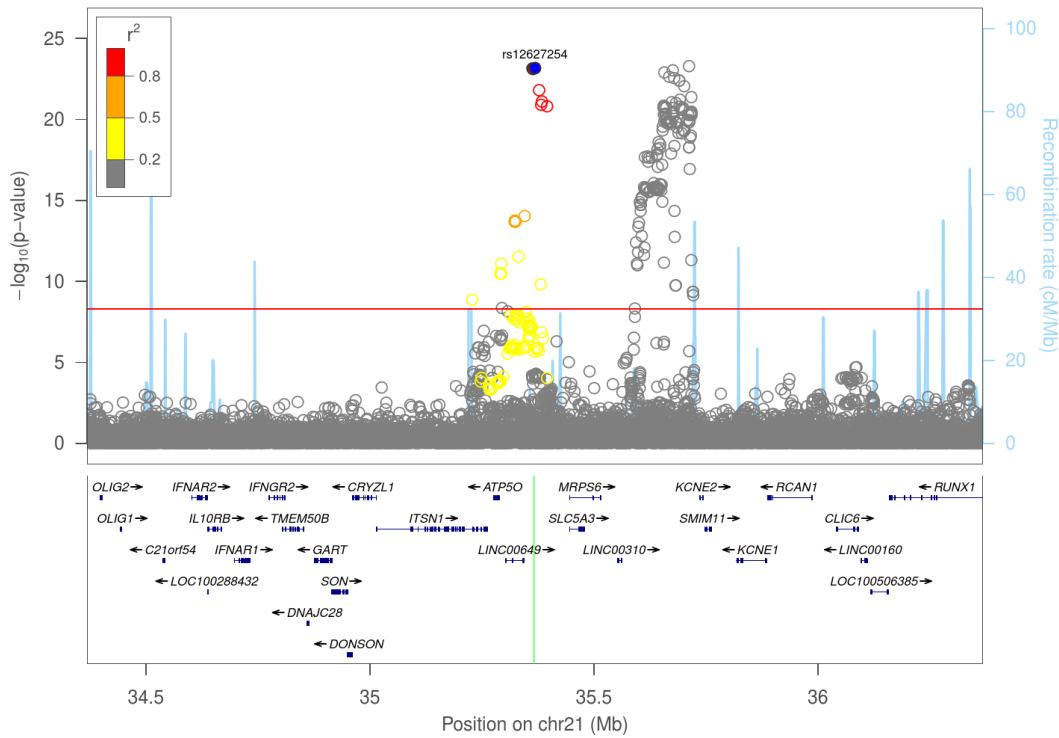
ZFP82, FVC tier 2 rs2967516



LOC101929395, FEV₁ tier 2 rs6032942



LINC00649, FEV₁/FVC tier 1 rs12627254



PPP6R2, FEV₁ tier 2 rs113111175

