

## Supplementary Figures

# Multi-ancestry genome-wide gene-sleep interactions identify novel loci for blood pressure

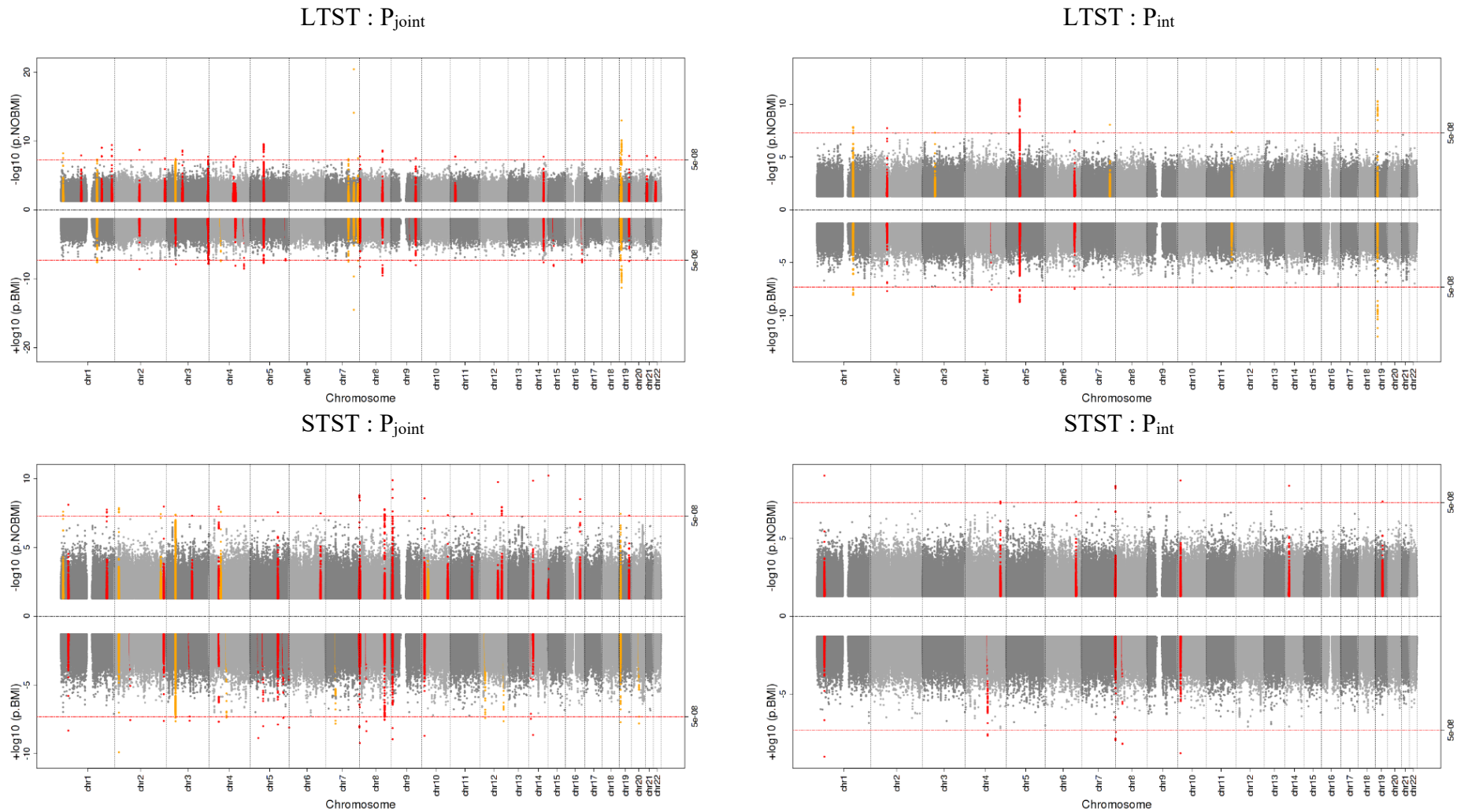
Wang H, et al.

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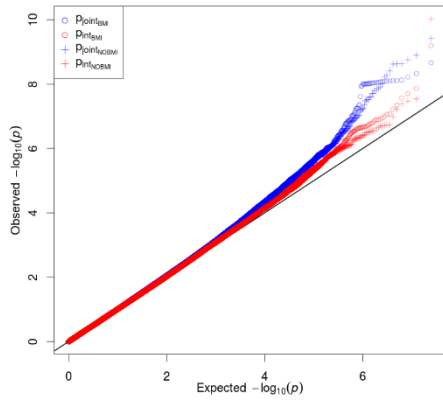
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Upper panel are results without adjusting for BMI, lower panel are results adjusted for BMI. The minimum p-values across four BP traits were used to generate each plot. The orange points correspond to significant loci ( $P < 5 \times 10^{-8}$ ) overlapped with previously reported BP loci (distance  $< 1\text{Mb}$ ), and the red points correspond to novel significant loci.

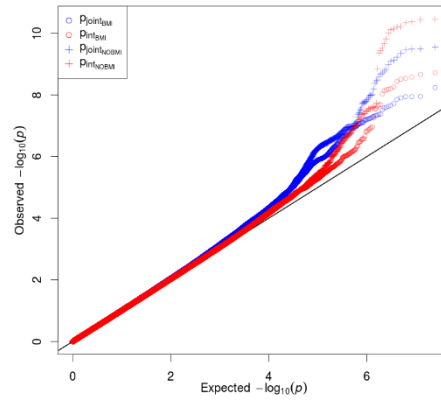


Supplementary Fig. 2. QQ plots of genome-wide gene-sleep interaction analyses in stage 1 MULTI ancestry group.

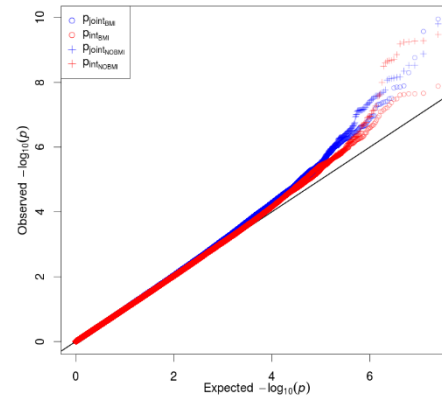
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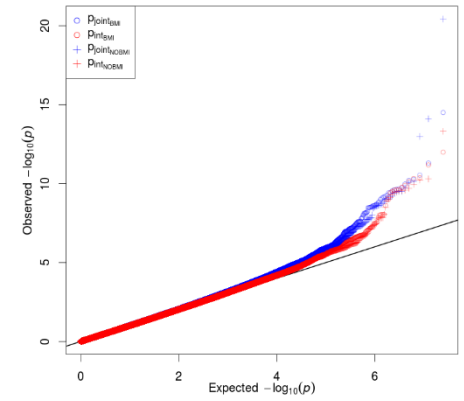
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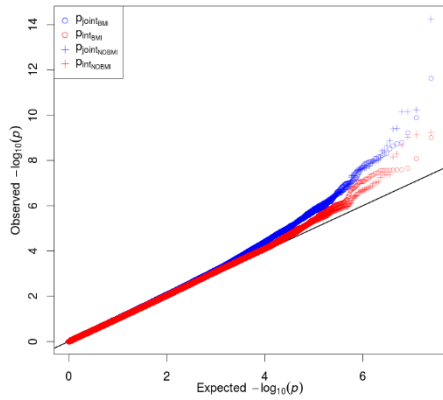
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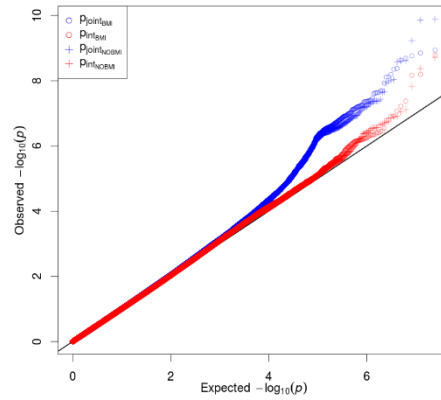
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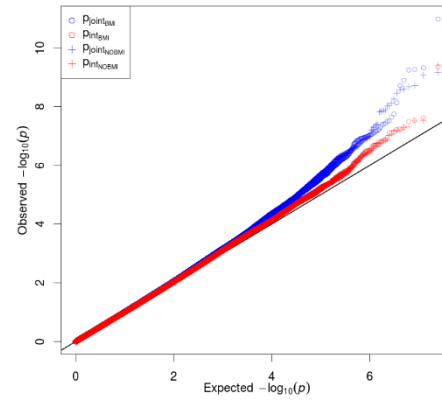
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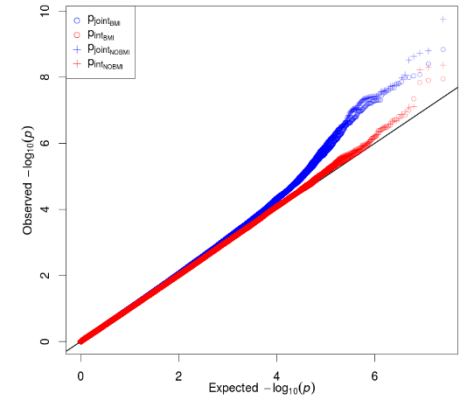
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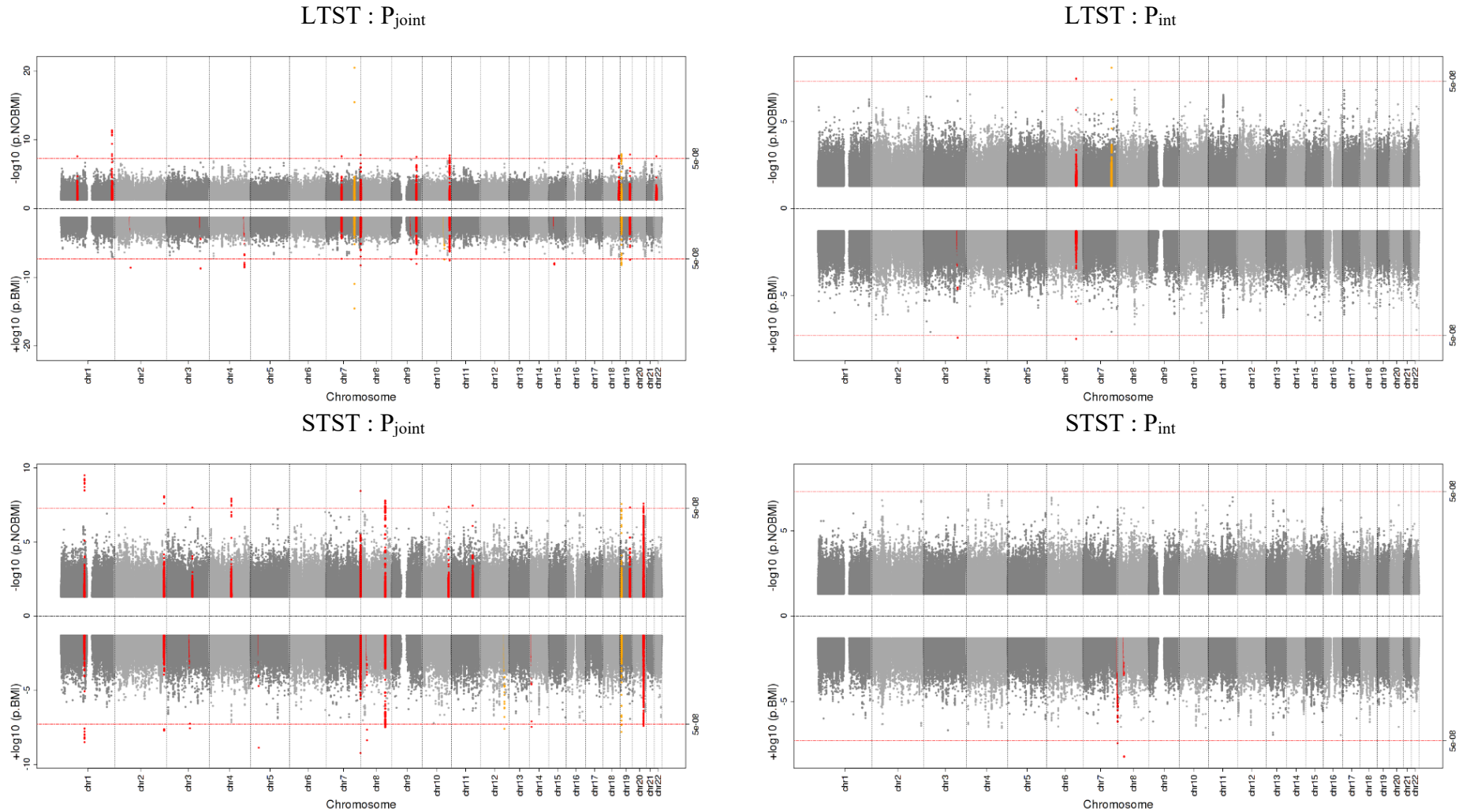


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### Supplementary Fig. 3. Miami plots of genome-wide gene-sleep interaction analyses in stage 1 EUR group.

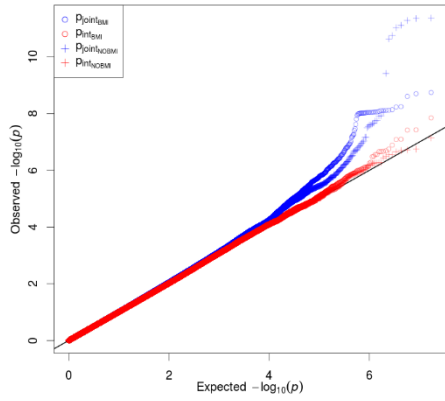
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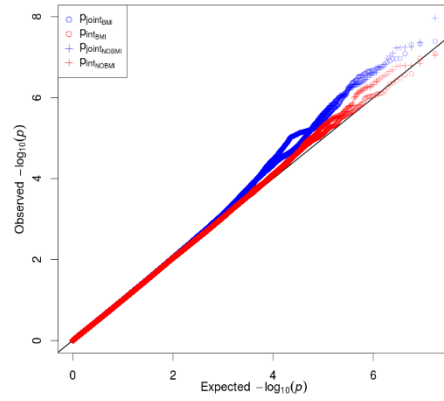


Supplementary Fig. 4. QQ plots of genome-wide gene-sleep interaction analyses in stage 1 EUR group.

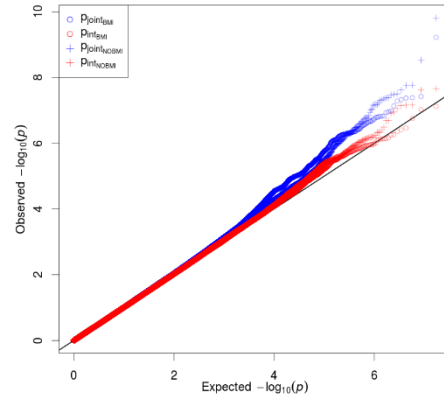
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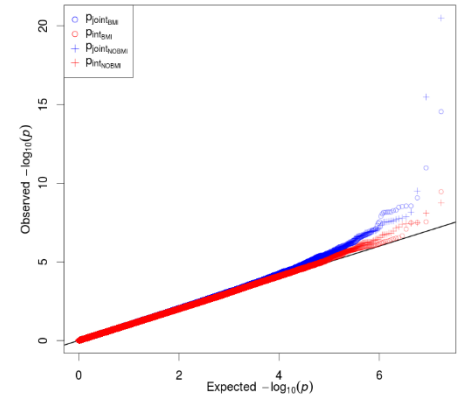
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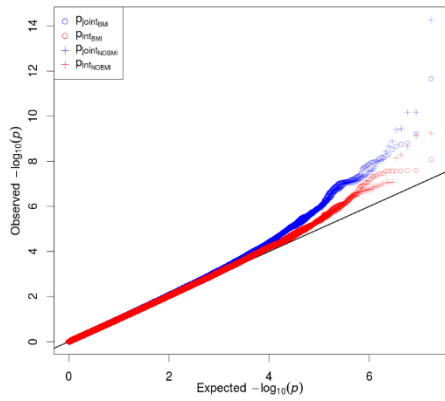
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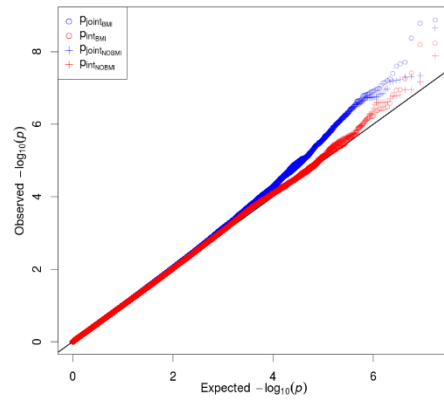
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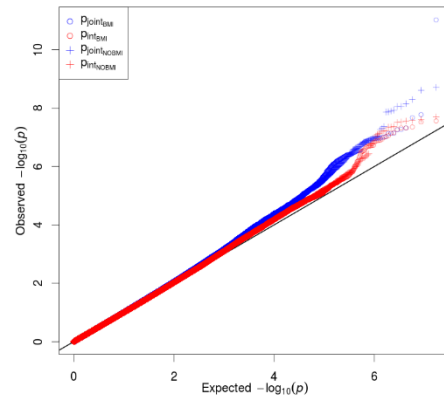
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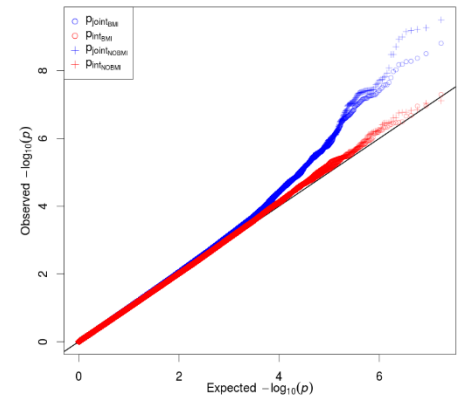
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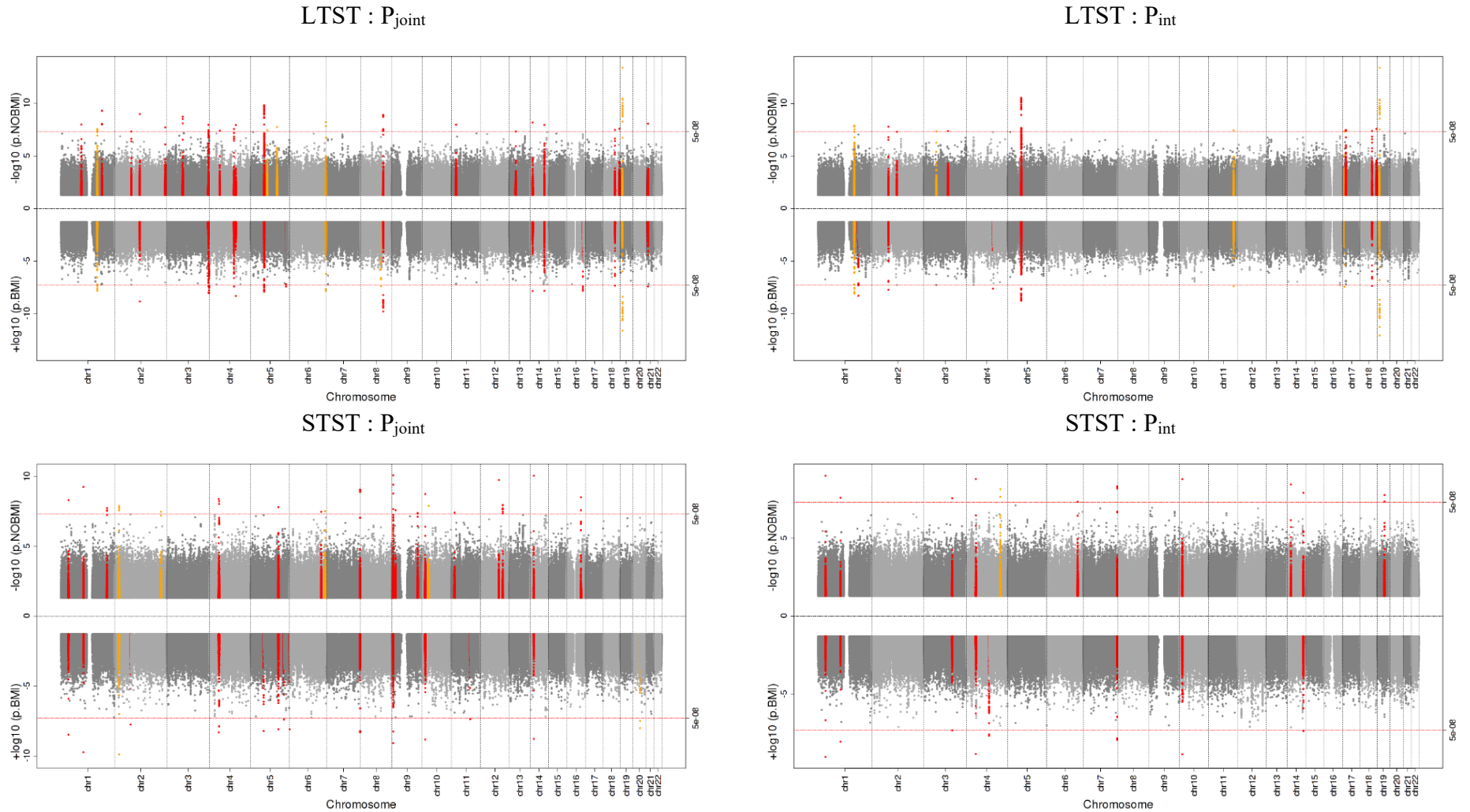


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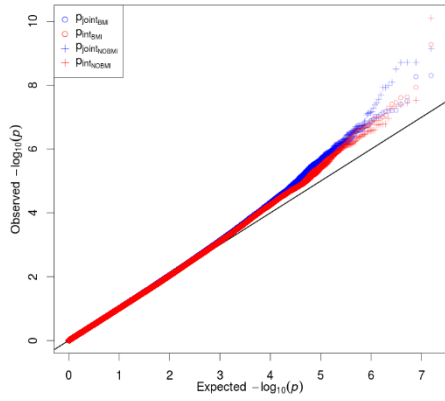
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Upper panel are results without adjusting for BMI, lower panel are results adjusted for BMI. The minimum p-values across four BP traits were used to generate each plot. The orange points correspond to significant loci ( $P < 5 \times 10^{-8}$ ) overlapped with previously reported BP loci (distance  $< 1\text{Mb}$ ), and the red points correspond to novel significant loci.

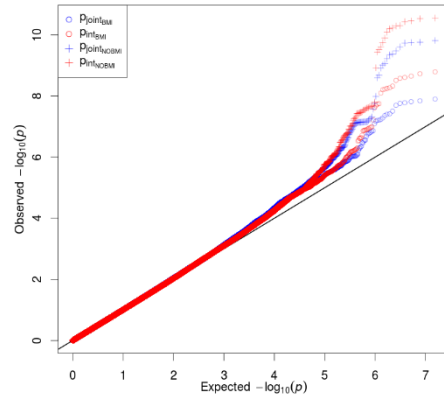


Supplementary Fig. 6. QQ plots of genome-wide gene-sleep interaction analyses in stage 1 AFR group.

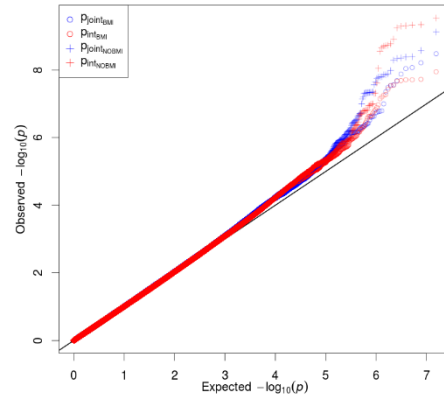
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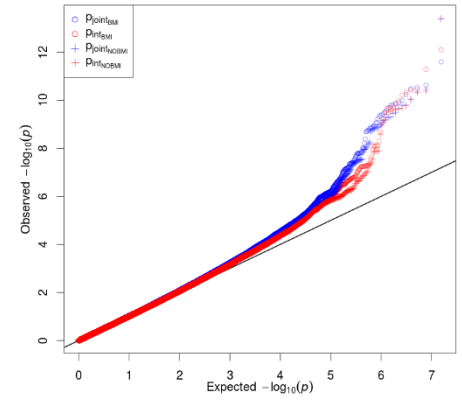
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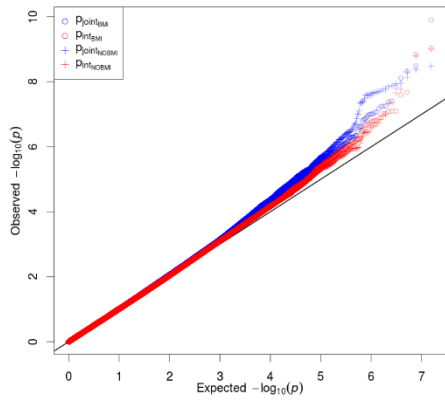
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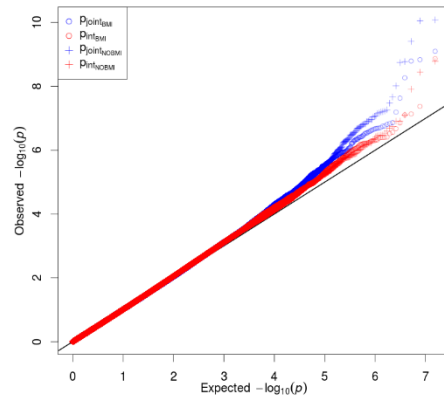
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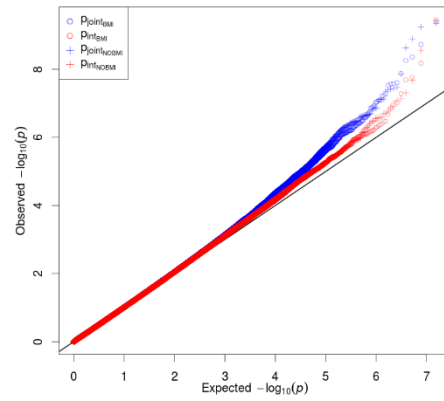
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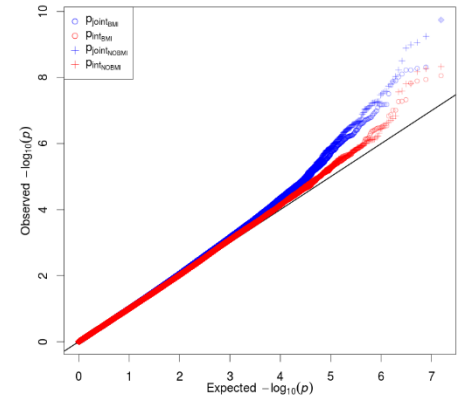
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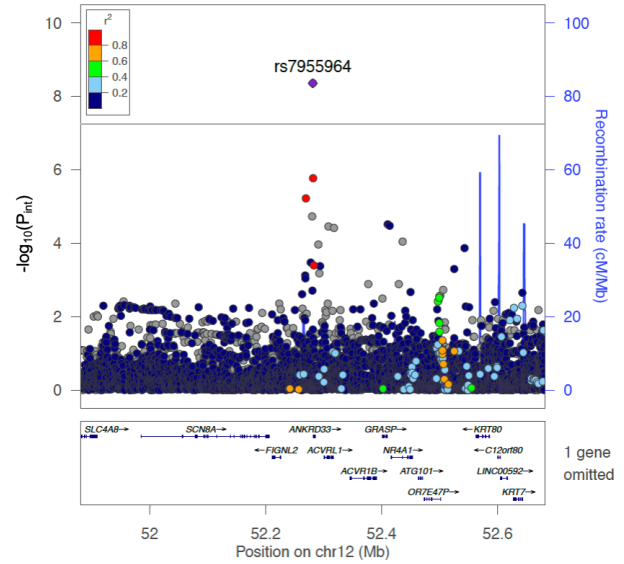
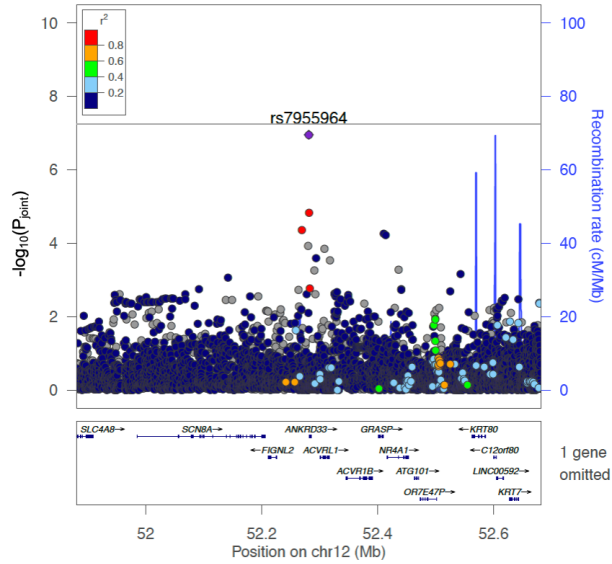


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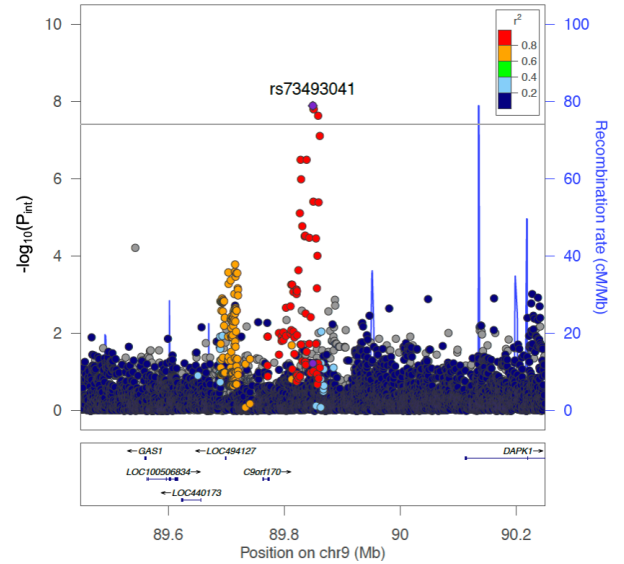
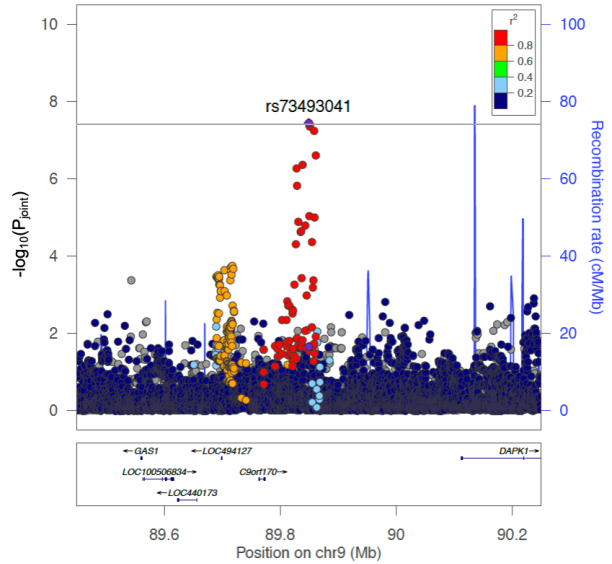


Supplementary Fig. 7. Regional association plots of  $P_{\text{joint}}$  and  $P_{\text{int}}$  for 3 replicated novel loci in stage 1+2 MULTI ancestry analyses (Table 1).

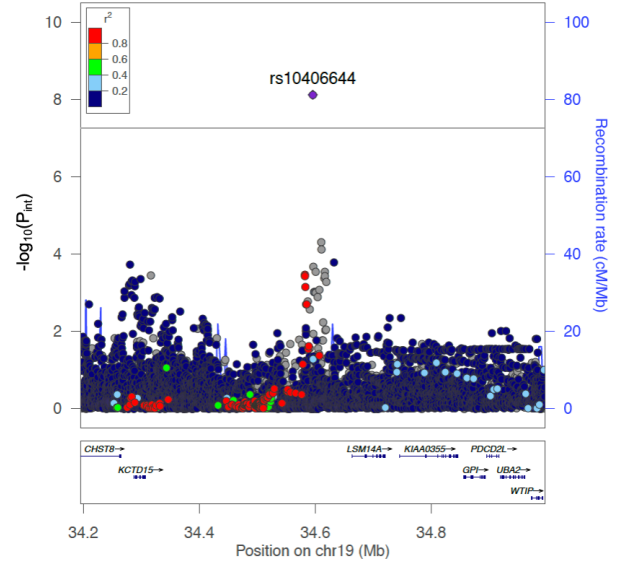
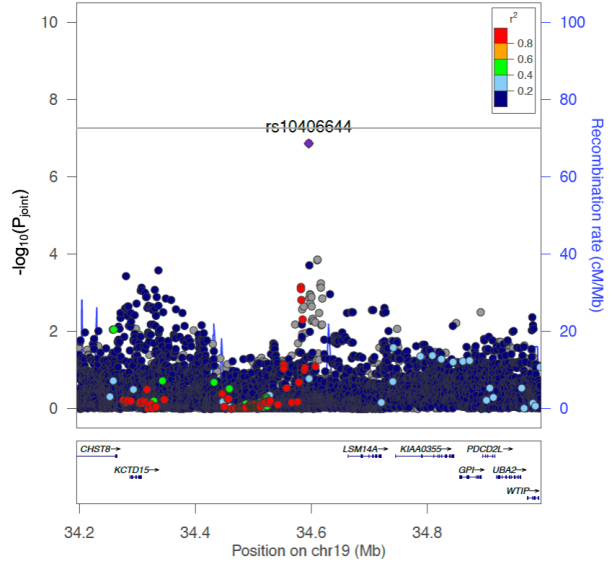
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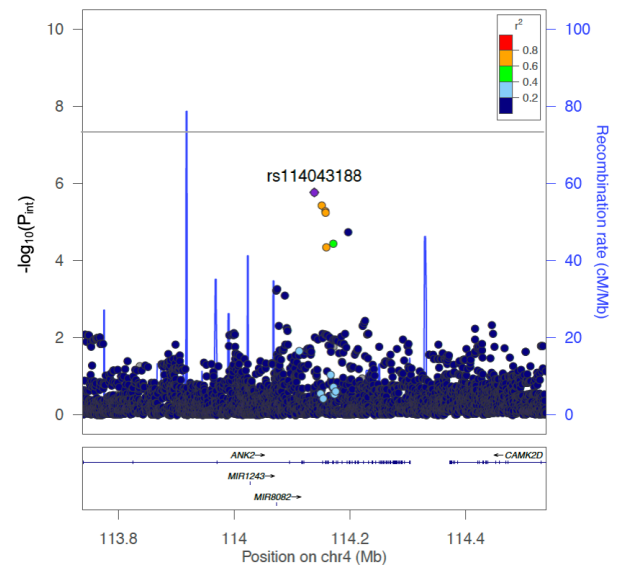
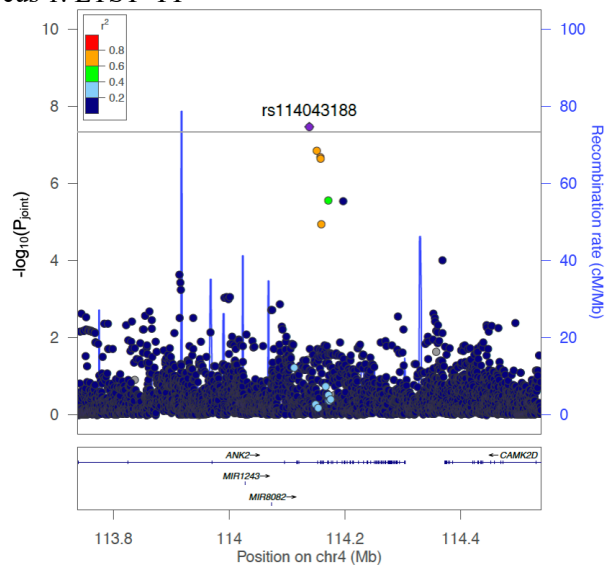


Locus 3. STST- PP

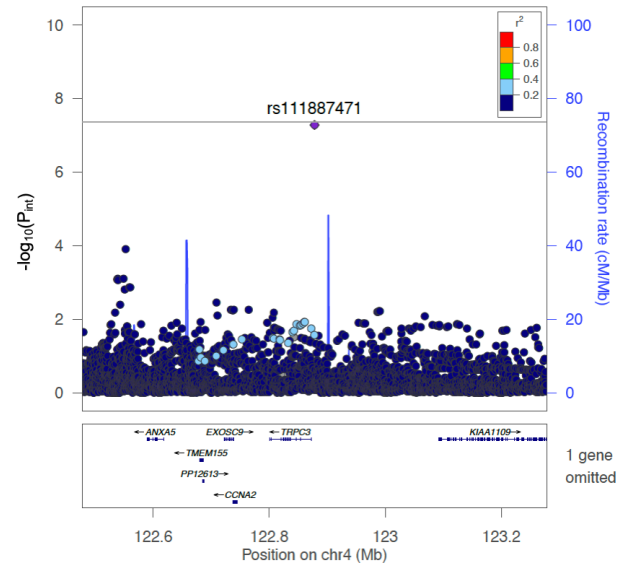
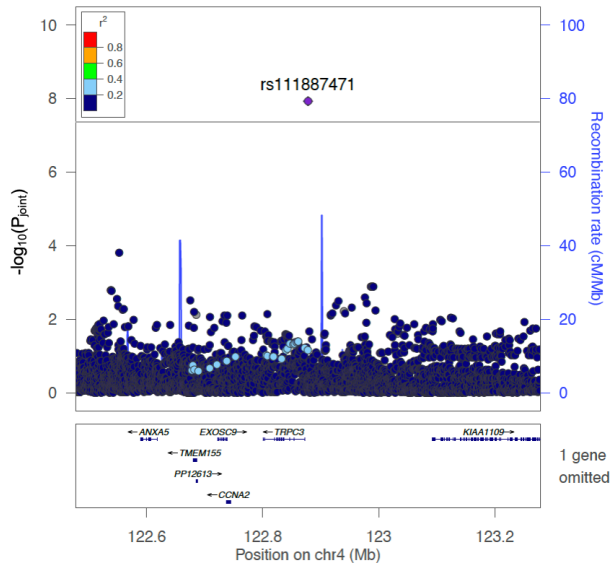


Supplementary Fig. 8. Regional association plots of  $P_{\text{joint}}$  and  $P_{\text{int}}$  for 3 novel loci in stage 1 AFR analyses (Supplementary Table 8).

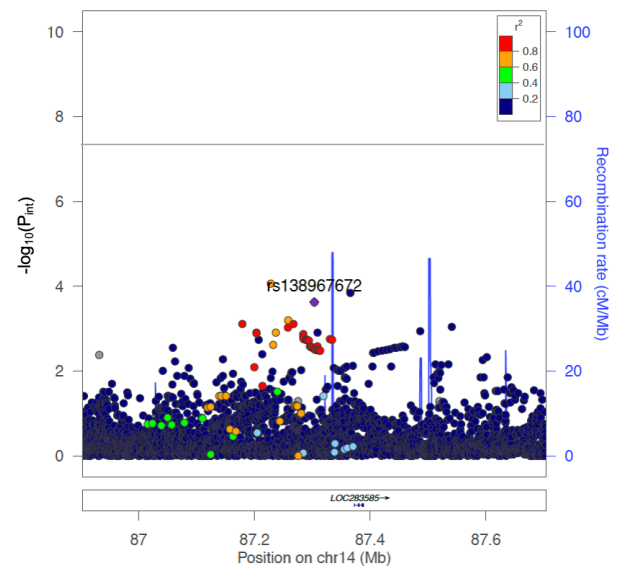
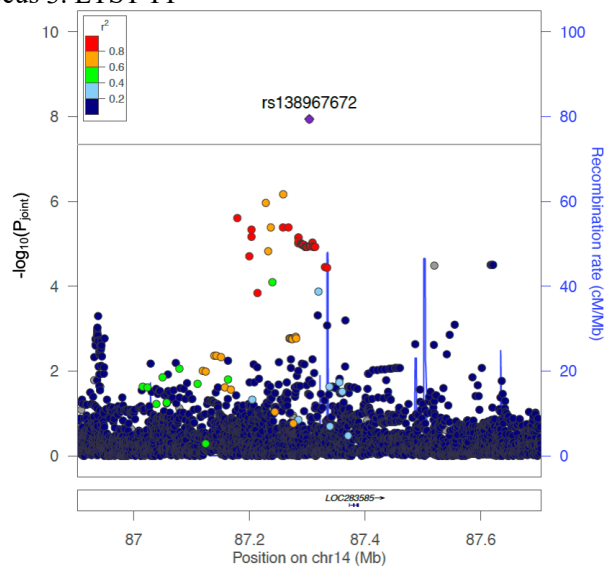
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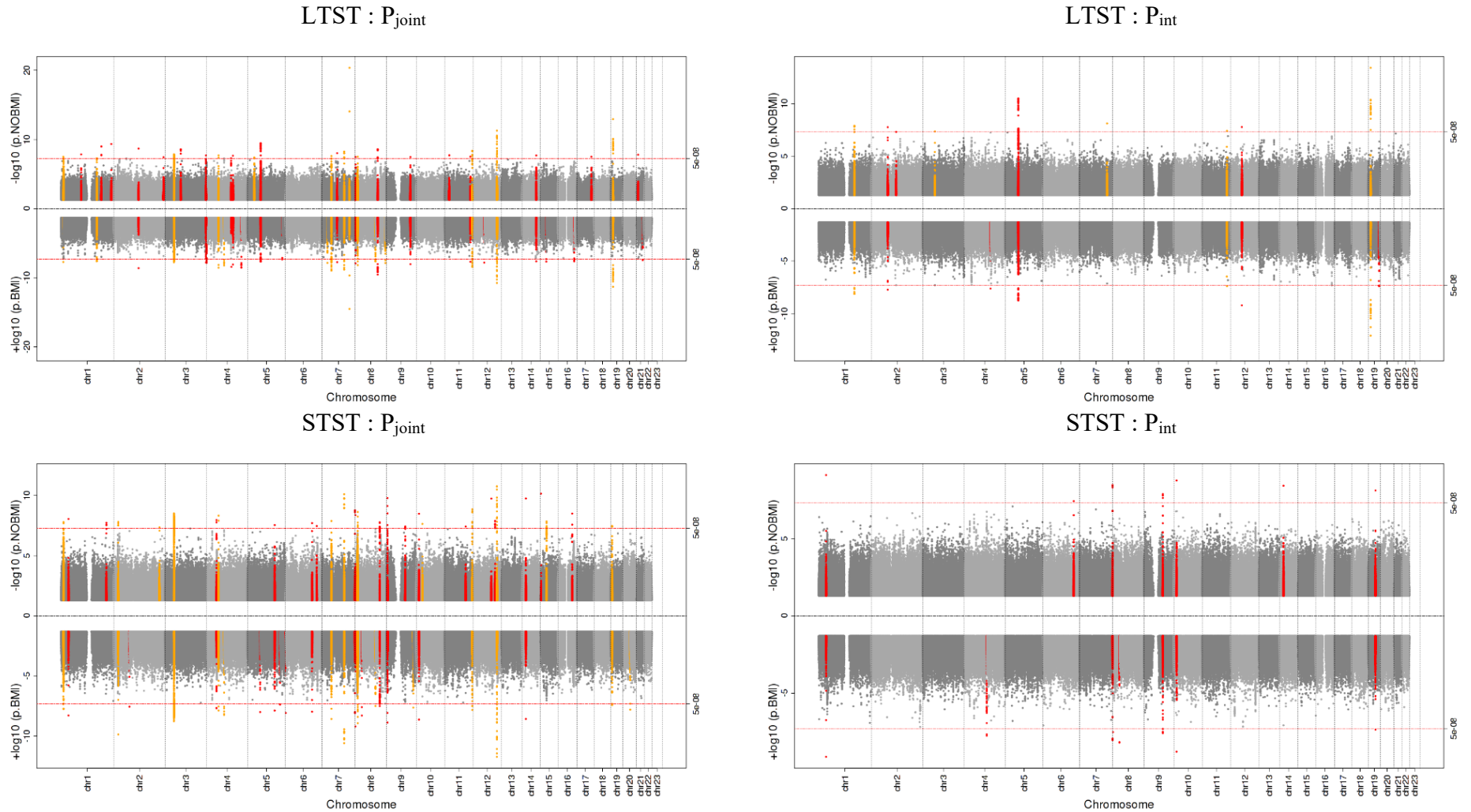


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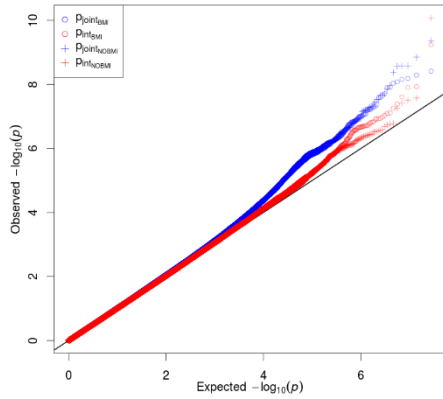
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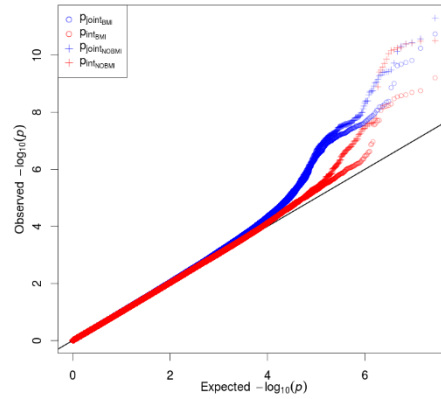


Supplementary Fig. 10. QQ plots of genome-wide gene-sleep interaction analyses in stage 1+2 MULTI ancestry group.

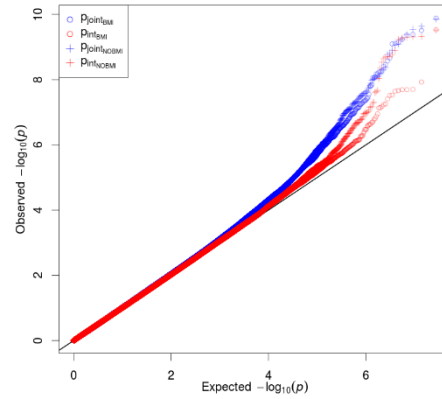
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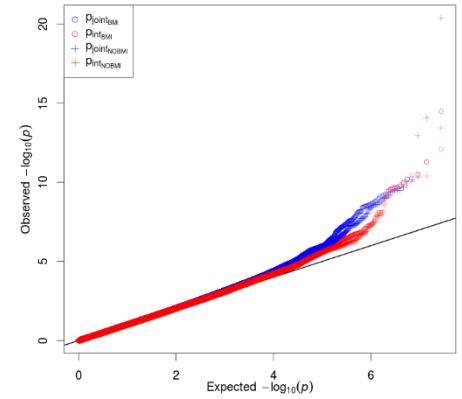
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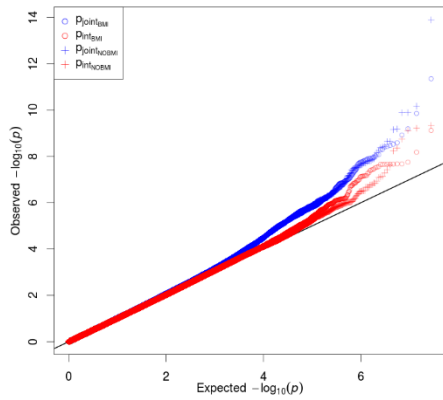
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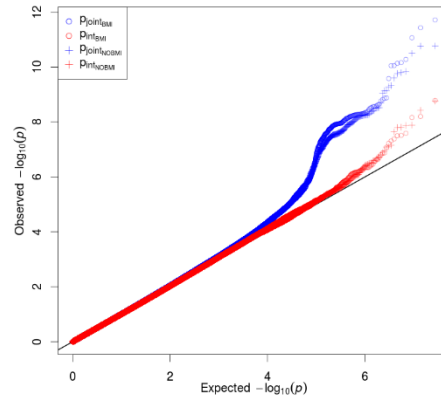
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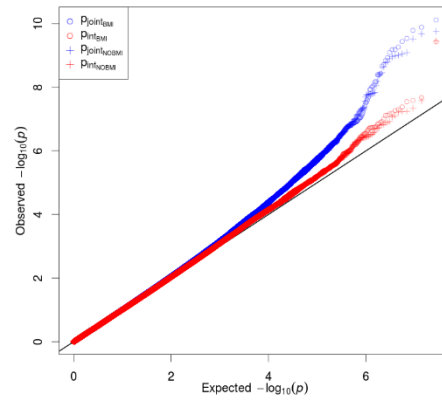
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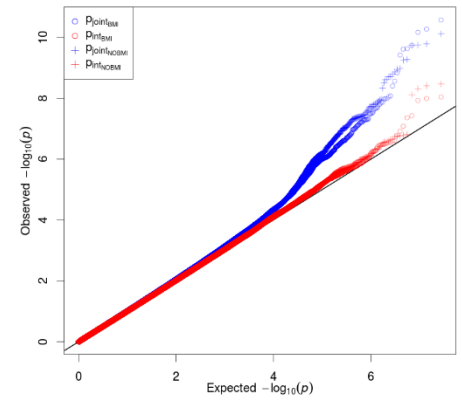
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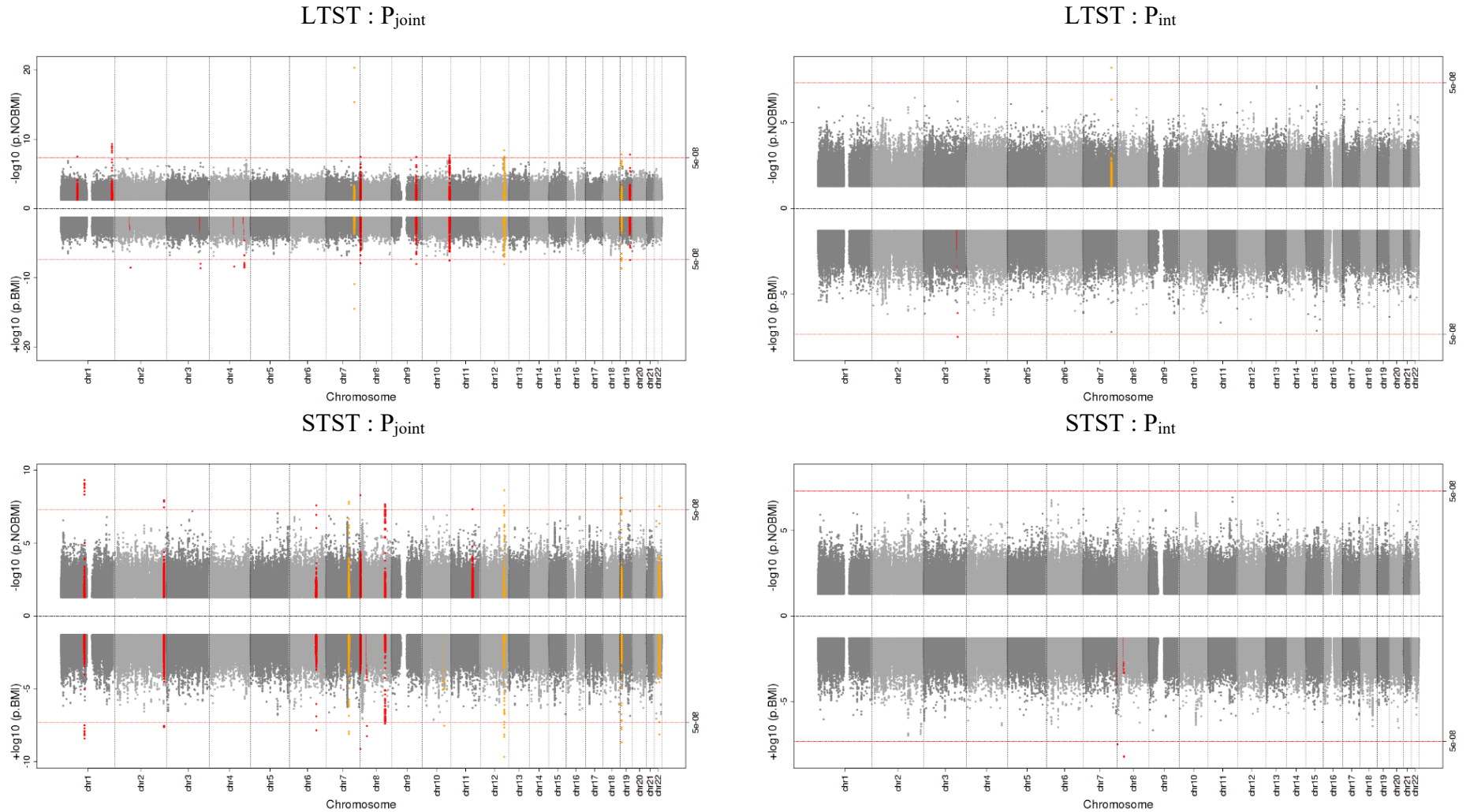
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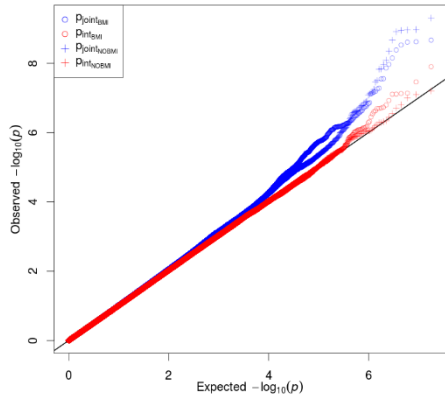
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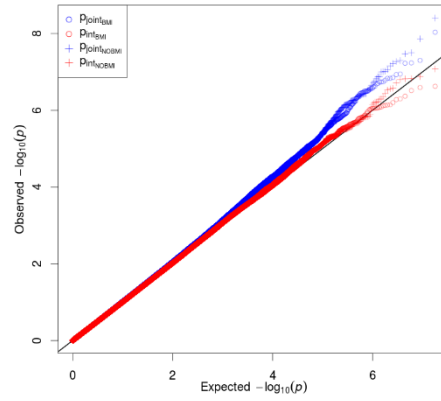


Supplementary Fig. 12. QQ plots of genome-wide gene-sleep interaction analyses in stage 1+2 EUR group.

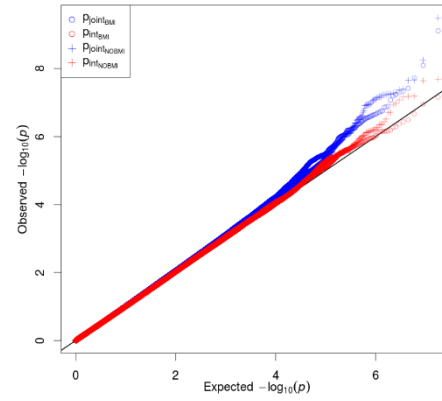
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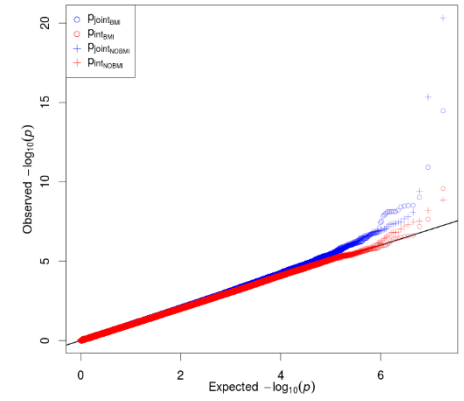
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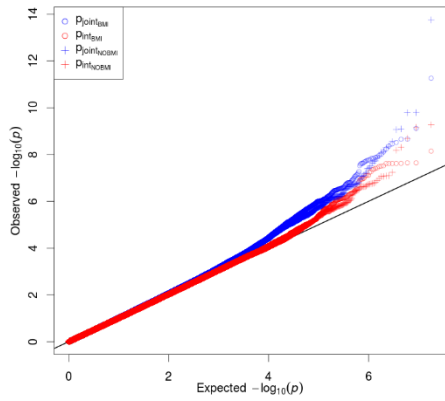
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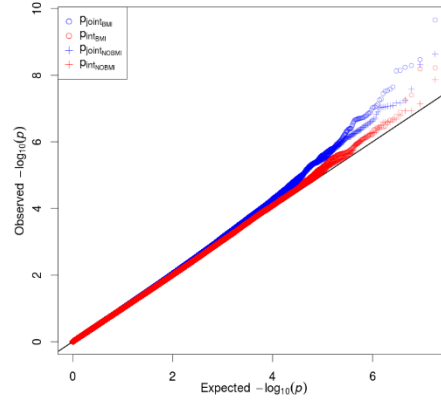
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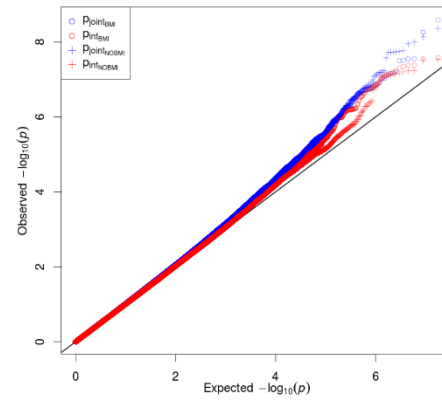
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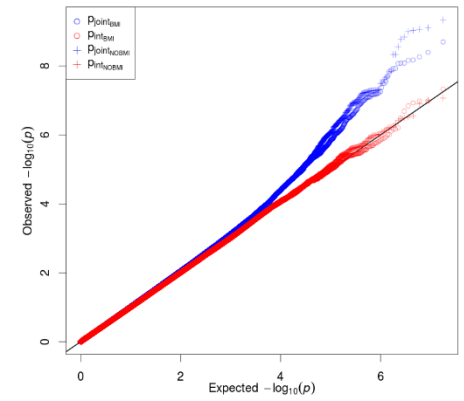
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STST-MAP

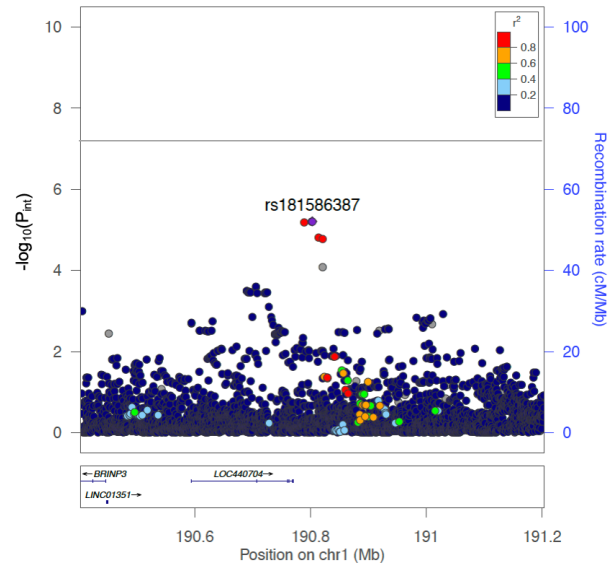
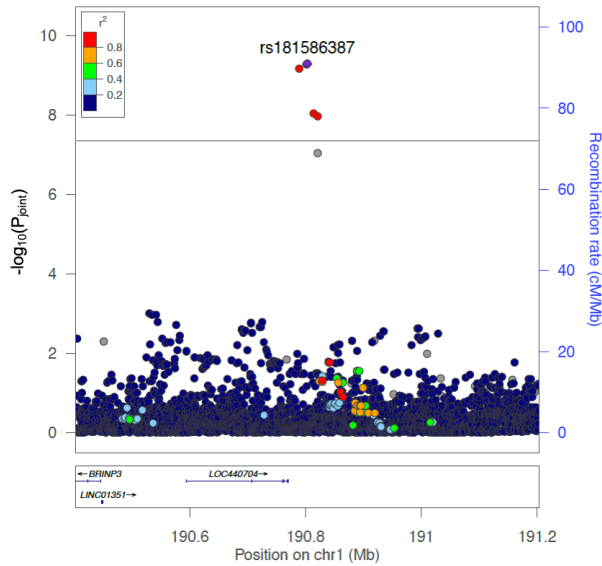


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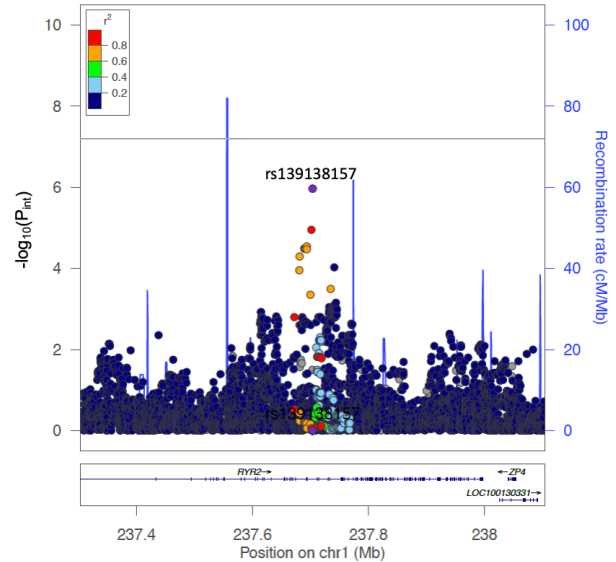
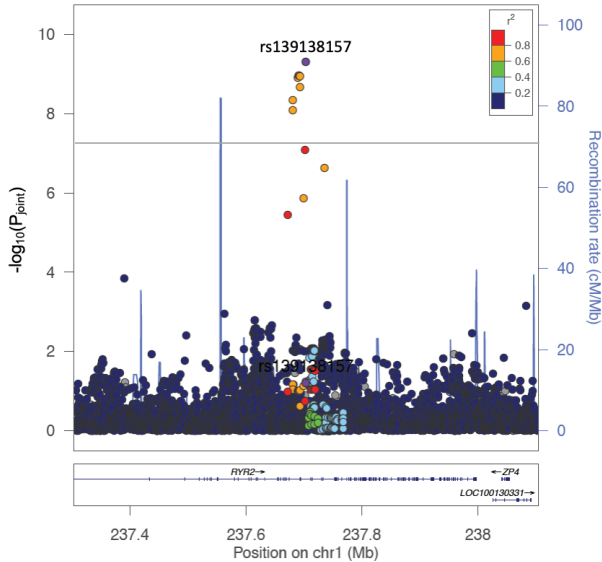


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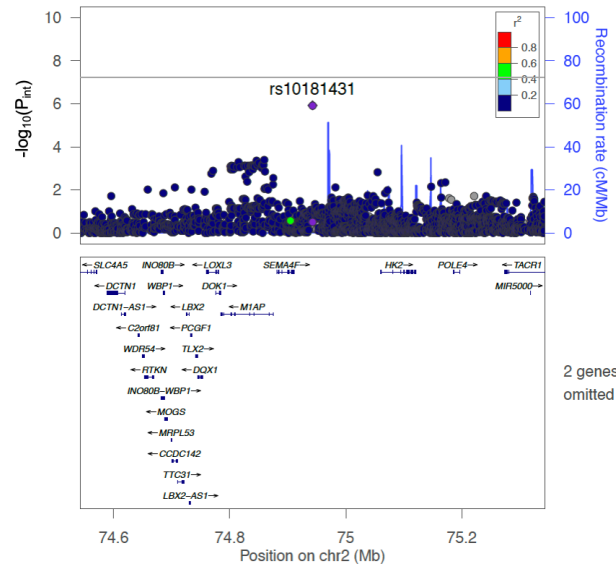
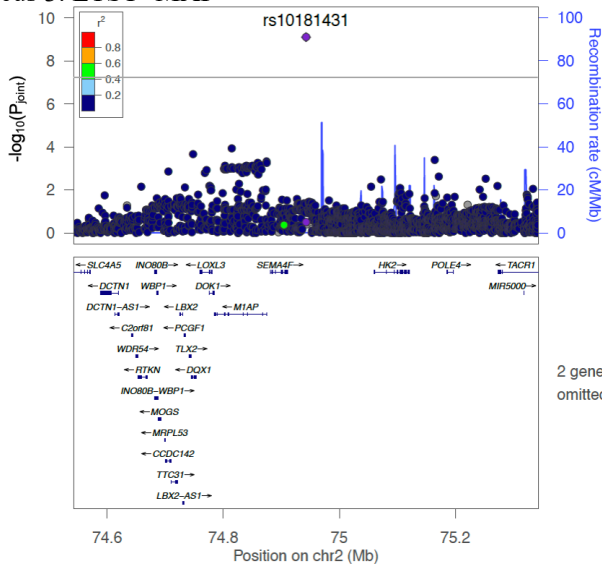
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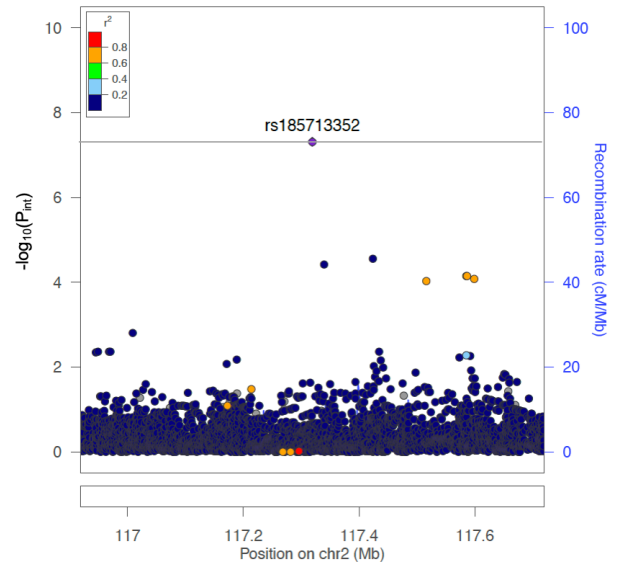
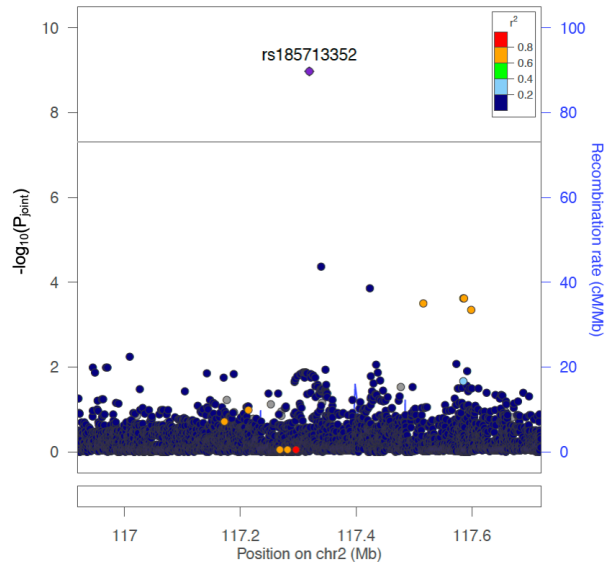
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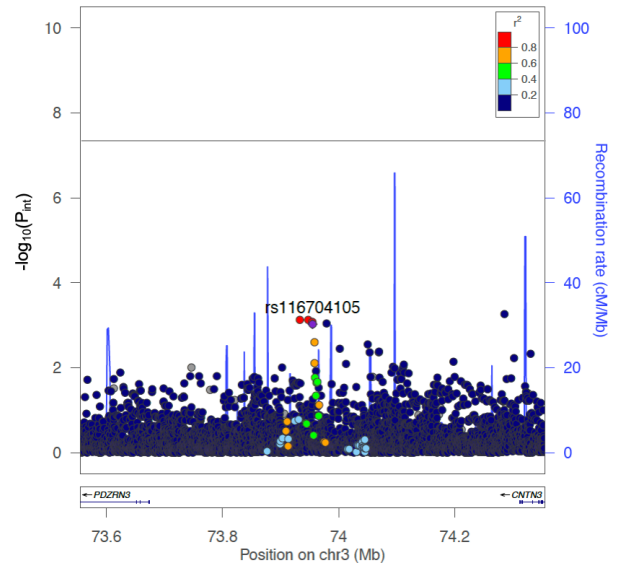
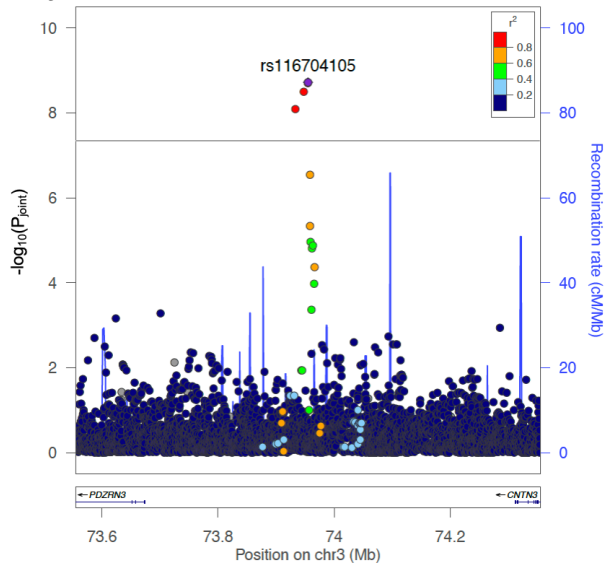
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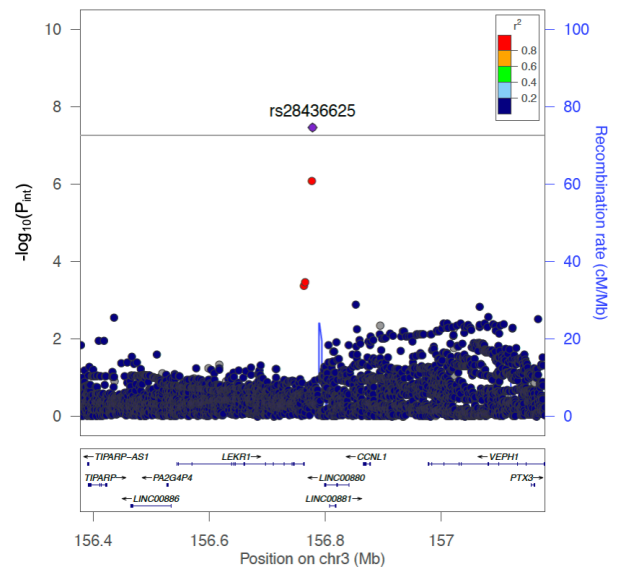
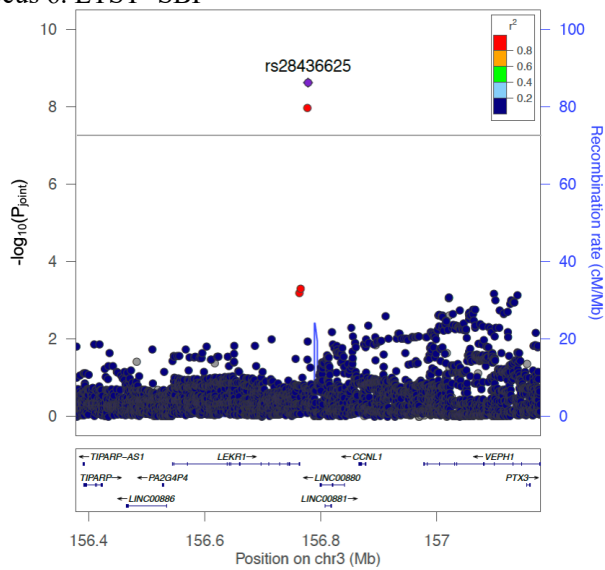
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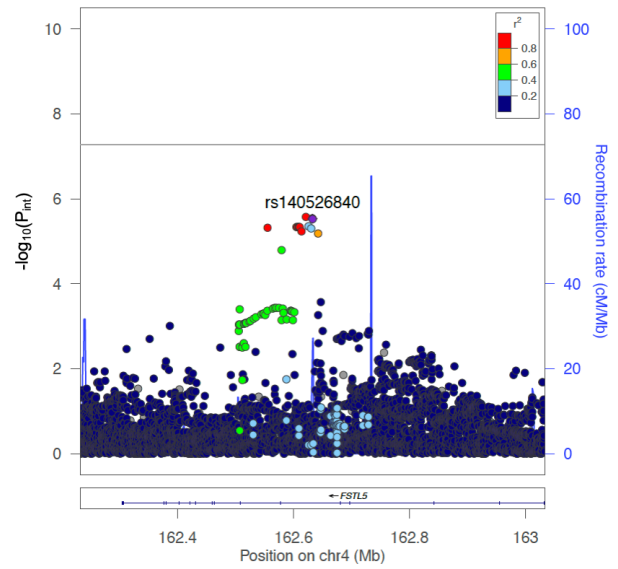
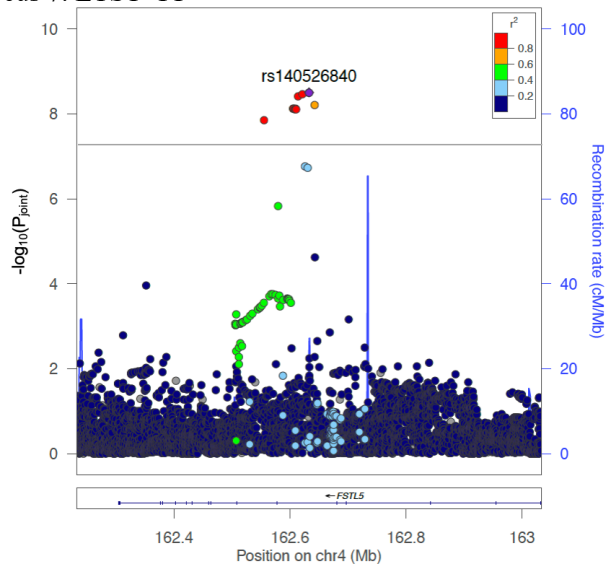
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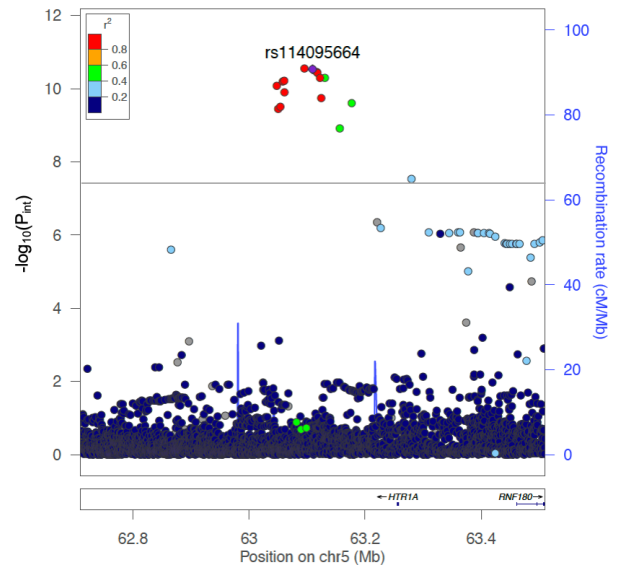
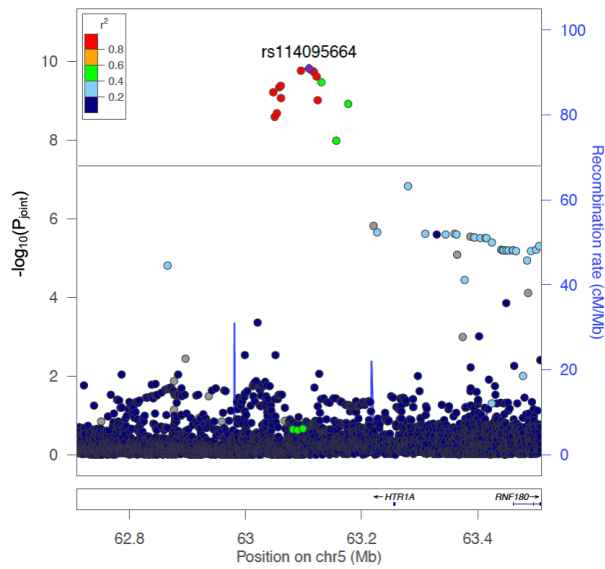
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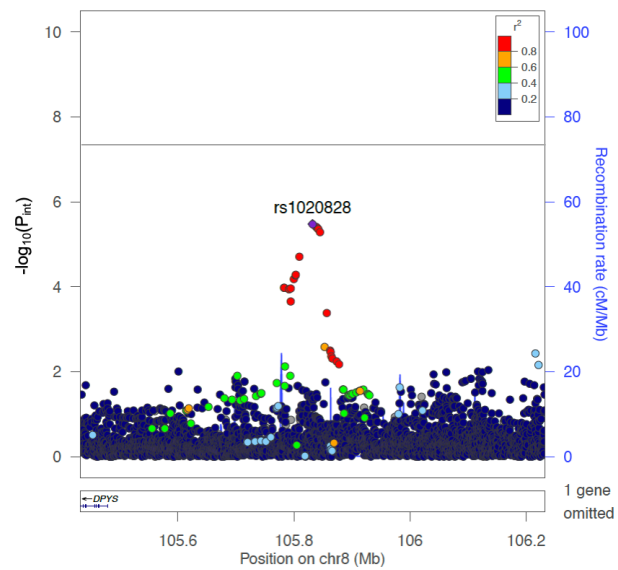
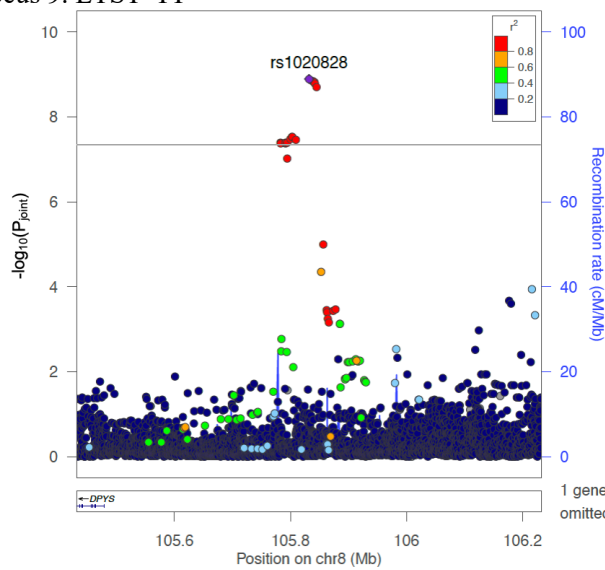
Locus 7. LTST- PP



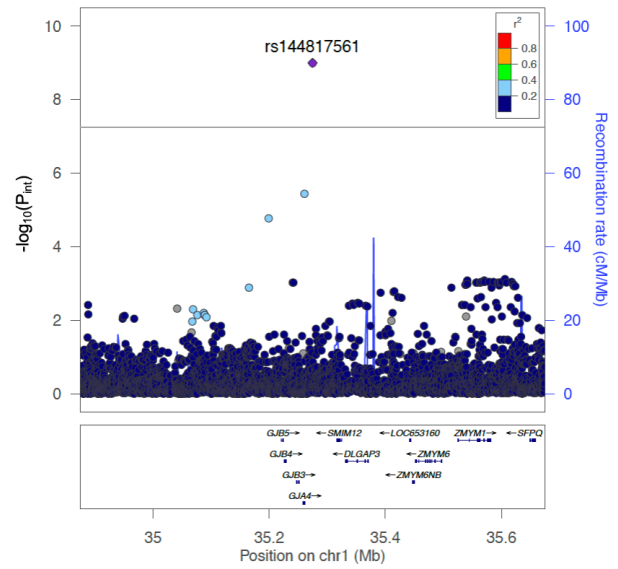
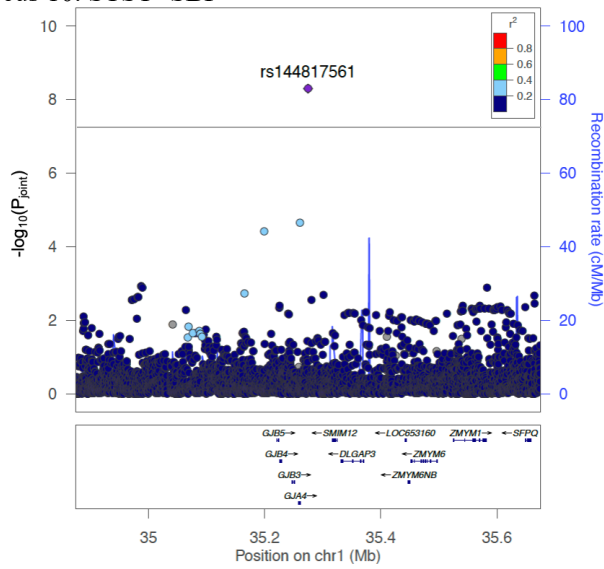
Locus 8. LTST- DBP



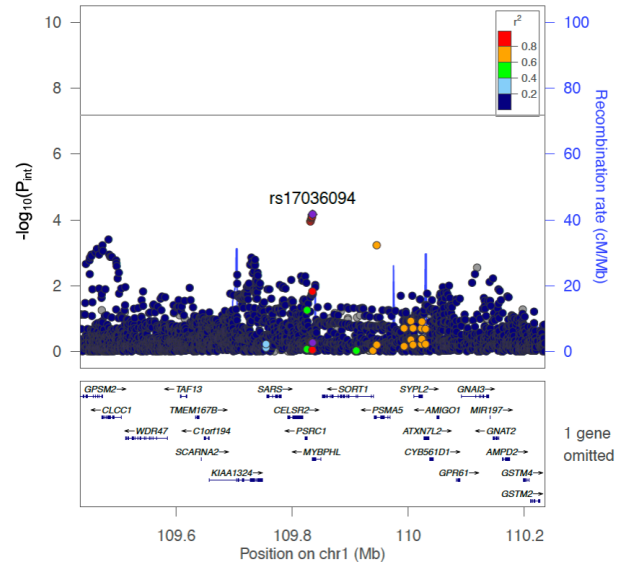
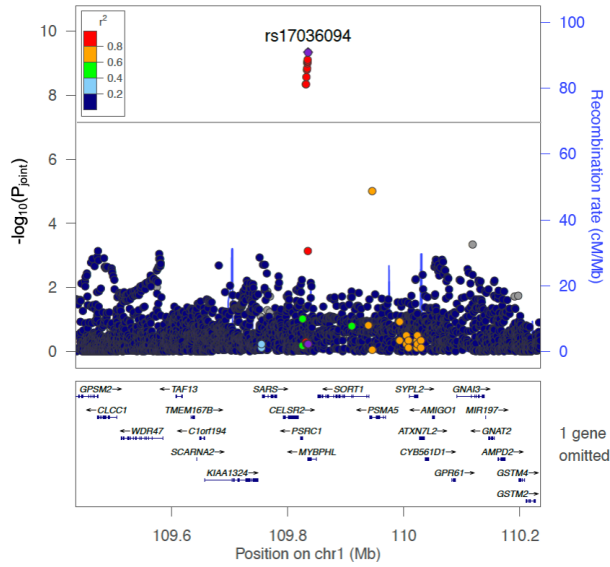
Locus 9. LTST- PP



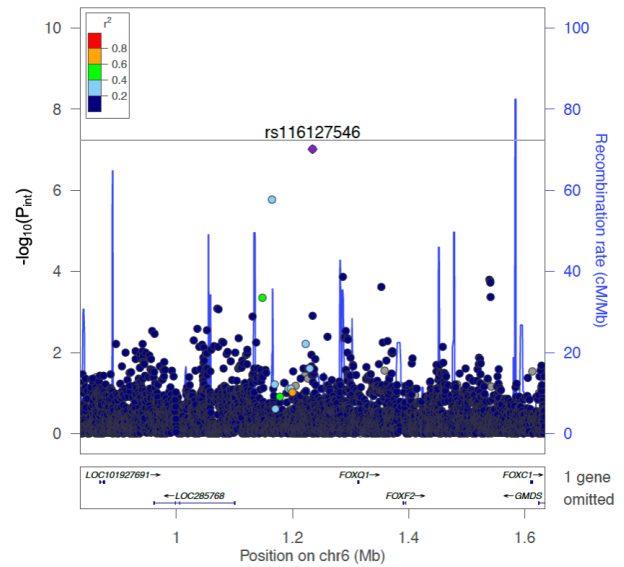
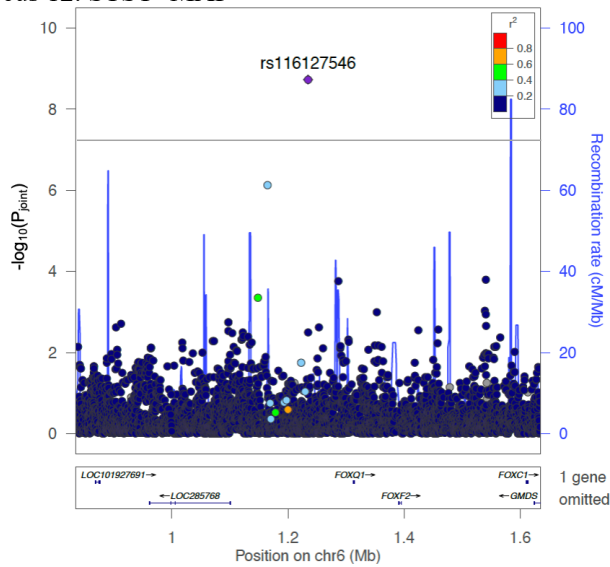
Locus 10. STST- SBP



Locus 11. STST- PP

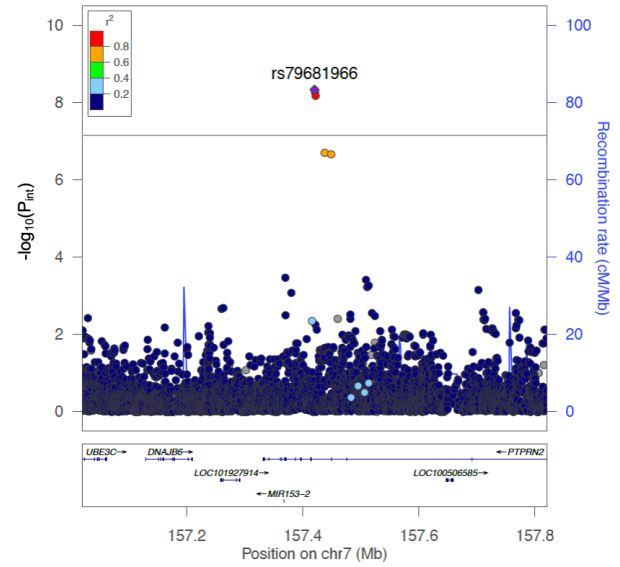
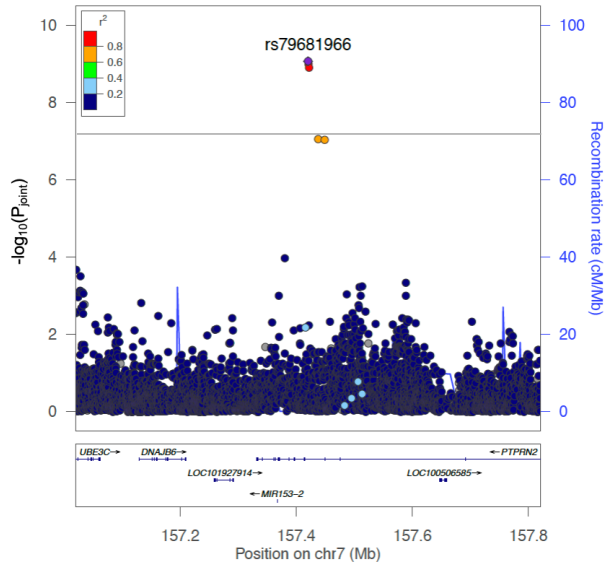


Locus 12. STST- MAP

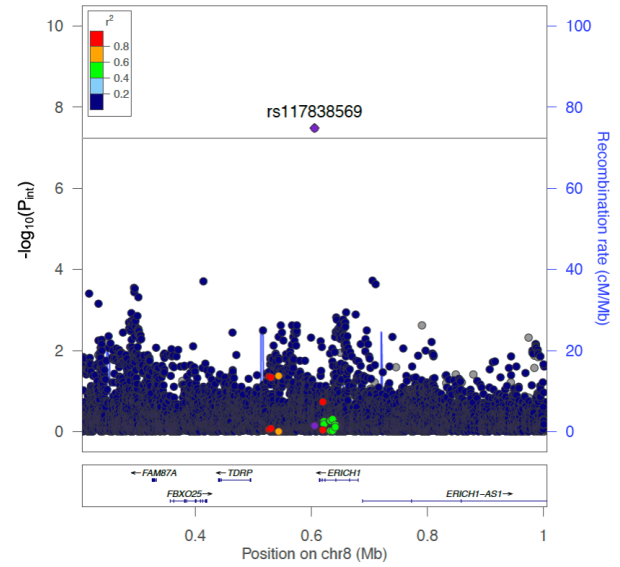
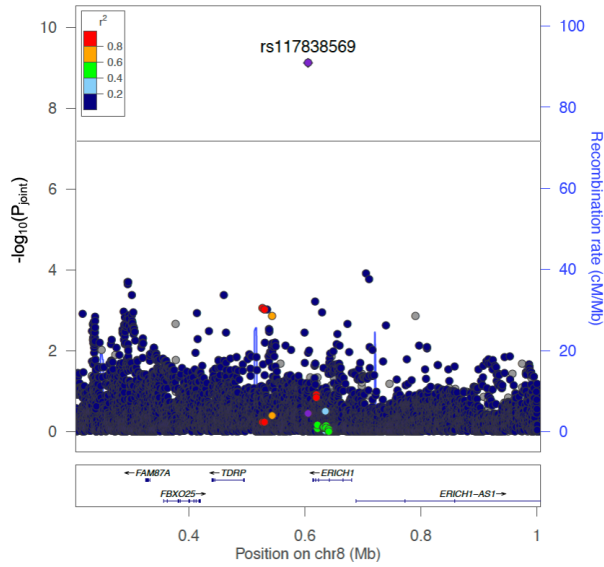




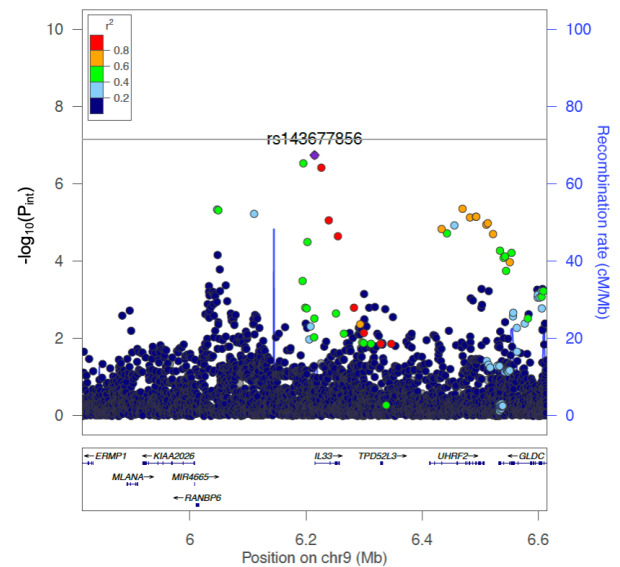
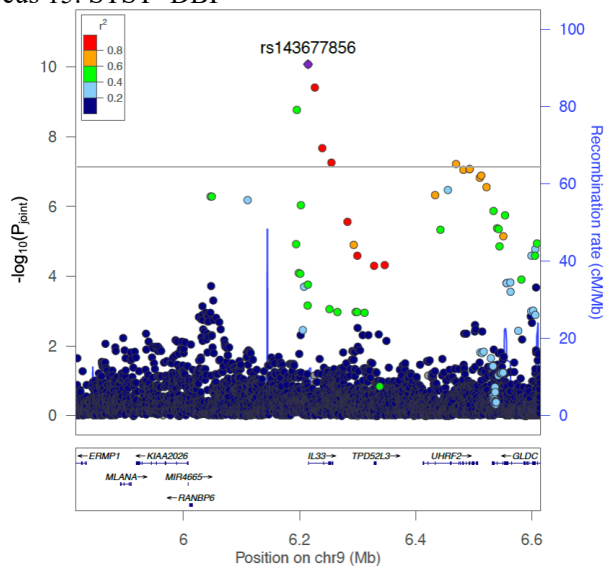
Locus 13. STST- PP



Locus 14. STST- SBP

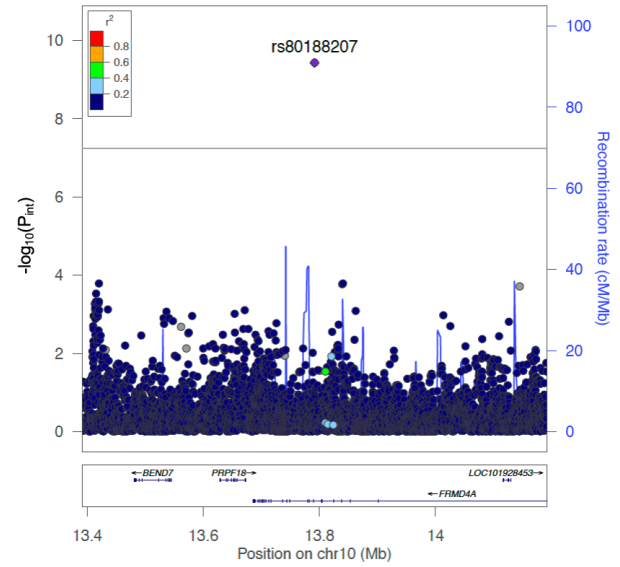
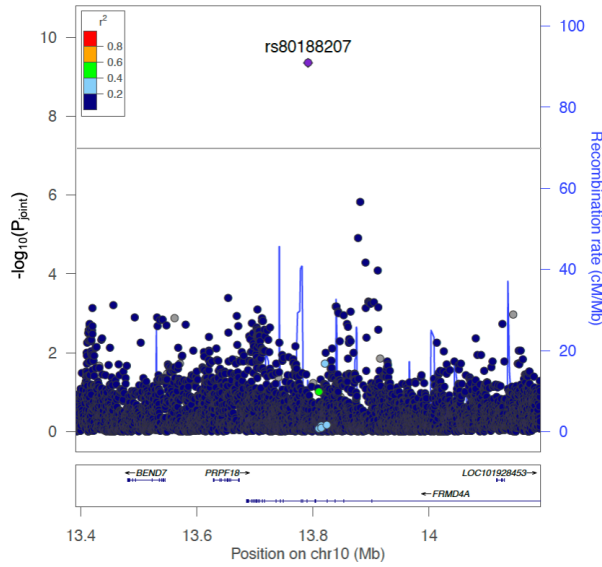


Locus 15. STST- DBP

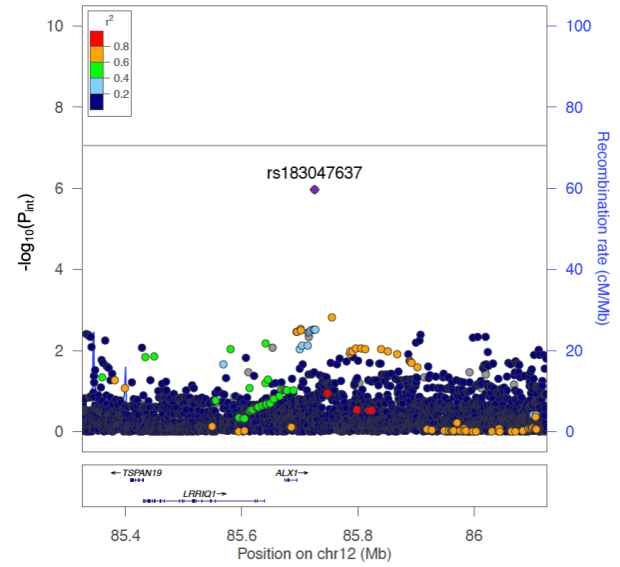
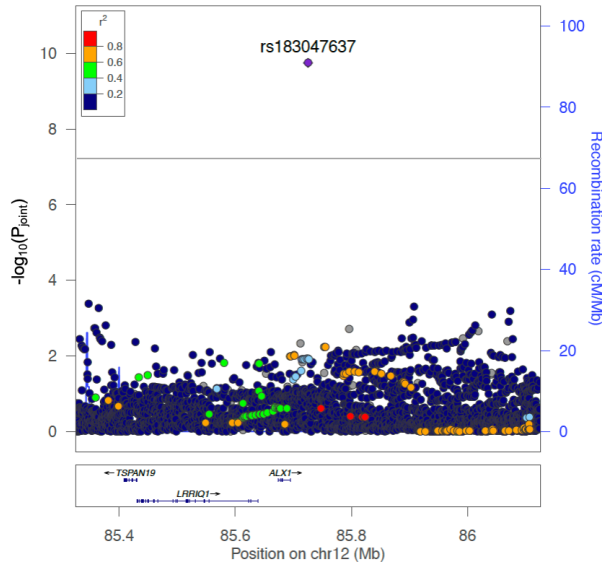




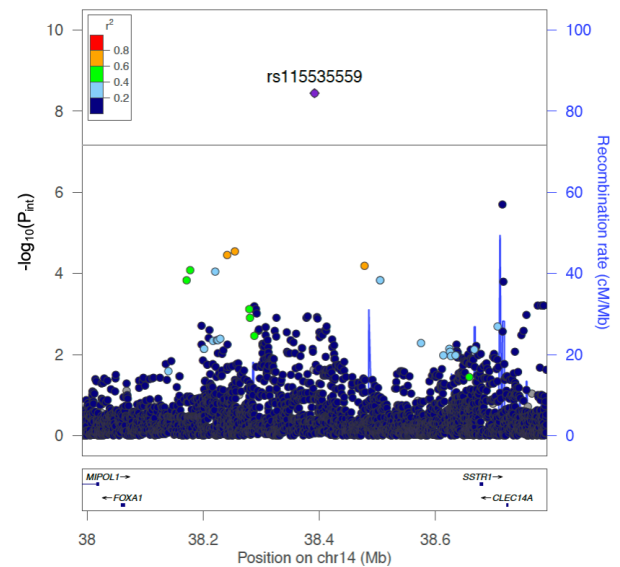
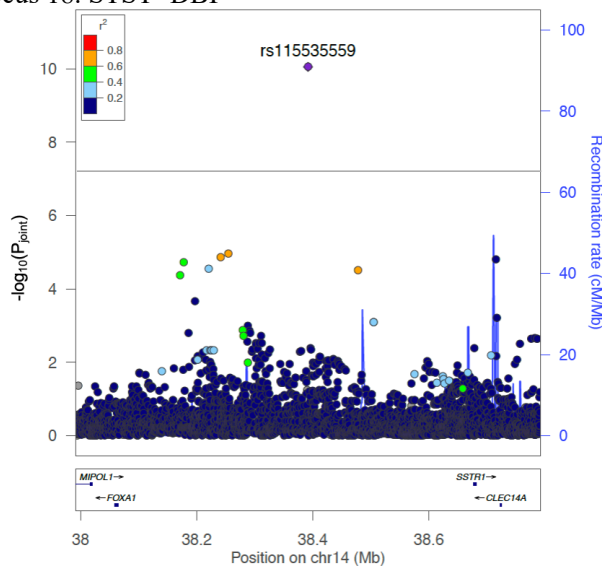
Locus 16. STST- MAP



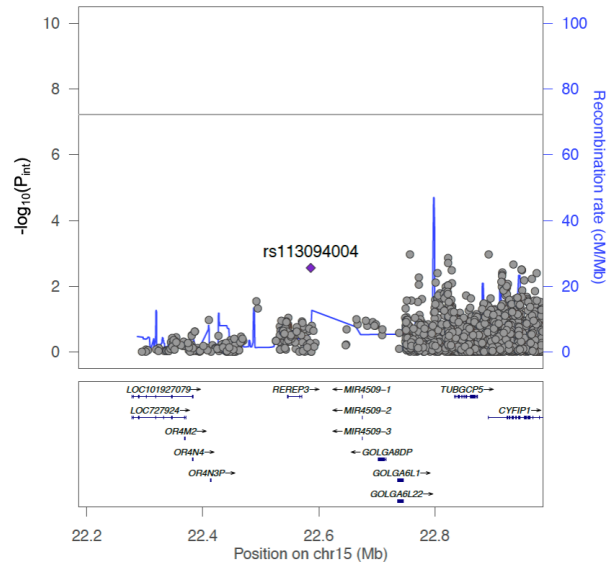
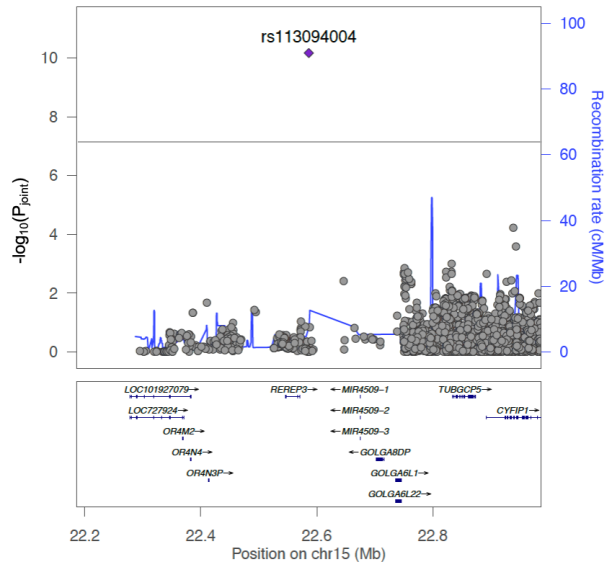
Locus 17. STST- PP



Locus 18. STST- DBP



Locus 19. STST- SBP



Locus 20. STST- PP

