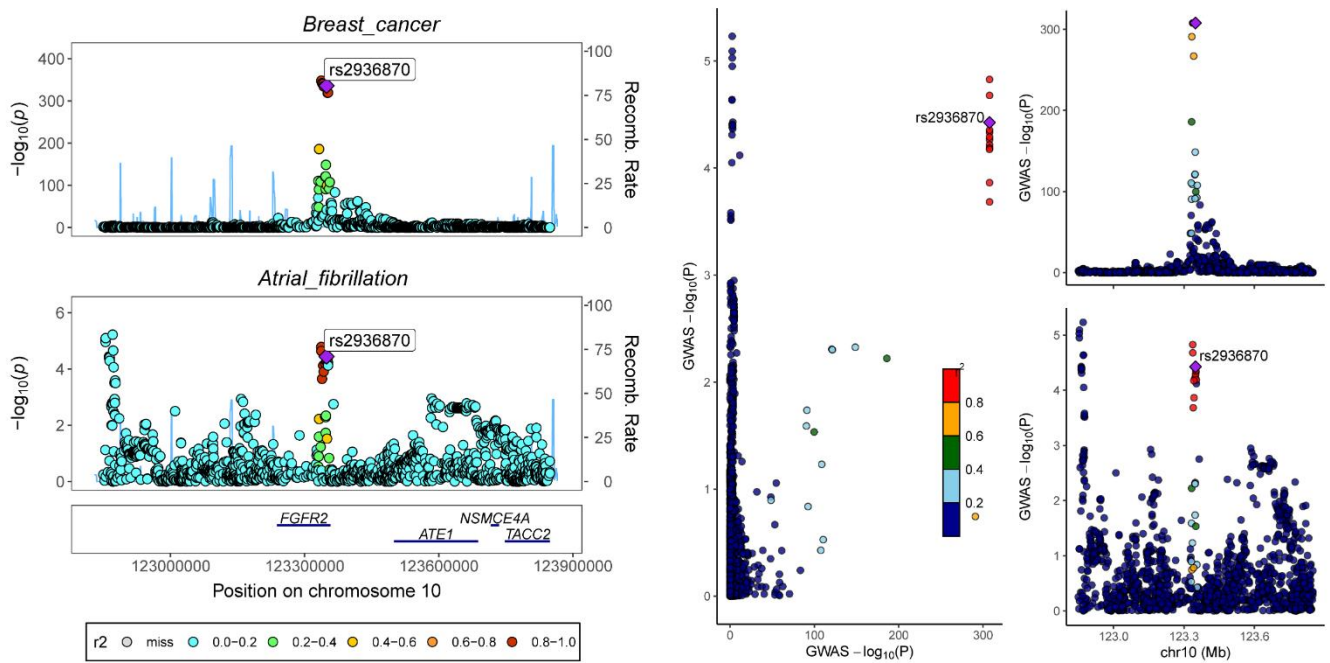
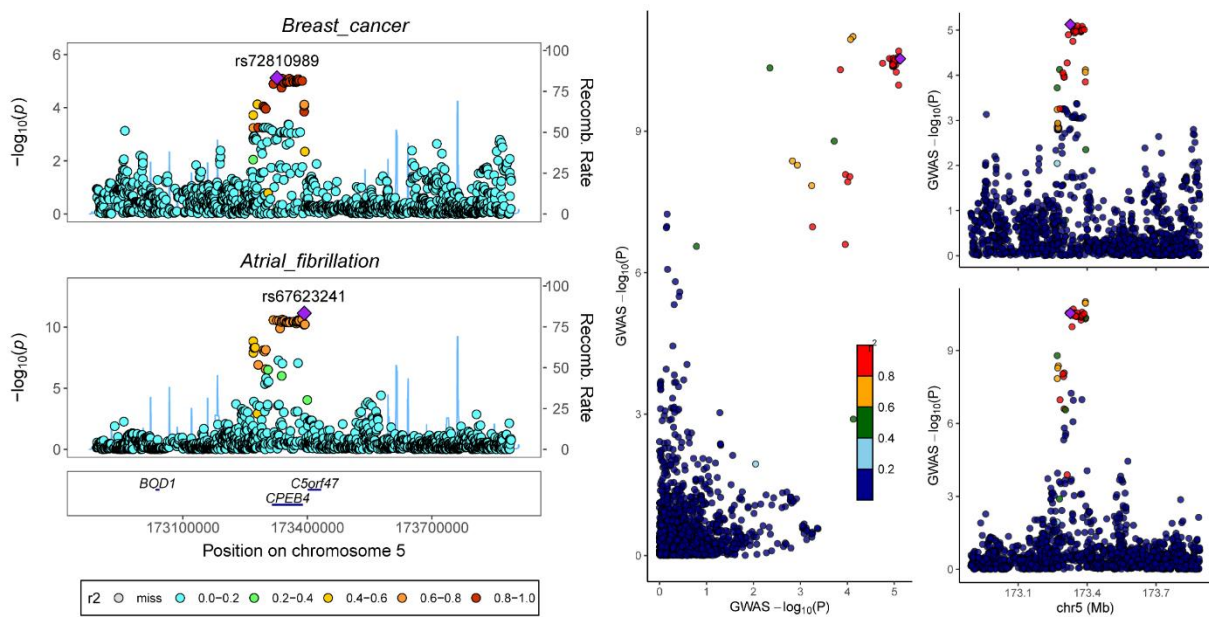


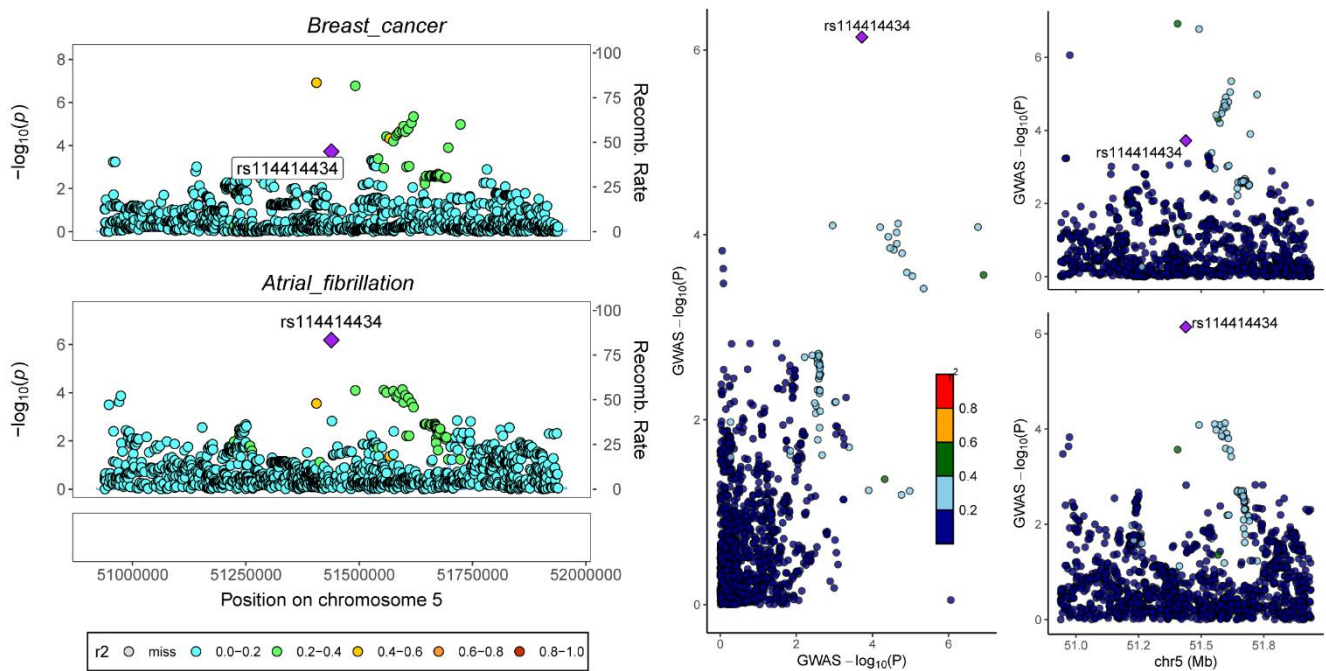
Supplementary Figure 1: Visualization of the colocalization results for the pleiotropic SNP rs1973765.



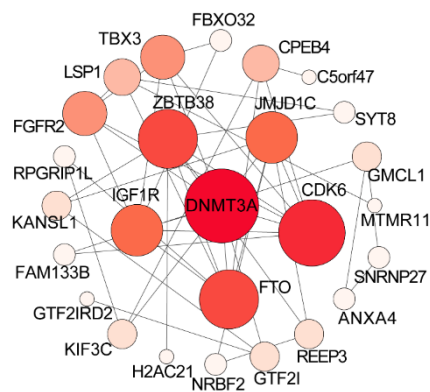
Supplementary Figure 2: Visualization of the colocalization results for the pleiotropic SNP rs2936870.



Supplementary Figure 7: Visualization of the colocalization results for the pleiotropic SNP rs56180201.



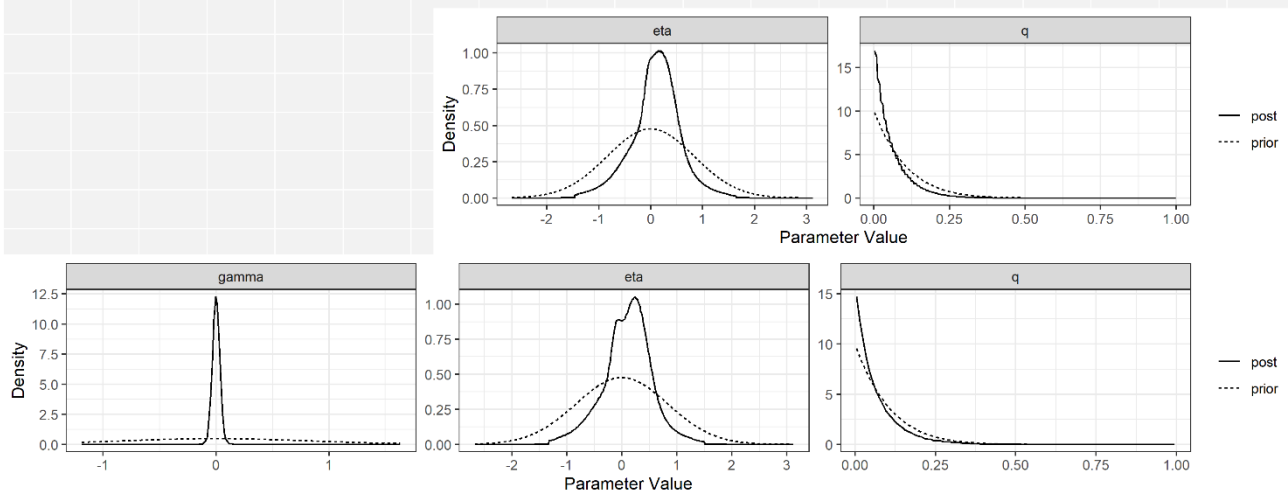
Supplementary Figure 8: Visualization of the colocalization results for the pleiotropic SNP rs114414434.



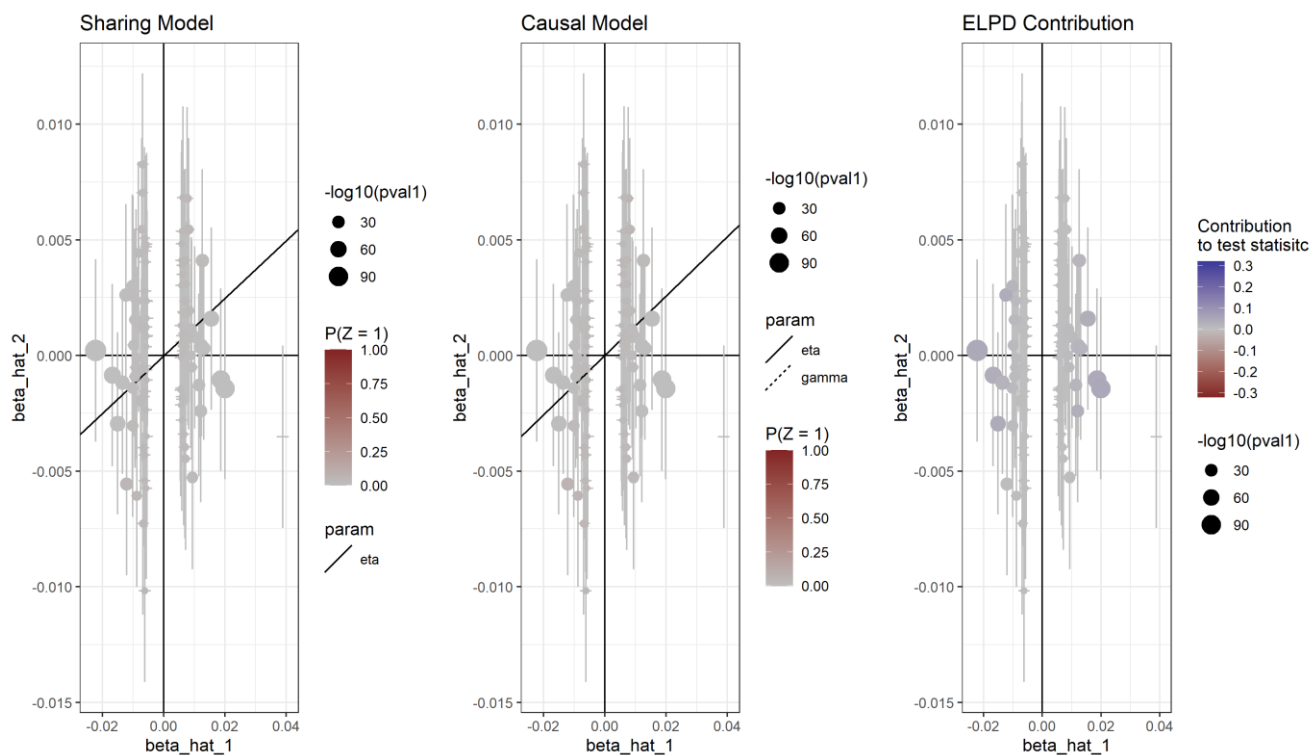
Supplementary Figure 9: PPI network diagram of 30 genes with potential pleiotropy, including 26 protein-coding genes. The size of the circles is proportional to the number of interactions between proteins. The lines between the circles represent the interactions between proteins.

	model1	model2	delta_elpd	se_delta_elpd	z	p
1	null	sharing	0.4	0.29	1.4	0.91
2	null	causal	1.4	0.50	2.8	1.00
3	sharing	causal	1.0	0.34	3.0	1.00

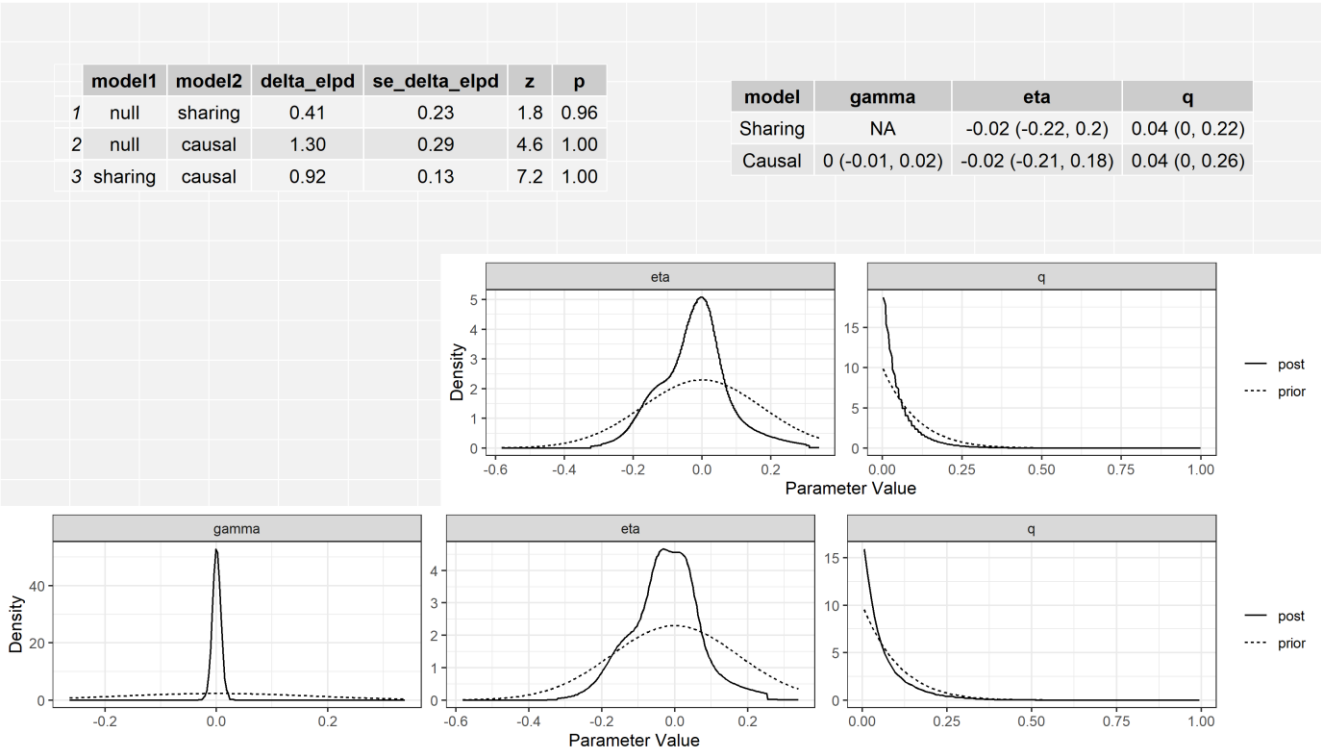
model	gamma	eta	q
Sharing	NA	0.12 (-0.97, 1.06)	0.04 (0, 0.22)
Causal	0 (-0.07, 0.07)	0.13 (-0.9, 1)	0.05 (0, 0.26)



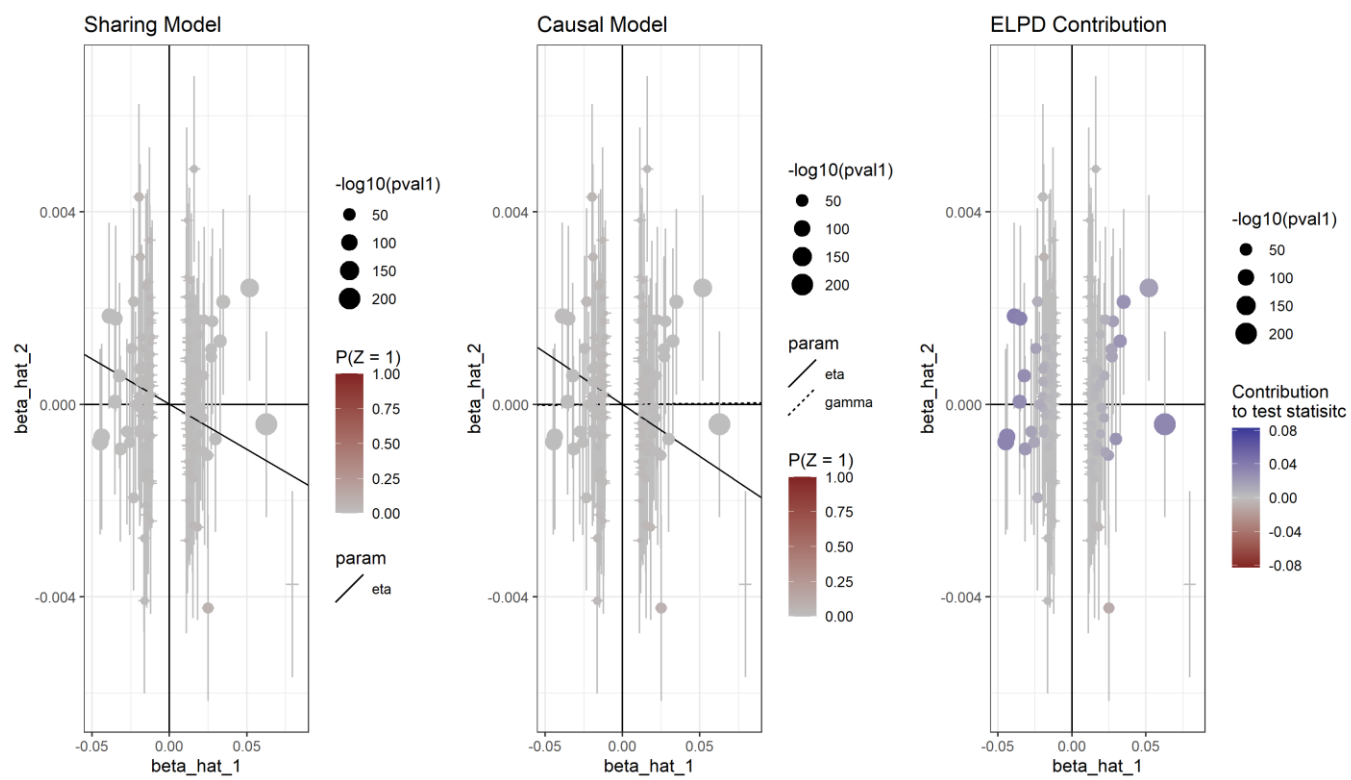
Supplementary Figure 10: Univariate MR analysis of atrial fibrillation on breast cancer, results of the CAUSE model.



Supplementary Figure 11: Scatter plot of the CAUSE model results from the univariate MR analysis of atrial fibrillation on breast cancer.



Supplementary Figure 12: Univariate MR analysis of breast cancer on atrial fibrillation, results of the CAUSE model.



Supplementary Figure 13: Scatter plot of the CAUSE model results from the univariate MR analysis of breast cancer on atrial fibrillation.