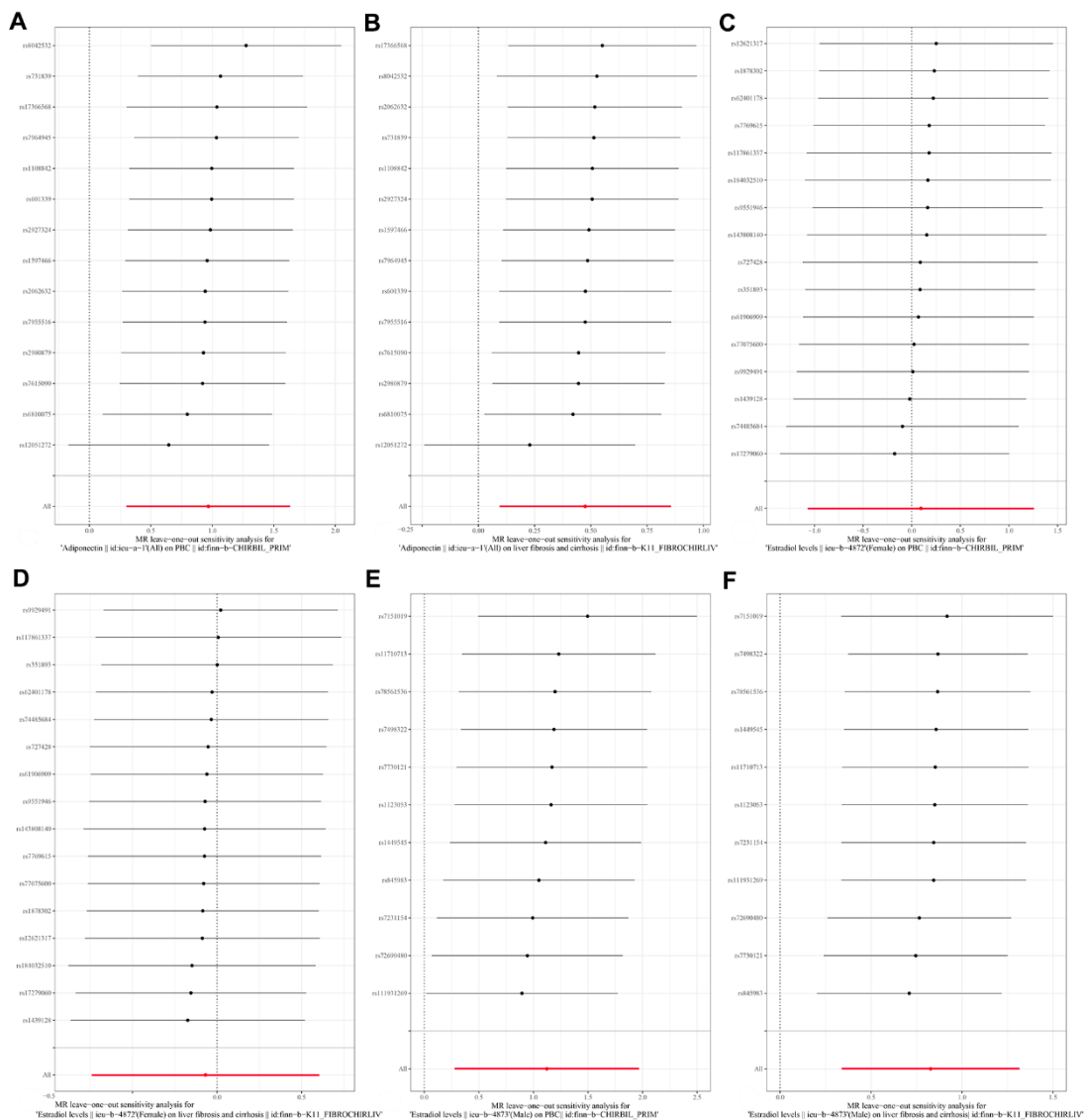


## Supplementary material

# Investigating the potential impact of sex hormones and adiponectin on the risk of liver fibrosis and cirrhosis: a Mendelian randomization study

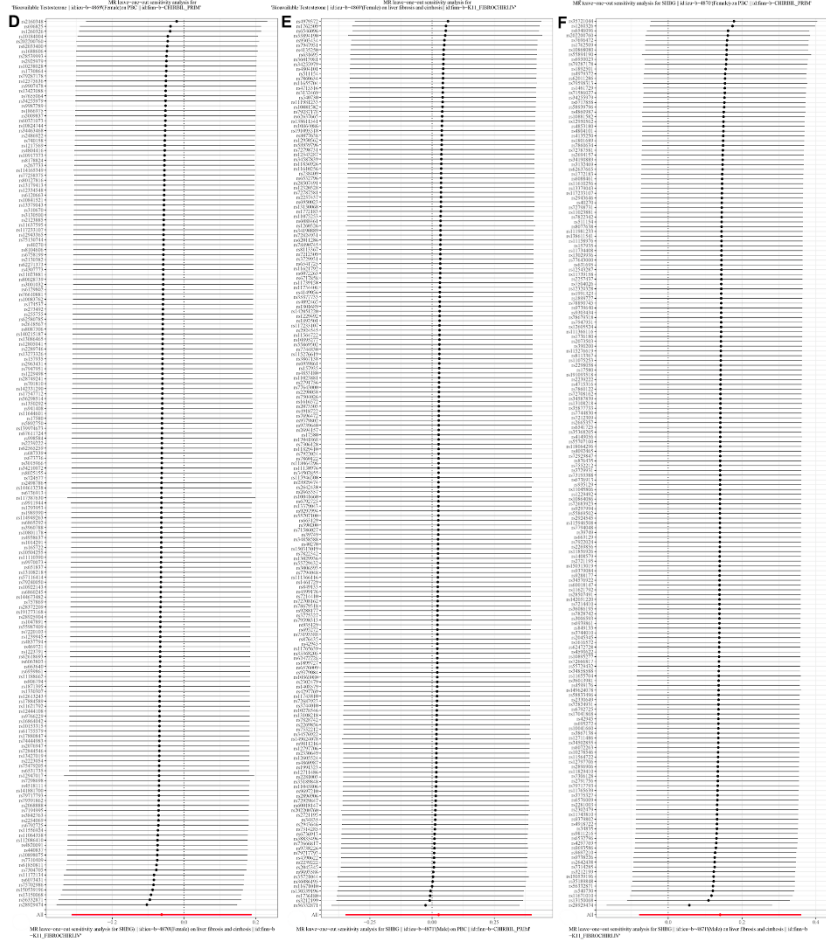
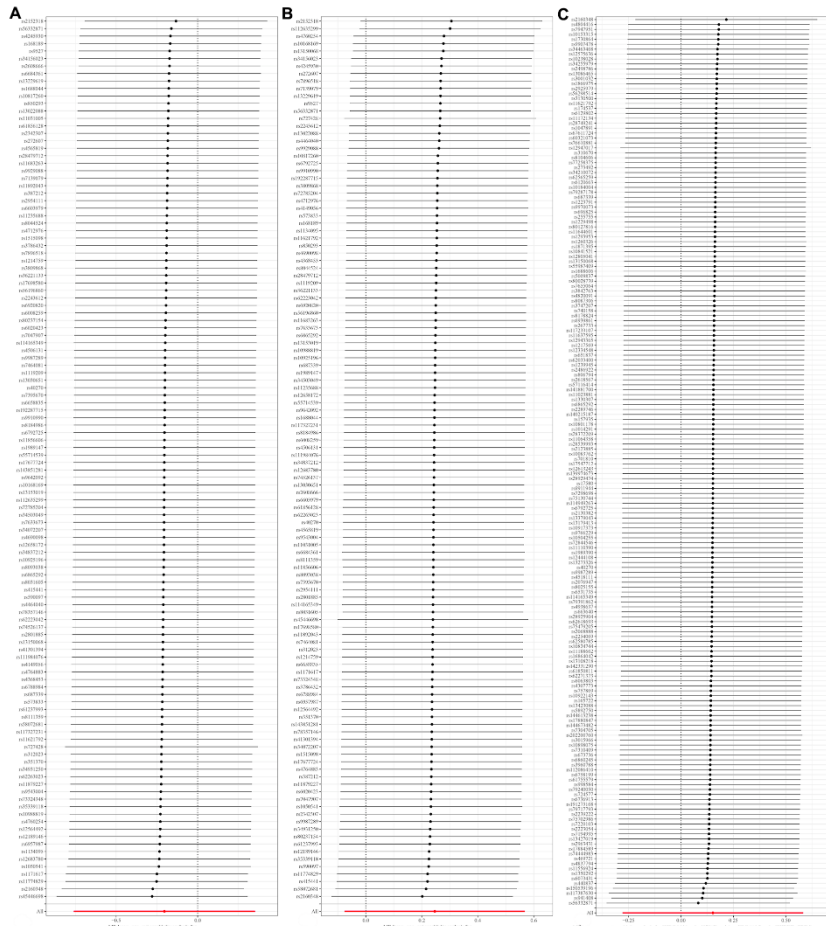
**Supplementary Figure S1.** Leave-one-out sensitivity tests of the causal relationship between: **(A)** adiponectin and PBC; **(B)** adiponectin and liver fibrosis and cirrhosis; **(C)** estradiol and PBC in females; **(D)** estradiol and liver fibrosis and cirrhosis in females; **(E)** estradiol and PBC in males; **(F)** estradiol and liver fibrosis and cirrhosis in males



**Supplementary Figure S2.** Leave-one-out sensitivity tests of the causal relationship between: (A) total testosterone and PBC in females; (B) total testosterone and liver fibrosis and cirrhosis in females; (C) total testosterone and PBC in males; (D) total testosterone and liver fibrosis and cirrhosis in males; (E) bioavailable testosterone and PBC in males; (F) bioavailable testosterone and liver fibrosis and cirrhosis in males



**Supplementary Figure S3.** Leave-one-out sensitivity tests of the causal relationship between: (A) bioavailable testosterone and PBC in females; (B) bioavailable testosterone and liver fibrosis and cirrhosis in females; (C) sex hormone-binding globulin and PBC in females; (D) sex hormone-binding globulin and liver fibrosis and cirrhosis in females; (E) sex hormone-binding globulin and PBC in males; (F) sex hormone-binding globulin and liver fibrosis and cirrhosis in males



Genomic tracks showing read alignments and coverage for different samples. Each track includes a list of genomic coordinates on the left and a corresponding plot of read depth on the right. The tracks are labeled with 'A', 'B', 'C', 'D', 'E', and 'F' at the top left.

**Supplementary Table SI.** GWAS summary data regarding exposure and outcomes

<b>Phenotype</b>	<b>Populations</b>	<b>Sample size (case)</b>	<b>Consortium</b>	<b>GWAS ID</b>
<b>Exposures</b>				
Estradiol		17134	IEU GWAS database	ieu-b-4873
Bioavailable testosterone	Males	184205	IEU GWAS database	ieu-b-4868
Total testosterone		199569	IEU GWAS database	ieu-b-4865
SHBG		185221	IEU GWAS database	ieu-b-4871
Estradiol		53391	IEU GWAS database	ieu-b-4872
Bioavailable testosterone	Females	180386	IEU GWAS database	ieu-b-4869
Total testosterone		199569	IEU GWAS database	ieu-b-4864
SHBG		214989	IEU GWAS database	ieu-b-4870
Adiponectin	All	39883	ADIPOGen	ieu-a-1
<b>Outcomes</b>				
Liver fibrosis and cirrhosis	All	214403 (811)	FinnGen R10	finn-b-K11_FIBROCHIRLIV
PBC		176861 (271)	FinnGen R10	finn-b-CHIRBIL_PRIM

GWAS – genome-wide association studies, FIBROCHIRLIV – liver fibrosis and cirrhosis, CHIRBIL\_PRIM – primary biliary cholangitis, PBC – primary biliary cirrhosis, SHBG – sex hormone-binding globulin

**Supplementary Table SII.** Other MR method about the causal effect of genetically predicted sex hormones, adiponectin and the risk of liver fibrosis, cirrhosis and PBC

Outcome	Exposure	Populations	Methods	SNPs	OR (95% CI)	<i>p</i>
Liver fibrosis and cirrhosis	Estradiol	Males	MR Egger	11	1.488 (0.403–5.503)	0.566
			Weighted median	11	1.835 (0.944–3.566)	0.073
			Simple mode	11	1.747 (0.684–4.457)	0.27
			Weighted mode	11	1.775 (0.801–3.938)	0.188
	Total testosterone		MR Egger	148	1.108 (0.638–1.923)	0.717
			Weighted median	148	1.106 (0.693–1.765)	0.672
			Simple mode	148	0.971 (0.378–2.489)	0.95
			Weighted mode	148	0.971 (0.606–1.555)	0.901
	Bioavailable testosterone	MR Egger	72	1.008 (0.371–2.736)	0.988	
		Weighted median	72	1.134 (0.603–2.133)	0.697	
		Simple mode	72	0.885 (0.258–3.038)	0.846	
		Weighted mode	72	0.885 (0.371–2.109)	0.783	
	SHBG	MR Egger	186	0.990 (0.651–1.505)	0.963	
		Weighted median	186	0.953 (0.643–1.414)	0.812	
		Simple mode	186	1.037 (0.460–2.337)	0.931	
		Weighted mode	186	0.840 (0.539–1.310)	0.443	
Estradiol	Females	MR Egger	16	1.168 (0.305–	0.824	

				4.468)	
		Weighted median	16	0.977 (0.394–2.423)	0.96
		Simple mode	16	1.067 (0.276–4.128)	0.927
		Weighted mode	16	1.082 (0.327–3.573)	0.899
		MR Egger	101	1.868 (0.988–3.532)	0.057
Total testosterone		Weighted median	101	1.312 (0.717–2.402)	0.379
		Simple mode	101	2.849 (0.736–11.033)	0.133
		Weighted mode	101	1.377 (0.771–2.462)	0.283
		MR Egger	118	1.454 (0.706–2.993)	0.312
Bioavailable testosterone		Weighted median	118	1.366 (0.802–2.326)	0.251
		Simple mode	118	2.374 (0.804–7.014)	0.12
		Weighted mode	118	1.365 (0.764–2.440)	0.295
		MR Egger	179	1.019 (0.632–1.645)	0.938
SHBG		Weighted median	179	0.960 (0.619–1.487)	0.854
		Simple mode	179	0.865 (0.344–2.177)	0.758
		Weighted mode	179	0.907 (0.576–1.428)	0.674
		MR Egger	14	2.066 (1.182–3.611)	0.026
Adiponectin	All	Weighted median	14	1.451 (0.855–2.463)	0.168
		Simple mode	14	1.450 (0.553–	0.463

				3.800)	
		Weighted mode	14	1.796 (1.090–2.958)	0.039
		MR Egger	11	1.747 (0.161–18.949)	0.657
	Estradiol	Weighted median	11	1.450 (0.455–4.620)	0.53
		Simple mode	11	1.507 (0.235–9.673)	0.674
		Weighted mode	11	1.353 (0.374–4.896)	0.655
		MR Egger	149	1.150 (0.473–2.796)	0.759
	Total testosterone	Weighted median	149	1.175 (0.556–2.483)	0.672
		Simple mode	149	1.559 (0.367–6.624)	0.549
		Weighted mode	149	1.031 (0.470–2.262)	0.94
PBC	Males	MR Egger	72	1.510 (0.282–8.072)	0.632
	Bioavailable testosterone	Weighted median	72	2.271 (0.759–6.801)	0.143
		Simple mode	72	1.021 (0.083–12.508)	0.987
		Weighted mode	72	1.983 (0.313–12.551)	0.469
		MR Egger	187	1.137 (0.579–2.234)	0.71
	SHBG	Weighted median	187	1.041 (0.576–1.883)	0.893
		Simple mode	187	1.023 (0.301–3.476)	0.971
		Weighted mode	187	1.023 (0.568–1.841)	0.939
	Estradiol	MR Egger	16	0.904 (0.079–	0.937

				10.408)	
		Weighted median	16	0.718 (0.158–3.263)	0.668
		Simple mode	16	1.047 (0.109–10.058)	0.969
		Weighted mode	16	0.811 (0.124–5.298)	0.829
		MR Egger	101	1.813 (0.605–5.428)	0.29
Total testosterone		Weighted median	101	1.254 (0.421–3.739)	0.685
		Simple mode	101	0.883 (0.105–7.446)	0.909
		Weighted mode	101	1.090 (0.384–3.096)	0.872
		MR Egger	118	1.677 (0.526–5.346)	0.384
Bioavailable testosterone		Weighted median	118	0.942 (0.374–2.372)	0.899
		Simple mode	118	0.929 (0.151–5.715)	0.937
		Weighted mode	118	1.023 (0.368–2.839)	0.966
		MR Egger	180	1.919 (0.894–4.120)	0.096
SHBG		Weighted median	180	1.122 (0.545–2.309)	0.754
		Simple mode	180	0.884 (0.213–3.674)	0.866
		Weighted mode	180	1.178 (0.527–2.632)	0.69
	All	MR Egger	14	2.832 (0.924–8.682)	0.094
Adiponectin		Weighted median	14	2.514 (0.973–6.492)	0.057
		Simple mode	14	4.429 (0.816–	0.108

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			24.045)	
	Weighted mode	14	2.650 (1.200– 5.854)	0.031

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SHBG – sex hormone-binding globulin, PBC – primary biliary cirrhosis, MR – mendelian randomization, SNPs – single-nucleotide polymorphisms, OR – odds ratio, CI – confidence intervals.