Descriptions of Supplementary Figures and Tables

Figure S1 The leave-one-out analysis for gut microbiota on aortic stenosis.

Figure S2 The scatter plots show overall effect of blood metabolites on aortic stenosis.

Figure S3 The leave-one-out analysis for inflammatory proteins on aortic stenosis.

Figure S4 The scatter plots show overall effect of gut microbiota on aortic stenosis.

Figure S5 The leave-one-out analysis for blood metabolites on aortic stenosis.

Figure S6 The scatter plots show overall effect of inflammatory proteins on aortic stenosis.

Figure S7 The leave-one-out analysis for immune cells on aortic stenosis.

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Table S1. Detailed information for genome-wide association studies involved in the present Mendelian randomization study.

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Figure S1. The leave-one-out analysis for gut microbiota on aortic stenosis.



0.6

Figure S2. The scatter plots show overall effect of blood metabolites on aortic stenosis.

0.1









Figure S3. The leave-one-out analysis for inflammatory proteins on aortic stenosis.





0.10 0.15 0.20 SNP effect on PWY 7323 (superpathway of GDP mannose derived O antigen building blocks biosynthesis)

























Figure S5. The leave-one-out analysis for blood metabolites on aortic stenosis.



Figure S6. The scatter plots show overall effect of inflammatory proteins on aortic stenosis.



Figure S7. The leave-one-out analysis for immune cells on aortic stenosis.



Figure S8. The scatter plots show overall effect of immune cells on aortic stenosis.