



## OPEN Lipid metabolism and gallstone disease risk: a multicenter study

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This study aimed to explore the associations between various lipid parameters and the risk of gallstone disease (GSD). A multicenter cross-sectional study was conducted to explore the associations between nine lipid parameters and the risk of GSD. Multivariable logistic regression models, adjusted for covariates, were employed to evaluate these associations. Stratified analyses by age and sex were performed to assess population-specific effects. Data from multiple centers were integrated using meta-analysis to generate pooled odds ratios with 95% confidence intervals. Trial Sequential Analysis (TSA) was implemented to determine the sufficiency of the sample size and the robustness of the associations under cumulative evidence conditions. This multicenter study included 580,935 participants, with a GSD prevalence of 7.43%. After adjusting for covariables, the combined multicenter results showed that total cholesterol, high-density and low-density lipoprotein cholesterol, and non-HDL cholesterol were negatively associated with GSD risk. In contrast, the Castelli risk index I and II, atherogenic coefficient, and atherogenic index of plasma were positively associated with the risk of GSD. Subgroup analyses and sensitivity analyses yielded consistent results. Stratified analysis revealed distinct lipid-gallstone associations between people with cholecystectomy and gallbladder stones. TSA revealed that the required amount of information has been achieved, and the results are conclusive. This study suggests significant associations between lipid metabolism and the risk of GSD. There are differences in the relationship between blood lipid indexes and cholecystectomy and gallstones. These findings not only enhance our understanding of the relationship between lipid metabolism and GSD but also provide a novel perspective for risk assessment and prevention strategies.

**Keywords** Blood lipids, Cholesterol, Non-high-density lipoprotein, Castelli risk index, Atherosclerosis index, Gallstones

### Abbreviations

GSD	Gallstone disease
TC	Total cholesterol
TG	Triglycerides
LDL-C	Low-density lipoprotein cholesterol
HDL-C	High-density lipoprotein cholesterol
non-HDL-C	Non-high-density lipoprotein-cholesterol
CRI I	Castelli risk index-I
CRI 2	Castelli risk index-II
AC	Atherogenic coefficient

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AIP	Atherogenic index of plasma
FBG	Fasting blood glucose
SBP	Systolic blood pressure
DBP	Diastolic blood pressure
T-bil	Total bilirubin
D-bil	Direct bilirubin
ALT	Alanine transaminase
AST	Aspartate transaminase
GGT	Gamma-glutamyl transferase
Cr	Creatinine
UA	Uric acid
UN	Urea nitrogen
ALB	Albumin
GLB	Globulin
(A/G)	Albumin-to-globulin ratio

Gallstone disease (GSD) is one of the most prevalent disorders within the digestive system<sup>1</sup>. Its global prevalence varies significantly by region, with the highest rates observed in the Americas, followed by Africa, Europe, and Asia<sup>2,3</sup>. Notably, in China, the prevalence of GSD is significantly higher than the overall level in Asia and is increasing rapidly<sup>4</sup>. This trend is partly attributed to the acceleration of urbanization and the rise in unhealthy lifestyle patterns, such as high-fat diets and insufficient physical activity, which have contributed to the annual increase in GSD incidence<sup>5,6</sup>. Although the mortality rate associated with GSD is relatively low, failure to promptly treat and manage GSD and its associated complications can result in severe consequences, including liver damage, liver failure, biliary tract infections (e.g., cholangitis), pancreatitis, diabetes, and increased all-cause mortality<sup>7–11</sup>. Consequently, early detection, preventive measures, and timely intervention for GSD are of substantial public health significance.

Despite significant advances in hepatobiliary pathophysiology, the precise mechanisms underlying GSD formation remain incompletely understood. Numerous studies have demonstrated that disruptions in serum sterol metabolism, imbalances in bile components, insulin resistance (IR), and other factors are intricately associated with the onset of GSD<sup>12</sup>. Current epidemiological investigations have predominantly focused on conventional lipid parameters, including total cholesterol (TC), triglycerides (TG), low-density lipoprotein cholesterol (LDL-C), and high-density lipoprotein cholesterol (HDL-C), for GSD risk stratification<sup>13</sup>. However, these isolated measurements may inadequately capture the complex lipid metabolic derangements associated with GSD risk.

With growing recognition of the association between abnormal lipid profiles and various diseases, non-traditional lipid indicators—such as non-high-density lipoprotein cholesterol (non-HDL), the Castelli risk index, and the atherogenic index—have garnered attention as complementary measures to traditional lipid metrics<sup>14–16</sup>. Previous studies have explored the associations between these non-traditional lipid parameters and the risk of GSD<sup>17–19</sup>. However, due to limitations in data and other factors, the relationships between lipid-related indicators and the risk of GSD still require in-depth validation using large sample sizes. Moreover, most prior research has focused on populations in the United States, with relatively fewer studies conducted on Chinese populations.

Given these gaps, this study conducts a large-scale, multicenter, cross-sectional investigation using population-based data to examine the relationship between both traditional and non-traditional lipid indicators and the risk of GSD formation. This research aims to fill the existing knowledge gaps and provide a more comprehensive understanding of the role of lipid metabolism in GSD, particularly in the Chinese population. Trial Sequential Analysis (TSA) was also implemented to determine the sufficiency and the robustness of the results.

## Result

### Baseline characteristics of subjects in four hospitals

This cross-sectional study recruited a total of 580,935 participants from four hospitals, including 43,170 individuals diagnosed with gallstone disease (GSD) and 537,765 individuals without GSD. The baseline characteristics and laboratory findings of the participants are presented in Table 1.

Among the 142,088 participants enrolled from the hospital in Liangjiang New Area, 7.66% (10,881 cases) were diagnosed with GSD. In the hospital in Kaizhou District, 8.81% (25,735 cases) of the 292,210 participants were identified with GSD. For the Hospital in Qianjiang District (70,615 participants) and the hospital in Beijing (76,022 participants), the detection rates of GSD were 3.41% (2,413 cases) and 5.45% (4,141 cases), respectively.

Compared with the non-GSD group, the GSD group exhibited significantly advanced age, higher level of urea nitrogen (UN), uric acid (UA), globulin (GLB), and higher proportion of abnormal blood glucose, kidney stones, fatty liver, and hypertension. In contrast, albumin levels and the albumin-to-globulin ratio (A/G) were significantly higher in the healthy control group. Among the nine lipid profile indicators, all were significantly elevated in the GSD group compared with the non-GSD group, except for high-density lipoprotein cholesterol (HDL-C) (all  $P < 0.001$ ).

### The associations between lipid profile indexes and the risk of GSD

When lipid indicators were analyzed as categorical variables, taking the results from the hospital in Liangjiang New Area as an example, the multivariable logistic regression analysis revealed that, higher level of TC, HDL-C, LDL-C, non-HDL were negatively associated the risk of GSD. In contrast, the CRI1, CRI2, AC, and AIP were

	Hospital in Liangjiang New Area, Chongqing (n = 142,088)				Hospital in Kaizhou District, Chongqing (n = 292,210)			
	GSD (n = 10,881)	non-GSD (n = 131,207)	statistic	P-value	GSD (n = 25,735)	non-GSD (n = 266,475)	statistic	P value
Age (year)	52 (42,61)	38 (31,50)	-76.58	<0.001	52 (44,66)	44 (34,52)	-96.696	<0.001
Female [n(%)]	5093 (46.8)	54,428 (41.5)	116.99	<0.001	14,259 (55.4)	120,290 (45.1)	995.542	<0.001
BMI (kg/m <sup>2</sup> )	24.43 (22.35,28.62)	23.12 (20.93,25.36)	-39.146	<0.001	25.0 (22.9,27.2)	23.8 (21.6,26)	-56.989	<0.001
Cr	69 (69,81)	71 (59,81)	-4.042	<0.001	67 (57.3,79.5)	69.7 (58.7,81.2)	-16.506	<0.001
UN (mmol/L)	5.1 (4.3,6.1)	5.0 (4.2,5.8)	-13.9	<0.001	5.2 (4.3,6.2)	5.1 (4.2,6.0)	-12.85	<0.001
UA (μmol/L)	341 (281,408)	339 (275,409)	-2.96	<0.001	328.1 (272.8,391.7)	327.5 (268.9,393.5)	-2.14	0.032
SBP (mmHg)	128 (116, 141)	121 (111, 133)	-37.262	<0.001	NA	NA	NA	NA
DBP (mmHg)	77 (70,86)	73 (66,81)	-31.112	<0.001	NA	NA	NA	NA
ALB	46 (45,48)	47 (45,49)	-29.161	<0.001	44.6 (41.9,46.5)	45.1 (42.3,47.1)	-22.802	<0.001
GLB	30 (27,32)	29 (26,32)	-11.481	<0.001	30.2 (27.9,32.7)	29.6 (27.4,32)	-23.244	<0.001
A/G	1.6 (1.4, 1.8)	1.6 (1.5, 1.8)	-20.183	<0.001	1.5 (1.37, 1.64)	1.55 (1.42, 1.68)	-33.932	<0.001
TP	76 (73,79)	76 (73,79)	-7.248	<0.001	74.5 (70.5,77.8)	74.6 (70.6,77.8)	-0.982	0.326
TBIL (μmol/L)	12.8 (10.3, 16.3)	12.7 (10.1, 16.1)	-3.417	<0.001	NA	NA	NA	NA
DBIL (μmol/L)	3.7 (2.9,4.8)	3.8 (3.0,4.9)	-4.981	<0.001	NA	NA	NA	NA
AST	21 (18,26)	21 (17,26)	-10.99	<0.001	24 (20,29)	23 (19,29)	-13.615	<0.001
ALT	22 (16,33)	21 (14,32)	-12.438	<0.001	23 (16,34)	21 (15,32)	-18.579	<0.001
GGT	26 (17,42)	22 (15,37)	-22.902	<0.001	27 (18,45)	23 (16,39)	-29.286	<0.001
ALP	NA	NA	NA	NA	88 (73, 106)	84 (69, 101)	-29.76	<0.001
TC	4.97 (4.34,5.60)	4.81 (4.23,5.44)	-15.137	<0.001	5.06 (4.45,5.73)	4.94 (4.34,5.60)	-18.144	<0.001
TG	1.49 (1.04,2.17)	1.22 (0.86, 1.85)	-32.237	<0.001	1.55 (1.08,2.28)	1.31 (0.92,1.96)	-43.337	<0.001
HDL-C	1.32 (1.12, 1.56)	1.37 (1.16, 1.61)	-13.164	<0.001	1.31 (1.13,1.54)	1.35 (1.16,1.58)	-18.271	<0.001
LDL-C	3.01 (2.49,3.57)	2.87 (2.36,3.42)	-15.727	<0.001	2.62 (2.22,3.05)	2.53 (2.14,2.97)	-19.129	<0.001
non-HDL-C	3.60 (3.00,4.25)	3.40 (2.82,4.04)	-20.267	<0.001	3.71 (3.11,4.35)	3.55 (2.94,4.21)	-25.069	<0.001
CRI1	3.74 (3.10,4.47)	3.50 (2.87,4.25)	-22.099	<0.001	3.84 (3.22,4.54)	3.63 (3.00,4.38)	-29.179	<0.001
CRI2	2.31 (1.77,2.87)	2.12 (1.59,2.73)	-19.708	<0.001	2.01 (1.61,2.45)	1.89 (1.48,2.36)	-26.825	<0.001
AC	2.74 (2.10,3.47)	2.50 (1.87,3.25)	-22.099	<0.001	2.84 (2.22,3.54)	2.63 (2.00,3.38)	-29.179	<0.001
AIP	0.05 (-0.15,0.26)	-0.05 (-0.26,0.18)	-29.652	<0.001	0.07 (-0.13,0.28)	-0.02 (-0.22,0.20)	-40.801	<0.001
Hypertension [n(%)]	4007 (36.8)	25,963 (19.8)	1752.47	<0.001	8470 (32.9)	57,276 (21.5)	1754.768	<0.001
Fatty liver [n(%)]	4516 (41.5)	33,838 (25.8)	1258.959	<0.001	10,610 (41.2)	70,678 (26.5)	2527.162	<0.001
Renal calculus [n(%)]	599 (5.5)	5568 (4.2)	38.502	<0.001	949 (3.7)	8623 (3.2)	15.108	<0.001
Abnormal blood glucose [n(%)]	1483 (13.6)	6510 (5.0)	1860.984	<0.001	3174 (12.3)	14,006 (5.3)	3191.447	<0.001
	Hospital in Qianjiang District, Chongqing (n = 70,615)				Hospital in Beijing (n = 76,022)			
	GSD (n = 2413)	non-GSD (n = 68,202)	statistic	P value	GSD (n = 4141)	non-GSD (n = 71,881)	statistic	P value
Age (year)	51 (44,61)	33 (22,46)	-48.761	<0.001	54 (45,67)	42 (33,51)	-52.155	<0.001
Female [n(%)]	1304 (54.0)	30,855 (45.2)	72.77	<0.001	1711 (41.3)	29,037(40.4)	1.384	0.239
BMI (kg/m <sup>2</sup> )	24.63 (22.52,26.82)	23.69 (21.45,25.96)	-13.794	<0.001	25.95 (23.73,28.26)	24.97 (22.44,27.48)	-16.448	<0.001
Cr	NA	NA	NA	NA	NA	NA	NA	NA
UN (mmol/L)	4.94 (4.06,5.92)	4.63 (3.77,5.59)	-11.405	<0.001	NA	NA	NA	NA
UA (μmol/L)	NA	NA	NA	NA	337 (281,397)	325 (266,388)	-8.337	<0.001
SBP (mmHg)	129 (117, 141)	124 (113, 135)	-14.814	<0.001	125 (115, 137)	119 (110, 130)	-23.071	<0.001
DBP (mmHg)	79 (71,87)	76 (69,84)	-10.291	<0.001	76 (70,82)	74 (67,80)	-10.848	<0.001
ALB	NA	NA	NA	NA	NA	NA	NA	NA
GLB	28.3 (25.1,31.2)	27.7 (24.7,30.6)	-6.342	<0.001	NA	NA	NA	NA
A/G	NA	NA	NA	NA	NA	NA	NA	NA
TP	NA	NA	NA	NA	NA	NA	NA	NA
TBIL (μmol/L)	13.3 (10.6, 16.9)	13.6 (10.3, 17.7)	-1.598	0.110	NA	NA	NA	NA
DBIL (μmol/L)	NA	NA	NA	NA	NA	NA	NA	NA
AST	23.0 (19.1,28.4)	22.1 (18.0,28.2)	-5.439	<0.001	21 (18,25)	20 (17,24)	-8.004	<0.001
ALT	23.0 (16.2,33.9)	19.4 (13.0,32.0)	-12.18	<0.001	19 (15,28)	19 (13,28)	-4.212	<0.001
GGT	28.4 (18.0,88.0)	33.3 (17.1,61.6)	-4.253	<0.001	NA	NA	NA	NA
ALP	76 (61.0,93.0)	74.0 (57.9,91.9)	-4.851	<0.001	NA	NA	NA	NA
TC	5.02 (4.39,5.63)	4.91 (4.31,5.53)	-4.844	<0.001	4.88 (4.25,5.56)	4.73 (4.15,5.39)	-8.357	<0.001
Continued								

	Hospital in Qianjiang District, Chongqing (n = 70,615)				Hospital in Beijing (n = 76,022)			
	GSD (n = 2413)	non-GSD (n = 68,202)	statistic	P value	GSD (n = 4141)	non-GSD (n = 71,881)	statistic	P value
TG	1.77 (1.23,2.50)	1.70 (1.06,2.61)	-4.945	<0.001	1.38 (0.98,2.00)	1.24 (0.84,1.90)	-11.833	<0.001
HDL-C	1.31 (1.10,1.52)	1.36 (1.13,1.59)	-6.767	<0.001	1.28 (1.10,1.50)	1.31 (1.12,1.54)	-5.683	<0.001
LDL-C	2.70 (2.24,3.18)	2.62 (2.14,3.12)	-4.704	<0.001	3.05 (2.50,3.57)	2.93 (2.43,3.46)	-7.730	<0.001
non-HDL-C	3.69 (3.08,4.27)	3.53 (2.94,4.16)	-7.094	<0.001	3.56 (2.95,4.19)	3.39 (2.78,4.05)	-10.447	<0.001
CRI1	3.83 (3.18,4.59)	3.60 (2.99,4.41)	-8.865	<0.001	3.80 (3.15,4.51)	3.62 (2.94,4.37)	-10.262	<0.001
CRI2	2.08 (1.63,2.58)	1.94 (1.48,2.50)	-7.777	<0.001	2.38 (1.86,2.94)	2.25 (1.71,2.85)	-8.760	<0.001
AC	2.83 (2.18,3.59)	2.60 (1.99,3.41)	-8.865	<0.001	2.80 (2.15,3.51)	2.62 (1.94,3.37)	-10.262	<0.001
AIP	0.14 (-0.07,0.33)	0.10 (-0.14,0.32)	-6.423	<0.001	0.04 (-0.15,0.24)	-0.02 (-0.24,0.21)	-11.260	<0.001
Hypertension [n(%)]	747 (31.0)	12,430 (18.2)	248.904	<0.001	1931 (46.6)	18,618 (25.9)	853.079	<0.001
Fatty liver [n(%)]	1066 (44.1)	8074 (11.8)	2163.018	<0.001	1963 (47.4)	26,172 (36.4)	202.998	<0.001
Renal calculus [n(%)]	63 (2.6)	667 (1.0)	60.737	<0.001	290 (7.0)	2302 (3.2)	171.735	<0.001
Abnormal blood glucose [n(%)]	NA	NA	NA	NA	812 (19.6)	6395 (8.9)	588.703	<0.001

**Table 1.** Baseline characteristics of GSD and non-GSD subjects recruited at each hospital. BMI = Body Mass Index; Cr = Creatinine; UN = Urea Nitrogen; UA = Uric Acid; SBP = Systolic Blood Pressure; DBP = Diastolic Blood Pressure; ALB = Albumin; GLB = Globulin; A/G = Albumin-to-globulin Ratio; TP = Total Protein; TBIL = Total Bilirubin; DBIL = Direct Bilirubin; AST = Aspartate Transaminase; ALT = Alanine Transaminase; GGT = Gamma-glutamyl Transferase; ALP = Alkaline Phosphatase; TC = Total Cholesterol; TG = Triglycerides; HDL-C = High-density Lipoprotein Cholesterol; LDL-C = Low-density Lipoprotein Cholesterol; non-HDL-C = Non-high-density Lipoprotein-Cholesterol; CRI1 = Castelli Risk Index—I; CRI2 = Castelli Risk Index—II; AC = The Atherogenic Coefficient; AIP = The Atherogenic Index of Plasma.

positively associated the GSD risk. No significant associations were found between TG and GSD risk (Table 2). The results were not entirely consistent across other centers.

When lipid indicators were analyzed as continuous variables, again taking the results from the hospital in Liangjiang New Area as an example, each unit or standard deviation (SD) increase in TC, HDL-C, LDL-C, non-HDL were negatively associated with GSD risk. Conversely, each unit or SD increase in TG, CRI1, CRI2, AC, and AIP were positively associated with the risk of GSD (Table 2). The results were not completely consistent across other centers.

### The pooled associations between lipid profile indexes and GSD risk

The meta-analysis pooled results from four centers demonstrated significant protective associations between elevated levels of conventional lipid markers (TC, HDL-C, LDL-C, and non-HDL cholesterol) and gallstone disease risk. Conversely, higher values of non-traditional indices (CRI1, CRI2, atherogenic coefficient and AIP) demonstrated significant positive associations with GSD risk. Triglyceride levels, however, showed no statistically significant association with gallstone risk in categorical analyses (Fig. 1).

When analyzed as continuous variables, consistent protective effects were observed in TC, HDL-C, LDL-C, and non-HDL cholesterol, with each unit increment corresponding to significantly reduced GSD risk. In contrast, per-unit increases in TG, CRI1, AC and AIP were significantly associated with elevated GSD risk. No significant association was detected for continuous CRI2 measurements (pooled OR = 1.010; 95%CI: 0.980–1.041).

The direction and significance of associations remained robust when scaled by standard deviation increments. Each standard deviation (SD) increase in conventional lipid parameters persistently correlated with reduced GSD risk, while SD increments in CRI1, AC and AIP maintained significant positive associations. Neither TG nor CRI2 exhibited statistically significant relationships with GSD risk when analyzed per SD increase.

These associations remained consistent when stratified by age and sex (Table 3).

### Comparison of results between cholecystectomy and gallstones and sensitivity analysis excluding specific metabolic diseases structure

We further stratified the outcomes into cholecystectomy and gallbladder stones. Due to the unavailability of stratified outcome data from Hospital in Qianjiang District, the analysis was restricted to the three remaining centers, with detailed results presented in Table SW. Meta-analysis of the three centers revealed that when lipids were treated as categorical variables, TC, HDL-C, CRI1, CRI2, AC, and AIP were significantly associated with gallstone formation, with TC and HDL-C acting as protective factors and the remainder as risk factors. These associations were largely consistent with those observed in the cholecystectomy population; however, in the latter, we also identified a positive association between TG and cholecystectomy, as well as inverse associations for LDL-C and non-HDL-C.

To evaluate the robustness of our findings, we conducted a series of sensitivity analyses by systematically excluding specific subpopulations. As detailed in Table SY, exclusion of individuals with fatty liver did not materially alter the observed associations. Similarly, when obese individuals (BMI  $\geq$  28 kg/m<sup>2</sup>) were excluded (Table SZ), and when participants with dysglycemia (fasting blood glucose > 7.0 mmol/L) were excluded (Table

	Hospital in Liangjiang New Area, Chongqing		Hospital in Kaizhou District, Chongqing		Hospital in Qianjiang District, Chongqing		Hospital in Beijing	
Index	OR (95%)	P-value	OR (95%)	P-value	OR (95%)	P-value	OR (95%)	P-value
TC, mmol/L								
< 3.1	ref		ref		ref		ref	
3.1–5.7	<b>0.652 (0.554,0.768)</b>	<0.001	<b>0.730 (0.646,0.824)</b>	<0.001	0.815 (0.592,1.122)	0.210	0.820 (0.636,1.057)	0.126
> 5.7	<b>0.589 (0.498,0.697)</b>	<0.001	<b>0.633 (0.560,0.717)</b>	<0.001	0.720 (0.518,1.001)	0.051	<b>0.764 (0.587,0.993)</b>	0.044
TG, mmol/L								
< 0.4	ref		ref		ref		ref	
0.4–1.7	0.724 (0.504,1.041)	0.081	1.099 (0.733,1.649)	0.648	<b>3.915 (2.206,6.948)</b>	<0.001	1.115 (0.608,2.045)	0.724
> 1.7	0.838 (0.582,1.206)	0.341	1.249 (0.832,1.875)	0.284	<b>3.695 (2.081,6.563)</b>	<0.001	1.127 (0.613,2.073)	0.701
HDL-C, mmol/L								
< 0.9	ref		ref		ref		ref	
0.9–2.0	<b>0.838 (0.768,0.915)</b>	<0.001	<b>0.875 (0.820,0.933)</b>	<0.001	1.104 (0.940,1.297)	0.226	<b>0.834 (0.730,0.953)</b>	0.008
> 2.0	<b>0.622 (0.544,0.712)</b>	<0.001	<b>0.684 (0.623,0.752)</b>	<0.001	0.900 (0.671,1.207)	0.482	<b>0.732 (0.587,0.913)</b>	0.006
LDL-C, mmol/L								
< 2.07	ref		ref		ref		ref	
2.07–3.1	<b>0.910 (0.851,0.973)</b>	<0.001	<b>0.916 (0.884,0.950)</b>	<0.001	0.943 (0.842,1.055)	0.302	0.932 (0.830,1.046)	0.232
> 3.1	<b>0.853 (0.797,0.913)</b>	<0.001	<b>0.839 (0.804,0.876)</b>	<0.001	<b>0.855 (0.754,0.970)</b>	0.015	0.907 (0.807,1.019)	0.100
non-HDL, mmol/L								
< P25	ref		ref		ref		ref	
IQR	<b>0.939 (0.889,0.993)</b>	0.026	<b>0.955 (0.922,0.989)</b>	0.011	0.911 (0.815,1.019)	0.102	0.951 (0.872,1.038)	0.260
> P75	<b>0.926 (0.870,0.985)</b>	0.015	<b>0.858 (0.824,0.894)</b>	<0.001	<b>0.851 (0.749,0.966)</b>	0.013	<b>0.881 (0.798,0.972)</b>	0.011
CRI 1								
< P25	ref		ref		ref		ref	
IQR	<b>1.196 (1.130,1.266)</b>	<0.001	<b>1.125 (1.084,1.167)</b>	<0.001	1.054 (0.937,1.185)	0.383	1.037 (0.947,1.135)	0.439
> P75	<b>1.239 (1.158,1.326)</b>	<0.001	<b>1.116 (1.068,1.166)</b>	<0.001	1.023 (0.893,1.171)	0.747	1.010 (0.908,1.124)	0.851
CRI 2								
< P25	ref		ref		ref		ref	
IQR	<b>1.163 (1.099,1.230)</b>	<0.001	<b>1.116 (1.077,1.158)</b>	<0.001	1.082 (0.963,1.215)	0.186	1.018 (0.932,1.113)	0.691
> P75	<b>1.183 (1.108,1.264)</b>	<0.001	<b>1.107 (1.060,1.155)</b>	<0.001	1.022 (0.894,1.169)	0.748	1.002 (0.903,1.111)	0.970
AC								
< P25	ref		ref		ref		ref	
IQR	<b>1.196 (1.130,1.266)</b>	<0.001	<b>1.125 (1.084,1.167)</b>	<0.001	1.054 (0.937,1.185)	0.383	1.037 (0.947,1.135)	0.439
> P75	<b>1.239 (1.158,1.326)</b>	<0.001	<b>1.116 (1.068,1.166)</b>	<0.001	1.023 (0.893,1.171)	0.747	1.010 (0.908,1.124)	0.851
AIP								
< P25	ref		ref		ref		ref	
IQR	<b>1.404 (1.323,1.491)</b>	<0.001	<b>1.301 (1.252,1.351)</b>	<0.001	<b>1.175 (1.045,1.321)</b>	0.007	<b>1.225 (1.114,1.347)</b>	<0.001
> P75	<b>1.556 (1.448,1.671)</b>	<0.001	<b>1.444 (1.379,1.512)</b>	<0.001	1.048 (0.912,1.204)	0.511	<b>1.227 (1.094,1.375)</b>	<0.001
TC, per unit	<b>0.937 (0.917,0.958)</b>	<0.001	<b>0.914 (0.902,0.927)</b>	<0.001	<b>0.942 (0.900,0.986)</b>	0.010	<b>0.942 (0.909,0.975)</b>	<0.001
TG, per unit	<b>1.036 (1.024,1.049)</b>	<0.001	<b>1.030 (1.021,1.038)</b>	<0.001	1.018 (0.990,1.047)	0.198	0.975 (0.948,1.002)	0.069
HDL-C, per unit	<b>0.697 (0.650,0.748)</b>	<0.001	<b>0.741 (0.711,0.774)</b>	<0.001	<b>0.873 (0.763,0.998)</b>	0.047	<b>0.791 (0.706,0.885)</b>	<0.001
LDL-C, per unit	<b>0.942 (0.919,0.967)</b>	<0.001	<b>0.907 (0.889,0.926)</b>	<0.001	<b>0.926 (0.874,0.980)</b>	0.008	<b>0.946 (0.907,0.987)</b>	<0.001
TC, per SD	<b>0.941 (0.922,0.961)</b>	<0.001	<b>0.915 (0.903,0.928)</b>	<0.001	<b>0.946 (0.907,0.987)</b>	0.010	<b>0.946 (0.916,0.977)</b>	<0.001
TG, per SD	<b>1.056 (1.037,1.075)</b>	<0.001	<b>1.045 (1.032,1.058)</b>	<0.001	1.025 (0.987,1.065)	0.198	0.967 (0.932,1.003)	0.069
HDL, per SD	<b>0.885 (0.864,0.906)</b>	<0.001	<b>0.895 (0.881,0.909)</b>	<0.001	<b>0.953 (0.910,0.999)</b>	0.047	<b>0.925 (0.892,0.960)</b>	<0.001
LDL, per SD	<b>0.953 (0.934,0.973)</b>	<0.001	<b>0.938 (0.926,0.951)</b>	<0.001	<b>0.945 (0.905,0.985)</b>	0.008	<b>0.958 (0.927,0.989)</b>	0.010
non-HDL, per unit	<b>0.968 (0.946,0.991)</b>	0.006	<b>0.941 (0.927,0.954)</b>	<0.001	<b>0.953 (0.909,0.999)</b>	0.046	<b>0.959 (0.925,0.995)</b>	0.025
CRI 1, per unit	<b>1.025 (1.007,1.042)</b>	0.005	<b>1.018 (1.004,1.032)</b>	0.009	1.001 (0.982,1.021)	0.908	1.007 (0.975,1.039)	0.676
CRI 2, per unit	<b>1.036 (1.012,1.061)</b>	0.003	<b>1.028 (1.005,1.052)</b>	0.015	<b>0.943 (0.893,0.996)</b>	0.034	1.004 (0.964,1.046)	0.842
AC, per unit	<b>1.025 (1.007,1.042)</b>	0.005	<b>1.018 (1.004,1.032)</b>	0.009	1.001 (0.982,1.021)	0.908	1.007 (0.975,1.039)	0.676
AIP, per unit	<b>1.586 (1.473,1.708)</b>	<0.001	<b>1.492 (1.419,1.569)</b>	<0.001	<b>1.159 (1.015,1.325)</b>	0.030	1.112 (0.983,1.258)	0.090
non-HDL, per SD	<b>0.970 (0.950,0.991)</b>	0.006	<b>0.942 (0.929,0.956)</b>	<0.001	<b>0.957 (0.916,0.999)</b>	0.046	<b>0.962 (0.929,0.995)</b>	0.025
CRI 1, per SD	<b>1.033 (1.010,1.057)</b>	0.005	<b>1.019 (1.005,1.034)</b>	0.009	1.002 (0.965,1.040)	0.908	1.008 (0.973,1.044)	0.676
CRI 2, per SD	<b>1.031 (1.011,1.053)</b>	0.003	<b>1.018 (1.004,1.033)</b>	0.015	<b>0.926 (0.862,0.994)</b>	0.034	1.004 (0.969,1.039)	0.842
AC, per SD	<b>1.033 (1.010,1.057)</b>	0.005	<b>1.019 (1.005,1.034)</b>	0.009	1.002 (0.965,1.040)	0.908	1.008 (0.973,1.044)	0.676
Continued								

Index	Hospital in Liangjiang New Area, Chongqing		Hospital in Kaizhou District, Chongqing		Hospital in Qianjiang District, Chongqing		Hospital in Beijing	
	OR (95%)	P-value	OR (95%)	P-value	OR (95%)	P-value	OR (95%)	P-value
AIP,per SD	<b>1.161 (1.134,1.189)</b>	<0.001	<b>1.132 (1.115,1.150)</b>	<0.001	<b>1.058 (1.006,1.114)</b>	0.030	1.035 (0.995,1.076)	0.090
	The ORs were adjusted for gender, age, BMI, fatty liver, renal calculus, hypertension, blood glucose level, AST, GGT, TBIL, UA,ALB, A/G		The ORs were adjusted for Age, gender,BMI, ALT, AST, GGT, ALP, ALB, A/G, UA, Cr, fatty liver, renal calculus, hypertension, blood glucose level		The ORs were adjusted for age, gender, BMI, ALT, GGT, fatty liver, renal calculus, hypertension		The ORs were adjusted for age,gender,BMI,AST,ALT,UA,blood glucose level,hypertension,renal calculus,fatty liver	

**Table 2.** Association between lipid profile and GSD with multivariable analysis by logistic regression in four hospitals. TC = Total Cholesterol; TG = Triglycerides; HDL-C = High-density Lipoprotein Cholesterol; LDL-C = Low-density Lipoprotein Cholesterol; non-HDL-C = Non-high-density Lipoprotein-Cholesterol; CRI1 = Castelli Risk Index—I; CRI2 = Castelli Risk Index—II; AC = The Atherogenic Coefficient; AIP = The Atherogenic Index of Plasma.

SX), the results remained largely consistent with those derived from the full population. Hospital in Qianjiang District was excluded from the dysglycemia sensitivity analysis due to lack of glucose data. The consistency of findings across these sensitivity analyses reinforces the robustness of our primary results.

### Results of trial sequential analysis (TSA) of lipid indices

The TSA plot revealed that all nine lipid profile indicators crossed both the conventional boundary and the TSA monitoring boundary, achieving the required information size (Fig. 2). This demonstrates that the current study provides robust and conclusive evidence, eliminating the need for additional studies to further validate these findings.

### Discussion

This study employed a multicenter cross-sectional design, enrolling 580,935 participants from four hospitals. Of note, significant variations were observed in both the detection rate of gallstones and the association between blood lipid indicators and gallstone presence across the participating hospitals, with Hospital of Qianjiang district exhibiting the most distinct pattern. These disparities may be attributable to regional epidemiological differences—specifically, a lower incidence of gallstone disease in northern China compared with southwestern China<sup>20</sup>. Furthermore, the unique dietary habits prevalent among ethnic minority populations in the Qianjiang region may contribute to the differential gallstone incidence observed there. It is also noteworthy that across all study sites, the number of female gallstone patients was significantly higher than that of males, a finding consistent with previously published epidemiological data<sup>20–22</sup>. Because of the heterogeneity across multiple centers, we did not use direct pooling of data for analysis. Instead, a meta-analysis was subsequently conducted to pool the results, aiming to comprehensively and systematically elucidate the associations between traditional and non-traditional lipid profiles and the risk of GSD. The pooled results indicated that TC, LDL-C, and HDL-C were inversely associated with GSD risk, while TG showed no significant overall association. This lack of association may be due to the fact that TG does not directly participate in GSD formation. However, subgroup analyses revealed a positive correlation between TG and GSD in individuals over 60 years of age and in females, which is consistent with previous findings<sup>23</sup>. This phenomenon may be related to estrogen secretion in women and slower lipid metabolism observed in the elderly population.

The specific mechanisms through which lipids contribute to GSD formation remain unclear. The relationships between TC, LDL-C, and GSD are particularly controversial. Some studies suggest a positive association, while others report an inverse association, and some find no association at all<sup>4,19,24–27</sup>. A study by Zhang et al. demonstrated that higher TC and LDL-C levels were significantly associated with a reduced risk of GSD, aligning with our findings<sup>13</sup>. Similarly, Chen et al. identified lower TC levels and higher TG levels as independent risk factors for GSD, potentially due to compensatory cholesterol secretion by the liver and reduced bile acid secretion<sup>19</sup>. The inverse associations between TC, LDL-C, and gallstone formation observed in this study are consistent with findings from multiple cross-sectional and Mendelian randomization studies<sup>18,28</sup>. The large sample size in the present analysis further strengthens the robustness of these results. However, the underlying mechanisms by which TC and LDL-C may confer protection against gallstone development remain unclear. One plausible explanation is that intrahepatic cholesterol levels exert a more direct influence on gallstone pathogenesis than serum cholesterol concentrations<sup>19</sup>. Additionally, different biological pathways through which plasma LDL-C is reduced may have divergent—and sometimes opposing—effects on gallstone risk<sup>29</sup>. HDL-C is a well-established protective factor against GSD, facilitating the reverse transport of cholesterol from peripheral tissues (including the gallbladder) to the liver, thereby reducing cholesterol deposition in bile and lowering the risk of gallstone formation<sup>4,30,31</sup>.

Emerging evidence suggests that non-traditional lipid indices, derived from conventional lipid measurements, demonstrate superior predictive capacity for metabolic disorders compared to individual lipoprotein assessments<sup>32,33</sup>. These composite biomarkers integrate multifaceted lipid metabolism information, yet their association with cholelithiasis pathogenesis remains insufficiently characterized. Our multi-center pooled analyses demonstrated significant associations between novel atherogenic indices and GSD formation risk,

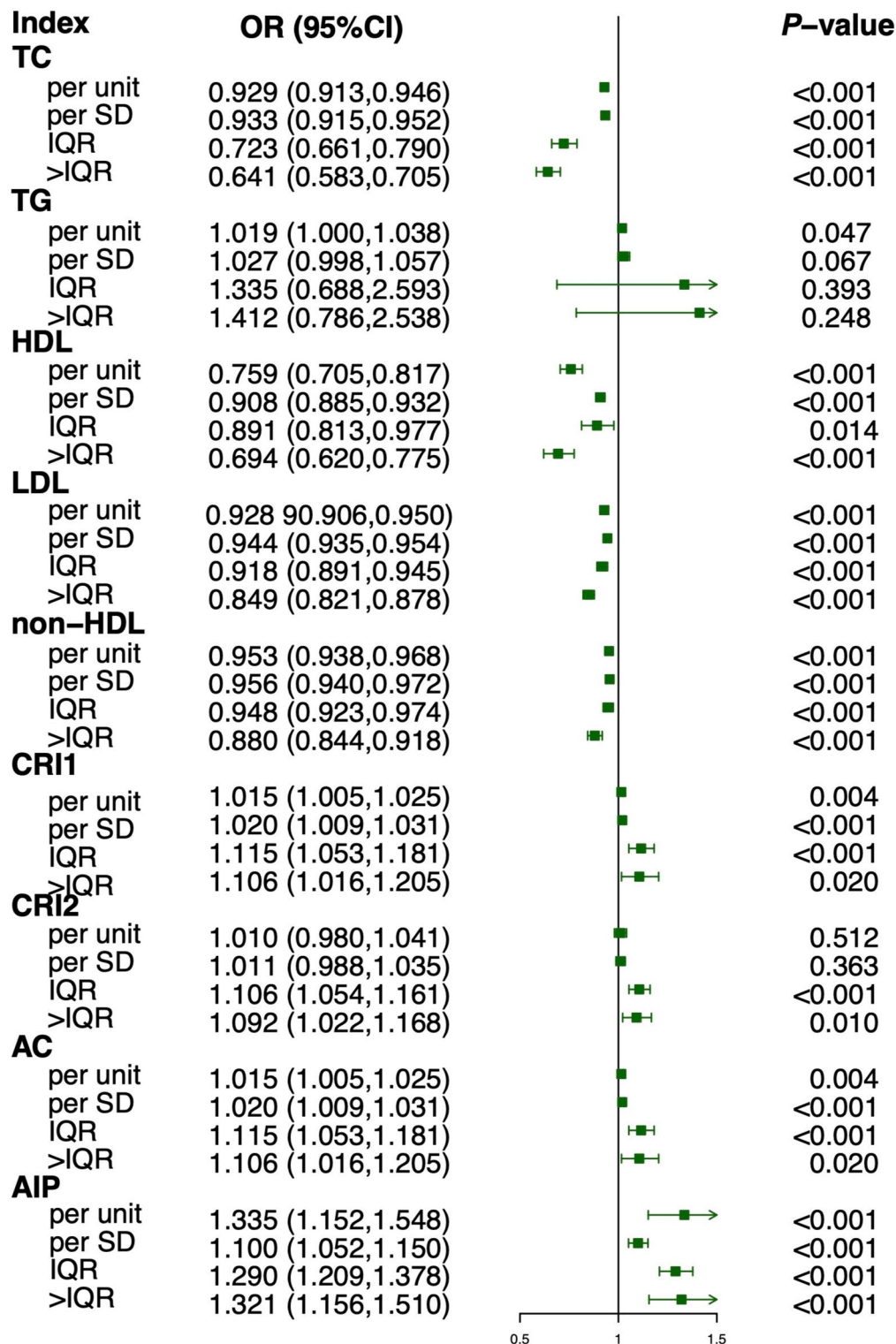


Fig. 1. The forest plot for the pooled associations between nine lipid parameters and the risk of GSD.

providing critical insights into the pathophysiological interplay between metabolic dysregulation and biliary stone development.

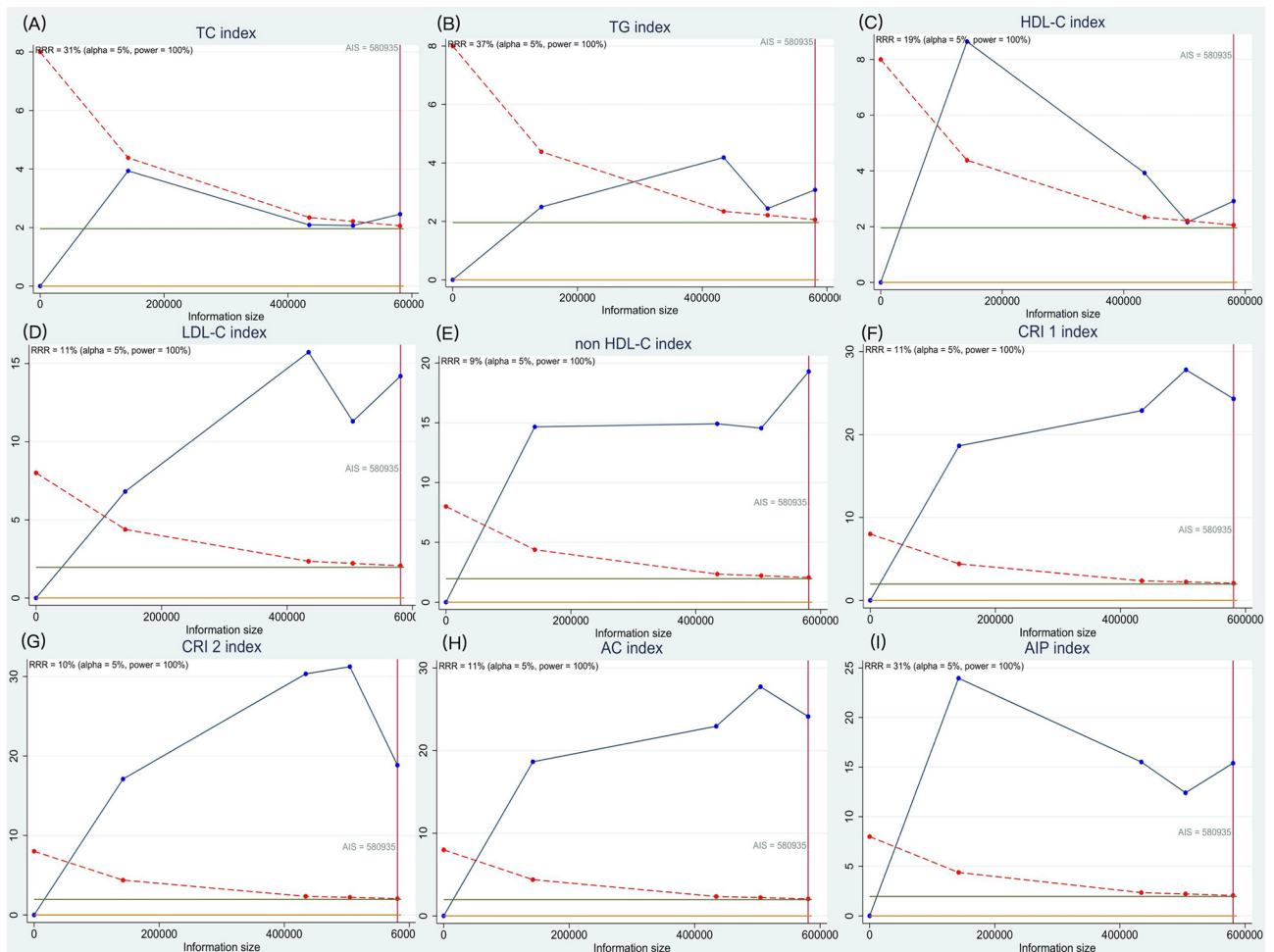
Notably, non-high-density lipoprotein cholesterol (non-HDL-C), calculated as total cholesterol minus HDL-C and encompassing atherogenic lipoproteins including LDL-C, VLDL-C, IDL-C, and Lp(a), exhibited an inverse correlation with cholelithiasis incidence, potentially attributable to its predominant LDL-derived composition ( $\approx 70\%$ ) modulating cholesterol trafficking and crystallization dynamics.

Index	Subgroup	OR(95%CI)	P-value	Index	Subgroup	OR(95%CI)	P-value		
Per unit				Per SD					
TC	Age	<40	0.912 (0.856,0.973)	0.005	TC	Age	<40	0.917 (0.863,0.974)	0.005
		40–60	0.948 (0.916,0.981)	0.002			40–60	0.951 (0.918,0.985)	0.005
		>60	0.902 (0.884,0.922)	0.000			>60	0.909 (0.887,0.932)	0.000
	Gender	Male	0.904 (0.884,0.926)	0.000		Gender	Male	0.910 (0.887,0.933)	0.000
		Female	0.966 (0.939,0.993)	0.015			Female	0.967 (0.941,0.994)	0.018
TG	Age	<40	1.016 (0.980, 1.054)	0.375	TG	Age	<40	1.021 (0.968, 1.078)	0.441
		40–60	1.018 (0.996, 1.040)	0.117			40–60	1.025 (0.993, 1.059)	0.126
		>60	1.024 (1.006,1.043)	0.008			>60	1.035 (1.009,1.063)	0.009
	Gender	Male	1.017 (0.995, 1.039)	0.132		Gender	Male	1.024 (0.992, 1.057)	0.147
		Female	1.045 (1.006,1.085)	0.023			Female	1.065 (1.006,1.127)	0.030
HDL	Age	<40	0.712 (0.610,0.832)	0.000	HDL	Age	<40	0.888 (0.842,0.937)	0.000
		40–60	0.754 (0.682,0.834)	0.000			40–60	0.905 (0.876,0.935)	0.000
		>60	0.799 (0.726,0.880)	0.000			>60	0.926 (0.891,0.963)	0.000
	Gender	Male	0.713 (0.667,0.762)	0.000		Gender	Male	0.889 (0.874,0.905)	0.000
		Female	0.835 (0.736,0.946)	0.005			Female	0.939 (0.897,0.983)	0.007
LDL	Age	<40	0.938 (0.880,0.999)	0.047	LDL	Age	<40	0.954 (0.912,0.999)	0.046
		40–60	0.954 (0.917,0.993)	0.022			40–60	0.965 (0.941,0.989)	0.005
		>60	0.885 (0.860,0.910)	0.000			>60	0.915 (0.897,0.934)	0.000
	Gender	Male	0.888 (0.861,0.917)	0.000		Gender	Male	0.915 (0.901,0.928)	0.000
		Female	0.980 (0.949, 1.011)	0.209			Female	0.985 (0.964, 1.006)	0.149
non-HDL	Age	<40	0.936 (0.862, 1.017)	0.117	non-HDL	Age	<40	0.940 (0.870, 1.016)	0.118
		40–60	0.974 (0.943, 1.007)	0.121			40–60	0.976 (0.945, 1.008)	0.136
		>60	0.915 (0.895,0.935)	0.000			>60	0.919 (0.900,0.939)	0.000
	Gender	Male	0.930 (0.910,0.951)	0.000		Gender	Male	0.935 (0.913,0.957)	0.000
		Female	0.987 (0.957, 1.018)	0.408			Female	0.988 (0.960, 1.017)	0.417
CRI1	Age	<40	1.013 (0.961, 1.067)	0.643	CRI1	Age	<40	1.022 (0.961, 1.086)	0.493
		40–60	1.014 (1.004,1.025)	0.006			40–60	1.018 (1.006,1.031)	0.004
		>60	0.984 (0.958, 1.011)	0.244			>60	0.980 (0.949, 1.012)	0.220
	Gender	Male	1.007 (0.997, 1.016)	0.176		Gender	Male	1.008 (0.996, 1.020)	0.201
		Female	1.029 (0.999, 1.061)	0.055			Female	1.037 (1.002,1.074)	0.036
CRI2	Age	<40	1.031 (0.959, 1.109)	0.410	CRI2	Age	<40	1.031 (0.976, 1.090)	0.277
		40–60	1.025 (1.003,1.048)	0.023			40–60	1.021 (1.007,1.036)	0.002
		>60	0.949 (0.919,0.979)	0.001			>60	0.958 (0.927,0.990)	0.010
	Gender	Male	0.992 (0.971, 1.014)	0.480		Gender	Male	0.995 (0.977, 1.013)	0.590
		Female	1.035 (0.983, 1.089)	0.188			Female	1.031 (0.993, 1.071)	0.110
AC	Age	<40	1.015 (1.006,1.025)	0.001	AC	Age	<40	1.020 (1.009,1.031)	0.000
		40–60	1.007 (0.997,1.016)	0.176			40–60	1.008 (0.996,1.020)	0.202
		>60	1.030 (1.016,1.044)	0.000			>60	1.042 (1.024,1.061)	0.000
	Gender	Male	1.040 (1.018,1.063)	0.000		Gender	Male	1.053 (1.026,1.082)	0.000
		Female	1.014 (1.004,1.025)	0.006			Female	1.018 (1.006,1.031)	0.004
AIP	Age	<40	1.377 (1.045,1.815)	0.023	AIP	Age	<40	1.111 (1.022,1.207)	0.013
		40–60	1.329 (1.159,1.523)	0.000			40–60	1.098 (1.053,1.146)	0.000
		>60	1.297 (1.194,1.408)	0.000			>60	1.087 (1.059,1.116)	0.000
	Gender	Male	1.302 (1.179,1.437)	0.000		Gender	Male	1.089 (1.054,1.126)	0.000
		Female	1.411 (1.126,1.769)	0.003			Female	1.119 (1.047,1.194)	0.001
IQR				>IQR					
TC	Age	<40	0.734 (0.566,0.951)	0.019	TC	Age	<40	0.612 (0.453,0.826)	0.001
		40–60	0.779 (0.674,0.899)	0.001			40–60	0.702 (0.606,0.812)	0.000
		>60	0.693 (0.541,0.888)	0.004			>60	0.616 (0.465,0.816)	0.001
	Gender	Male	0.697 (0.622,0.780)	0.000		Gender	Male	0.585 (0.521,0.658)	0.000
		Female	0.796 (0.693,0.914)	0.001			Female	0.750 (0.652,0.863)	0.000
Continued									

Index	Subgroup	OR(95%CI)	P-value	Index	Subgroup	OR(95%CI)	P-value		
TG	Age	<40	1.358 (0.677,2.723)	0.389	TG	Age	<40	1.373 (0.808,2.335)	0.242
		40–60	1.813 (0.980,3.354)	0.058			40–60	<b>1.917 (1.129,3.256)</b>	0.016
		>60	0.639 (0.307, 1.331)	0.232			>60	0.708 (0.345, 1.452)	0.347
	Gender	Male	1.108 (0.452,2.718)	0.822		Gender	Male	1.190 (0.485,2.919)	0.704
		Female	1.399 (0.798,2.452)	0.241			Female	<b>1.501 (1.005,2.242)</b>	0.047
HDL	Age	<40	<b>0.885 (0.798,0.981)</b>	0.020	HDL	Age	<40	<b>0.667 (0.561,0.792)</b>	0.000
		40–60	<b>0.865 (0.793,0.945)</b>	0.001			40–60	<b>0.680 (0.586,0.790)</b>	0.000
		>60	1.020 (0.819, 1.269)	0.862			>60	<b>0.766 (0.612,0.958)</b>	0.019
	Gender	Male	<b>0.869 (0.778,0.970)</b>	0.012		Gender	Male	<b>0.651 (0.580,0.732)</b>	0.000
		Female	0.955 (0.851, 1.073)	0.441			Female	<b>0.768 (0.647,0.911)</b>	0.003
LDL	Age	<40	0.973 (0.918, 1.031)	0.349	LDL	Age	<40	0.888 (0.762, 1.036)	0.130
		40–60	<b>0.911 (0.874,0.948)</b>	0.000			40–60	<b>0.879 (0.822,0.941)</b>	0.000
		>60	<b>0.875 (0.821,0.933)</b>	0.000			>60	<b>0.771 (0.720,0.826)</b>	0.000
	Gender	Male	<b>0.886 (0.848,0.926)</b>	0.000		Gender	Male	<b>0.779 (0.742,0.818)</b>	0.000
		Female	0.961 (0.919, 1.005)	0.080			Female	0.958 (0.894, 1.026)	0.244
non-HDL	Age	<40	0.973 (0.875, 1.082)	0.615	non-HDL	Age	<40	0.876 (0.714, 1.075)	0.204
		40–60	<b>0.961 (0.925,0.998)</b>	0.037			40–60	<b>0.909 (0.845,0.979)</b>	0.011
		>60	<b>0.874 (0.824,0.926)</b>	0.000			>60	<b>0.805 (0.756,0.857)</b>	0.000
	Gender	Male	<b>0.919 (0.882,0.957)</b>	0.000		Gender	Male	<b>0.830 (0.776,0.888)</b>	0.000
		Female	0.989 (0.953, 1.026)	0.554			Female	0.964 (0.889, 1.046)	0.384
CRI1	Age	<40	1.101 (0.962, 1.259)	0.161	CRI1	Age	<40	1.120 (0.940, 1.335)	0.206
		40–60	<b>1.155 (1.111,1.202)</b>	0.000			40–60	<b>1.163 (1.086,1.246)</b>	0.000
		>60	1.015 (0.958, 1.076)	0.604			>60	0.949 (0.887, 1.015)	0.124
	Gender	Male	<b>1.088 (1.035,1.143)</b>	0.001		Gender	Male	<b>1.089 (1.034,1.147)</b>	0.001
		Female	<b>1.115 (1.065,1.167)</b>	0.000			Female	1.079 (0.961, 1.211)	0.197
CRI2	Age	<40	<b>1.131 (1.017,1.258)</b>	0.023	CRI2	Age	<40	<b>1.157 (1.031,1.299)</b>	0.013
		40–60	<b>1.147 (1.103,1.192)</b>	0.000			40–60	<b>1.142 (1.092,1.195)</b>	0.000
		>60	0.983 (0.928, 1.040)	0.544			>60	<b>0.912 (0.831,1.000)</b>	0.050
	Gender	Male	1.049 (0.990, 1.111)	0.104		Gender	Male	1.046 (0.994, 1.100)	0.082
		Female	<b>1.121 (1.082,1.161)</b>	0.000			Female	<b>1.112 (1.034,1.195)</b>	0.004
AC	Age	<40	<b>1.129 (1.097,1.161)</b>	0.000	AC	Age	<40	<b>1.128 (1.090,1.167)</b>	0.000
		40–60	<b>1.088 (1.035,1.143)</b>	0.001			40–60	<b>1.089 (1.034,1.147)</b>	0.001
		>60	<b>1.123 (1.084,1.164)</b>	0.000			>60	<b>1.126 (1.074,1.181)</b>	0.000
	Gender	Male	<b>1.162 (1.095,1.233)</b>	0.000		Gender	Male	<b>1.208 (1.114,1.310)</b>	0.000
		Female	<b>1.155 (1.111,1.202)</b>	0.000			Female	<b>1.157 (1.105,1.211)</b>	0.000
AIP	Age	<40	<b>1.240 (1.036,1.485)</b>	0.019	AIP	Age	<40	1.250 (0.931, 1.677)	0.138
		40–60	<b>1.334 (1.281,1.390)</b>	0.000			40–60	<b>1.374 (1.247,1.513)</b>	0.000
		>60	<b>1.207 (1.136,1.283)</b>	0.000			>60	<b>1.287 (1.175,1.410)</b>	0.000
	Gender	Male	<b>1.208 (1.146,1.273)</b>	0.000		Gender	Male	<b>1.329 (1.255,1.407)</b>	0.000
		Female	<b>1.284 (1.184,1.392)</b>	0.000			Female	<b>1.273 (1.047,1.549)</b>	0.016

**Table 3.** Pooled-analysis results of subgroup analysis by age and sex in the multi-center cross-sectional study. Note: TC = Total Cholesterol; TG = Triglycerides; HDL-C = High-density Lipoprotein Cholesterol; LDL-C = Low-density Lipoprotein Cholesterol; non-HDL-C = Non-high-density Lipoprotein-Cholesterol; CRI1 = Castelli Risk Index—I; CRI2 = Castelli Risk Index—II; AC = The Atherogenic Coefficient; AIP = The Atherogenic Index of Plasma.

Multivariable regression analyses identified four atherogenic indices as independent risk predictors: CRI 1, CRI 2, AC, and AIP, with AIP demonstrating the strongest positive correlation. A large-scale study using the NHANCE database demonstrated a significant non-linear association between elevated AIP and cholelithiasis incidence (adjusted OR: 1.45; 95% CI: 1.09–1.93), with diabetes mellitus identified as a mediating factor<sup>33</sup>. Meanwhile, Cheng Q et al. found that elevated AC (non-HDL to HDL ratio) was associated with an increased incidence of cholelithiasis using the NHANCE database being consistent with our findings<sup>32</sup>. The mechanistic convergence of atherogenic indices and GSD formation appears rooted in shared metabolic dysregulation. Multiple plausible mechanisms may account for the positive association between AIP and AC with GSD pathogenesis. First, dyslipidemia induces systemic metabolic perturbations involving hepatic and biliary systems. Specifically, hyperlipidemia promotes hepatic cholesterol hypersecretion while reducing biliary phospholipid content, thereby disrupting cholesterol solubility and creating supersaturated bile conducive to lithogenesis<sup>34</sup>.



**Fig. 2.** Trial Sequential Analysis plot for the associations between nine lipid parameters and the risk of GSD. The plot for TC (A), TG (B), HDL-C (C), LDL-C (D), non-HDL-C (E), CRI 1 (F), CRI 2 (G), AC (H), AIP (I).

Second, insulin resistance emerges as a critical intermediary in this pathophysiological axis. Recent studies have demonstrated a nonlinear dose–response relationship between insulin resistance and AIP levels, characterized by progressive elevation of AIP with declining insulin sensitivity<sup>35</sup>. This metabolic state enhances biliary cholesterol saturation through dual pathways: (1) hyperlipidemia-induced inactivation of FoxO1 transcription factor upregulates hepatic cholesterol transporters ABCG5/G8, augmenting biliary cholesterol excretion; (2) insulin resistance directly promotes ABCG5/ABCG8 expression, further increasing biliary cholesterol concentration and accelerating crystal nucleation. Concurrently, insulin resistance-mediated downregulation of farnesoid X receptor (FXR) signaling impairs bile acid biosynthesis, exacerbating cholesterol supersaturation and lithogenicity<sup>36,37</sup>. Collectively, these interrelated pathways establish a mechanistic framework linking dyslipidemia, insulin resistance, and gallstone formation.

Interestingly, stratified analyses in our study revealed distinct associations of TG, LDL-C, non-HDL-C, and atherogenic coefficient (AC) between patients who underwent cholecystectomy and those with gallbladder stones. These differences may be attributed to alterations in lipid metabolism profiles following gallbladder removal, as supported by previous evidence<sup>38,39</sup>. Moreover, both gallstone patients and those with a history of cholecystectomy exhibited significantly higher atherogenic index of plasma (AIP) compared with healthy controls, suggesting AIP may serve as a valuable and potential predictive marker in both clinical populations.

Notably, existing investigations into the association between novel lipid biomarkers and GSD have been exclusively based on U.S. populations, with no studies conducted in Chinese cohorts<sup>31,32</sup>. Moreover, prior research has primarily utilized the NHANCE database characterized by limited sample sizes. This large-scale, multicenter study in Chinese populations addresses this critical gap by providing the first evidence from an East Asian cohort. Using TSA, we dynamically monitored cumulative evidence for nine lipid biomarkers. The results demonstrated that all cumulative Z-curves for odds ratios (ORs) crossed both the conventional  $\alpha$ -boundary and the pre-specified information boundary, validating the robustness of the findings.

However, several limitations merit acknowledgment. First, the cross-sectional nature of this investigation inherently limits our ability to determine causal associations between GSD and lipid biomarkers, as this design cannot establish temporality or exclude reverse causation. Second, residual confounding from

unmeasured variables (e.g., genetic predispositions) persists; critically, lifestyle factors (diet, physical activity), detailed dietary patterns and drugs related to lipid metabolism were unavailable in our retrospective dataset, preventing adjustment. Third, the retrospective nature of the ultrasound data collection imposed constraints: standardized sonographer training protocols and prospective quality control measures for diagnostic criteria were not implemented, potentially introducing inter-observer variability. Future longitudinal cohort studies and randomized controlled trials (RCTs) are necessary to confirm these associations and explore underlying mechanisms.

## Method

### Study participant

This multicenter cross-sectional study, titled "A Multicenter Cohort Study on the Risk Factors for Gallstone Disease in the Physical Examination Population" (Registration number: ChiCTR2500095891), was conducted across four tertiary care hospital in China from January 2015 to May 2020. The participating centers included a tertiary hospital in Qianjiang District, Chongqing; a tertiary hospital in Kaizhou District, Chongqing; a tertiary hospital in Liangjiang New Area, Chongqing; and a tertiary hospital in Beijing city.

Eligible participants were systematically recruited from the health screening centers of these institutions using strict inclusion criteria: (a) completion of abdominal ultrasonography in accordance with standardized diagnostic protocols; and (b) availability of comprehensive demographic, anthropometric, and biochemical data. The dataset encompassed information on age, sex, height, weight, blood glucose levels, diastolic blood pressure (DBP) and systolic blood pressure (SBP), liver and kidney function indicators, as well as medical histories of conditions such as fatty liver and kidney stones. For participants who had undergone multiple health examinations, only the most recent data were used to ensure the currency and relevance of the information.

Ethical approval for this study was obtained from the ethics committee of West China Fourth Hospital and West China School of Public Health, Sichuan University (approval number: Gwl2021055). The study strictly adhered to the ethical guidelines stipulated in the 1964 Declaration of Helsinki and all subsequent amendments, ensuring the highest standards of ethical conduct in human-subject research. This study is a retrospective analysis of non-identifiable researcher-collected data, human consent to participate statements do not apply. We confirm that: The research does not involve any human experimentation; No identifiable subject data or biological materials were utilized; No personal privacy concerns or commercial interests are implicated in this work.

### Laboratory examinations and indices

Fasting blood samples were collected from study participants and analyzed within one hour in the laboratories of the participating hospitals. Using biochemical analyzers, a comprehensive panel of serum biochemical parameters was carefully measured. These parameters included high-density lipoprotein cholesterol (HDL-C), low-density lipoprotein cholesterol (LDL-C), total cholesterol (TC), triglycerides (TG), fasting blood glucose (FBG), systolic blood pressure (SBP), diastolic blood pressure (DBP), total bilirubin (T-bil), direct bilirubin (D-bil), alanine transaminase (ALT), aspartate transaminase (AST), gamma-glutamyl transferase (GGT), creatinine (Cr), uric acid (UA), urea nitrogen (UN), albumin (ALB), globulin (GLB), and the albumin-to-globulin ratio (A/G). However, there are differences in the examination items among different centers, and a few indicators are missing. Each hospital strictly followed internationally recognized standard protocols for biochemical measurements, ensuring the accuracy, reliability, and comparability of the data. Subsequently, several lipid-related indices were calculated using well-established formulas:

Non-high-density lipoprotein-cholesterol (non-HDL-C) = (TC—HDL-C);

Castelli risk index—I (CRI 1) = (TC/HDL-C);

Castelli risk index—II (CRI 2) = (LDL-C/HDL-C);

The atherogenic coefficient (AC) = ((TC—HDL—C)/HDL—C).;

The atherogenic index of plasma (AIP) = (log(TG/HDL—C)).

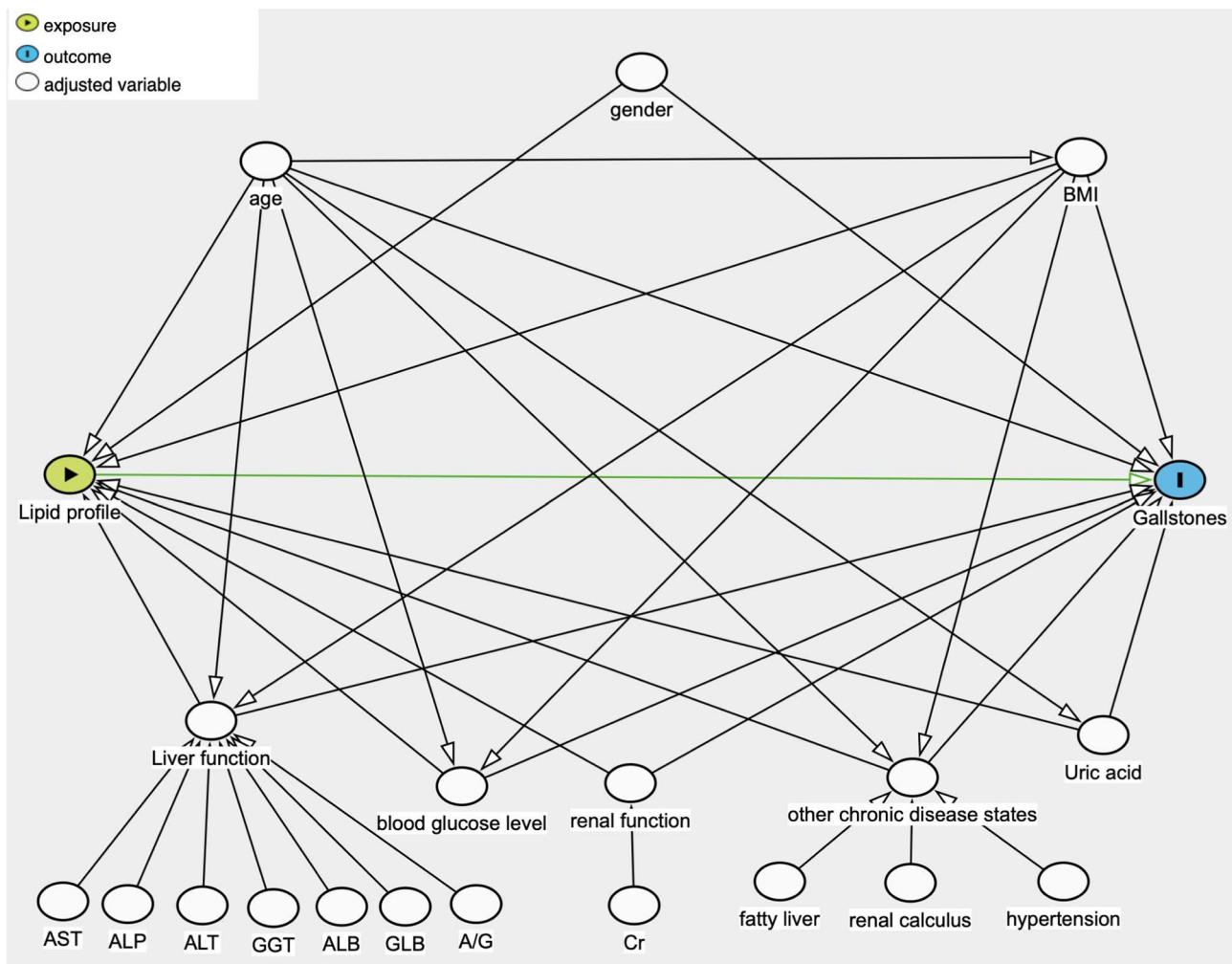
### Definitions

Ultrasonography examinations were performed by highly experienced radiologists. The diagnosis of GSD was established based on one or both of the following criteria: (i) The presence of one or more hyperechoic structures within the gallbladder or biliary system, which exhibited acoustic shadowing or demonstrated gravity-dependent movement; (ii) In patients who had previously undergone cholecystectomy due to gallstones, the absence of gallbladder visualization on ultrasonography was considered evidence of a prior GSD diagnosis<sup>13</sup>.

In this cross-sectional study, we defined the fasting blood glucose variable > 7 mmol/L as an abnormal blood glucose level. And lipid markers were categorized into three distinct groups according to the clinical diagnostic criteria. The classification details are as follows: TG (mmol/L): > 5.7, 3.1–5.7, and < 3.1; TG (mmol/L): > 1.7, 0.4–1.7, and < 0.4; LDL-C (mmol/L): > 3.1, 2.07–3.1, and < 2.07; HDL-C (mmol/L): > 2.0, 0.9–2.0, and < 0.9. Additionally, non-HDL cholesterol, CRI1, CRI2, AC, and AIP were divided into three groups based on the interquartile range: > IQR, IQR, and < IQR.

### Statistical analysis

In this cross-sectional study, statistical analysis was performed using SPSS software version 19 (IBM, USA). Missing data were addressed using multiple imputation (MI). All continuous variables exhibited non-normal distributions and were thus expressed as medians and interquartile ranges (IQR), denoted as P50 (P25, P75). Comparisons between the GSD and non-GSD groups were conducted using the Wilcoxon rank-sum test.



**Fig. 3.** Directed acyclic graphs between blood lipid indexes and gallstones.

Categorical variables were presented as counts and percentages, with intergroup comparisons performed using the chi-square test.

The adjustment set was selected based on a pre-specified Directed Acyclic Graph (DAG), which was developed by integrating established clinical knowledge with the variables actually available in our retrospective multicenter dataset (Fig. 3). Then a multivariable logistic regression model was employed to evaluate the associations between nine key lipid indicators and the risk of GSD. Lipid parameters were analyzed both as categorical variables and as continuous variables to assess their relationships with the risk of GSD, with subgroup analyses conducted by age and gender. Comprehensive Meta-Analysis (CMA) software was used to pool the results across four hospitals, generating pooled odds ratios (OR) and 95% confidence intervals (CI). Heterogeneity was assessed using Cochran's Q test, with the magnitude of heterogeneity quantified by the  $I^2$  statistic. When significant heterogeneity was present ( $I^2 \geq 50\%$ ), the DerSimonian and Laird random-effects model was applied; otherwise, a fixed-effects model was used. Forest plots were generated using R version 4.3. Trial Sequential Analysis (TSA) was conducted using Stata 17.0 to determine the sufficiency of the sample size and the robustness of the results. TSA was performed based on the pooled data from multiple centers, incorporating both higher-level and highest-level groups. A  $2 \times 2$  contingency table was constructed, and the relative risk reduction (RRR) values were calculated using the combined odds ratios (OR). The statistical power analysis was conducted with a type I error rate ( $\alpha$ ) of 0.05 and a type II error rate ( $\beta$ ) of 0.2.

## Conclusion

This study underscores the pivotal role of lipid metabolic dysregulation in GSD pathogenesis, revealing that TC, HDL-C, LDL-C, and non-HDL-C are negatively correlated with the risk of GSD. Conversely, elevated levels of the CRI1, CRI2, AC, and AIP are significantly associated with an increased risk of GSD. The above results were different between cholecystectomy and gallbladder stones. These findings not only elucidate the underlying mechanisms of GSD, but also establish a comprehensive risk stratification framework that integrates lipid homeostasis biomarkers. This framework holds substantial potential for clinical translation, offering a novel and practical approach for early risk assessment, targeted prevention, and personalized management of GSD. By

integrating these lipid parameters into clinical practice, our study provides actionable insights to enhance the precision of GSD prevention and improve patient outcomes.

### Data availability

The datasets generated during and analysed during the current study are not publicly available due to the confidentiality of multi-center collaboration but are available from the corresponding author on reasonable request.

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## Author contributions

XW, YJ, LZ: conceptualization, and investigation, writing original draft, revision. YJ, HYL, MLH: methodology, assisted in the conceptualization, contributed to the draft writing. YJ: software, imaging analysis, and revision. YJ, CLW, XYZ, HYL, WQY, XW, XB, YTX, LJX: contributed to the methodology, writing, and data curation. MLH, XW, WQY, HYL, GHJ, JL: contributed to the conception, and draft writing. XW, YJ, YY, YML: writing, review, and editing. XW, CLW, XYZ, GCL, XFS, GHJ, LZ, FL, YML, YY: data collection, revision. All authors reviewed the final draft and agreed on its content and conclusions. All authors had full access to all the data in the study and had final responsibility for the decision to submit for publication. XW, CLW, YML, FL, YY and LZ accessed and verified the data.

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## Declarations

### Competing interests

The authors declare no competing interests.

### Ethics approval and consent to participate

Ethical approval for this study was obtained from the ethics committee of West China Fourth Hospital and West China School of Public Health, Sichuan University (approval number: Gwll2021055). This study is a retrospective analysis of non-identifiable researcher-collected data. The requirement for informed consent from the study subjects was waived by the IRB of Ethical Approval of Ethics Committee of West China Fourth Hospital and West China School of Public Health, Sichuan University due to the retrospective study design. We confirm that: The research does not involve any human experimentation; No identifiable subject data or biological materials were utilized; No personal privacy concerns or commercial interests are implicated in this work.

### Additional information

**Supplementary Information** The online version contains supplementary material available at <https://doi.org/10.1038/s41598-026-37603-x>.

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