



# Inverse Associations Between Circulating Secreted Frizzled Related Protein 2 (sFRP2) and Cardiometabolic Risk Factors

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**Background:** Secreted frizzled-related protein 2 (sFRP2) plays an important role in metabolic syndrome and cardiovascular diseases (CVDs); However, its relevance with cardiometabolic diseases remains to be elucidated. We aimed to determine the serum levels of sFRP2 in patients at different stages of heart failure (HF) with or without type 2 diabetes mellitus (T2DM), and assess the correlation between circulating sFRP2 levels and cardiometabolic risk factors.

**Methods:** In this study, serum samples from 277 patients visiting Zhongshan Hospital affiliated to Fudan University were collected. These patients were clinically diagnosed and categorized as five groups, including the control group, pre-clinical HF group, pre-clinical HF+T2DM group, HF group and HF+T2DM group. Serum sFRP2 levels were measured with enzyme-linked immunosorbent assay (ELISA) tests and the clinical characteristics of each patient were recorded. Spearman rank correlation analysis and multiple stepwise linear regression analysis were conducted. Univariate and multivariate logistic regression analysis were performed to screen risk factors for HF in patients with CVDs.

**Results:** Serum sFRP2 levels were significantly lower in the HF+T2DM group compared with the other four groups. Spearman rank correlation analysis showed that sFRP2 was negatively correlated with parameters including patients' age, fasting plasma glucose (FPG), glycated hemoglobin A1c (HbA1c), cardiac troponin T (cTNT), N-terminal pro-B-type natriuretic peptide (NT-proBNP), high-sensitivity C-reactive protein (hs-CRP), left atrial dimension (LAD) and left ventricular posterior wall (LVPW), and positively correlated with hemoglobin, estimated glomerular filtration rate (eGFR), albumin, total cholesterol (TC), low-density lipoprotein cholesterol (LDL-C), and left ventricular ejection fraction (LVEF). However, in multiple regression analysis, significant associations with ln(sFRP2) were observed only in FPG, hs-CRP and LAD. Higher serum sFRP2 was significantly linked to lower odds of HF in patients with CVDs.

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**Conclusion:** sFRP2 progressively decreased when glucose homeostasis and cardiac function deteriorated. sFRP2 acted as a risk factor for HF in patients with CVDs, especially in those with concomitant T2DM.

Keywords: secreted frizzled-related protein 2 (sFRP2), cardiometabolic risk factors, heart failure (HF), type 2 diabetes (T2DM), cross-section study

# INTRODUCTION

Despite numerous advances in diagnosis and treatment, cardiovascular diseases (CVDs) remain the leading cause of global mortality. Heart failure (HF), characterized by dyspnea or limited activity due to fluid retention and/or reduced cardiac output, is the end stage of most CVDs and affects at least 23 million people worldwide (1). To emphasize the importance of early intervention and guide therapy, the American College of Cardiology Foundation/American Heart Association (ACCF/AHA) started to promote the concept of HF stages in 2001 (2) and have categorized HF into four stages. Stage A is defined as high risk for HF, including hypertension, metabolic syndrome, tobacco use, and known cardiotoxic agents. Stage B is defined as heart disease without signs or symptoms of HF. Stage C refers to HF with prior or current symptoms, and stage D refers to refractory HF requiring specialized interventions (3). The China Medical Association (CMA) has adapted this concept since 2007 and has classified HF into pre-HF, pre-clinical HF, clinical HF and refractory HF stages (4), which correspond to stages A to D in the American guideline, respectively. The progression of HF stages is irreversible and is associated with a decrease in 5-year survival (5). The rational for treating each stage of HF varies from controlling risk factors, treating structural heart disease to reducing morbidity and mortality.

In the past decade, the prevalence of type 2 diabetes (T2DM) has increased by 30% globally (6), with the number of influenced Chinese adults increasing from 128 million in 2007 to 155 million in 2017 (7). Previous studies have consistently demonstrated a 2- to 4-fold increased risk of HF in individuals with T2DM compared with their age- and sex-matched non-diabetic counterparts, even after adjustment for other cardiovascular risk factors (8, 9). The blood glucose control level also influences the development and progression of HF (10–12). Moreover, HF patients with concomitant T2DM exhibit significantly worse outcomes compared with those without T2DM (13). Therefore, investigating the T2DM-related characteristics in a HF patient cohort may give insight into our understanding of how the metabolic disorders incorporate with CVDs.

Secreted frizzled-related protein (sFRP) family consists of 5 members (sFRP1-5) that structurally resemble the Wnt frizzled receptors and play important roles in cardiac development and various cardiovascular pathophysiological conditions (14). Among these proteins, sFRP2 is considered to be the most potent modulator of the Wnt signaling (15). Previous studies have shown divergent roles of sFRP2 in myocardial fibrosis (16–18), hypertrophy (19), angiogenesis (20), cardiac cell death (21), and regeneration (22, 23) *via* bi-directional regulation of

the canonical Wnt pathway, activation of non-canonical Wnt pathway or modulation of other signaling pathways (24). The expression of sFRP2 in cardiac tissues was increased in the early stage of HF and subsequently decreased with the progression of HF (19). In patients with severe decompensated HF, serum sFRP2 levels were higher in those experiencing primary outcome events (25). However, data on serum sFRP2 are still lacking in healthy subjects and in patients with cardiac abnormalities that have not developed into HF. As an adipokine, sFRP2 expression in the white adipose tissue (WAT) was upregulated in obesity (26) and downregulated with moderate weight loss (27). Serum sFRP2 was positively associated with circulating insulin, homeostasis model assessment of  $\beta$  cell secretory capacity (HOMA-β) and insulin resistance (HOMAIR), body mass index (BMI) and triglycerides (TG), and was increased in patients with abnormal glucose tolerance (AGT) (28). Circulating sFRP4 (29-32) and sFRP5 (33-36) levels were associated with the risk of prediabetes/T2DM, obesity and CVDs. Lower levels of serum sFRP5 was associated with worse prognosis in patients with HF, especially in those with concomitant T2DM (37). However, it is obscure whether sFRP2 is involved in the comorbidity of HF and T2DM.

In this study, we investigated the serum levels of sFRP2 among patients in different HF stages with or without T2DM and analyzed its association with fasting blood glucose (FBG), glycated hemoglobin (HbA1c), blood lipid levels, BMI and other cardiometabolic risk factors. We also explored whether sFRP2 could serve as an indicator of the comorbidity of HF and T2DM.

# **METHODS**

### **Study Population**

This study complied with the principles of the Declaration of Helsinki and was approved by the Ethics Committee of Zhongshan Hospital, Fudan University (B2020-078R). During the period of August 2020 to August 2021, a total of 425 serum samples from patients admitted to Zhongshan Hospital were initially collected. Those with T1DM, history of malignancy, autoimmune diseases, severe infection, severe renal failure (estimated glomerular filtration rate (eGFR) <30 mL/min/1.73 m<sup>2</sup> or under renal replacement therapy), severe hepatic disease (bilirubin > 3× the upper limit of normal, or aspartate aminotransferase/alanine aminotransferase/alkaline phosphatase > 5× the upper limit of normal, or cirrhosis), active bleeding or severe anemia (Hb < 60 g/L) were excluded from the study.

Finally, 277 of these serum samples were included for the following analysis. Patients were classified and grouped according to the Guideline for the diagnosis and treatment of heart failure

in China (2018 edition) (38) and the guideline for T2DM (2020 edition) (39). The healthy subjects and pre-HF patients (stage A) were pooled as control group (n = 53) and none of them was diagnosed with T2DM. The pre-clinical HF patients (stage B) were subdivided into pre-clinical HF group without T2DM (n = 57) and pre-clinical HF+T2DM group (n = 54), depending on their medical history of T2DM. The HF patients, including both clinical HF (stage C) and refractory HF (stage D), were also stratified by the medical history of T2DM and thus subdivided into HF group without T2DM (n = 63) and HF+T2DM group (n = 50).

The cardiac conditions of these patients were further determined following the latest guidelines for hypertension (40), ischemic heart disease (41, 42), valvular heart disease (43), cardiomyopathy (44), and congenital heart disease (45). Based on echocardiography, patients with left ventricular ejection fraction (LVEF) < 40%, 40–49%, and  $\geq$ 50% were subclassified as HF with reduced ejection fraction (HFrEF), midrange EF (HFmrEF), and preserved EF (HFpEF), respectively (46).

# Clinical Characteristics and sFRP2 Detection

Medical records were carefully documented, including age, gender, BMI, smoking habit, history of hypertension and CVDs, medication, New York Heart Association (NYHA) functional class, biochemistry tests and echocardiographic parameters.

Fasting venous blood was collected with a serum separation tube, followed by stratification at room temperature for 15 min. Serum were separated and then stored at  $-80^{\circ}$ C until analysis.

An enzyme-linked immunosorbent assay (ELISA) kit purchased from Shanghai Yu Bo Biotech Co., Ltd was used to measure the levels of serum sFRP2 in accordance with the manufacturer's instruction manual. The intra-assay and inter-assay variations were 5.4% and 7.5%, respectively.

# **Statistical Analysis**

Statistical analyses were performed in R software (Version 4.0.4) and GraphPad Prism Software (Version 8.3.0). Continuous variables were presented as mean  $\pm$  standard error of mean or median and interquartile range. Categorical variables were exhibited as numbers or proportions. Differences among groups were compared using one-way ANOVA test for normally distributed continuous variables with homogeneous variance, otherwise Kruskal-Wallis test along with Dunn post-hoc test were used. Pearson's chi-squared test or Fisher's exact test was used for categorical variables. Spearman rank correlation analysis identified factors correlated with sFRP2. The levels of sFRP2 were log<sub>e</sub>-transformed (ln-transformed) for further analyses to optimize the fitted equation. Cross-sectional associations between ln(sFRP2) and cardiometabolic risk factors were estimated by multiple stepwise linear regression analysis. Multicollinearity was considered present for a variance inflation factor (VIF) > 2.5. Univariate logistic regression analysis was performed to screen risk factors for HF and the variables with significance were further investigated in the multivariate logistic stepwise regression analysis. All statistical tests were two-sided and a *P*-value < 0.05 was considered as statistically significant.

TABLE 1   Demographic and laboratory information of the study population.						
	Control ( $n = 53$ )	Pre-HF ( <i>n</i> = 57)	Pre-HF +T2DM ( <i>n</i> = 54)	HF ( <i>n</i> = 63)	HF + T2DM ( <i>n</i> = 50)	P-value
Demographic characte	ristics					
Age (years)	62 (56, 66)	65 (54, 67)	66 (56, 71)	64 (52, 72)	68 (62, 74)	0.15
Gender (male%)	32 (60.4)	40 (70.2)	41 (75.9)	46 (73.0)	31 (62.0)	0.33
BMI (kg/m <sup>2</sup> )	$23.8\pm0.6$	$24.2\pm0.5$	$24.8\pm0.4$	$25.1 \pm 0.6$	$24.7\pm0.6$	0.16
Smoking [n (%)]	15 (28.3)	19 (33.3)	19 (35.1)	17 (27.0)	8 (18.0)	0.22
Biochemistry indicators	S					
Hemoglobin (g/L)	$142.8\pm1.6$	$130.0\pm2.5$	$128.7 \pm 2.5$	$121.0 \pm 1.9$	$111.0 \pm 3.9$	<0.001
Alb (g/L)	$46.8\pm0.5$	$40.8\pm0.5$	$40.5\pm0.7$	$38.4\pm0.6$	$36.9\pm0.7$	<0.001
eGFR (mL/min/1.73 m <sup>2</sup> )	$87.3\pm1.9$	$83.6 \pm 2.2$	$77.6\pm2.7$	$75.3 \pm 3.0$	$62.8\pm3.8$	<0.001
FPG (mmol/L)	4.9 (4.6, 5.1)	5.5 (4.9, 6.1)	6.6 (5.6, 8.2)	5.4 (4.7, 6.4)	8.5 (6.1, 10.6)	<0.001
HbA1c (%)	5.7 (5.5, 5.9)	5.6 (5.4, 5.8)	6.6 (5.9, 7.8)	5.9 (5.4, 6.4)	7.6 (6.4, 8.1)	<0.001
cTNT (ng/mL)	0.005 (0.004, 0.006)	0.019 (0.007, 0.077)	0.015 (0.009, 0.056)	0.082 (0.025, 0.192)	0.055 (0.028, 0.142)	<0.001
NT-proBNP (pg/mL)	36.3 (27.5, 43.4)	310.0 (64.6, 698.0)	284.0 (77.1, 1410.0)	1263.0 (676.0, 2759.0)	2261.0 (647.0, 5270.0)	<0.001
TC (mmol/L)	4.97 (4.47, 5.48)	3.32 (2.88, 3.88)	3.63 (2.98, 4.88)	3.76 (3.14, 4.26)	3.32 (2.83, 3.93)	<0.001
TG (mmol/L)	1.47 (1.10, 2.21)	1.33 (0.94, 1.70)	1.58 (1.02, 2.03)	1.25 (0.93, 1.87)	1.51 (1.15, 2.12)	0.09
LDL-C (mmol/L)	2.82 (2.46, 3.32)	1.56 (1.26, 2.17)	1.64 (1.32, 2.56)	2.01 (1.57, 2.62)	1.56 (1.25, 2.02)	<0.001
HDL-C (mmol/L)	1.24 (1.08, 1.48)	1.08 (0.86, 1.31)	1.08 (0.83, 1.31)	0.99 (0.85, 1.13)	0.98 (0.80, 1.16)	<0.001
hs-CRP (mg/L)	1.0 (0.4, 1.3)	1.0 (0.1, 2.8)	1.4 (0.6, 5.4)	3.8 (1.0, 8.8)	3.5 (1.2, 26.2)	<0.001

Continuous variables are presented as mean ± standard error of mean or median (interquartile range). Categorical variables are expressed as number (percentages). Statistically significant values are indicated in italic. BMI, body mass index; Alb, albumin; eGFR, estimated glomerular filtration rate; FPG, fasting plasma glucose; HbA1c, glycated hemoglobin; cTNT, cardiac troponin T; NT-proBNP, N-terminal pro-B-type natriuretic peptide; TC, total cholesterol; TG, triglyceride; LDL-C, low-density lipoprotein cholesterol; HDL-C, high-density lipoprotein cholesterol; high-sensitivity C-reactive protein.

# RESULTS

# Clinical Characteristics of the Study Population

The 277 participants were divided into 5 groups: control group (n = 53), pre-clinical HF group (n = 57), preclinical HF+T2DM group (n = 54), HF group (n = 63) and HF+T2DM group (n = 50). There was no significant difference in demographic characteristics such as age, gender, BMI and smoking habit among these groups. The preclinical HF+T2DM group and HF+T2DM group had higher FPG and HbA1c than their nondiabetic counterparts and the control group (Table 1). Compared with the control group, the levels of cardiac troponin T (cTNT), N-terminal pro-B-type natriuretic peptide (NT-proBNP) and left atrial dimension (LAD) increased in the pre-clinical HF group and further elevated in the HF group (Tables 1, 2). Similar results were obtained in the pre-clinical HF+T2DM group and the HF+T2DM group. Patients with CVDs exhibited reduced LVEF and elevated pulmonary artery pressure (PAP) compared with the control group. Echocardiographic examinations showed enlarged left ventricular end-diastolic dimension (LVDd) and left ventricular end-systolic dimension (LVDs), as well as thickened interventricular septal thickness (IVS) and left ventricular posterior wall (LVPW) in the HF group and HF+T2DM group compared with the control group, pre-clinical HF group and pre-clinical+T2DM group (**Table 2**).

## Serum sFRP2 Levels in Different Groups

Next, we compared the serum sFRP2 levels among the patient groups (**Figure 1**). There was no significant difference between the control group and the pre-clinical HF group (P = 0.94) or the HF group (P = 0.43). However, compared with the control group, serum sFRP2 level decreased significantly in the HF+T2DM group (P < 0.0001). Although no significant difference was found between the pre-clinical HF group and the pre-clinical HF+T2DM group (P = 0.62), the HF+T2DM group had lower levels of serum sFRP2 than the HF group (P < 0.0001). Our results showed that serum sFRP2 levels reduced

TABLE 2   Medical history and echocardiography of the study population.						
	Control ( $n = 53$ )	Pre-HF ( <i>n</i> = 57)	Pre-HF+T2DM ( <i>n</i> = 54)	HF ( <i>n</i> = 63)	HF+T2DM ( <i>n</i> = 50)	P-value
History of cardiovascular diseases						
IHD [n (%)]	/	47 (82.6)	51 (94.4)	30 (47.6)	27 (54.0)	<0.001
VHD [n (%)]	/	4 (7.0)	6 (11.1)	19 (30.2)	14 (28.0)	<0.01
Cardiomyopathy [n (%)]	/	2 (3.5)	2 (3.7)	8 (12.7)	5 (10.0)	0.16
CHD [n (%)]	/	3 (5.3)	3 (5.6)	6 (9.5)	3 (6.0)	0.83
Hypertension [n (%)]	19 (35.8)	33 (57.9)	45 (83.3)	27 (42.9)	40 (80.0)	<0.001
Chronic medication						
ACEI/ARBs [n (%)]	12 (22.6)	16 (28.1)	17 (31.5)	14 (22.2)	18 (36.0)	0.45
CCBs [n (%)]	11 (20.8)	15 (26.3)	17 (31.5)	13 (20.6)	22 (44.0)	<0.01
Beta-blockers [n (%)]	1 (1.9)	28 (49.1)	22 (40.7)	20 (31.7)	21 (42.0)	<0.001
Diuretics [n (%)]	3 (5.7)	6 (10.5)	7 (13.0)	29 (46.0)	19 (38.0)	<0.001
Digoxin [n (%)]	/	0 (0.0)	1 (1.9)	7 (11.1)	4 (8.0)	<0.05
Nitrate esters [n (%)]	/	8 (14.0)	11 (20.4)	11 (20.7)	16 (32.0)	0.12
Antiplatelet [n (%)]	/	40 (70.2)	28 (51.9)	31 (49.2)	28 (56)	0.10
Anticoagulant [n (%)]	/	9 (15.8)	3 (5.6)	15 (23.8)	4 (8.0)	<0.05
Insulin [n (%)]	/	/	9 (16.7)	/	17 (34.0)	<0.05
OAD [n (%)]	/	/	49 (90.7)	/	19 (38.0)	<0.001
Statins [n (%)]	7 (13.2)	33 (57.9)	19 (35.2)	13 (20.6)	24 (48.0)	<0.001
Echocardiographic parameters						
LVEF (%)	67 (63,69)	62 (56, 66)	63 (57, 66)	60 (35, 64)	58 (44, 65)	<0.001
ARD (mm)	33 (30, 34)	34 (30, 36)	33 (31, 36)	33 (31, 37)	34 (32, 38)	0.17
LAD (mm)	36 (33, 38)	40 (37, 43)	41 (38, 44)	47 (41, 51)	44 (42, 48)	<0.001
LVDd (mm)	46 (44, 47)	46 (44, 51)	47 (44, 53)	53 (46, 58)	51 (47, 59)	<0.001
LVDs (mm)	30 (28, 30)	30 (29, 34)	30 (28, 35)	35 (30, 47)	32 (30, 43)	<0.001
IVS (mm)	9 (8, 10)	10 (9,11)	10 (9, 13)	10 (9, 10)	10 (9, 10)	<0.001
LVPW (mm)	8 (8, 9)	9 (9, 10)	9 (9, 10)	10 (9, 11)	10 (9, 11)	<0.01
PAP (mmHg)	30 (29, 30)	31 (30, 35)	33 (31, 38)	36 (31, 44)	35 (31, 45)	<0.001

Continuous variables are presented as mean ± standard error of mean or median (interquartile range). Categorical variables are expressed as number (percentages). Statistically significant values are indicated in italic. IHD, ischemic heart disease; VHD, valvular heart disease; CHD, congenital heart disease; ACEI/ARB, angiotensin-converting enzyme inhibitors or angiotensin II receptor blockers; CCB, calcium channel blocker; OAD, oral antidiabetic drugs; LVEF, left ventricular ejection fraction; ARD, aortic root dimension; LAD, left atrial dimension; LVDd, Left ventricular end-diastolic dimension; LVDs, left ventricular end-systolic dimension; IVS, interventricular septal thickness; LVPW, left ventricular posterior wall; PAP, pulmonary artery pressure.



in the co-occurrence of HF and T2DM, rather than in either individual condition.

# Association of sFRP2 With Clinical Variables

The correlation between sFRP2 and clinical variables were analyzed with Spearman rank correlation analysis using data from all the 277 participants. The presence and absence of diseases were assigned as 1 and 0, respectively. No, past and current tobacco use were assigned as 0,1,2, respectively. The detailed data are shown in **Table 3**. The Spearman correlation coefficient between sFRP2 and FPG was <-0.5, suggesting a moderate inverse correlation. SFRP2 had weak negative correlations with HbA1c, cTNT, NT-proBNP and weak positive correlations with hemoglobin and eGFR, as evidenced by the coefficients between -0.4 and -0.2 or 0.2 and 0.4. Although the *P*-values were <0.05, the correlation coefficients of sFRP2 with age, albumin, TC, LDL-C, high-sensitivity C-reactive protein (hs-CRP), LVEF, LAD and LVPW were between -0.2 and 0.2, indicating that the correlations were marginal.

With ln(sFRP2) serving as the dependent variable and the clinical variables correlated with sFRP2 in Spearman correlation analyses as independent variables, a multiple stepwise regression analysis was performed using data from all the 277 participants. During the stepwise regression, age, hemoglobin, Alb, eGFR, HbA1c, cTNT, NT-proBNP, LDL-C and LVPW were removed to reach the lowest Akaike information criterion (AIC) level. Potential confounders such as medications (with or without drugs were assigned as 1 and 0, respectively) were also added as independent variables, but no significant associations were found (data not shown). Finally, FPG, TC, hs-CRP, LVEF and LAD were

**TABLE 3** | Spearman's correlation of sFRP2 with clinical variables.

Indicator	ρ	P-value
Age	-0.1591595	<0.01
Gender	-0.0567796	0.35
BMI	-0.0980334	0.16
Smoking	-0.0538868	0.38
IHD	-0.0252443	0.68
VHD	-0.0699490	0.25
Hypertension	-0.1141791	0.06
Hemoglobin	0.2127307	<0.001
Alb	0.1946989	<0.01
eGFR	0.2082241	<0.001
FPG	-0.5345865	<0.001
HbA1c	-0.3071817	<0.001
cTNT	-0.2118334	<0.001
NT-proBNP	-0.3009578	<0.001
TC	0.1492633	<0.05
TG	-0.0672510	0.29
LDL-C	0.1639592	<0.01
HDL-C	0.1214094	0.06
hs-CRP	-0.1856972	<0.01
LVEF	0.1175641	<0.05
ARD	-0.0618573	0.32
LAD	-0.1438706	<0.05
LVDd	-0.1023572	0.10
LVDs	-0.116702	0.06
IVS	-0.093653	0.13
LVPW	-0.1213423	<0.05
PAP	-0.0072189	0.91

Statistically significant values are indicated in italic. BMI, body mass index; IHD, ischemic heart disease; VHD, valvular heart disease; Alb, albumin; eGFR, estimated glomerular filtration rate; FPG, fasting plasma glucose; HbA1c, glycated hemoglobin; cTNT, cardiac troponin T; NT-proBNP, N-terminal pro-B-type natriuretic peptide; TC, total cholesterol; TG, triglyceride; LDL-C, low-density lipoprotein cholesterol; HDL-C, high-density lipoprotein cholesterol; hS-CRP, high-sensitivity C-reactive protein; LVEF, left ventricular ejection fraction; ARD, aortic root dimension; LAD, left atrial dimension; LVDd, Left ventricular end-diastolic dimension; LVDs, left ventricular posterior wall; PAP, pulmonary artery pressure.

selected as independent variables in the optimized model. The *P*-value of the model was <2.2e-16 and the adjusted *R*-squared was 0.5109, which indicated 51.09% of the changes in ln(sFRP2) could be attributed to these factors (**Table 4**). VIFs were <1.5 in all analyses, suggesting that no collinearity was observed. As shown in **Figure 2**, residuals were equally distributed and approximately accorded with a normal distribution.

# Relationship Between sFRP2 and the Risk of HF

To investigate the risk factors of HF in patients with CVDs, a univariate logistic regression analysis was performed between pre-clinical HF patients (the pre-clinical HF group and preclinical HF+T2DM group) and HF patients (the HF group and HF+T2DM group). In total, 224 of the 277 participants were included in this analysis. As shown in **Table 5**, hemoglobin, albumin, eGFR and ln(sFRP2) were negatively associated with the odds of HF, whereas FPG, HbA1c, and hs-CRP showed positive associations with the odds of HF. The 7 variables with significance were further investigated in the multivariate logistic stepwise regression analysis. Finally, hemoglobin, albumin, eGFR and ln(sFRP2) were included in the logistic regression equation (**Table 6**).

# DISCUSSION

In our study, sFRP2 decreased significantly in the HF+T2DM group compared with the control group and the pre-clinical HF+T2DM group. Serum sFRP2 was negatively correlated with cTNT, NT-proBNP, LAD, LVPW, and positively correlated with LVEF. In addition, higher sFRP2 was significantly associated to lower odds of HF in patients with CVDs. These results suggest that the decreased circulating sFRP2 may be a risk factor for HF. Several experimental studies supported our observations and revealed the potential mechanisms. sFRP2 attenuated

	Estimate	Std. error	t value	P-value
(Intercept)	7.361267	0.414689	17.751	<0.001
FPG	-0.179817	0.015973	-11.258	<0.001
TC	0.058419	0.030584	1.910	0.06
hs-CRP	-0.006420	0.001744	-3.682	<0.001
LVEF	0.005843	0.003451	-1.693	0.07
LAD	-0.014792	0.006302	-2.347	<0.05

Statistically significant values are indicated in italic. FPG, fasting plasma glucose; TC, total cholesterol; hs-CRP, high-sensitivity C-reactive protein; LVEF, left ventricular ejection fraction; LAD, left atrial dimension.

cardiac fibrosis and HF induced by pressure overload (19) and autoimmune myocarditis (16) *via* the Wnt/ $\beta$ -catenin and TGF- $\beta$  pathways, respectively. sFRP2 prevented cardiomyocyte apoptosis (21) and pathological hypertrophy (19). *SFRP2* 

 TABLE 5 | Risk factors for HF in 244 cardiac patients: univariate logistic regression.

	OR (95% CI)	P-value
Age	1.0166780 (0.9958878, 1.0379020)	0.12
Gender	0.8284024 (0.4743534, 1.4467070)	0.51
BMI	1.0403330 (0.9532089, 1.1354200)	0.38
Smoking	0.7870194 (0.5334641, 1.1610890)	0.23
Hypertension	0.8865784 (0.5183912, 1.5162700)	0.66
Hemoglobin	0.9744476 (0.9620449, 0.9870103)	<0.001
Alb	0.8674282 (0.8141866, 0.9241514)	<0.001
eGFR	0.9779109 (0.9655423, 0.9904378)	<0.01
FPG	1.1219660 (1.0124180, 1.2433680)	<0.05
HbA1c	1.2742450 (1.0220770, 1.5886280)	<0.05
cTNT	1.0953920 (0.8179942, 1.4668600)	0.54
TC	1.0374960 (0.8384569, 1.2837850)	0.74
TG	0.9669338 (0.7421804, 1.2597490)	0.80
LDL-C	1.0332420 (0.7431301, 1.4366110)	0.85
HDL-C	0.5222179 (0.2337020, 1.1669200)	0.11
hs-CRP	1.0210930 (1.0052670, 1.0371690)	<0.01
In(sFRP2)	0.5360921 (0.3668849, 0.7833375)	<0.01

Statistically significant values are indicated in italic. BMI, body mass index; Alb, albumin; eGFR, estimated glomerular filtration rate; FPG, fasting plasma glucose; HbA1c, glycated hemoglobin; cTNT, cardiac troponin T; TC, total cholesterol; TG, triglyceride; LDL-C, low-density lipoprotein cholesterol; HDL-C, high-density lipoprotein cholesterol; hs-CRP, high-sensitivity C-reactive protein.



TABLE 6   Risk factors for HF in 244	cardiac patients: multivariate logistic
regression.	

	OR (95% CI)	P-value
Hemoglobin	0.9856071 (0.9714623, 0.9999579)	<0.01
Alb	0.9123722 (0.8501815, 0.9791121)	<0.05
eGFR	0.9835357 (0.9700838, 0.9971742)	<0.01
In(sFRP2)	0.6633438 (0.4362906, 0.9985590)	<0.05

Statistically significant values are indicated in italic. Alb, albumin; eGFR, estimated glomerular filtration rate.

transgene mice exhibited higher LVEF and reduced infarct size due to the increased angiogenesis mediated by ATF6 signaling (20). In cardiac regeneration therapy, sFRP2 also improved bone marrow stromal cells (BMSCs) transplantation (22, 23) and enhanced the differentiation of cardiac progenitor cells (CPCs) (47). However, Yang et al. reported that in HF patients, sFRP2 was positively associated with extracellular volume (ECV) fraction, a parameter in cardiovascular magnetic resonance (CMR) imaging used to assess myocardial fibrosis. During follow-up periods, patients with primary cardiac events had higher levels of sFRP2 (25). Since they only enrolled patients with severe decompensated HF and NYHA functional class III-IV, the disparity may be attributed to different stages of HF and the elevated sFRP2 in their study may be a compensatory factor to counteract myocardial fibrosis. Indeed, an initial compensatory increase of cardiac sFRP2 level followed by a substantial decrease of sFRP2 during the progression of HF has already been reported in an animal model (19).

Our study also revealed that sFRP2 was lower in the HF+T2DM group than in the HF group and was negatively correlated with FPG and HbA1c. A previous study showed higher sFRP2 levels in patients with AGT (28). Since the diabetic patients in our study was subdivided from patients with CVDs, the effect of CVDs on sFRP2 may be the main reason for the difference. This discrepancy may also be attributed to their enrollment of pre-diabetes patients. Prediabetes is also associated with increased risk of CVDs (48) and HF (49). Further studies are needed to explore whether sFRP2 is associated with prediabetes and whether this association, if any, is involved in CVDs and HF.

It is intriguing that serum sFRP2 levels reduced in the cooccurrence of HF and T2DM, rather than in either individual disease. We performed a univariate logistic regression analysis specifically in patients with CVDs and T2DM (the pre-clinical HF+T2DM group and the HF+T2DM group, data not shown), and found a more obvious inverse association between ln(sFRP2) and HF (OR: 0.42, 95% CI: 0.24-0.74, P < 0.01) compared with that in all the patients with CVDs (OR: 0.54, 95% CI: 0.37–0.78, P < 0.05). Patients with the comorbidity of HF and T2DM have distinct biomarkers compared to those without T2DM (50) and sFRP2 is potentially one of them. A falling heart undergoes metabolic perturbations characterized by impaired lipid metabolism and a substrate preference switch toward glycolysis and ketone body oxidation (51). The hyperglycemia, insulin resistance, and hyperinsulinemia in T2DM trigger a cascade of deleterious effects that contribute to the development of HF (52). sFRP2 was reported to be associated with lipogenesis and insulin resistance (53), which indicated that sFRP2 might modulate heart function *via* regulating metabolism.

In our study, the level of hs-CRP in the HF+T2DM group was higher than that in other groups and sFRP2 was negatively correlated with hs-CRP. This is unsurprising since HF and T2DM are accompanied by chronic low-grade systemic inflammation (54). Previous studies have reported the association between sFRP2 and inflammation. sFRP2 was downregulated in inflammation-induced muscle atrophy and prevented inflammatory muscle atrophy (55). sFRP2 enhanced the osteo/odontogenic differentiation and paracrine potentials under inflammation conditions via inhibiting canonical Wnt/βcatenin and nuclear factor kappa B (NF-kB) signaling pathways (56). However, Zhou et al. demonstrated that serum sFRP2 was significantly upregulated in patients with COPD and knockdown of sFRP2 in peripheral blood mononuclear cells (PBMCs) attenuated airway inflammation (57). Thus, further studies are mandated to explore the detailed roles of sFRP2 in cardiovascular inflammation.

The main strengths of our study are the inclusion of people with different HF stages, which allow us to explore the association of sFRP2 and severity of HF. Furthermore, we collected and adjusted multiple cardiovascular risk factors. However, several limitations should be noted. First, the crosssectional design precluded us from drawing casual conclusions and further studies are required to define the role of sFRP2 in cardiac function and glucose homeostasis. Second, there were significant differences in etiologies between each group, although there were no significant differences of sFRP2 between ischemic and non-ischemic heart diseases in each group (data not shown) and Spearman rank correlation analysis confirmed no significant association between sFRP2 and etiologies. Third, the measurement of serum sFRP2 by ELISA could not reflect its sources and targets. SFRP2 gene is ubiquitously expressed in adipose tissue, small intestine, colon, heart, skeletal muscle and other organs, with the highest expression in adipose tissue (58, 59). As an adipokine, circulating sFRP2 is not linearly correlated with SFRP2 gene expression in WAT (27), indicating that other sources of sFRP2 production should be investigated. Since the cardiac sFRP2 expression increased in the early stage of HF and subsequently decreased with the development of HF (19), less sFRP2 production from heart may explain the declined circulating sFRP2. Previous studies focused on the autocrine and paracrine function, while the changes in serum sFRP2 may indicate its endocrine function. SFRP2 exerts effects on cardiomyocytes, cardiac fibroblast cells, endothelial cells (24) and adipocytes (28) and its function exhibits a high degree of cell specificity. More researches are necessary to explore why serum sFRP2 reduced in the comorbidity of HD and T2DM and the relevant mechanism.

# CONCLUSIONS

The level of sFRP2 was negatively correlated with age, FPG, HbA1c, cTNT, NT-proBNP, hs-CRP, LAD and LVPW, and positively correlated with hemoglobin, eGFR, albumin, TC, LDL-C and LVEF. Higher serum sFRP2 was significantly linked to lower odds of HF in patients with CVDs. sFRP2 is a promising

risk factor for the comorbidity of HF and T2DM, which might pave novel ways for the diagnosis and treatment of clinical HF from the perspective of metabolism.

## DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

# ETHICS STATEMENT

The studies involving human participants were reviewed and approved by Ethics Committee of Zhongshan Hospital affiliated to Fudan University. The patients/participants provided their written informed consent to participate in this study.

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# **AUTHOR CONTRIBUTIONS**

PG and MC designed the study. PG, MC, HW, and XW performed the experiments. WL and XJ contributed to the interpretation of the results. MC analyzed the data and wrote the first draft of the manuscript. WG, PG, and YZ supervised the project and revised the paper. All authors contributed to the article and approved the submitted version.

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