

# Associations between insulin-like growth factor-1 standard deviation score and overall nutritional parameters in patients with maintenance hemodialysis: a cross-sectional study

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# Abstract

**Background** This study investigated the association between insulin-like growth factor-1 and nutritional status indicators in patients undergoing maintenance hemodialysis (MHD).

**Methods** Patients undergoing MHD for > 3 months were included in this single-center cross-sectional study in March 2021. Clinical, demographic, and body mass data and blood samples were collected before the hemodialysis sessions. Serum insulin-like growth factor-1 (IGF-1) levels were measured using a radioimmunoassay, and serum IGF-1 standard deviation score (SDS) was calculated for MHD patients according to age and sex. The nutritional status of patients was assessed using serum albumin, serum prealbumin, handgrip strength, pinching strength, upper arm muscle circumference, lean body mass, phase angle, seven-point subjective global assessment (SGA) score, and geriatric nutritional risk index (GNRI). The patients were divided into groups according to tertiles of serum IGF-1 SDS levels. Spearman correlation analyses and univariate and multivariate binary logistic regression analyses were used to determine the association between serum IGF-1 SDS and nutritional status parameters.

**Results** A total of 155 MHD patients (male: female = 90:65) were enrolled in the study, with a median dialysis vintage of 28.0 (11.0, 55.0) months, and an average age of 66 (65.5 ± 13.0) years. The median of IGF-1 SDS was -0.1 (-0.6 to 0.6). Compared to patients with higher IGF-1 SDSs, patients with lower IGF-1 SDSs had lower levels of serum ceruloplasmin (341.0 [287.5, 416.0] vs 395.0 [327.0, 451.0] vs 409.0 [349.5, 507.5], p=0.002), serum albumin (34.7 ± 3.0 vs 37.0 ± 3.1 vs 37.8 ± 2.6, p < 0.001), serum prealbumin (270.3 [233.7, 327.8] vs 326.0 [279.3, 355.6] vs 363.0 [324.2, 398.2], p < 0.001), handgrip strength (13.8 [10.0, 20.7] vs 17.7 [10.7, 22.5] vs 23.3 [16.6, 27.8], p < 0.001), pinch strength (4.6 [3.9, 6.0] vs 4.9 (3.9, 6.9) vs 6.5 [4.7, 8.7], p=0.002), phase angle (3.3 [3.0, 3.8] vs 3.9 [3.4, 4.7] vs 4.3 [3.6, 5.2, p < 0.001), modified Creatinine Index (83.1 ± 19.7 vs 93.1 ± 23.4 vs 113.9 ± 24.3, p < 0.001), intracellular water (14.5 ± 4.4 vs 16.1 ± 4.9 vs 16.9 ± 4.4, p=0.031), higher extracellular water (26.9 ± 5.8 vs 25.7 ± 5.5 vs 25.1 ± 3.1, p=0.042), and higher malnutrition risk as defined by GNRI (49.0% vs 15.7% vs 11.5%, p < 0.001) and SGA (53.9% vs 23.5% vs 7.7%, p < 0.001). **Conclusions** Lower IGF-1 SDSs are independently associated with higher malnutrition risk in patients with MHD.

**Keywords** Insulin-like growth factor-1 standard deviation score  $\cdot$  Seven-point subjective global assessment  $\cdot$  Geriatric nutritional risk index  $\cdot$  Protein-energy wasting  $\cdot$  Hemodialysis  $\cdot$  End-stage renal disease

	Abbreviations		
	MHD	Maintenance hemodialysis	
	CKD	Chronic kidney disease	
Tingting Xing and Yao Xu have contributed equally to this work.	ESRD	End-stage renal disease	
	SGA	Seven-point subjective global assessment	
🖂 Yun Liu	GNRI	Geriatric Nutritional Risk index	
316463862@qq.com	GH	Growth hormone	
🖂 Rongshao Tan	IGFBPs	Insulin-like growth factor bind proteins	
tanrongshao@126.com	IGF-1	Insulin-like growth factor-1	

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IGF-1 SDS	Insulin-like growth factor-1 standard devia-
	tion score
PEW	Protein-energy wasting
DM	Diabetes mellitus
IL-6	Interleukin-6
hs-CRP	Hypersensitive C-reactive protein
mCI	Modified Creatinine Index
ECW	Extracellular body water
ICW	Intracellular body water

# Introduction

Patients with chronic kidney disease (CKD) are prone to malnutrition or protein-energy wasting (PEW), which is a strong risk factor for poor prognoses in patients undergoing maintenance hemodialysis (MHD) [1]. The worldwide prevalence of PEW ranges from 11 to 54% in patients with CKD stages 3–5 [2], and between 28 and 54% in patients undergoing dialysis [3]. The prevalence of PEW varies in different studies, depending on the assessment method. The Seven-Point Subjective Global Assessment (SGA) [4] and Geriatric Nutritional Risk Index (GNRI) [5] are widely used tools for PEW assessment. Abnormalities in growth hormone (GH)/ insulin-like growth factor-1 (IGF-1) are considered significant factors in the development of skeletal muscle wasting in patients with CKD via inflammation [6]. Studies have shown that lower serum IGF-1 levels are significantly associated with biochemical and anthropometric markers of malnutrition [7, 8]. Researchers have also observed an inverse association between serum IGF-1 levels and nutritional status assessed by traditional SGA in patients undergoing MHD [9]. Since serum IGF-1 levels are strongly related to age or sex, serum IGF-1 SDSs were calculated using the method described by Isojima et al. to minimize the bias of these confounding factors [10].

To the best of our knowledge, no study has analyzed the relationship between standardized IGF-1 levels and indicators of comprehensive nutritional assessment such as the 7-point SGA and GNRI, in patients with MHD. Hence, we designed the present cross-sectional study to explore the relationship between IGF-1 SDS levels and the overall nutritional status indicators in these patients.

### Methods and materials

#### Population and study design

This was a single-center, observational, cross-sectional study of patients undergoing MHD at Guangzhou Red Cross Hospital in March 2021. Patients included (1) had end-stage renal disease (ESRD); (2) performed maintenance hemodialysis regularly for more than 3 months, 3 times a week, and for four hours each time; (3) aged 18–80 years; and (4) provided informed consent. The exclusion criteria were: (1) neurological or mental illness or inability to complete the questionnaire; (2) history of acute heart failure, severe infection, or malignant tumor within 3 months; (3) history of surgery within 3 months; (4) pacemaker use; and (5) refusal to participate in the study.

#### **Research methodology**

#### Demographic, clinical, and laboratory parameters

Data on the following demographic and clinical parameters were collected: history of diabetes, sex, age, and dialysis vintage. Venous blood samples were collected by nurses shortly before the hemodialysis sessions. They were sent to the clinical laboratory of Guangzhou Red Cross Hospital within 2 h. Routine blood tests, serum biochemical tests, and enzyme immunity measurements were performed. Further, the following parameters were measured: Serum creatinine, Kt/V, serum albumin, serum prealbumin, serum IGF-1, serum interleukin-6 (IL-6), serum ceruloplasmin, and serum hypersensitive C-reactive protein (hs-CRP).

## Measurement and calculation of IGF-1

Serum IGF-1 was measured by high-performance liquid chromatography-mass spectrometry using a Thermo Q Exactive Focus instrument (Thermo Fisher Scientific, Waltham, MA, USA). The assay was calibrated using standards prior to testing. All reagents used in this study (including inoculated strains, reagent wedges, calibrators, and dilutions) were obtained from the same batch. The intra-group coefficients of variation ranged from 2.4% to 6.3%, and the inter-group coefficients of variation ranged from 3.0% to 7.6%. The sensitivity was 20  $\mu$ g/L, and the upper limit of detection was 1600  $\mu$ g/L. The SDS of serum IGF-1 was calculated using the following formula [10]:

SDS = (measurement - mean)/standard deviation (SD).

#### **Nutritional indicators**

**Body mass** A body composition analyzer (Bodystat5000, UK) was used to measure the body composition of patients. According to the manufacturer's guidelines, measurements were performed in the supine position, with electrodes attached to the hands and feet on the side without a fistula for hemodialysis. Resistance (R in ohms) and reactance (Xc in ohms) values were recorded at 50 kHz. Patients with implantable electronic devices (e.g., electronic heart pace-

makers) were excluded. Lean body mass, phase angle, extracellular water (ECW), ECW%, intracellular water (ICW), ICW%, and the ECW/ICW ratio were measured during the process.

**Calculation of nutritional indices** According to the 7-point SGA, a total score of 6 to 7 was classified as normal nutritional status, 3 to 5 as mild to moderate malnutrition, and 1 to 2 as severe malnutrition. A 7-point SGA score  $\leq$  5 points was used as the diagnostic criterion for malnutrition or PEW [11].

The GNRI was calculated according to the following formula:

GNRI =1.489 × serum albumin (g/L) + 41.7 × (actual body weight/ideal body weight)

where the ideal body weight was calculated as  $22 (kg/m^2) \times height$  [12]. If the actual body weight was above the ideal body weight, the value of "(actual body weight/ideal body weight)" was set to 1. At present, some studies suggest that a GNRI in MHD patients of <91.2 can be defined as a risk of malnutrition [13].

The mCI was calculated using parameters including sex, age, spKt/V (for urea clearance), and pre-hemodialysis creatinine level, using the following formula [14]:

$$\begin{split} \text{MCI} \; (\text{mg/kg/day}) &= 16.21 + 1.12 \times (0 \text{ for women; } 1 \text{ for men}) \\ &\quad - 0.06 \times \text{age (years)} - 0.08 \times SPKt/V \text{ for urea} \\ &\quad + 0.009 \times \text{pre-hemodialysis creatinine (umol/L)}. \end{split}$$

**Grip strength and pinch strength** Grip strength and pinch strength were measured using the BASELINE digital Grip Force Tester (12-0091, Fabrication Enterprises Inc., USA) and the BASELINE digital Pinch Force tester (12-0081, Fabrication Enterprises Inc., USA). The participants were instructed to apply as much grip or pinch force as possible on the instruments with the dominant hand or the hand without fistulae. The measurement was repeated thrice, and the maximum value was recorded.

**Anthropometry** Upper arm circumference was measured according to the criteria developed by Frisancho [15]. The upper arm muscle circumference was calculated according to the following formula:

upper arm muscle circumference (cm)

- = upper arm circumference (cm) 3.14
  - × triceps skinfold thickness (cm).

#### Parameters of hemodialysis

Patients in this study were treated on a Braun Dialog + (B. Braun Co., Ltd., Melsungen, Germany) dialysis machine using a REXEED-15L high-throughput polysulfone membrane dialyzer (Asahi Kasei Corp., Tokyo, Japan) with a membrane area of  $1.5 \text{ m}^2$ , dialysis blood flow rate of 200–300 ml/min, dialysis fluid flow rate of 500 ml/min, and dialysis duration of 4 h.

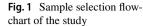
#### **Statistical analysis**

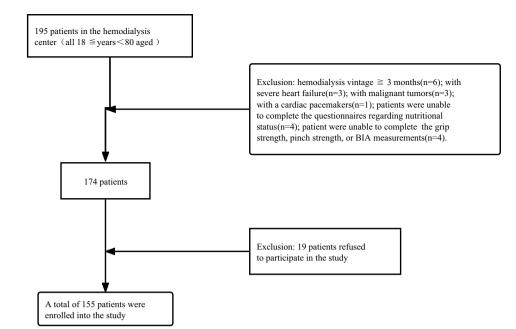
Continuous variables with a normal distribution were described as  $\overline{x} \pm s$ , and variables with a non-normal distribution were described as the median (25-75% interquartile range). Categorical variables were described as percentages. Patients were divided into three groups based on their serum IGF-1 SDS levels. Differences in clinical and demographic data among the three groups were assessed by one-way analysis of variance, Kruskal–Wallis test, or  $\chi^2$  test, as appropriate. Spearman analyses were used to evaluate the correlation between the IGF-1 SDS and nutritional indicators. Univariate analyses were conducted to select potential explanatory variables with dependent variables of malnutrition defined as SGA <5 and GNRI <91.2. In multivariate binary logistic regression analyses, potentially relevant variables or known to be important in the physiology of malnutrition: sex, age, dialysis vintage, Kt/V ( $\geq$  1.2), DM, IL-6, phosphorus, and 25-OH-D values, were included to determine whether IGF-1 SDS levels was independently associated with malnutrition. SPSS (version 22.0; IBM Corp, Armonk, NY, USA) was used for statistical analyses. Significance was set at p < 0.05.

# Results

#### Characteristics of the study population

A total of 155 patients with MHD (Fig. 1) were included (male/female = 90:65), with a median dialysis vintage of 28.0 (11.0, 55.0) months and an average age of 66 ( $65.5 \pm 13.0$ ) years. The sample selection flowchart is shown in Fig. 1. The baseline characteristics of the total study population and those after stratification by serum IGF-1 tertile are presented in Table 1. The median serum IGF-1 level was 186.0 (140.0, 248.7) ng/mL. The median of IGF-1 SDS levels were levels - 0.1 (- 0.6 to 0.6). Malnourished patients, as defined by the 7-point SGA and the GNRI, accounted for 28.4% (n=44) and 25.3% (n=39) of the study population, respectively. Patients with lower serum IGF-1 levels were older and had lower levels of serum creatine, serum ceruloplasmin, serum albumin, serum prealbumin, handgrip strength, pinch strength, upper arm circumference, upper





arm muscle circumference, phase angle, modified Creatinine Index (mCI), ICW, higher levels of ECW%, serum IL-6, and higher malnutrition risk, as defined by the SGA and GNRI values (all p < 0.05).

# Correlations between IGF-1 SDS and nutritional indicators

In Spearman's analyses, we observed that the levels of serum IGF-1 SDS were positively correlated with the values of grip strength (r=0.32, p<0.001), pinch strength (r=0.28, p<0.001), phase angle (r=0.33, p<0.001), lean body mass (r=0.30, p<0.001), serum albumin (r=0.42, p<0.001), serum prealbumin (r=0.52, p<0.001), mCI (r=0.52, p<0.001), ICW (r=0.21, p=0.012), and were negatively correlated with the levels of ECW% (r=-0.22, p=0.010) (Fig. 2). There were no associations between ECW (r=0.13, p=0.141), ICW% (r=0.02, p=0.800), and ECW/ICW (r=-0.11, p=0.180).

# Associations between the serum IGF-1 and nutrition risk defined by different nutritional indicators

The results of univariate binary logistic regression analysis are shown in Table 2. We found that the lowest tertile of IGF-1 SDS levels (SGA  $\leq$  5: OR = 13.71, 95% CI = 4.31–43.61, p < 0.001; GNRI  $\leq$  91.2: OR = 7.21, 95% CI = 2.62–19.87, p < 0.001), mCI, and ICW values were related to malnutrition defined by the 7-point SGA and GNRI. Higher levels of serum hs-CRP and IL-6 were related to the malnutrition as defined by the GNRI. The multivariate binary logistic regression results are presented in Table 3. After partial or full adjustment for sex, age, dialysis vintage, Kt/V ( $\geq$  1.2), diabetes mellitus(DM), serum IL-6, serum 25 hydroxyvitamin D<sub>3</sub>, and serum phosphorus levels, we found that the lowest tertile of IGF-1 SDS levels (SGA  $\leq$  5: OR = 9.69, 95% CI = 2.49–37.62, p=0.001; GNRI  $\leq$  91.2: OR = 5.71, 95% CI = 1.64–19.89, p=0.006) were independently positively associated with malnutrition defined as SGA  $\leq$  5 and GNRI  $\leq$  91.2.

# Discussion

Our study showed that the patients with lower serum IGF-1 SDS levels had lower serum levels of creatine, ceruloplasmin, albumin, and prealbumin; higher serum levels of IL-6; lower values of grip strength, pinch strength, upper arm circumference, upper arm muscle circumference, phase angle, and modified Creatinine Index (mCI); and higher malnutrition defined by the two nutrition indicators, SGA and GNRI. After adjusting for confounding factors in multivariate binary logistic regression models, we found that the lowest quartile of IGF-1 SDS levels were independently associated with malnutrition defined by SGA and GNRI.

Disorder of the GH/IGF-1 axis may occur at any stage of CKD. Notably, IGF-1 is mainly produced in the liver. Circulating insulin-like growth factor-binding proteins (IGFBPs), which are transport proteins, modulate IGF-1 bioavailability, prolong its half-life, and regulate its activity in target tissues [16]. Normal total IGF levels in uremia are thought to result from decreased IGF degradation caused by enhanced IGFBP binding. Binding to IGFBP protects IGF1 from metabolic

Table 1 Baseline Characteristics of the study population an	d after stratification by tertiles of serum IGF-1 SDS
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Characteristics	Total ( <i>n</i> = 155)	Tertile 1 $(n=52)$	Tertile 2 $(n=51)$	Tertile 3 $(n=52)$	p value
DM, n (%)	86 (55.5)	27 (51.9)	25 (49.0)	34 (65.4)	0.2
Sex <i>n</i> (%)	90 (58.1)	29 (55.8)	24 (47.1)	37 (71.2)	0.04
Age	$65.5 \pm 13.0$	$73.1 \pm 8.9$	$65.5 \pm 12.4$	$57.9 \pm 12.7$	< 0.001
Dialysis vintage (months)	28.0 (11.0, 55.0)	31.0 (10.0, 46.0)	39.0 (12.0, 56.0)	26.0 (11.5, 61.0)	0.7
Creatinine µmol/L	$933.6 \pm 279.7$	$786.7 \pm 215.4$	$893.7 \pm 254.0$	$1116.7 \pm 262.5$	< 0.001
Kt/V	$1.4 \pm 0.4$	$1.3 \pm 0.3$	$1.4 \pm 0.7$	$1.3 \pm 0.2$	0.2
Albumin (g/L)	$36.5 \pm 3.1$	$34.7 \pm 3.0$	$37.0 \pm 3.1$	$37.8 \pm 2.6$	< 0.001
Prealbumin (mg/L)	326.0 (272.6, 365.5)	270.3 (233.7, 327.8)	326.0 (279.3, 355.6)	363.0 (324.2, 398.2)	< 0.001
IGF-1 (ng/mL)	186.0 (140.0, 248.7)	119.8 (89.5, 139.8)	186.0 (172.0, 197.7)	277.9 (248.8, 320.8)	< 0.001
IGF-1 SDS	- 0.1(- 0.6 to 0.6)	- 0.9 (- 1.3 to 0.7)	- 0.1(- 0.3 to 0.0)	0.9 (0.6 to 1.4)	< 0.001
IL-6 (pg/mL)	8.9 (6.7, 15.6)	11.1 (7.5, 20.9)	8.7 (5.2, 13.7)	8.0 (5.2, 11.0)	0.01
Ceruloplasmin (mg/L)	377.0 (324.0, 457.0)	341.0 (287.5, 416.0)	395.0 (327.0, 451.0)	409.0 (349.5, 507.5)	0.002
hs-CRP (mg/L)	3.20 (1.6, 8.6)	3.40 (1.7, 9.9)	3.6 (2.0, 8.3)	2.6 (1.0, 7.4)	0.1
Grip strength (kg)	17.7 (11.8, 24.4)	13.8 (10.0, 20.7)	17.7 (10.7, 22.5)	23.3 (16.6, 27.8)	< 0.001
Pinch strength (kg)	5.0 (4.2, 6.9)	4.6 (3.9, 6.0)	4.9 (3.9, 6.9)	6.5 (4.7, 8.7)	0.002
Upper arm circumference (cm)	23.5 (22.0, 26.1)	22.9 (21.0, 24.0)	24.0 (21.1, 25.7)	25.6 (23.1, 28.0)	< 0.001
Upper arm muscle circumference (cm)	21.0 (19.1, 23.1)	19.7 (18.3, 21.6)	20.6 (18.9, 22.9)	22.4 (20.6, 24.5)	< 0.001
Lean body mass (kg)	75.3 (66.8, 81.6)	73.1 (63.0, 79.3)	74.2 (66.0, 81.6)	77.4 (68.5, 81.5)	0.3
Phase Angle	3.9 (3.2, 4.7)	3.3 (3.0, 3.8)	3.9 (3.4, 4.7)	4.3 (3.6, 5.2)	< 0.001
GNRI	$94.5 \pm 5.1$	$91.5 \pm 4.7$	$95.3 \pm 5.2$	$96.7 \pm 4.0$	< 0.001
mCI	$96.8 \pm 25.9$	$83.1 \pm 19.7$	$93.1 \pm 23.4$	$113.9 \pm 24.3$	< 0.001
ECW (lt)	$15.6 \pm 3.5$	$16.0 \pm 2.8$	$15.5 \pm 3.6$	$15.1 \pm 4.3$	0.5
ECW%	$25.9 \pm 4.9$	$26.9 \pm 5.8$	$25.7 \pm 5.5$	$25.1 \pm 3.1$	0.042
ICW (lt)	$15.9 \pm 4.6$	$14.5 \pm 4.4$	$16.1 \pm 4.9$	$16.9 \pm 4.4$	0.031
ICW%	$26.6 \pm 7.8$	$26.6 \pm 8.4$	$26.8 \pm 8.8$	$26.7 \pm 6.2$	0.7
ECW/ICW ratio	$1.0 \pm 0.2$	$1.0 \pm 0.3$	$1.0 \pm 0.2$	$0.9 \pm 0.2$	0.2
SGA score $\leq 5$ , $n$ (%)	44 (28.4)	28 (53.9)	12 (23.5)	4 (7.7)	< 0.001
GNRI≤91.2, <i>n</i> (%)	39 (25.3)	25 (49.0)	8 (15.7)	6 (11.5)	< 0.001

Continuous variables were expressed as mean  $\pm$  standard deviation for normally distributed variables and median (interquartile range) for variables with a skewed distribution. Differences between groups were assessed using the *t* test, Mann–Whitney *U* test, or chi-square analysis, as appropriate. A *p* < 0.05 was considered statistically significant

*Kt/v* the urea clearance index, was used to quantify dialysis treatment adequacy (K, dialyzer clearance of urea, *t* dialysis time, *V* volume of distribution of urea, approximately equal to the total body water of the patient), *DM* diabetes mellitus, *IGF-1* insulin-like growth factor-1, *IGF-1 SDS* insulin-like growth factor-1 standard deviation score, *IL-6* interleukin-6, *hs-CRP* hypersensitive C-reactive protein, *GNRI* Geriatric Nutritional Risk Index, *mCI* Modified Creatinine Index, *ECW* extracellular body water, *ICW* intracellular body water

degradation, but could also inhibit IGF1 interaction with its receptor, resulting in impaired IGF1 bioactivity [17]. It has been shown that serum IGF-I levels vary with age, gender, puberty, physiological status, and ethnicity [18]. In one study, serum IGF-1 was evaluated as the SDS of IGF-1 levels based on a Japanese population reference range established according to age and sex [10].

GH and IGF-1 resistance may adversely affect metabolism and nutritional status in adults with ESRD [19]. In another study, researchers observed that protein catabolism was also associated with impaired IGF-I signaling pathways in ESRD-related sarcopenia [20]. Serum IGF-1 is a positive marker of skeletal muscle strength and mass in patients undergoing hemodialysis [21]. In addition, phase angle (PhA) is a parameter obtained from direct measurements of bioelectrical impedance analysis (BIA). It is widely used as a marker of cellular health and has been recognized as a valuable measure for nutritional assessment, reflecting both muscle mass and muscle function [22, 23]. Extracellular body water (ECW) and intracellular body water (ICW) measured using BIA have also been introduced as markers, and studies have shown that the ECW/ICW ratio is associated with malnutrition [24]. The IGF-deficient group has been found to have a lower PhA, ICW%, and higher ECW values [25]. Other researchers have also found a correlation between serum IGF-1 and biochemical and anthropological parameters of malnutrition in patients with MHD [26, 27]. Our results showed that serum

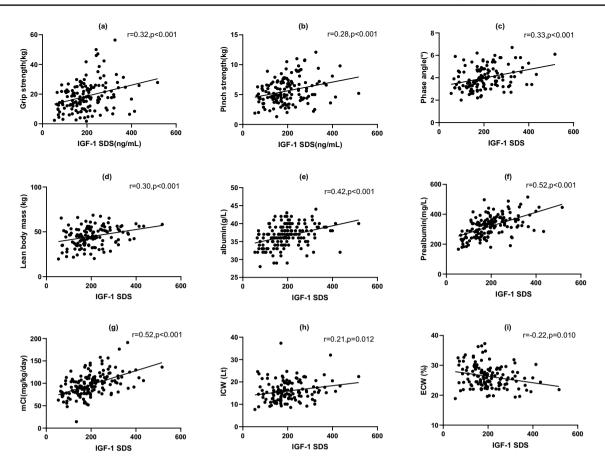


Fig. 2 Spearman's correlation analyses between levels of grip strength (a), pinch strength (b), phase angle (c), lean body mass (d), serum albumin (e), serum prealbumin (f), mCI (g), ICW (h), ECW% (i), and serum levels of IGF-1 SDS

Clinical parameters	Malnutrition by SGA $\leq 5$			Malnut	Malnutrition by GNRI $\leq 91.2$		
	OR	95% CI	<i>p</i> _value	OR	95% CI	<i>p</i> _value	
IGF-1 SDS	0.25	0.13~0.46	< 0.001	0.39	0.23~0.64	< 0.001	
IGF-1 T3	Ref	Ref	Ref				
IGF-1 T1	13.71	4.31~43.61	< 0.001	7.21	2.62~19.87	< 0.001	
IGF-1 T2	3.52	1.05~11.79	0.04	1.36	0.44~4.25	0.59	
hs-CRP mg/L	1.01	0.98~1.04	0.56	1.05	1.01~1.09	0.006	
IL-6 pg/mL	1.02	0.99~1.05	0.06	1.04	1.01, 1.07	0.01	
mCI	0.95	0.93~0.97	< 0.001	0.96	$0.95 \sim 0.98$	< 0.001	
ECW (Lt)	0.87	$0.78 \sim 0.98$	0.02	0.90	0.81~1.01	0.07	
ECW%	1.06	0.97~1.16	0.20	1.07	0.97~1.17	0.16	
ICW(Lt)	0.84	$0.75 \sim 0.94$	0.002	0.88	0.80~0.98	0.01	
ICW%	0.99	0.94~1.04	0.79	1.0	0.96~1.06	0.86	
ECW/ICW ratio	2.79	0.65~11.74	0.16	4.17	0.97~18.16	0.06	

T1 Tertile 1, T2 Tertile 2, T3 Tertile 3, Ref take the third quartile as a reference, IGF-1 insulin-like growth factor-1, hs-CRP hypersensitive C-reactive protein, IL-6 interleukin-6, mCI Modified Creatinine Index, ECW extracellular body water, ICW intracellular body water

Table 2Correlations betweennutritional indicators andclinical parameters

Table 3 Associations between the serum IGF-1 and malnutrition defined by different nutritional indicators

	Malnutrition by SGA $\leq 5$			Malnutrition by GNRI $\leq$ 91.2			
	IGF-1 T3 OR (95% CI) P	IGF-1 T1 OR (95% CI) p	IGF-1 T2 OR (95% CI) p	IGF-1 T3 OR (95% CI) <i>p</i>	IGF-1 T1 OR (95% CI) p	IGF-1 T2OR (95% CI) p	
Model 1	Ref	13.71 4.31~43.61<0.001	3.52 1.05~11.79 0.041	Ref	7.21 2.62~19.87 <0.001	1.36 0.44~4.25 0.593	
Model 2	Ref	17.96 3.1~103.97 0.001	2.95 0.54 ~ 16.21 0.213	Ref	6.81 1.5~30.81 0.013	1.36 0.27~6.85 0.711	
Model 3	Ref	14.25 2.7~75.15 0.002	2.3 0.46~11.6 0.312	Ref	7.06 1.57~31.85 0.011	1.72 0.37~8.06 0.489	
Model 4	Ref	11.5 2.23 ~ 59.16 0.003	1.7 0.34~8.58 0.52	Ref	5.74 1.29~25.54 0.022	1.35 0.28~6.38 0.707	
Model 5	Ref	14.82 2.62~84 0.002	2.25 0.41~12.3 0.351	Ref	5.71 1.27~25.7 0.023	1.15 0.22~5.95 0.867	
Model 6	Ref	13.66 2.68~69.52 0.002	2.2 0.45~10.67 0.327	Ref	6.82 1.56~29.74 0.011	1.67 0.37~7.55 0.508	
Model 7	Ref	9.69 2.49~37.62 0.001	2.74 0.68 ~ 11.13 0.158	Ref	5.71 1.64~19.89 0.006	1.88 0.22~7.46 0.85	

T1 Tertile 1, T2 Tertile 2, T3 Tertile 3, Ref take the third quartile as a reference

Model 1: Unadjusted

Model 2: Adjusted for sex, age, dialysis vintage, Kt/V (≥1.2), DM, Serum IL-6 levels

Model 3: Adjusted for sex, age, dialysis vintage, Kt/V (≥1.2), DM, Serum phosphorus

Model 4: Adjusted for sex, age, dialysis vintage, Kt/V ( $\geq$  1.2), DM, Serum 25 hydroxyvitamin D3

Model 5: Adjusted for sex, age, dialysis vintage, Kt/V (≥1.2), Serum IL-6, Serum 25 hydroxyvitamin D3

Model 6: Adjusted for sex, age, dialysis vintage, Kt/V (≥1.2)+DM

Model 7: Adjusted for sex, age, dialysis vintage, DM, serum IL-6 level, and serum phosphorus level

Kt/v urea clearance index, DM diabetes mellitus, IL-6 interleukin-6

albumin, serum prealbumin, grip strength, pinch strength, body phase angle, lean body mass, mCI, and ICW were all significantly and positively correlated with serum IGF-1 SDS, and ECW% was negatively correlated with serum IGF-1 SDS, which is consistent with the results of a previous study [24–27]. However, our study did not find a correlation between serum IGF-1 SDS and the ECW/ICW ratio.

However, the underlying factors associated with these findings are complex. First, IGF-1 induces protein synthesis and myogenesis by activating the Akt/mTOR pathway, resulting in the growth and repair of skeletal muscles [20]. Together with our results, these findings indicate that serum IGF-1 levels may reflect protein anabolism and skeletal muscle function in MHD patients. Second, lower serum IGF-1 levels may reflect inflammatory status. Chronic inflammation has been reported to disrupt the GH/IGF-1 axis through relative GH and/or IGF-1 insufficiency, peripheral resistance to GH/IGF-1 receptors, inhibition of GH/IGF-1 signaling, dysregulation of IGF-binding proteins, reduced IGF-1 bioavailability, and altered gene regulation through the micro-RNA system [28–32].

Clinical data showed significantly higher levels of circulating IL-6 and IL-6 receptors in older individuals [33], with IL-6 levels being a significant predictor of sarcopenia [34]. Lastly, there is a negative correlation between plasma levels of CRP and IL-6 and rates of mixed muscle and myosin heavy chain protein synthesis in a population-based study of the general community, further illustrating the likelihood of inflammation contributing to reduced protein synthesis in sarcopenia [35]. Researchers have observed that inflammation and malnutrition in patients with ESRD often coexist and are mutually causal, a condition termed malnutrition–inflammation complex syndrome (MICS) [36]. Possible causes of MICS include comorbidities, oxidative and carbonyl stress, nutrient loss through dialysate, anorexia, loss of appetite, uremic toxins, decreased clearance of inflammatory cytokines, and volume overload [37]. In conclusion, a possible mechanism for attenuated skeletal muscle protein synthesis in CKD may be suppression of insulin/IGF-1 signaling due to metabolic acidosis, upregulated pro-inflammatory cytokine expression, and malnutrition due to anorexia [38].

Finally, in persistent systemic and tissue inflammation, the impairment of IGF-I-mediated signaling pathways preferentially increases muscle protein catabolism and inhibits muscle protein anabolism, resulting in excessive muscle protein loss, high hospitalization rates, and high rates of cardiovascular events and mortality among patients with CKD and ESRD [37, 39]. In our study, we also found that the group with a lower IGF-1 SDS had higher IL-6 levels; therefore, the results support the above notion.

This study has some limitations. First, this was a crosssectional observational study, and causal conclusions could not be drawn. Second, given that our study had a small sample size, bias cannot be ruled out. Third, our results cannot exclude the possibility of residual confounding. Lastly, we did not measure serum IGFBPs and growth hormone levels; therefore, the relationship between the GH/IGF-1 axis and nutritional status in patients with MHD remains unclear.

Although a crossover clinical trial and randomized controlled trial have shown that the application of recombinant human growth hormone and IGF-1 could improve protein synthesis and metabolism in patients with ESRD [40, 41], these studies included only a small number of patients with short follow-up periods. However, there are currently no prospective studies with large enough sample sizes or sufficient evidence to determine the potential long-term side effects of recombinant IGF-1 or GH supplementation in patients with MHD.

Our study confirmed that higher serum IGF-1 SDS levels are significantly associated with better nutritional status as assessed by biochemical markers, anthropometric measures, body composition parameters, and comprehensive nutrition assessment. Future observational and interventional studies with larger sample sizes and multicenter are still needed to provide a theoretical basis for routine clinical application of IGF-1 preparations to improve nutritional status in patients with ESRD and dialysis.

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**Data availability statement** The data that support the findings of this study are available from the corresponding author upon reasonable request.

#### Declarations

**Conflict of interest** The authors have declared that no conflict of interest exists.

**Ethical approval** This study was approved by the Ethics Committee of Guangzhou Red Cross Hospital [No. 2021-066-01]. This study adhered to the tenets of the Declaration of Helsinki and the Guidance on Sample Collection of Human Genetic Diseases by the Ministry of Public Health of China and does not contain any studies with animals performed by any of the authors.

**Informed consent** Informed consent was obtained from all individual participants included in the study.

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