

A decorative border at the top of the page featuring various food icons such as fish, peppers, mushrooms, and fruits in a colorful, repeating pattern.

NUTRITIONAL ASPECTS OF KIDNEY DISEASE

EDITED BY: Alice Sabatino and Carla Maria Avesani
PUBLISHED IN: Frontiers in Nutrition





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ISSN 1664-8714

ISBN 978-2-88974-005-5

DOI 10.3389/978-2-88974-005-5

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NUTRITIONAL ASPECTS OF KIDNEY DISEASE

Topic Editors:

Alice Sabatino, University of Parma, Italy

Carla Maria Avesani, Karolinska Institutet, Sweden

Citation: Sabatino, A., Avesani, C. M., eds. (2021). Nutritional Aspects of Kidney Disease. Lausanne: Frontiers Media SA. doi: 10.3389/978-2-88974-005-5

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Editorial: Nutritional Aspects of Kidney Disease

Alice Sabatino^{1*} and Carla Maria Avesani²

¹ Dipartimento di Medicina e Chirurgia, Università di Parma, Parma, Italy, ² Medical Unit Clinical Nutrition, Division of Renal Medicine and Baxter Novum, Department of Clinical Science, Technology and Intervention, Karolinska Institutet, Karolinska University Hospital, Stockholm, Sweden

Keywords: kidney disease, nutrition, chronic kidney disease, acute kidney injury, body composition, end-stage kidney disease, dialysis

Editorial on the Research Topic

Nutritional Aspects of Kidney Disease

The kidneys are homeostatic organs that remove waste products from the metabolism and regulate the concentration of mineral, ions and the volume of body fluids. In the presence of kidney disease (KD) this function is compromised, leading to accumulation of water, minerals and waste products in the body. A close nutritional follow-up which comprises periodical nutritional assessment and supervised dietary modification is key for the management of KD (1, 2), playing an important role in reducing the risk for cardiovascular disease and managing the clinical effects of the reduced glomerular filtration rate, while keeping an optimal nutritional status (3).

This recent Research Topic on “Nutritional aspects of kidney disease” represents a collection of 11 articles, including 8 original research articles, ranging from pathophysiological-related mechanisms to body composition assessment, dietary patterns, and clinical counseling in the field of chronic kidney disease (CKD) and acute kidney injury (AKI). A common theme across the Research Topic is the assessment of nutritional status and body composition of patients with KD. This is an important topic for investigation since protein energy wasting (PEW) and sarcopenia are common in patients with KD, and are associated with adverse clinical outcomes (4, 5). In the current issue Macedo et al. showed that older adults on hemodialysis can concomitantly have malnutrition and sarcopenia, and that both conditions can be independently present in this patient group even if overweight is present. In addition, the authors also demonstrated that malnutrition and sarcopenia together have a worse quality of life and survival.

Many are the factors involved in the worsening of nutritional status of patients with CKD, including unsupervised dietary modifications, which in conjunction with the loss of appetite often observed in this patient group, may lead to spontaneous reduction in energy and nutrient intake (6). Moreover, the catabolic effects of kidney replacement therapy (KRT), metabolic and hormonal derangements, the presence of systemic inflammation and comorbid conditions, and also reduced physical activity contribute to a state of negative energy and protein balance both in patients with AKI and patients with CKD/ESKD (6, 7). A recent hypothesis is that patients undergoing dialysis are more “anabolic resistant,” characterized by blunted stimulation of muscle protein synthesis to common anabolic stimuli such as food intake and physical activity. Garibotto et al. thoroughly discuss this hypothesis in a very interesting and in-depth review, bringing awareness to a problem that is far from being solved.

Unique aspects of KD, such as fluid overload and consequent body weight fluctuations, hinder a reliable assessment of nutritional status, and consequently, the management of these conditions. This topic gathers much interest and most papers from this special collection discuss or investigate different tools to assess body composition in patients with KD. Sabatino et al. presented the

OPEN ACCESS

Edited and reviewed by:

Maurizio Muscaritoli,
Sapienza Università di Roma, Italy

*Correspondence:

Alice Sabatino
alice.sabatino86@gmail.com

Specialty section:

This article was submitted to
Clinical Nutrition,
a section of the journal
Frontiers in Nutrition

Received: 27 October 2021

Accepted: 29 October 2021

Published: 17 November 2021

Citation:

Sabatino A and Avesani CM (2021)
Editorial: Nutritional Aspects of Kidney
Disease. *Front. Nutr.* 8:803351.
doi: 10.3389/fnut.2021.803351

results of an observational study that investigated the role of ultrasound (US) in the monitoring of quadriceps muscle mass of patients with AKI. They observed an important reduction in quadriceps muscle mass in the first 5 days in the ICU stay, which is expected given the strong catabolic stimuli present in this clinical setting. Overall, this paper provides evidence that US is emerging as a sensitive alternative for the body composition assessment of patients with KD. Aligned with this idea, Reis et al. investigated the role of electrical bioimpedance (BIA) in patients on peritoneal dialysis, showing how single frequency BIA seems to be more accurate than multi-frequency BIA in this clinical setting. Finally, Bellafronte et al. showed that predictive equations applied to estimate skeletal muscle mass from BIA parameters can be useful for the identification of patients with low muscle mass. In addition, Sabatino et al. discuss in a narrative review the usefulness of different bedside tools currently studied and/or applied in the clinical practice for patients on hemodialysis. An emphasis is given to simplified creatinine index [SCI], bioimpedance spectroscopy (BIS) and muscle US, all suitable for clinical practice use. From these, US is the only method that measures muscle in different dimensions, while BIS and SCI are dependent on either theoretical assumptions or on the use of population specific regression equations, which may lead to errors when fluid alteration (over or dehydration) is present, a condition often observed in patients with KD.

A decline in physical function is also of much concern in this clinical setting, being related with increased mortality, falls, fractures, disability and low quality of life (8). Although malnutrition, sarcopenia and low physical function partially overlap, they do not develop in the same rate nor necessarily at the same time (4). In this regard, Mota Silva et al. demonstrated how low lean mass as assessed by phase angle from BIA, but not inflammation and overhydration, has a moderate but statistically significant correlation with worse physical function in patients on peritoneal dialysis.

Another important challenge in the current nutritional management of patients with end stage kidney disease (ESKD) on hemodialysis is the achievement of a good dietary quality pattern

(9). The fear of hyperkalemia often precludes clinicians from recommending patients to eat fruits and vegetables, which results in poor fiber intake. A low-fiber dietary intake seems to associate with the development of low grade systemic inflammation through the promotion of intestinal dysbiosis and should not be overlooked (10). Other unwanted consequences, especially in this clinical setting, may arise from a pro-inflammatory dietary habit. Qin et al. show how pro-inflammatory dietary patterns, as assessed by the dietary inflammatory index (DII), positively correlated with parathyroid hormone (PTH) and increased the risk of hyperparathyroidism in patients with early stages of CKD. Although it is a challenge to improve diet quality; tabus, pre-conceptions and non-evidence based dietary counseling regarding the role of fruits and vegetables in the development of hyperkalemia must be resolved, and dietitians should be called into action. That is the theme of a very interesting article published in the current Research Topic. Chan et al. evaluates the impact of individualized nutritional counseling in improving diet quality in patients with CKD, showing encouraging results.

Finally, complementing this special collection, Wang et al. describe the risk factors for renal impairment in adult patients with short bowel syndrome; and Graidis et al. discuss the role of vitamin D in the development and prognosis of AKI as well as a therapeutic adjuvant.

To summarize, we understand that this volume brings a conjunction of papers that makes up a translational point of view for the medical nutritional therapy of kidney diseases—all aligned with the unanswered questions pointed as topics requiring investigation by the KDOQI 2020 Clinical Practice Guideline for Nutrition in Chronic Kidney Disease (1). We are certain that this reading will add new information for clinicians and researchers in the field of nutrition and kidney diseases.

AUTHOR CONTRIBUTIONS

All authors listed have made a substantial, direct and intellectual contribution to the work, and approved it for publication.

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Risk Factors for Renal Impairment in Adult Patients With Short Bowel Syndrome

Peng Wang^{1†}, Jianbo Yang^{2†}, Yupeng Zhang¹, Li Zhang¹, Xuejin Gao¹ and Xinying Wang^{1*}

¹ Department of General Surgery, Jinling Hospital, Medical School of Nanjing University, Nanjing, China, ² Department of General Surgery, The First Affiliated Hospital of USTC, Division of Life Sciences and Medicine, University of Science and Technology of China, Hefei, China

OPEN ACCESS

Edited by:

Carla Maria Avesani,
Karolinska Institutet, Sweden

Reviewed by:

Christiane Ramos,
Federal University of Sao Paulo, Brazil
Ailema González-Ortiz,
National Autonomous University of
Mexico, Mexico
Maria Ines Barreto Silva,
Rio de Janeiro State Federal
University, Brazil

*Correspondence:

Xinying Wang
wangxinying@nju.edu.cn

[†]These authors have contributed
equally to this work

Specialty section:

This article was submitted to
Clinical Nutrition,
a section of the journal
Frontiers in Nutrition

Received: 18 October 2020

Accepted: 14 December 2020

Published: 18 January 2021

Citation:

Wang P, Yang J, Zhang Y, Zhang L,
Gao X and Wang X (2021) Risk
Factors for Renal Impairment in Adult
Patients With Short Bowel Syndrome.
Front. Nutr. 7:618758.
doi: 10.3389/fnut.2020.618758

Renal impairment is a common complication in patients with intestinal failure that is mostly caused by short bowel syndrome (SBS) and is associated with adverse outcomes that severely affect the quality of life or even survival. The prevalence and risk factors for renal impairment in patients with SBS remain unclarified. Therefore, we aimed to determine the prevalence of renal impairment and identify potential risk factors for renal impairment in adult patients with SBS. We retrospectively identified 199 patients diagnosed with SBS admitted to the Department of General Surgery between January 1, 2012 and January 1, 2019, from a prospectively maintained database. Overall, 56 patients (28.1%) with decreased renal function (eGFR < 90 mL/min/1.73 m²). The median duration of SBS was 7 months (IQR, 3–31 months) and the mean eGFR was 103.1 ± 39.4 mL/min/1.73 m². Logistic regression modeling indicated that older age [odds ratio (OR), 1.074; 95% CI, 1.037–1.112, *P* < 0.001], kidney stones (OR, 4.887; 95% CI, 1.753–13.626; *P* = 0.002), decreased length of the small intestine (OR, 0.988; 95% CI, 0.979–0.998; *P* = 0.019), and prolonged duration of SBS (OR, 1.007; 95% CI, 1.001–1.013; *P* = 0.046) were significant risk factors for renal impairment. This is the largest study that has specifically explored the risk factors for renal impairment in a large cohort of adults with SBS. The present study showed that renal function should be closely monitored during treatment, and patients should be given prophylactic interventions if necessary. This retrospective study is a part of clinical study NCT03277014, registered in ClinicalTrials.gov PRS. And the PRS URL is <http://register.clinicaltrials.gov>.

Keywords: short bowel syndrome, selective digestive decontamination, risk factors, intestinal failure, renal impairment

INTRODUCTION

Short bowel syndrome (SBS) is a rare disease usually caused by the removal of the small intestine due to a variety of underlying diseases, and it accounts for 74.4% of chronic intestinal failure (1). In patients with intestinal failure who are treated with long-term parenteral nutrition (PN) support therapy, impaired kidney function has also been reported (2–5).

Renal injury is a serious life-threatening complication that beginning with renal impairment. Worldwide, increasing numbers of patients are affected by chronic kidney disease (6–8). Two factors that have been reported are important for end-stage renal disease (ESRD): aging and type 2 diabetes mellitus (8). The gold standard for evaluating renal function involves inulin or radiolabeled markers to measure the glomerular filtration rate (GFR) (9, 10). However, considering the cost and complexities of measuring GFR, estimated GFR (eGFR) is widely used in clinical work.

In patients with SBS, studies showed that chronic renal failure (CRF) was related to intestinal failure and long-term PN (3, 11). Infections, dehydration, hypovolemic state, and age are also reported were risk factors for renal impairment in patients with SBS; however, these results were not consistent among studies (2–4). In summary, the risk factors of renal impairment ($\text{eGFR} < 90 \text{ mL/min/1.73 m}^2$) in adult patients with SBS remain uncharacterized.

Small intestinal bacterial overgrowth (SIBO) occurs commonly in SBS. A study found that among patients with SBS 63% (27 of 43) of patients had SIBO (12). This means that the gut microbiota was disrupted in patients with SBS. A study showed that in patients with SBS, both fecal and colonic biopsy samples were found to have a high prevalence of *Lactobacillus*, with an associated depletion of *Clostridia* and *Bacteroidetes* (13). The crosstalk between the gut microbiota and the host has attracted considerable attention owing to its involvement in various diseases. Many papers report that chronic kidney disease (CKD) is associated with gut dysbiosis and altered host-microbiota crosstalk (14–16). One paper also suggested that the overgrown bacteria translocated from the gut to the blood, where they contributed to the development of CKD (16). Selective digestive decontamination (SDD) is a treatment for SIBO and the principal method that uses broad-spectrum and other antibiotics including those that cover anaerobic bacteria and antifungals (17–19). However, whether there is a crosstalk between gut flora and renal function in patients with SBS remains unclear.

This study aimed to determine the prevalence of renal impairment in clinical practice and to identify potential risk factors for renal impairment in adult patients with SBS.

MATERIALS AND METHODS

Participants

We retrospectively identified consecutive SBS patients from a prospectively maintained database at the Clinical Nutrition therapy center, Jinling Hospital, from January 1, 2012, to January 1, 2019. The prospective database was maintained to enable the guidance of patients' home nutritional support. We included patients who underwent small bowel resection with the length of remaining small bowel $< 200 \text{ cm}$ and were diagnosed with SBS. Patients were excluded if (1) $\text{eGFR} < 90 \text{ mL/min/m}^2$ before SBS diagnosed; (2) their serum creatinine and cystatin C level increased significantly in the last 3 months; (3) their remaining intestinal length and the anatomical structure was not clear; (4) the patients were at the acute stage of SBS (< 4 weeks after intestinal resection surgery); (5) age < 18 years. The present study

was approved by the Ethics Committee of Jingling Hospital. The ethics committee registration number was 2015ZFYJ-010.

Demographic data at admission were collected from the database, including age, sex, body mass index (BMI), nutritional risk screening 2002 (NRS-2002) score (≥ 3.0 means nutritional risk) (20), subjective global assessment (SGA) score (A means well-nourished; B means moderately malnourished; C means severely malnourished) (21), diabetes mellitus, hypertension, and serum concentrations of albumin, pre-albumin, creatinine, cystatin C, hemoglobin, and C-reactive protein. Unintentional weight loss in recent months (3–6 months), and underlying diseases were recorded. The intestinal anatomy (including small bowel length, anatomy type, presence of an intact ileocecal valve, and colon-in-continuity) were also recorded. Parenteral nutrition (PN) dependence, whether patients received SDD treatment, presence of nosocomial catheter-related bloodstream infections (CRBSIs) were evaluated in hospitalization for intestinal rehabilitation after SBS diagnosis in our center.

Definitions

The diagnosis of SBS was made in our clinical nutrition center by two gastroenterologists, three surgeons, and one dietitian. The anatomy of the remaining small bowel was used to classify SBS into type 1 (end-jejunostomy with no colon-in-continuity), type 2 (jejunocolic anastomosis with partial colon-in-continuity), and type 3 (jejunoleal anastomosis with an intact colon). The *duration of SBS* was defined as the time interval between the diagnosis of SBS and the previous hospital visit. *Malnutrition* was retrospectively diagnosed by two clinical nutritional physicians, according to the ESPEN diagnostic criteria of 2015 (22). *Kidney stones* were diagnosed based on abdominal imaging (CT, MRI, or ultrasound).

Creatinine and cystatin C were used to calculate the eGFR, using the CKD-EPI equation (23, 24). According to the KIDGO Guideline for the evaluation and management of chronic kidney disease (CKD) which published in 2012, the prognosis of CKD graded by eGFR categories was divided into five grades (G1, normal or high; G2, mildly decreased; G3a, mildly to moderate decreased; G3b, moderate to severe decreased; G4, severely decreased; G5, kidney failure). Hence, we defined the *Renal impairment* as an eGFR of $< 90 \text{ mL/min/1.73 m}^2$ (11, 25). The diagnosis of SIBO in patients with SBS mainly depends on two methods; the culture of duodenal fluid was widely used and was considered the gold standard (26), and the other is the hydrogen breath test (27). Nevertheless, both tests raise several issues regarding repeatability, accuracy, and optimal cut-off values (28–30). For these reasons, there is no consensus regarding the optimal test for SIBO. The management of patients with SIBO should consider gastroenterological, surgical, microbiologic, pharmacologic, nutritional, and metabolic factors (31). So, we developed an SDD treatment process combining the etiology, symptomology, and the results of the fecal smear test to determine whether SIBO was present in patients with SBS. We empirically diagnosed patients with SIBO according to (1) the pathogenesis [proton pump inhibitors used for more than 3 weeks; intestinal motility impaired; intestinal anatomy changed (32, 33)], (2) symptoms reported in the literature [diarrhea/stool

smelly, abdominal pain, bloating, weight loss, unexpected plateau during the weaning of PN (12, 34, 35)], and (3) the results of the smear test of fecal bacteria (proportion of bacteria was abnormal, and the appearance of other abnormal bacteria). Patients diagnosed with SIBO were then treated with SDD (oral amikacin/vancomycin/gentamicin with metronidazole and fluconazole three times per day for 3 days).

Statistical Analysis

Numerical data that were normally distributed were expressed as mean \pm standard deviation (SD); others were expressed as median (first-to-third interquartile range). Categorical variables were expressed as numbers and percentages. Binary data were presented as 0 (no) or 1 (yes). Data between groups were compared using Student's *t*-test for normally distributed values and the Mann-Whitney *U*-test was used to compare median values, while Pearson's chi-square test was used for categorical variables. Potential risk factors for renal impairment were evaluated by univariate analysis, and risk factors with $P < 0.1$ were included in multivariate binary logistic stepwise regression analysis. In the multivariate analysis statistical significance was set at two-sided $P < 0.05$. Statistical analyses were performed using SPSS software (version 20.0 Win; SPSS Inc., Chicago, IL, USA).

RESULTS

Patient Characteristics

A total of 239 patients were diagnosed with SBS in our center during 2012–2019. However, according to our exclusion criteria, data from 40 patients were not used in the present study because they were either in the acute stage of the SBS (20 patients) or $\text{eGFR} < 90 \text{ mL/min/1.73 m}^2$ before SBS diagnosed (6 patients) or age < 18 years old (14 patients). Eventually, we identified 199 patients for the study (Figure 1). The mean age was 48.8 ± 17.3 years, and 138 (69.3%) were male. The mean BMI was $18.1 \pm 4.6 \text{ kg/m}^2$. Regarding the SGA score, there were 12 patients in level A (well nourished), 113 patients in level B (moderately malnourished), and 74 patients in level C (severely malnourished), as for the NRS2002 score there are 96 (48.5%) patients with nutritional risk ($\text{NRS2002} \geq 3$). The median duration of SBS was 7 months (interquartile range 3–31). There were 118 (59.2%) patients requiring PN at admission; 141 (71.4%) patients were malnourished according to ESPEN diagnostic criteria of 2015. There were 34 (17.1%) patients with kidney stones; 56 (28.1%) patients with bowel symptoms who received SDD treatment, while 34 (17.1%) contracted nosocomial CRBSIs. The characteristics of the study population at admission are outlined in Table 1.

The mean small bowel length of the patients was $85.6 \pm 45.5 \text{ cm}$; the median was 80.0 cm ; 53 (26.6%) patients had type 1, 69 (34.7%) had type 2, and 77 (38.7%) had type 3 SBS. A total of 146 (73.4%) patients had continuity of the colon. The ileocecal valve had been removed in 77 (38.7%) patients (Table 2). The most common underlying diseases were mesenteric ischemia (41.7%), volvulus (23.6%), surgical

complications (19.6%), radiation enteritis (6.0%), and others (9.0%) (Table 3).

The mean eGFR of patients with SBS was $103.1 \pm 39.4 \text{ mL/min/1.73 m}^2$. There were 56 (28.1%) patients decreased the eGFR . According KDIGO guideline, in the 56 patients with renal impairment, 25 (44.6%) of them were classified as G2 ($\text{eGFR} 60\text{--}89 \text{ mL/min/1.73 m}^2$), 10 (17.9%) of them were classified as G3a ($\text{eGFR} 45\text{--}59 \text{ mL/min/1.73 m}^2$), 6 (10.7%) of them were classified as G3b ($\text{eGFR} 30\text{--}44 \text{ mL/min/1.73 m}^2$), 6 (10.7%) of them were classified as G4 ($\text{eGFR} 15\text{--}29 \text{ mL/min/1.73 m}^2$) and 9 (16.1%) of them were classified as G5 ($\text{eGFR} < 15 \text{ mL/min/1.73 m}^2$) and four of them required dialysis therapy. The incidence of renal impairment was 23.2% in type 1 SBS, 41.1% in type 2 SBS, and 35.7% in type 3 SBS ($P = 0.962$).

Patients were divided into a renal impairment group (decreased eGFR , $\text{eGFR} < 90 \text{ mL/min/1.73 m}^2$) and a normal renal function group ($\text{eGFR} \geq 90 \text{ mL/min/1.73 m}^2$). Compared with patients in the normal renal function group, patients with renal impairment tend to be older (58.6 ± 12.5 years vs. 44.9 ± 17.4 years, $P < 0.001$), having longer duration of SBS (12.0 vs. 5.0 months, $P = 0.051$), and lower hemoglobin concentration (97.8 ± 27.6 vs. $107.2 \pm 20.1 \text{ g/L}$, $P = 0.031$) and albumin concentration (34.7 ± 7.3 vs. $36.9 \pm 6.6 \text{ g/L}$, $P = 0.046$). Compared with patients in the normal renal function group, in the renal impairment group there were more patients with kidney stones (28.6 vs. 12.6%, $P = 0.008$), receiving SDD treatment (53.5 vs. 18.1%, $P < 0.001$), with nosocomial CRBSIs (25.0 vs. 13.9%, $P = 0.067$) and hypertensive (23.2 vs. 6.9%, $P = 0.001$). Regarding small bowel length, the renal impairment group was shorter than the other group (75.8 ± 41.2 vs. $89.4 \pm 46.6 \text{ cm}$, $P = 0.068$). The two groups were similar regarding sex, BMI, NRS-2002 score, diabetes, PN dependence, and the biochemical indicators of nutritional status (all $P > 0.1$).

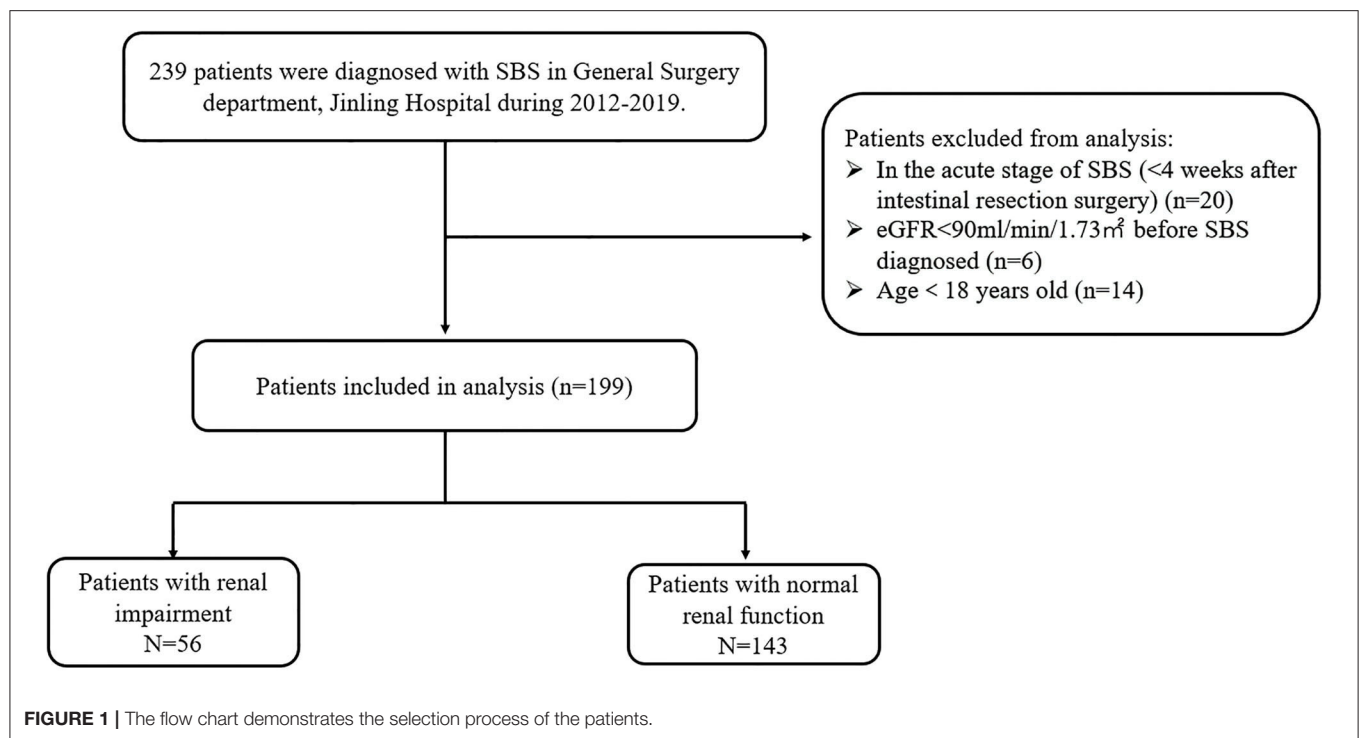
Independent Risk Factors for Renal Impairment in Patients With Short Bowel Syndrome

We incorporated potential variables with $P < 0.1$ in the univariate analysis into a multivariate logistic regression analysis. The results indicated that the age ($P < 0.001$), SBS duration ($P = 0.046$), small bowel length ($P = 0.019$), and existing kidney stones ($P = 0.002$) were independent risk factors for reduced eGFR . SBS duration had an OR of 1.007 per month resulting in a 10-year cumulative risk of 2.3 folds (1.007^{120}) (Table 4).

DISCUSSION

To the best of our knowledge, this is the largest study that has specifically explored the risk factors for renal impairment in a large cohort of adults with SBS ($n = 199$). The incidence of renal impairment in adults with SBS was 28.1%, and the median duration from SBS diagnosis to the development of renal impairment was 12 months. Renal impairment in adults with SBS was influenced by age, shorter length of the small bowel, kidney stones, and prolonged duration of SBS.

Renal impairment has been widely reported in patients with SBS. To date, Buchman et al. (3) reported decreased GFR $> 20\%$



in 52.5% of patients, and 3.5% decline per year of creatinine clearance and a reduced kidney function in ~50% of patients was observed; Lauverjat et al. (4) reported that decreased GFR was found in 9 of 16 patients with SBS also receiving home parenteral nutrition (HPN). Some risk factors have been reported, including infections, dehydration, hypovolemic state, duration of HPN, and patient age; however, these results were not consistent among studies (2–4). Both studies had results similar to ours. Ylinen et al. (11) found that small bowel length was a risk factor for renal impairment in children. Agostini et al. (36) found that age, urologic disease, sepsis, and SBS significantly influenced the development of CKD in patients with HPN, and that eGFR comparable with the median decreased -2.4 to -7.3% per year. Because our study absorbed all patients either receiving PN support therapy, or weaned off PN, and receiving enteral nutrition support therapy, the incidence of renal impairment only was 28.1%.

Patients with SBS need long-term PN. However, with the progress of treatment the amount of fluid and basal metabolism required by patients changed significantly. Most patients do not adjust their PN support plan timeously, resulting in them becoming chronically dehydrated for long periods. A retrospective study of 33 patients with long-term PN found that the GFR decreased by $3.5\% \pm 6.3\%$ per year (3). Similarly, in our study, we found that the cumulative risk of eGFR was 2.3 fold (1.007^{120}) after 10 years. A decrease in renal function with aging is a natural phenomenon.

In patients with SBS, renal impairment is mainly due to systemic fluid imbalance caused by excessive loss of ostomy fluid, CRBSIs, chronic dehydration, and electrolyte imbalance

(37). The small bowel length is an important factor that can affect nutrients and electrolyte absorption, as well as renal function (11). Similar to the results of previous reports, the length of the small intestine in our study was an important factor for determining renal impairment. In the present study, nosocomial CRBSIs are not considered independent risk factors for renal impairment because we did not analyze the occurrence of CRBSIs during HPN, and CRBSIs are well-controlled during hospitalization. However, in the renal impairment group, a higher percentage of patients suffered CRBSIs (25.0 vs. 13.9%).

The ileocecal valve plays an important role in the alimentary canal since it can influence nutrients absorbed and keep electrolytes balanced. Although in the present study the renal dysfunction has no difference in patients with or without the ileocecal valve, we found that the eGFR of SBS in various types were significantly different (type 1 SBS was 107 ± 33.5 mL/min/1.73 m², type 2 SBS was 94.1 ± 43.7 mL/min/1.73 m² and type 3 SBS was 108.4 ± 38.1 mL/min/1.73 m², type 2 vs. type 3, $P = 0.037$). The eGFR was significantly lower in patients with type 2 SBS than in type 3 SBS. Although type 1 SBS includes patients with stomas who are prone to water and electrolyte loss, no significant reduction in eGFR was found. This phenomenon may be related to the acute phase of the disease, since most of the acute phase occurs during the hospitalization period when doctors pay more attention to water and electrolyte balance.

Kidney stones have been reported in patients with SBS, resulting in adverse clinical sequelae (38, 39). The prevalence of kidney stones is increased in patients who have undergone intestinal surgery, especially in patients with jejunostomy or ileostomy, with an occurrence of 5–15% (40–42). Patients

TABLE 1 | Characteristics of adult patients with SBS with and without renal impairment.

Variables	All patients	Patients with renal impairment*	Patients with normal renal function	P-value
No. of patients	199	56	143	
Age, year	48.8 ± 17.3	58.6 ± 12.5	44.9 ± 17.4	<0.001
Male, n (%)	138 (69.3)	43 (76.8)	95 (66.4)	0.154
Diabetes mellitus, n (%)	7 (3.5)	1 (1.8)	6 (4.2)	0.682
Hypertension, n (%)	23 (11.6)	13 (23.2)	10 (6.9)	0.001
Kidney stones, n (%)	34 (17.1)	16 (28.6)	18 (12.6)	0.008
BMI, kg/m ²	18.1 ± 4.6	18.0 ± 4.7	18.1 ± 4.6	0.889
Malnutrition, n (%)	141 (71.4)	42 (76.8)	99 (69.9)	0.291
NRS2002 ≥ 3, n (%)	96 (48.5)	26 (46.4)	70 (49.0)	0.832
<3, n (%)	103 (51.5)	30 (53.6)	73 (51.0)	
SGA class (A; B; C)	12/113/74	4/27/25	8/86/49	0.311
Hemoglobin (g/L)	104.5 ± 23.5	97.8 ± 29.6	107.2 ± 20.1	0.031
Albumin (g/L)	36.3 ± 6.8	34.7 ± 7.3	36.9 ± 6.6	0.046
Serum creatinine (μmol/L)	56.1 (43.0–76.0)	116.0 (89.7–185.5)	50.0 (38.5–61.0)	<0.001
Cystatin c (mg/L)	0.8 (0.6–1.0)	1.3 (1.1–1.9)	0.7 (0.6–0.8)	0.010
eGFR (mL/min/1.73 m ²)	103.1 ± 39.4	51.8 ± 27.8	123.2 ± 20.4	<0.001
≥ 90, n (%)	143 (71.9)	–	143 (100)	
60–89, n (%)	25 (12.5)	25 (44.6)	–	
45–59, n (%)	10 (5.0)	10 (17.9)	–	
30–44, n (%)	6 (3.0)	6 (10.7)	–	
15–29, n (%)	6 (3.0)	6 (10.7)	–	
<15, n (%)	9 (4.5)	9 (16.1)	–	
ALT (U/L)	33.0 (18.8–58.3)	31.5 (15.8–58)	33.0 (19.0–58.8)	0.547
AST (U/L)	28.5 (18.0–43.0)	25.0 (17.3–59.5)	29.0 (18.0–41.5)	0.124
CRP (mg/L)	3.9 (0.9–19.6)	8.8 (2.0–30.0)	3.0 (0.8–10.9)	0.191
PN requirement, n (%)				0.947
Yes	118 (59.2)	33 (58.9)	85 (59.4)	
Weaned	81 (40.7)	23 (41.1)	58 (40.6)	
Duration of SBS, (months)	7.0 (3.0–31.0)	12.0 (5.3–46.0)	5.0 (2.0–16.0)	0.051
SDD, n (%)	56 (28.1)	30 (53.5)	26 (18.1)	<0.001
Nosocomial CRBSIs, n (%)	34 (17.1)	14 (25.0)	20 (13.9)	0.067

Values were presented as n (%), or mean ± SD, or median (first-to-third interquartile range).

BMI, body mass index; NRS2002, Nutrition Risk Screening 2002; Scr, Serum creatinine; ALT, Alanine aminotransferase; AST, Aspartate aminotransferase; CRP, C-reactive protein; PN, Parenteral Nutrition; SBS, Short Bowel Syndrome; SDD, Selective decontamination of the digestive tract; CRBSIs, catheter-related blood stream infections. Duration of SBS: the time interval between the diagnosis of SBS and the previous hospital visit.

*Renal impairment was defined as eGFR < 90 mL/min/1.73 m² calculated by the CKD-EPI equation. P-value < 0.05 were bolded and italicized.

with kidney stones have significantly increased incidences of adverse renal outcomes, including chronic renal failure (43). Our previous studies also found that patients with kidney stones have lower eGFR than patients without kidney stones (44). In the present study, we found that patients with kidney stones may be prone to renal impairment. Nevertheless, the mechanisms linking nephrolithiasis to impairment of renal function remain unclear. The intestinal surgery alters the anatomy of intestinal results in malabsorption of bile acids and fatty acids (45). Normally, the oxalate in dietary binds to calcium in luminal forming calcium oxalate which cannot be absorbed. At the same time, unabsorbed fatty acids bind to luminal calcium and form insoluble calcium soaps which inhibit the binding of luminal oxalate to calcium and increase its absorption. Besides these, the bile acids enrichment in colon and increases the absorption of oxalate, results in

hyperoxalate and increase the formation of oxalate stones (38). There is also a study found that the plasma oxalate increase at lower GFR levels even among those without enteric or primary hyperoxaluria and established the relationships between plasma oxalate, GFR and urine oxalate among patients with routine urinary stone disease. In present study, we found that the patients with kidney stones may be prone to renal dysfunction, may be the decreased of GFR resulted the kidney stones (46).

SDD is an infection preventive measure for ICU patients that was proposed more than 30 years ago (47). It has been reported to have a favorable effect on mortality in adult patients in general intensive care units (48). In patients with SBS, a review showed that cyclical use (1 week per month) of broad-spectrum antibiotics was the mainstay of therapy for small-intestinal bacterial overgrowth (49).

TABLE 2 | The intestinal anatomy of adult patients with SBS with and without renal impairment.

Variables	All patients	Patients with renal impairment*	Patients with normal renal function	P-value
No. of patients, <i>n</i>	199	56	143	
Small bowel length (cm)	85.6 ± 45.5	75.8 ± 41.2	89.4 ± 46.6	0.068
Median (cm)	80.0 (57.5–117.5)	77.5 (50.0–100.0)	90 (60.0–120.0)	
Anatomy type, <i>n</i> (%)				0.962
I	53 (26.6)	13 (23.2)	40 (28.0)	
II	69 (34.7)	23 (41.1)	46 (32.2)	
III	77 (38.7)	20 (35.7)	57 (39.9)	
Ileocecal valve intact, <i>n</i> (%)	77 (38.7)	20 (35.7)	57 (39.9)	0.591
Colon in continuity, <i>n</i> (%)	146 (73.4)	43 (76.8)	103 (72.0)	0.495

Values were presented as *n* (%), or mean ± SD, or median (first-to-third interquartile range).

Type I: end-jejunostomy with no colon-in-continuity.

Type II: jejunocolic anastomosis with partial colon-in-continuity.

Type III: jejunoleal anastomosis with an intact colon.

*Renal impairment was defined as eGFR < 90 mL/min/1.73 m² calculated by the CKD-EPI equation.

TABLE 3 | The underlying diseases in adult patients with SBS with and without renal impairment.

Variables	All patients	Patients with renal impairment*	Patients with normal renal function	P-value
No. of patients, <i>n</i>	199	56	143	
Mesenteric ischemia	83 (41.7)	30 (53.5)	53 (37.1)	0.266
Radiation enteritis	12 (6.0)	3 (5.4)	9 (6.3)	1.000
Surgical complications	39 (19.6)	9 (16.1)	30 (20.1)	1.000
Volvulus	47 (23.6)	8 (14.3)	39 (27.3)	0.269
Others	18 (9.0)	6 (10.7)	12 (8.4)	1.000

Values were presented as *n* (%).

*Renal impairment was defined as eGFR < 90 mL/min/1.73 m² calculated by the CKD-EPI equation.

TABLE 4 | Multivariate analysis of the factors associated with renal impairment in adults with SBS.

Variables	β	OR	95%CI	P-value
Age	0.071	1.074	1.037–1.112	<0.001
Kidney stones	1.587	4.887	1.753–13.626	0.002
Length of SI	−0.012	0.988	0.979–0.998	0.019
Duration of SBS	0.007	1.007	1.001–1.013	0.046
Hypertension	0.140	1.150	0.338–3.910	0.823
Hemoglobin	−0.016	0.984	0.961–1.008	0.183
Albumin	0.041	1.042	0.955–1.136	0.441
SDD	0.820	2.272	0.897–5.755	0.356
CRBSIs	−0.088	0.916	0.313–2.679	0.837

SBS, Short bowel syndrome; SI, Small intestine; SDD, Selective decontamination of the digestive tract.

Duration of SBS: the time interval between the diagnosis of SBS and the previous hospital visit. P-value < 0.05 were bolded and italicized.

We diagnosed patients with SIBO using our SDD management scheme. Many papers reported that disordered gut flora could influence the renal function of the host. Vaziri et al. (50) found that there were 190 bacterial OTUs in the stool, with marked differences in abundance between patients with ESRD and normal controls. In addition, Wang et al. performed a clinical trial that showed that bacterial DNA was detected in the blood of six (20%) patients with ESRD. The bacterial genera found in the blood that were overgrown in the intestines were

mainly *Klebsiella* spp., *Proteus* spp., *Escherichia* spp., *Enterobacter* spp., and *Pseudomonas* spp. (16). This paper suggests that the overgrown bacteria translocated from the gut to the blood, where they contributed to the development of CKD. Hence, SIBO may be an independent risk factor for renal impairment in patients with SBS. However in the present study, 56 (28.1%) patients received SDD treatment (may with SIBO), less than the previously reported value (63%) (12). The reason for our low result is that we can only rely on symptoms and experience to

diagnose SIBO. And SIBO is not an independent risk factor for renal impairment in patients with SBS. The relationship between intestinal flora disorder and SBS-related kidney damage needs further study.

The present study had several limitations. First, it was a retrospective single-institution study performed at a tertiary-care referral center. Second, the dietary habits, home PN formulation, and total time that patients required PN support therapy were not available. Third, the incidence of CRBSIs during HPN could not be recorded. Fourth, because most patients with SBS in our center were adults, we only evaluated the risk factors for renal impairment in adults. Fifth, it was a retrospective study performed at a tertiary-care referral center, the data of urine test (hematuria and proteinuria) was lacking. Therefore, the incidence of renal impairment may have been underestimated. Large, multicenter, prospective studies focusing on the reason for renal impairment in both pediatric and adult patients with SBS should be conducted in the future.

The strengths of the present study. Firstly, it is the largest study that has specifically explored the risk factors for renal dysfunction in a large cohort of adults with SBS ($n = 199$). In addition, we found that the duration of SBS, kidney stones, old age, and the length of the small bowel are the independent risk factors for renal dysfunction in adult patients with SBS. And, SBS duration had an OR of 1.007 per month resulting in a 10-year cumulative risk of 2.3 fold (1.007^{120}). Given that patients with SBS have increased risk for renal impairment and associated adverse outcomes, close monitoring of renal function and prophylactic interventions should be conducted routinely in clinical practice.

CONCLUSION

Renal function impairment is common in adults with SBS and can result in adverse clinical consequences. In our study, 56 (28.1%) in 199 patients developed renal function impairment with $\text{eGFR} < 90 \text{ mL/min/1.73 m}^2$, 34 (17.1%) developed nephrolithiasis. The risk of renal impairment is increased in patients with older age, kidney stones, shorter length of remaining small intestine, and prolonged duration of SBS. In the chronic stage of SBS, renal impairment is primarily related to water and electrolyte loss, kidney stones, old age, and the length of the small bowel. Given that patients with SBS have increased

risk for renal impairment and associated adverse outcomes, close monitoring of renal function to avoid dehydration, and prophylactic interventions should be conducted.

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 the Ethics Committee of Jingling Hospital. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

PW and XW contributed to conception, design of the research, and critically revised the manuscript. PW and JY contributed to acquisition, analysis, and interpretation of the data. XG, LZ, and YZ drafted the manuscript. All authors agreed to be fully accountable for ensuring the integrity and accuracy of the work, read, and approved the final manuscript.

FUNDING

This study was supported by the National Natural Science Foundation of China (81470797 and 81770531), the Science Foundation of Outstanding Youth in Jiangsu Province (BK20170009), Military Medical Innovation Project (18CXZ031), the National Science and Technology Research Funding for Public Welfare Medical Projects (201502022), and The 13th Five-Year Plan Foundation of Jiangsu Province for Medical Key Talents (ZDRCA2016091).

ACKNOWLEDGMENTS

We thank Editage China for editing the English text of a draft of this manuscript.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Vitamin D and Acute Kidney Injury: A Two-Way Causality Relation and a Predictive, Prognostic, and Therapeutic Role of Vitamin D

Spyridon Graidis^{1†}, Theodosios S. Papavramidis^{2†} and Maria Papaioannou^{1*}

¹ Laboratory of Biological Chemistry, Faculty of Health Sciences, School of Medicine, Aristotle University of Thessaloniki, Thessaloniki, Greece, ² 1st Propedeutic Department of Surgery, Faculty of Health Sciences, School of Medicine, AHEPA University Hospital, Aristotle University of Thessaloniki, Thessaloniki, Greece

OPEN ACCESS

Edited by:

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*Correspondence:

Maria Papaioannou
mpapaioannou@auth.gr
orcid.org/0000-0003-3149-6523

†ORCID:

Spyridon Graidis
orcid.org/0000-0003-0356-850X
Theodosios S. Papavramidis
orcid.org/0000-0002-7097-9889

Specialty section:

This article was submitted to
Clinical Nutrition,
a section of the journal
Frontiers in Nutrition

Received: 18 November 2020

Accepted: 31 December 2020

Published: 04 March 2021

Citation:

Graidis S, Papavramidis TS and
Papaioannou M (2021) Vitamin D and
Acute Kidney Injury: A Two-Way
Causality Relation and a Predictive,
Prognostic, and Therapeutic Role of
Vitamin D. *Front. Nutr.* 7:630951.
doi: 10.3389/fnut.2020.630951

Background: Acute kidney injury (AKI) constitutes a multi-factorially caused condition, which significantly affects kidney function and can lead to elevated risk of morbidity and mortality. Given the rising scientific evidence regarding vitamin D's (VitD's) multisystemic role, the connection between AKI and VitD is currently being studied, and the complex relation between them has started to be unraveled.

Methods: A systematic review had been conducted to identify the pathogenetic relation of VitD and AKI and the potential role of VitD as a biomarker and therapeutic–renoprotective factor.

Results: From 792 articles, 74 articles were identified that fulfilled the inclusion criteria. Based on these articles, it has been found that not only can VitD disorders (VitD deficiency or toxicity) cause AKI but, also, AKI can lead to great disruption in the metabolism of VitD. Moreover, it has been found that VitD serves as a novel biomarker for prediction of the risk of developing AKI and for the prognosis of AKI's severity. Finally, animal models showed that VitD can both ameliorate AKI and prevent its onset, suggesting its renoprotective effect.

Conclusion: There is a complex two-way pathogenetic relation between VitD disorders and AKI, while, concomitantly, VitD serves as a potential novel predictive–prognostic biomarker and a treatment agent in AKI therapy.

Keywords: vitamin D, acute kidney injury, pathogenesis, biomarker, prediction, prognosis, therapeutic agent

INTRODUCTION

Acute kidney injury (AKI), acute kidney failure (AKF), or acute renal failure (ARF) is a multi-factorially caused condition, characterized by sudden and rapid loss of kidney function. Acute kidney injury results in further systemic disorders and is related to elevated morbidity and mortality (1). There are many different definitions for AKI, characterized by different biochemical, physiological, and clinical cutoff points. The most preponderant ones are based on serum creatinine (sCr), glomerular filtration rate (GFR), and urine output (UO). According to the Kidney Disease

Improving Global Guidelines (KDIGO) Clinical Practice Guidelines for Acute Kidney Injury definition, there are three stages of AKI. Stage 1 includes one of the following: (a) 1.5–1.9 times baseline sCr, (b) ≥ 0.3 mg/dl increase of baseline sCr, or (c) <0.5 ml/kg/h UO for 6–12 h; Stage 2 includes one of the following: (a) 2–2.9 times baseline sCr or (b) <0.5 ml/kg/h for ≥ 12 h; and Stage 3 includes one of the following: (a) 3 times baseline sCr, (b) ≥ 4.0 mg/dl increase, (c) initiation of renal replacement therapy (RRT), (d) in patients <18 years old, a decrease of eGFR < 35 ml/min/1.73 m², (e) <0.3 ml/kg/h for >24 h, or (f) anuria ≥ 12 h (2).

The pathophysiology of AKI can be classified into: (1) prerenal azotemia (disruption of kidneys' blood inflow, without cellular or structural damage), (2) intrinsic or intrarenal AKI (direct damage to kidney's structure) due to acute tubular necrosis (ATN), vascular damage, and glomerular damage, (3) interstitial AKI (damage of the interstitial tissue), or (4) postrenal AKI (obstruction of urinary tract) (3). Moreover, AKI can be induced by septic shock, cardiogenic shock, pharmaceutical administration, hepatorenal syndrome, obstructive uropathy, and hypovolemia (4).

Due the necessity to specify the causes and the risk factors inducing AKI, many efforts have been made in order to identify sensitive, specific, and cost-efficient biomarkers, which may contribute to early detection and disease prognosis. In this regard, research has been also focused on the relation between AKI and vitamin D (VitD).

Apropos of VitD, it is an important part of the physiological function of various systems. VitD is, mainly, produced in the skin *via* UV radiation-induced transformation of 7-dehydroxycholesterol (7DHC) to cholecalciferol, which then undergoes consecutive hydroxylations in the liver and kidneys, accordingly, which leads to the synthesis of 1,25(OH)₂D (5, 6). However, 1,25(OH)₂D can be synthesized into many extrarenal tissues, which possess the CYP27B1 enzyme, which is essential for the process (7). VitD's effects are classified into endocrine, paracrine, and autocrine, while the mechanism of action is both genomic [based on the action of VitD receptor (VDR) on VitD response elements (VDREs)] and non-genomic (8, 9). Briefly, VitD affects, significantly, calcium-phosphate homeostasis, leads to bone reabsorption, enhances muscular contraction and proliferation, intensifies myocardial contracture and lowers blood pressure *via* affecting the renin-angiotensin-aldosterone system (RAAS), enhances innate immunity and changes the cytokine and chemokine profile of acquired immunity from proinflammatory to anti-inflammatory, and protects against autoimmune response and cancer (10).

The aim of the present systematic review is to unravel the relationship between AKI and VitD in both patients and experimental models. We assumed that this is a two-way relationship, insinuating that on the one hand, VitD levels may play a role in AKI, and inversely, AKI induces changes in VitD levels. In this perspective, we examined whether hypo- or hypervitaminosis causes AKI and, inversely, whether AKI causes hypo- or hypervitaminosis. It was actually pointless to search for active comparators. Moreover, we examined possible

implications of VitD as a predictive and prognostic marker for AKI, as well as its role in AKI treatment.

METHODS

In May 2020, a bibliographic search was conducted in PubMed, Scopus, and Embase regarding the relation between VitD and AKI. The terms employed were “vitamin D AND AKI,” “vitamin D AND ARF,” and “vitamin D AND acute kidney injury.” Additionally, a manual search of the reference lists of other studies had been conducted, to detect subsequent material that could be included in this systemic review.

Then, the articles were screened, based on:

1. Removing duplicates
2. Relativity to the subject
3. Date of publication
4. Validity of methods and conclusions

Articles that did not meet the above criteria were excluded.

RESULTS

There were 756 articles identified in the three databases. Moreover, 36 articles have been included from screening the reference lists of the articles. After the exclusion of 336 duplicates and 340 irrelevant studies, the remaining 117 articles were analyzed based on the eligibility for full access and the inclusion criteria. Finally, 74 articles matched our search criteria (**Figure 1**).

These articles reveal a two-way relation between VitD and AKI. Acute kidney injury can, *via* various mechanisms, lead to either D hypo- or hypervitaminosis. Moreover, both VitD deficiency and VitD toxicity can lead to the development of AKI.

Given the fact that our knowledge concerning the relationship between VitD and AKI comes from small series, cases, and experimental models, extrapolation of our results to the general population may be risky and biased. Moreover, introducing VitD in the treatment of AKI is also questionable on a clinical basis since there are no prospective randomized trials available and our knowledge is a result of animal experiments. Finally, more clinical data are available on the use of VitD levels as biomarkers concerning both the prediction and the prognosis of AKI. However, despite the fact that there are nine clinical studies on this matter, those studies are observational (either prospective or retrospective), and no data coming from randomized controlled trials are available.

DISCUSSION

The Development of AKI Is Associated With VitD Disorders

Acute kidney injury is the significant depletion of renal function, affecting the normal renal enzymatic activity, and consequently disrupts VitD. Usually, the decreased kidney function causes VitD deficiency, but in some cases, VitD toxicity is also observed (**Table 1**).

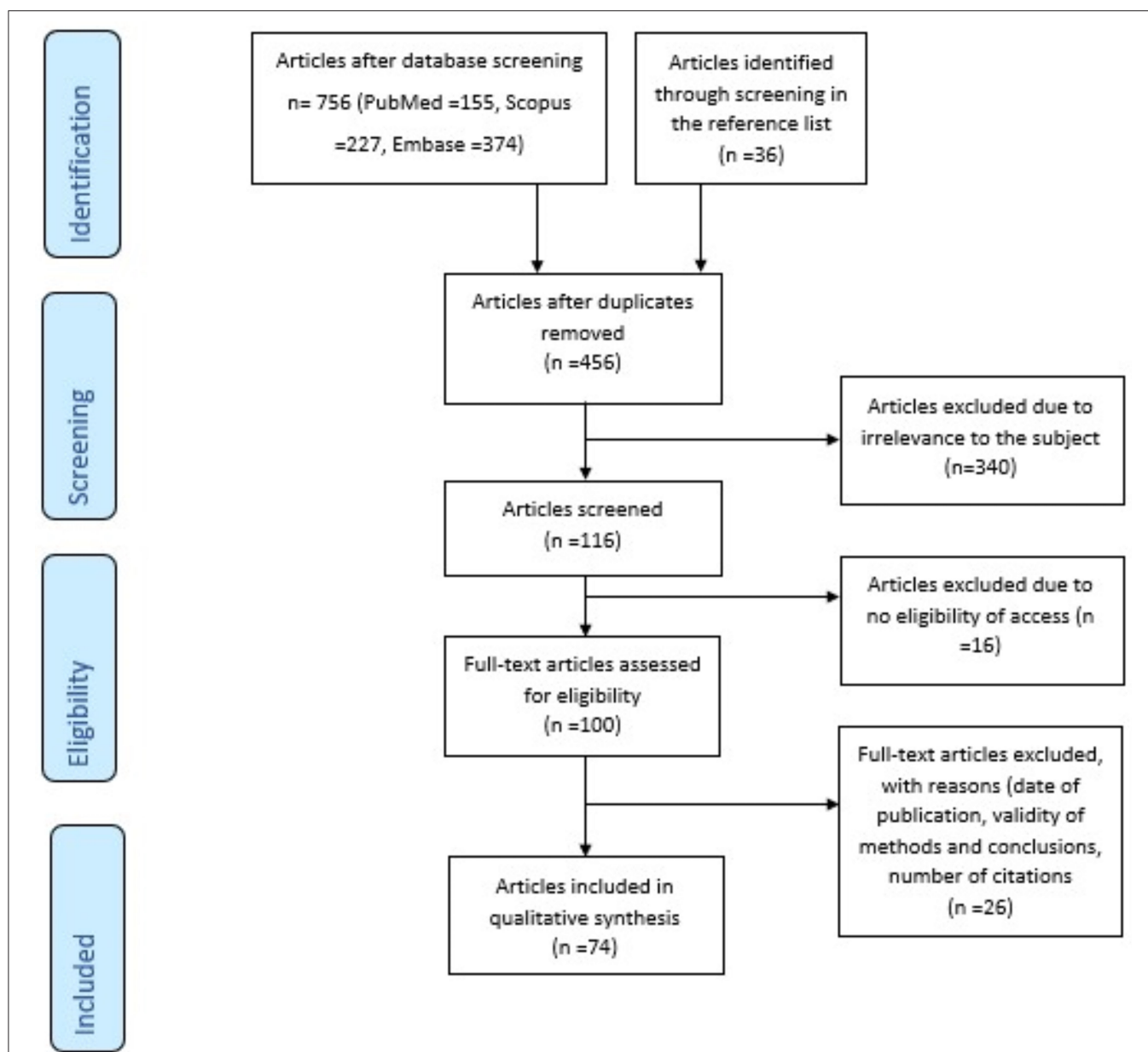


FIGURE 1 | Prisma flow diagram of the systematic review's methodology: This figure clearly depicts the methodology of the identification, screening, and exclusion of articles, in order to reach the final number ($n = 74$) of articles that are relevant to the topic and fulfill the inclusion criteria.

The progressive depletion of renal function during AKI leads to retention and accumulation of phosphate, due to the inability to excrete it. Phosphate acts as a negative regulator of 1-hydroxylase, an enzyme of $1,25(\text{OH})_2\text{D}$ synthesis, and therefore negatively contributes to VitD metabolism (11, 12). Moreover, as the final hydroxylation takes place in the kidney's active unit, the gradual loss of active nephrons leads to reduced ability of VitD synthesis (11) and to diminution of calcium's intestinal absorption. However, calcium's concentration depletion contributes to an increase of the parathyroid hormone (PTH). Due to secondary hyperparathyroidism and induction of CYP27B1, the levels of VitD can also be elevated (6, 11).

Fibroblast growth factor-23 (FGF-23), a phosphaturic factor, induces phosphaturia *via* acting on FGF-receptor (FGFR)/Klotho coreceptor complex, by (1) downregulating the luminal sodium/phosphate cotransporter in the proximal convoluted tubules, (2) suppressing kidneys' 1-hydroxylase, and (3) inducing 24-hydroxylase, which catalyzes the first step of VitD catabolism, causing hypophosphatemia and significantly lower levels of active VitD (13, 14). The soluble Klotho (sKl) can also regulate calcium and phosphate metabolism *via* an FGF-23-independent mechanism affecting the renal outer medullary potassium (ROMK) channel and the transient receptor potential vanilloid-5 (TRPV5). Moreover, sKl has a renoprotective effect

TABLE 1 | AKI as a cause of vitamin D (VitD) disruptions: AKI leads to the development of either hypovitaminosis D or hypervitaminosis D through various mechanisms.

Condition	Mechanism	Citation
Hypovitaminosis D	Retention and accumulation of phosphate.	(11, 12)
	Depletion of active nephrons, where the second hydroxylation occurs.	(11)
	Klotho deficiency leads to secondary increase of FGF-23 and deterioration of AKI.	(13, 14)
	Increase of FGF-23.	(13, 15)
Hypervitaminosis D	Due to calcium's diminution, the secondary hyperparathyroidism leads to induction of CYP27B1.	(6, 11)
	Rhabdomyolysis-induced mechanism.	(16)

This table presents the mechanisms via which AKI can lead to either VitD deficiency or VitD toxicity. FGF-23, fibroblast growth factor-23; AKI, acute kidney injury; CYP27B1, cytochrome P450 family 27 subfamily B member 1 or 1 α -hydroxylase.

via (1) the transforming growth factor- β 1 (TGF- β 1)-dependent anti-fibrotic mechanism and (2) the tumor necrosis factor (TNF)/nuclear factor- κ B (NF κ B)-dependent anti-inflammatory mechanism (14).

VitD deficiency can be caused by increased expression of FGF-23, which has been observed in AKI cases. Leaf et al. presented a case of a 45-year-old male hospitalized due to rhabdomyolysis and with a history of polysubstance abuse. The concurrent AKI was accompanied by elevated levels of FGF-23 and slightly decreased levels of 25(OH)D and 1,25(OH) $_2$ D (15). Additionally, FGF-23 can be secondarily elevated due to Klotho deficiency, which is prevalent, mostly, in chronic kidney disease (CKD) but also has been observed in AKI (13). Klotho's deficiency affects VitD levels, not only by upregulating FGF-23 but also through increasing the extent of AKI, due to depleted renoprotection, and leading, consequently, to greater VitD deficiency (14).

Moreover, Akmal et al. presented a study with patients suffering from rhabdomyolysis and AKI. The patients were initially hypocalcemic. However, during the diuretic phase, an elevation of 25(OH)D and 1,25(OH) $_2$ D was observed, which was greater in patients who developed hypercalcemia (16). These results suggest that the elevation of serum 1,25(OH) $_2$ D plays an important role in the development of hypercalcemia and that this increase may be due to extrarenal production or/and dysregulated renal production.

VitD Deficiency Induces AKI

VitD depletion, VDR knock-out, or disruption of VitD synthesis contributes to AKI development (Table 2) by leading to upregulation of RAAS and to elevated mRNA expression of renal-vascular renin. Due to obstruction and increased levels of extracellular matrix proteins (such as collagen I and fibronectin) and proinflammatory and profibrogenic factors [such as TGF- β , connective tissue growth factor, and monocyte chemoattractant protein-1 (MCP-1)], the renal injury becomes more severe. Moreover, epithelial-to-mesenchymal transition (EMT) was observed (17). Upon angiotensin I antagonist

administration, there was no difference between wild and VDR $^{-/-}$ mice, suggesting that angiotensin II is responsible for the increased renal damage. Contrarily, when VDR agonists were administered, proteinuria, podocytes' damage, mesangial dilation, macrophage infiltration, oxidative stress damage, proinflammatory and profibrogenic factors, and extracellular matrix protein and neutral lipid accumulation were reduced, proposing that VDR depletion worsens renal injury (17, 18).

Moreover, VitD deficiency can exacerbate pre-existing AKI [ischemia/reperfusion injury (IRI) induced AKI] by deteriorating the renal vascular condition and it can accelerate the AKI-to-CKD progression, via both an increased TGF- β 1 signaling and a decreased VDR and Klotho (19–21).

Buttar et al. presented a case of an 86-year-old woman with mild chronic disease and decreased levels of 25(OH)D, who developed rhabdomyolysis and secondary AKI after sitagliptin administration. The patient was on chronic atorvastatin therapy, and a possible interaction between these two drugs is proposed (22).

VitD as a Predictive and Prognostic Biomarker

In current clinical practice, most assays estimate total 25(OH)D, which cannot distinguish the three different forms of 25(OH)D [VitD binding protein (VDBP)-bound 25(OH)D, albumin-bound 25(OH)D, and free 25(OH)D]. This approach has many limitations since there are many variables that are influenced by physiologic and pathophysiologic conditions. The VDBP's affinity is affected by both hyperlipemic conditions and the three common variants of VDBP's gene, GC1F (group-specific component-1f), GC1S, and GC2 (23). Many efforts were made at inventing new assays that can directly, validly, efficiently, affordably, and quickly estimate free 25(OH)D, rather than calculating it using already-existing multi-factorial formulas. Free 25(OH)D measurement is considered to have more benefits than total 25(OH)D. So far, the studies have only proved the benefits of free 25(OH)D (1) in cases with differences in VDBP affinity, (2) in the elderly, (3) in pregnancy to detect VitD deficiency, (4) in liver diseases, (5) in kidney disorders (AKI, CKD), (6) in acromegaly, and (7) in allergies associated with atopy and pulmonary function in asthmatic children (23, 24). However, in many studies, no significant superiority of free 25(OH)D was observed. Still, there were many limitations in some of the studies, due to sample size and due to the use of monoclonal VDBP kits in multiracial/non-Caucasian populations, which affected the outcome (23, 24).

Regarding the predictive and prognostic role of VitD and its metabolites, mostly 25(OH)D but, also, 1,25(OH) $_2$ D can act as novel biomarkers of AKI (Supplementary Table 1).

Braun et al. found that 25(OH)D could serve as an independent predictor of AKI since serum 25(OH)D deficiency (<15 ng/ml) and insufficiency (15–30 ng/ml) are associated with greater risk of AKI. Moreover, serum 25(OH)D can also act as independent prognostic biomarker of 30-day mortality since its deficiency and insufficiency are linked to elevated risk of 30-day death (25).

TABLE 2 | VitD deficiency induces AKI through various mechanisms.

Condition	Type of model	Mechanism	Citation
VitD deficiency	Wild-type and VDR ^{-/-} mice	Upregulation of RAAS <i>via</i> lack of obstruction of angiotensin II receptor and mineral-corticoid receptor	(17)
	Diet-induced obesity mice	Elevation of extracellular matrix proteins and of profibrogenic and proinflammatory factors	(18)
	Wistar rats	Deterioration of already-existing renal vascular damage (ischemia/reperfusion injury induced AKI), which leads to accelerated AKI-to-CKD progression <i>via</i> increased TGF- β 1 signaling and <i>via</i> decreased VDR and Klotho expression	(19–21)
	Human case	In rhabdomyolysis-induced AKI, simultaneous administration of atorvastatin and sitagliptin	(22)

This table presents the mechanism through which VitD deficiency can lead to the development of AKI. RAAS, renin-angiotensin-aldosterone system; CKD, chronic kidney disease; TGF- β 1, tumor growth factor- β 1; VDR, VitD receptor.

Similarly, Zapatero et al. showed that serum 25(OH)D can act as a predictor of AKI, given that patients with serum 25(OH)D < 10.9 ng/ml had greater risk of AKI than VitD-sufficient patients. Moreover, lower levels of serum 25(OH)D were more common in non-survivor patients, and serum 25(OH)D acted as a prognostic biomarker of mortality (best cutoff value = 10.9 ng/ml) (26).

Leaf et al. found by using a two-multivariable-adjustment model that bioavailable 25(OH)D could serve as an independent prognostic biomarker of sepsis severity ($r = -0.45$) and mortality, while 25(OH)D can only correlate with sepsis severity ($r = -0.42$). Also, VDBP, 1,25(OH)₂D, and FGF-23 are predictive biomarkers of AKI, while FGF-23 served as biomarker of sepsis severity too ($r = 0.35$) (27).

Sahin et al. found that VitD acts as an independent predictive biomarker of contrast-induced nephropathy (CIN)-AKI, even in multivariable models (28).

Chaykovska et al. found that neither urinary VDBP (uVDBP) nor uVDBP/sCr could predict AKI development, but they served as biomarkers of need-of-dialysis, mortality, major adverse renal events (MAREs) (only uVDBP), and non-elective hospitalization. After adjustments, the predictive value of uVDBP was confirmed and was independent of well-known CIN risk factors, such as anemia, already-existing kidney injury, heart failure, and diabetes (29).

Vicente-Vicente et al. found that uVDBP can, potentially, predict the risk of gentamicin-induced AKI, in order to prevent its manifestation, given that increased uVDBP is associated with chronic proclivity to gentamicin nephrotoxicity (30).

Rebholz et al. found that VDBP, free 25(OH)D, bioavailable 25(OH)D, total 25(OH)D, and 1,25(OH)₂D (it is the only dependent biomarker) act as independent biomarkers of development of the ESRD stage, even after various multivariable models of adjustments (31).

Lai et al. studied the predictive and prognostic value of 1,25(OH)₂D, 25(OH)D, and [1,25(OH)₂D/25(OH)D] \times 1,000 ratio. As for the prediction of AKI, only 1,25(OH)₂D and [1,25(OH)₂D/25(OH)D] \times 1,000 ratio were significant, while as for the prognosis, these two biomarkers had been negatively correlated with AKI stage stratification (Risk, Injury, Failure). None of the markers could predict mortality, while 25(OH)D showed neither predictive nor prognostic effect (32).

Leaf et al. studied the prognostic value of human cathelicidin antimicrobial protein-18 (hCAP-18), free 25(OH)D and

bioavailable 25(OH)D, and the predictive value of hCAP-18. As for the prognosis, hCAP-18 measured on ICU day 1 is an independent biomarker of sepsis and 90-day mortality, as had been found *via* univariate and multivariate models of adjustments. However, hCAP-18 showed no significant predictive effect regarding AKI. As for the prognostic value of free and bioavailable 25(OH)D, only the former acted as a biomarker of 90-day mortality, while the latter showed no significant prognostic value (33).

Vitamin D in the Treatment of AKI

Many studies have been performed mainly with rats to define the potential use of VitD as treatment for AKI (Supplementary Table 2).

A dominant cause of AKI is IRI, induced by the actuation of inflammation and the increased expression of matrix metalloproteinases (MMPs). Ersan et al. studied the effect of paricalcitol on MMPs expression and, subsequently, on IRI progress. Pre-treatment with paricalcitol resulted in amelioration of IRI-AKI *via* an MMP-dependent inflammatory mechanism (34).

Hamzawy et al. studied the effect of pre-treatment with 22-oxacalcitriol (OCT) on IRI-AKI and found that it can ameliorate AKI through: (1) an anti-inflammatory mechanism *via* inhibition of Toll-like receptor-4 (TLR-4) and interferon- γ (IFN- γ), (2) a reduction of Na⁺/H⁺ exchanger-1 (NHE-1 exchanger), (3) a pro-autophagic action *via* elevating Beclin-1 expression and LC3II/LC3I ratio, (4) an anti-apoptotic action *via* elevating Bax/Bcl-2, cytochrome c and caspase-3 expression, and (5) an inhibitory action on G1 cell cycle arrest *via* reducing insulin-like growth factor-binding protein-7 (IGFBP-7) and tissue inhibitor of matrix metalloproteinases-2 (TIMP-2) expression (35).

In another study, Kapil et al. also investigated the protective role of VitD pre-treatment in IRI-AKI and showed a renoprotective effect in IRI against oxidation and lipid peroxidation mediated by peroxisome proliferator-activated receptor- γ (PPAR- γ) (36).

Additionally, Arfan et al. studied the impact of VitD on IRI-AKI and found that its administration can mitigate AKI through reducing the inflammation and the production of myofibroblasts, *via* reducing the expression TLR-4 and MCP-1 (37).

Also, Silva Barbosa studied the effect of estrogen sulfotransferase (SULT1E1) inhibition on IRI-AKI and found

that this inhibition attenuates AKI *via* elevating the VDR activation, as shown by the elevated Cyp24 α 1 and Ccnd1 and by the decreased Fgg expression (38). This effect has been found to be estrogen- and androgen-independent (38). Additionally, only in male mice, the liver expression of Sult1e1 has been found to be necessary for IRI-AKI development, thus proposing a tissue- and sex-specific relation between the expression of Sult1e1 and sensitivity to IRI-AKI (38).

Moreover, Lee et al. examined the effect of paracalcitol on IRI and demonstrated that it can attenuate AKI *via* an anti-inflammatory mechanism mediated by the inhibition of TLR-4 expression and the suppression of NF κ B signaling, by increasing IkB in a TNF- α -dependent way (39).

Xu et al. studied the effect of pre-treatment with VitD on liposaccharide (LPS)-induced AKI and found that it can attenuate AKI through: (1) an anti-oxidative mechanism *via* increasing glutathione (GSH), superoxide dismutase (SOD)-1, and SOD-2 and *via* decreasing nitric oxide synthase (iNOS), p47phox, and gp91phox (subunits of renal NADPH oxidase) and (2) an anti-apoptotic mechanism (40).

Du et al. studied the effect of paracalcitol on LPS-AKI and found that its administration ameliorates AKI through: (1) an anti-inflammatory mechanism *via* reduction of TLR-4 and (2) an anti-apoptotic mechanism *via* elevation of Bcl-2 (it is anti-apoptotic) and *via* decrease of caspase-3, PUMA (it is a pro-apoptotic member of the Bcl-2 family), and miR-155 (it targets Bcl-2 and blocks its expression) (41). Regarding the decrease of PUMA and miR-155, it has been found that this is due to the VitD-dependent inhibition of NF κ B expression by disrupting the IKK kinase complex or by blocking the p65/p50 nuclear translocation (41).

Park et al. investigated the impact of paricalcitol on cisplatin-induced AKI and found that it ameliorated AKI through: (1) inhibiting EMT, (2) reducing apoptosis *via* increasing Bcl-2 and *via* decreasing P-p53 and p21, p-Bad, Bax, and caspase-3 expression, and (3) increasing the cell proliferation *via* elevating the expression of p27^{kip1} and decreasing that of cyclin-dependent kinase-2 (CDK2) and Cyclin E. The underlying mechanism of these actions includes the inhibition of TGF- β 1 and p53 signaling pathways and the elevation of the p27^{kip1} signaling pathway (42).

Another study on the effect of paricalcitol on cisplatin-induced AKI demonstrated that the ameliorated AKI was correlated with reduced lipid peroxidation and ferroptotic cell death due to the direct binding of VDR to the glutathione peroxidase 4 (GPX4) promoter, which induces the expression of GPX4, affecting the induction of ferroptosis (43).

Also, Moneim et al. studied the impact of alfacalcidol and BQ-123, a selective endothelin receptor A blocker (ET_AR blocker), on cisplatin-induced AKI. Their administration led to attenuation of AKI *via* a VDR-, ET-1-, and ET_AR-dependent mechanism, which includes the signaling cascade of Pnf- κ Bp65, TNF- α , and TGF- β 1. The combined administration of both drugs led to an enhanced therapeutic effect. These results propose a merger of VitD and endothelin-1 signaling pathways, which is promising as a therapeutic option for cisplatin-induced AKI (44).

In another study, administration of 1,25(OH)₂D₃ on gentamicin-induced AKI had no beneficial effect on ameliorating

AKI (45). However, there was a decrease of the systolic blood pressure, to some extent, and an increase of the urine volume, probably due to its inhibitory role in RAAS. Also, VitD acted as an antioxidant factor by increasing GSH (45). So, VitD may be a promising therapeutic agent due to its RAAS- and GSH-related action.

Additionally, El-Boshy et al. studied the role of VitD in paracetamol-induced AKI and liver failure. They found that both the therapeutic and the prophylactic use of VitD can ameliorate AKI and liver failure through: (1) an anti-apoptotic mechanism *via* reducing caspase-3 and (2) a decrease of Cyp24 α 1 and VDBP expression and an increase of Cyp27b1, Cyp2R1, and VDR expression, (3) an anti-oxidative and anti-inflammatory mechanism *via* decreasing IL1- β , IL1R1 (IL-1 receptor 1), IL-6, IL6R (IL-6 receptor), IFN- γ , IFNGR1 (IFN- γ receptor 1), IL17A, and IL17RA (IL-17 receptor A) and *via* elevating GSH, chloramphenicol acetyltransferase (CAT), Gpx, IL10 (and its gene), IL22, and IL22RA (IL-22 receptor A) (46). The effect of the prophylactic use of VitD has been found to be greater than that of its use for treatment (46).

In another study, the administration of 500 and 1,000 IU/kg VitD can ameliorate AKI and liver failure induced by paracetamol. Interestingly, the dose of 500 IU/kg presented a greater protective role (47). The underlying mechanism includes: (1) an anti-oxidative action *via* the reduction of the expression of heme oxygenase 1 (HO-1) and its regulators, NrF2 and BACH1, and (2) an anti-inflammatory action *via* the reduction of NF κ B, TNF- α , and IL-10 (47).

Finally, Reis et al. studied the impact of calcitriol on rhabdomyolysis-induced AKI and found that AKI was ameliorated through: (1) an anti-inflammatory mechanism *via* decreasing NF κ B and Jun N-terminal kinase (p-JNK), MCP-1, and IL-1 β , (2) an anti-oxidative mechanism *via* elevating SOD and *via* decreasing 8-epi-PGF2 α and nitrotyrosine, (3) reducing vimentin, proliferating cell nuclear antigen (PCNA), and caspase-3, and (4) elevating CYP24 (48).

Vitamin D Toxicity Induced AKI

Although VitD seems to ameliorate AKI, its administration should be performed with extreme caution. Hypervitaminosis D can significantly afflict kidney function by inducing hypercalcemia and hyperphosphatemia. Specifically, hypercalcemia leads to the development of nephrogenic diabetes insipidus, which affects water homeostasis by causing polyuria and diuresis. Therefore, there is evident loss of water, leading to hypovolemia, which causes AKI and concomitantly deteriorates the hypercalcemia. So, there is a continuous circle of hypercalcemia causing hypovolemia-induced AKI and vice versa (49–51). Moreover, hypercalcemia and hypercalciuria can lead to deposition of calcium, thus causing nephrolithiasis and renal calcification, which can cause the development of AKI (49–51). Also, hypercalcemia can cause renal vasoconstriction, which subsequently leads to severe GFR decrease and AKI (49–51). As for the hyperphosphatemia, it can cause acute phosphate nephropathy due to the tubulointerstitial deposition of phosphate calcium, a condition that concomitantly worsens the already-existing hyperphosphatemia, thus leading to a

vicious circle of deterioration (49–51). As for the acute phosphate nephropathy, it can also be due to significant phosphate intake and the subsequent diarrhea-induced hypovolemia (52). Specifically, the hypovolemia leads to great reabsorption of water and sodium chloride in the proximal tubule and the descending limb of Henle's loop, while calcium and phosphate are not that easily reabsorbed (52). As a result, the greater concentration of these substances in the distal tubule and the collecting duct leads to significant deposition of calcium phosphate in these structures, a deposition that is exacerbated due to surface expression of hyaluronan and osteopontin as a result of hypovolemia-induced tubular injury (52). Among risk factors for the development of acute phosphate nephropathy are: advanced age, hypertension, CKD, drugs (such as angiotensin-converting enzyme inhibitors, angiotensin II receptor blockers, and loop- or thiazide-type diuretics), female gender, diabetes mellitus, and the use of non-steroidal anti-inflammatory drugs (NSAIDs) (52). As for the treatment of acute phosphate nephropathy, kidney replacement therapy (KRT) can be used to tackle this condition, and particularly, sustained low efficiency dialysis (SLED) has gained ground in treatment, given that it is a hybrid technique, which combines the advantages of both intermittent and continuous KRT (53).

VitD-mediated hypercalcemia can be due to: (1) excessive VitD₂ or D₃ ingestion/supplementation, (2) extravagant calcitriol ingestion or pharmaceutical administration, (3) elevated ectopic production of calcitriol (evident in granulomatous diseases, such as tuberculosis, sarcoidosis, leprosy, fungal infections, and others; in Hodgkin's and non-Hodgkin's lymphomas; in malignant

lymphoproliferative diseases), (4) milk-alkali syndrome (MAS), and (5) depleted catabolism of calcitriol due to mutations of CYP24A1 genes (54).

VitD toxicity due to immoderate administration of VitD supplements or overfortified milk is a global phenomenon (55), potentially affecting kidney function. Chowdry et al. presented a study of VitD toxicity incidents in a tertiary care center at the Sher-i-Kashmir Institute of Medical Sciences, in which 16 out of 19 patients were identified with hypervitaminosis D-induced AKI due to extravagant doses of VitD (median cumulative dose of VitD is 6,000,000 IU), in order to correct VitD deficiency. Not all the patients with toxic levels of 25(OH)D (>150 ng/ml) developed symptoms (56). Similarly, 13 patients in Brazil developed AKI due to intramuscular injection of veterinary supplements of vitamins A, D, and E for esthetic purposes (57).

Also, VitD toxicity-induced AKI has been observed due to anabolic steroid and VitD supplement abuse (58), dispensing errors (59), dosage malpractice and overcorrection of VitD deficiency (60), intramuscular injection of VitD after operation (61), overfortification of milk with VitD (62), over-the-counter supplements (63), and topical treatment with calcitriol analog in combination with oral calcium/VitD for psoriasis (64).

Additionally, in another study, 33 patients with osteoporosis treated with 0.75 µg/day eldecacitol developed hypercalcemia-induced AKI since the discontinuation of eldecacitol ameliorated the situation (65). Furthermore, 11 patients developed with hypercalcemia-induced AKI, due to alfacalcidol (9 patients) or calcitriol (2 patients) (65).

TABLE 3 | Cases of hypervitaminosis D as cause of AKI.

References	Article type	Sample size	Substance	Concentration
Chowdry et al. (56)	Case series	19 patients	25(OH)D	371 (190–988) ng/ml
Daher et al. (57)	Case series	16 patients	Vitamin D	135 ± 75 ng/ml
Daher et al. (58)	Case series	2 patients	Vitamin D	N/A
Nasri et al. (59)	Case	1 patient	Vitamin D	>400 nmol/L
Kaur et al. (60)	Case series	16 patients	Serum 25(OH)D	371 (175–1,161) ng/ml
Bansal et al. (61)	Case	1 patient	Serum 25(OH)D	150 ng/ml
Jacobus et al. (62)	Case series	8 patients	Serum 25(OH)D	293 ± 174 ng/ml
Koutkia et al. (63)	Case	1 patient	Serum 25(OH)D	487.3 ng/ml
Corden et al. (64)	Case	1 patient	Vitamin D	N/A
Aihara et al. (65)	Case series	43 patients	Eldecacitol (32 patients) Alfacalcidol (9 patients) Calcitriol (2 patients)	N/A
Tollit et al. (66)	Case	1 patient	Vitamin D	N/A
Lavender et al. (69)	Case	1 patient	1,25(OH) ₂ D ₃	318 pmol/L
Karmali et al. (70)	Case	1 patient	1,25(OH) ₂ D ₃	145 pg/ml
Altun et al. (71)	Case	1 patient	Calcitriol	N/A
Jeong et al. (72)	Case	1 patient	Calcitriol	N/A
Asghar et al. (73)	Case	1 patient	Vitamin D	119 ng/ml
Tsao et al. (74)	Case	1 patient	1,25(OH) ₂ D ₃	268 pmol/L

This table presents the different studies that have been done regarding vitamin D toxicity as a cause of AKI. It depicts information regarding the type of study conducted, the sample size, the vitamin D-related substance that has been calculated, and the concentration of this substance.

Moreover, VitD toxicity-induced AKI is related with granulomatous diseases (especially sarcoidosis), MAS, subclinical hyperparathyroidism, and immune reconstitution syndrome.

Tollitt and Solomon presented a case of a 38-year-old male and two other patients with a history of sarcoidosis. After treatment with high doses of cholecalciferol, the patient developed different symptoms of hypercalcemia, such as vomiting, nausea, muscle cramps, and constipation, and subsequent AKI (66). In these cases, the hypercalcemia is due to the extrarenal synthesis of $1,25(\text{OH})_2\text{D}$ by macrophages within the sarcoid granulomas, the process of which is without systemic regulation (67, 68). The sufficient administration of prednisolone inhibits macrophages' $1-\alpha$ hydroxylase activity and protects against hypercalcemia (66). A similar relation between VitD administration and VitD-induced hypercalcemia has been observed in tuberculosis and Hodgkin's lymphomas (69, 70).

MAS is a result of VitD, calcium carbonate, and bisphosphonate administration for osteoporosis, iatrogenic hypothyroidism, and idiopathic hypothyroidism treatment. Although there is limited knowledge regarding its pathophysiology, it affects patients who absorb more calcium than average. Among the risk factors are old age, medication that reduces GFR, and hypovolemia. MAS consists of hypercalcemia, metabolic alkalosis, and renal function disruption, and, particularly, hypercalcemia leads to AKI via renal vasoconstriction, polyuria, and GFR depletion (71, 72).

Asghar et al. presented a case of a 55-year-old female who had slightly elevated calcium and was found to be VitD-deficient. The prescription of VitD led later to the development of a giant cystic parathyroid adenoma and to manifestation of parathyroid crisis, accompanied by severe gastrointestinal symptoms and AKI. This suggests that VitD administration can unveil subclinical hyperparathyroidism (73).

Finally, Tsao et al. described a case of immune reconstitution inflammatory syndrome (IRIS) due to *Mycobacterium tuberculosis* lymphadenitis. It was manifested in a patient with history of silent human immunodeficiency virus-1 (HIV-1) infection, who was in highly active antiretroviral therapy (HAART) because of plasma viral load increase. This 48-year-old male patient, after the initiation of HAART, developed various hypercalcemia symptoms, accompanied by hypervitaminosis D [$1,25(\text{OH})_2\text{D}_3$, 268 pmol/L], which progressed to coma and AKI. IRIS can be due to many opportunistic infections, but the most common infection is *M. tuberculosis*, which is incriminated as the cause of VitD-induced hypercalcemia, which leads to AKI (74). All

cases of hypervitaminosis D-induced AKI are summarized in Table 3.

CONCLUSION

It can be concluded that there is a two-way relation between AKI and VitD. Specifically, disruptions of VitD—both hypovitaminosis and hypervitaminosis—can lead to the development of AKI, while also, AKI can contribute to dysregulation of VitD's homeostasis and function.

On this ground, due to this two-way causality relation, VitD is examined on whether its forms can act as a novel biomarker of AKI. Many studies have confirmed VitD's significant role as a predictive (it can help to determine the risk of developing AKI) and prognostic (it can help determine the stage, progress, clinical outcomes, and mortality risk of AKI) biomarker.

Finally, VitD is found to be a potentially important therapeutic factor for AKI due to its multisystemic functions, which include regulation of many enzymic mechanisms via genomic and non-genomic actions.

However, many prospective studies and trials need to be conducted, in order to: (1) fully determine the complex relation between VitD and AKI, (2) find the best cutoff points with the most significant statistic importance, and (3) determine the therapeutic protocols of VitD as treatment of AKI and its potential adverse effects.

Despite the necessity of future studies, VitD is a very promising biomarker and a potential treatment for AKI, which is one the most prominent health problems nowadays.

DATA AVAILABILITY STATEMENT

The original contributions generated in the study are included in the article/Supplementary Material, further inquiries can be directed to the corresponding author.

AUTHOR CONTRIBUTIONS

MP and TP: conceptualization and supervision. SG and MP: data curation and investigation. SG: formal analysis and writing—original draft. All authors methodology and writing—review and editing.

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fnut.2020.630951/full#supplementary-material>

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Evaluating the Impact of Goal Setting on Improving Diet Quality in Chronic Kidney Disease

Chi H. Chan^{1*}, Marguerite Conley², Marina M. Reeves³, Katrina L. Campbell^{4,5,6} and Jaimon T. Kelly^{5,6*}

¹ Faculty of Health Science and Medicine, Bond University, Gold Coast, QLD, Australia, ² Department of Nutrition and Dietetics, Princess Alexandra Hospital, Brisbane, QLD, Australia, ³ Faculty of Medicine, School of Public Health, The University of Queensland, Brisbane, QLD, Australia, ⁴ Allied Health Services, Metro North Hospital and Health Service, Brisbane, QLD, Australia, ⁵ Centre of Applied Health Economics, Griffith University, Brisbane, QLD, Australia, ⁶ Menzies Health Institute Queensland, Griffith University, Gold Coast, QLD, Australia

OPEN ACCESS

Edited by:

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(INCMNSZ), Mexico

*Correspondence:

Chi H. Chan
ianchan1230@gmail.com
Jaimon T. Kelly
jaimon.kelly@griffith.edu.au

Specialty section:

This article was submitted to
Clinical Nutrition,
a section of the journal
Frontiers in Nutrition

Received: 10 November 2020

Accepted: 05 February 2021

Published: 12 March 2021

Citation:

Chan CH, Conley M, Reeves MM,
Campbell KL and Kelly JT (2021)
Evaluating the Impact of Goal Setting
on Improving Diet Quality in Chronic
Kidney Disease.
Front. Nutr. 8:627753.
doi: 10.3389/fnut.2021.627753

Background: Improving diet quality in chronic kidney disease (CKD) is challenging due to a myriad of competing recommendations. Patient-centered goal setting can facilitate dietary behavior change; however, its role in improving diet quality in CKD has not been investigated.

Aim: The aim of the study is to evaluate the effects of goal setting on improving diet quality in stages 3–4 CKD.

Methods: Forty-one participants completed a 6-month dietitian-led telehealth (combined coaching calls and text messages) intervention as part of a larger RCT. Participants set one to two diet-related SMART goals and received weekly goal tracking text messages. Dietary intake was assessed using the Australian Eating Survey at baseline, 3, and 6 months, with diet quality determined using the Alternate Healthy Eating Index (AHEI).

Results: Significant improvements in AHEI (+6.9 points; 95% CI 1.2–12.7), vegetable (+1.1 serves; 95% CI 0.0–2.3) and fiber intake (+4.2 g; 95% CI 0.2–8.2) were observed at 3 months in participants setting a fruit and/or vegetable goal, compared with those who did not. However, no significant or meaningful changes were observed at 6 months. No other goal setting strategy appeared in effect on diet intake behavior or clinical outcomes in this group of CKD participants.

Conclusions: Patient-centered goal setting, particularly in relation to fruit and vegetable intake, as part of a telehealth coaching program, significantly improved diet quality (AHEI), vegetable and fiber intake over 3 months. More support may be required to achieve longer-term behavior change in stages 3–4 CKD patients.

Keywords: chronic kidney disease, diet quality, goal setting, self-management, telehealth

INTRODUCTION

Chronic kidney disease (CKD) is a public health issue affecting over 10% of the global population (1). Dietary intake and lifestyle behaviors are important elements in the self-management of CKD and its associated comorbidities (2, 3). Diet recommendations have historically focused on restricting nutrients such as protein, sodium, potassium, and phosphate to slow the progression

of CKD and manage its associated conditions such as hyperkalemia and hyperphosphatemia. However, the evidence base for these nutrient restrictions is conflicting, and people with CKD struggle to adhere to these recommendations for a long term (4). Recent evidence has hypothesized that food-based dietary pattern approaches may improve CKD outcomes (4). Observational data have shown that a healthy dietary pattern is associated with decreased risks of all-cause mortality in people with CKD (5) and reduces risks of CKD comorbidities such as cardiovascular disease and diabetes mellitus (6, 7). A small number of randomized controlled trials (RCTs) have also shown that people with CKD can improve their diet quality in the short term (8). However, the long-term adherence to diet pattern interventions is unknown and remains a critical factor to address in helping patients succeed in changing their dietary behavior in practice for a long term (9–11).

Goal setting as a strategy for behavior change has been shown to be effective in CKD comorbidities management and may facilitate long-term behavior change through improved self-efficacy (12, 13). For example, in diabetes and heart failure patients, goal setting has been found to be beneficial for behavioral changes in dietary intake and physical activity (14), and improves self-efficacy and self-management skills (15, 16). People who set a specific, measurable, achievable, realistic, and timed (SMART) goal are likely to attain a successful outcome (17), and it is hypothesized that this might help people with CKD to achieve better dietary intake. However, the effect of SMART goal setting on improving diet quality in CKD-specific populations has not been evaluated to date.

A recent pilot RCT of a telehealth coaching program in people with stages 3–4 CKD showed significant and meaningful improvements in diet quality following 3 months of intensive coaching (8, 18). However, these effects were attenuated after 3 months of less-intensive coaching. As coaching was tailored to participant goals in this study, this has prompted the need to investigate whether specific goal setting influenced the change in diet quality in this sample of individuals with CKD. Therefore, this current study aimed to investigate whether specific dietary goal setting was associated with improvements in diet quality and other dietary and health indicators in people with stages 3–4 CKD.

METHODS

This study is a single-arm secondary analysis of intervention participants recruited from the previous ENTICE-CKD trial. The previous ENTICE-CKD study was a pilot RCT originally conducted in 80 participants ($n = 41$ intervention; $n = 39$ control) across three sites with stages 3–4 CKD (eGFR < 60 ml/min/1.73 m²), who owned a mobile phone (18). All participants were metabolically stable and were cleared to participate by their treating nephrologists. Each participants' nephrologist maintained responsibility for the patients' medical management throughout the study period. A single-arm analysis within the intervention group was the most appropriate to answer our specific research aim as the control group participants

in the previous ENTICE-CKD study did not receive any intervention to set SMART goals nor had data collected on goal achievement throughout the study. This study conformed to the Declaration of Helsinki; all participants provided written informed consent, and the research protocol was approved by the Metro South Hospital and Health Service and Bond University Human Research Ethics Committees.

The previous ENTICE-CKD intervention is described in detail elsewhere (16, 18). Briefly, the previous ENTICE-CKD study was a telehealth intervention completed in two phases, phase 1 (baseline to 3 months) and phase 2 (3–6 months). **Figure 1** shows the study schema; the intervention participants received intensive telehealth coaching (telephone and text messages) fortnightly from the same Accredited Practising Dietitian (APD), with weekly tailored text messages to support the coaching content in phase 1. Each coaching call addressed improving diet quality in line with the Australian Dietary Guidelines (19), which was further tailored to suit individual participant-associated comorbidities and their set goals. A program workbook was provided to all participants, which included sections relevant to SMART goal setting strategies and self-monitoring practice. In the initial coaching call, the dietitian coach guided participants through the workbook and worked with participants to set one to two diet-related SMART goals. The setting of each SMART goal was done collaboratively between the dietitian and the participant, based on a comprehensive medical and nutrition assessment and the participant stage of change. The purpose of the intervention was to improve diet quality, and therefore, each SMART goal set by participants was designed to address an area of dietary intake in order to achieve this, rather than the improvement in a clinical outcome. The dietitian provided evidence-based nutrition counseling using motivational interviewing techniques (18), addressing barriers to goal achievement, and action planning for the following 2 weeks. Participants also determined their individual preference for the timing and frequency of receiving text messages, which supported the coaching content and reminded participants of their set goals (detailed in **Table 1**). In the five subsequent coaching calls, the dietitian checked participants' progress in regard to their set goals and provided counseling support as per the study protocol (18). In phase 2, the intervention participants received no further coaching calls, however, and were continued to be sent the same tailored text messages at a frequency determined by the participants (protocol detailed in **Table 1**).

The tailored text messages in both phases were sent to individual participants using a web-based, semi-automated text message management platform (Propelo, www.propelo.com.au), developed and administered by The University of Queensland's School of Public Health (20). Tailored text messages were used to remind and track participants' goals each week (see **Table 1**). Participants were asked to text back either "yes" or "no" regarding their weekly goal achievement.

Data were collected at three time-points throughout the study including baseline, end-of phase 1 (3 months), and end-of phase 2 (6 months). The primary outcome was the change in the Alternate Healthy Eating Index (AHEI)-2010. The AHEI-10

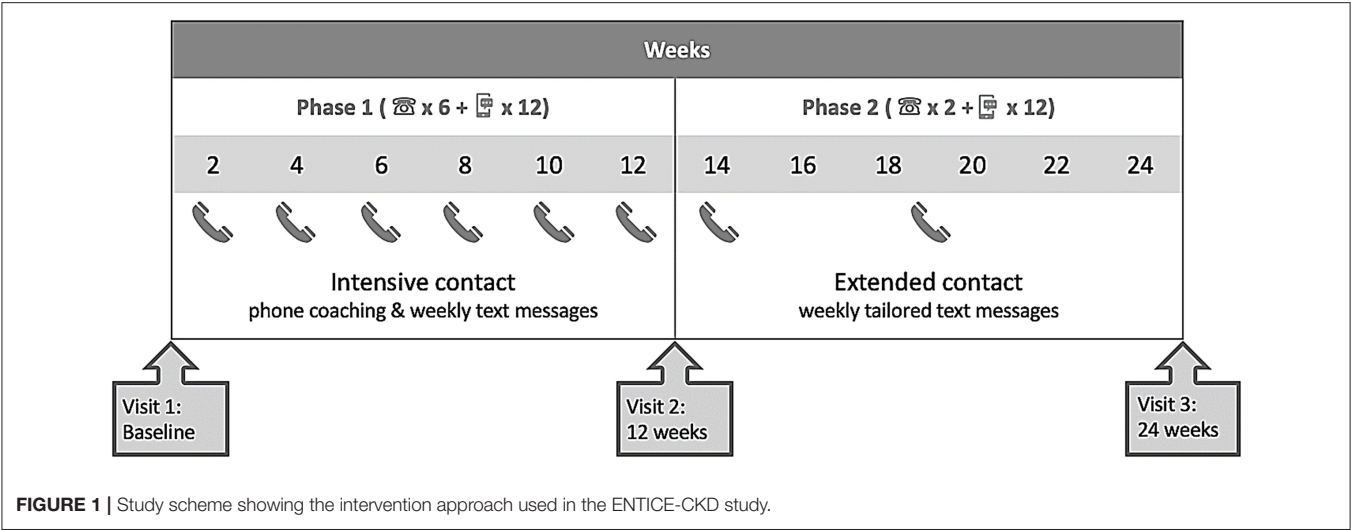


TABLE 1 | Text message type and examples.

Message type	Example text	Frequency
Educational	Dietary fiber intake reduces your cholesterol levels and controls your blood sugar. Include whole grain bread and cereals, fruits and vegetables regularly	1–4 every 2 weeks
Self-monitoring	Hi, (name). Are you keeping track of your fruit/vegetable intake every day? Remember your goal to meet at least 5 serves this week	1–4 every 2 weeks
Goal check	Hi, (name). Did you reach your goal to eat 5 fruits/vegetables 4 times this week? Text me back yes or no to let me know	2–4 every 2 weeks
Educational permutations	Choose high-fiber, low-potassium breakfast cereals. Good choices are Multigrain Weetbix, Rolled Oats, Guardian, Oatbritz, and Special K	0–2 every 2 weeks

score is derived from the consumption of 11 food groups or nutrients: (1) vegetables, (2) fruits, (3) whole grains, (4) sugar-sweetened beverages and fruit juice, (5) nuts and legumes, (6) red and processed meat, (7) fish, (8) sodium, (9) alcohol, (10) trans-fat, and (11) long-chain fats. Each of the 11 components is scored from 1 (poor adherence) to 10 (perfect adherence), with the total AHEI-2010 diet quality score therefore ranging from 0 (non-adherence) to 110 (perfect adherence), the higher AHEI score. Further information about the AHEI-2010 and how it was collected and applied in this study can be found elsewhere (8). We based the diet quality rating categories on previous studies showing approximations (<38 points indicating a “low quality,” 3–67 points as “intermediate,” and >68 points as a “high” diet quality) (21). Diet consumption data for the AHEI-2010 components 1–9 were collected on the Australian Eating Survey and calculated using the Australian food composition databases, AusNut 2011–2013 and AusFoods Revision 5 (Australian Government Publishing Service, Canberra) over a 3-month period. Data on the AHEI-2010 components 10 and 11 were calculated using NUTTAB 2010 (database developed by the Food Standards Australia and New Zealand) and FoodWorks (version 18, The Nutrition Company, Long Valley, NJ, USA). Secondary outcomes, which related to participants’ set goals, included fruit and vegetable intake, fiber intake, sodium intake, blood pressure, and body weight. Fruit and vegetable intake, fiber intake, and

sodium intake were measured using the Australian Eating Survey. Body weight and blood pressure were collected as part of usual care where possible or by a trained site investigator who was blind to the treatment allocation, following a standardized protocol (8).

Total goal attainment was determined based on the number of “yes” responses to the weekly goal check text messages. The number of goal achievements was measured between 0 (i.e., no goal achieved) to 12 (i.e., all goals achieved) in each 12-week phase. If participants did not respond to the goal check message, these data were recorded as goal not achieved.

For statistical analysis, participant goals were categorized into either a “no goal” (received all other text messages but elected not to set a goal outside the dietary intervention targets), a “fruit and/or vegetable” goal (as the majority of participants set both fruit and vegetable intake goals, rather than one of these food groups separately), or a “healthy eating” goal. These three dietary goals were the most frequently set goals by participants with enough data available to conduct an exploratory analysis. The baseline-observation-carried-forward (BOCF) approach was applied to impute missing data in the current study. Data were checked for normality. One-way analyses of covariance (ANCOVA; adjusted for baseline) were conducted to determine potential differences in the primary and secondary outcomes differed across the two groups defined as whether participants

TABLE 2 | Baseline characteristics.

Characteristic	Participant (n = 41)
Age, years	63 ± 12
Gender, % of male	63
Ethnicity, n (%)	
Caucasian	35 (85)
Asian	2 (5)
European	2 (5)
Indigenous	1 (2.5)
Other	1 (2.5)
Hypertension, n (%)	34 (83)
Diabetes, n (%)	16 (39)
Cardiovascular disease, n (%)	14 (34)
eGFR, ml/min/1.73 m ²	36 ± 12
Systolic blood pressure, mmHg	136 ± 18
Diastolic blood pressure, mmHg	80 ± 12
Weight, kg	96 ± 22
Body mass index, kg/m ²	33 ± 7
Alternate Healthy Eating Index	71.4 ± 11.8
Fruit, serves/day	1.5 ± 0.9
Vegetables, serves/day	4.2 ± 2.0
Sodium, mg	2,379 ± 1,392
Fiber, g	24.1 ± 9.1

Data were reported as mean ± SD.

eGFR, estimated glomerular filtration rate; g, grams; mg, milligrams; kg, kilograms.

had set one of the three listed goals or not, in each phase. Results from the analyses are presented as mean and 95% confidence interval (CI). A $p < 0.05$ (two sided) was considered statistically significant. However, given the exploratory nature of this pilot study, all results were also considered against minimal clinically important difference (MCID), which was informed by a thorough review of the literature and expert clinical guidance. For the purpose of this study, we defined MCID as 20% change in diet quality (AHEI score) (22), 5 grams (g) change in fiber intake (23), 0.5 serves in fruit intake (24), one serve in vegetable intake (24), 780 milligrams (mg) in sodium intake (25), 5 mmHg systolic blood pressure (SBP) (26), and 5% change in body weight (27). All statistical tests were performed using SPSS (version 26. Chicago: SPSS Inc.).

RESULTS

All 41 intervention participants from the previous ENTICE-CKD study were included in the analysis (63% male; mean age 63 ± 12 years; **Table 2**). Comorbidities were common across the population, 83% of participants having hypertension, 39% having diabetes, and 34% having cardiovascular disease.

At baseline (**Table 2**), the mean AHEI of participants was 71.4 points (rating in the “high” diet quality category). The average fruit and vegetable intake was 1.5 ± 0.9 serves and 4.2 ± 2.0 serves, respectively. The average fiber and sodium consumptions were 24.1 ± 9.1 g and 2,379 ± 1,392 mg, respectively. Participants

had a mean body mass index (BMI) and SBP of 33 ± 7 kg/m² and 136 ± 18 mmHg, respectively.

The most common goals set by participants included “general healthy eating” goal ($n = 15$ in phase 1; $n = 21$ in phase 2), “fruit and/or vegetable” goal ($n = 16$ in phase 1; $n = 13$ in phase 2), and “no goal” ($n = 5$ in phase 1; $n = 7$ in phase 2). A total of 944 goal check messages were sent to participants throughout the trial, 365 messages in phase 1 and 579 messages in phase 2. Participants’ response rates to goal check messages were 46.3% in phase 1 and 38.7% in phase 2, respectively, of which, 41.9% of participants stated they met their set goal in phase 1 and 34% in phase 2.

Table 3 shows the effect of goal setting on diet intake. Compared with participants who did not set a “fruit and/or vegetable” goal in phase 1 ($n = 25$), those that did ($n = 16$) had a statistically significant improvement in AHEI (+6.9 points, 95% CI 1.2–12.7; 10% increase), vegetable intake (1.1 serves per day, 95% CI 0.0–2.3), and fiber intake (+4.2 g, 95% CI 0.2–8.2). Fruit intake was the only variable shown to clinically meaningfully improve (+0.5 serves, 95% CI −0.1–1.1); however, it was not statistically significant. At 6 months, these associations were all attenuated; however, a significant (but unlikely clinically meaningful) increase in sodium intake (+413.7 mg, 95% CI 28–799.4) was observed in participants setting a “fruit and/or vegetable” goal ($n = 13$) compared with those who did not.

In those who set a “general healthy eating” goal (vs. those who did not), no statistically or clinically meaningful differences in any of the outcomes were observed at either 3 or 6 months. No associations were observed at 3 or 6 months in those who set “no goal” vs. those who set a goal.

There was no clinically or statistically significant effects observed for body weight or SBP (**Supplementary Table 1**).

DISCUSSION

This secondary analysis from the previous ENTICE-CKD study aimed to investigate the relationship between patient-centered goal setting and diet quality improvement and determine the specific dietary goals that might help people with CKD to improve their diet quality and clinical outcomes. Significant and potentially meaningful associations were primarily observed at 3 months. These results are exploratory and hypothesis generating and suggest that focusing on fruit and vegetable intake goals can help to promote a clinically significant short-term improvement in diet quality, vegetable and fruit intake, and dietary fiber intake with dietitian-led telehealth coaching.

Both diet quality and fiber intake are important in CKD self-management. Compared with other goal categories set by participants, “fruit and/or vegetable” goals appeared to be the strongest enabler of achieving an improved diet quality, vegetables and fiber intake in CKD patients, with the support of intensive telehealth coaching from a dietitian over 3 months. In one of the largest longitudinal analyses of continuous changes in the diet quality, a 20% improvement in AHEI was associated with an 8–17% reduced risk of all-cause mortality in people

TABLE 3 | Between group changes in AHEI and dietary intake across the two groups of participants who either set a specific goal compared with participants who did not, in each phase of the study.

Outcomes#Goal	Phase	AHEI (points)	Fruit (serves/day)	Vegetables (serves/day)	Fiber (g/day)	Sodium (mg/day)
Healthy eating goal ^a	Phase 1	-0.2 (-6.3, 5.9)	-0.3 (-1.0, 0.4)	0.6 (-0.6, 1.8)	0.6 (-3.7, 4.9)	109.8 (-407.9, 627.5)
	Phase 2	-1.0 (-7.3, 5.3)	-0.2 (-0.9, 0.4)	0.3 (-0.6, 1.3)	-0.3 (-3.6, 3.0)	-210.0 (-590.1, 170.2)
Fruit and/or vegetable goal ^b	Phase 1	6.9 (1.2, 12.7)*	0.5 (-0.1, 1.1)	1.1 (0.0, 2.3)*	4.2 (0.2, 8.2)*	-155.6 (-659.9, 348.6)
	Phase 2	-2.0 (-9.0, 5.0)	0.0 (-0.6, 0.7)	-0.5 (-1.5, 0.6)	0.4 (-3.1, 4.0)	413.7 (28.0, 799.4)*
No goal ^c	Phase 1	-4.5 (-13.4, 4.3)	-0.5 (-1.4, 0.5)	-0.7 (-2.4, 1.1)	-4.0 (-10.2, 2.1)	115.2 (-643.6, 874.0)
	Phase 2	1.1 (-7.4, 9.6)	0.0 (-0.8, 0.8)	0.2 (-1.1, 1.5)	-0.0 (-4.4, 4.3)	309.1 (-191.7, 810.0)

Data are reported as mean (95% CI).

^aPhase 1: "yes" *n* = 15; "no" *n* = 26. Phase 2: "yes" *n* = 21; "no" *n* = 20.

^bPhase 1: "yes" *n* = 16; "no" *n* = 25. Phase 2: "yes" *n* = 13; "no" *n* = 28.

^cPhase 1: "yes" *n* = 5; "no" *n* = 36. Phase 2: "yes" *n* = 7; "no" *n* = 34.

#Standard ANCOVA analysis reporting the mean (95% CI) change in the group of participants who set each specific goal, minus the change in participants who did not set each specific goal.

**p* < 0.05.

who ranked in the intermediate diet quality category (22). While the overall results of this exploratory study are far from conclusive, the AHEI improvement observed in phase 1 (+7 points; 10% improvement) is promising, particularly given the fact that our participants' baseline diet already ranked in the "high" diet quality category (22, 23). Similarly, with the noted clinical meaningful changes in fruit (0.5 serves) and vegetable intake (one serve), this placed participants mean intakes at two and five serves, respectively, in line with the Australian Dietary Guidelines (19). The fact that sodium intake increased by ~400 mg at 6 months is interesting and unexpected. It is important to note that at 3 months, sodium intake decreased by a non-significant 150 mg, in line with the significant improvement in diet quality. While we are not able to determine the exact reason why this occurred, the 6-month period of the study was typically (for the vast majority of participants) occurring over the winter and spring seasons in Australia. These seasons present well-known challenges for people with CKD in controlling their sodium intake, typically due to choosing more tinned vegetables and prepared soups (28). This also highlights the challenge that some participants had in maintaining their dietary changes after the initial 3-month coaching period, where they could not speak to a dietitian to discuss these potential issues. Future telehealth coaching studies should ensure specific education content (both telephone coaching, but likely more importantly, extended text message support), which considers assisting patients choose low-sodium pre-prepared foods and provide seasonal recipes.

Fruits and vegetables have many properties, which might make them an appealing intervention strategy for people with CKD trying to improve their diet quality. Fruits and vegetables release potassium salts, which generate bicarbonate naturally, and can decrease the kidney acid load. Current best-practice guidelines recommend that people with stages 1–4 CKD consume a diet higher in fruits and vegetables to effectively manage metabolic acidosis (4). Fruits and vegetables also have a low bioavailability of dietary phosphorus and calcium, which limits its bioavailability due to the presence of phytate in vegetable forms of phosphorous,

which can also promote better adherence to nutrient-restricted interventions (29).

Adequate daily intakes of fruits, vegetables, and fiber reflect a higher overall diet quality (30). Evidence shows that sufficient fiber intake reduces the risk of inflammatory reactions, which are directly linked with the progression of kidney diseases (31). Moreover, fiber is found to be related to inflammation metabolic acidosis and gut microbiota culture, which are also associated with CKD (32). Increasing dietary fiber intake may also attenuate imbalance in the gut microbiome, which is known to have a role in modification of protein-bound uremic toxins (33). A meta-analysis of 14 CKD controlled-feeding trials showed dietary fiber to significantly reduce serum urea and creatinine levels (34). As the most common dietary sources of fiber, increasing fruit and vegetable consumptions are likely to be the most practical way in which CKD patients can improve their fiber intake and overall diet quality.

No significant AHEI improvement was found at 6 months compared with the first 3 months of the study. However, this result is not conclusive due to the low participants' response rate to goal check text messages (<50% in phase 1 and approximately one third in phase 2). While it is critical to acknowledge the message response rate, it may also be important to consider the results in the context of the differences in intervention approaches used in each phase. Specifically, in phase 1, participants received coaching contact from the dietitian with six fortnightly telephone coaching and weekly tailored text messages for the first 3 months of the program, while in the second phase of the study, participants continued receiving the same tailored text messages but received no further telephone coaching with their dietitian. It was previously reported that previous ENTICE-CKD participants needed at least two coaching calls to start putting their set action plans in place (18). According to the achievement motivation theory, people tend to continue making positive changes once they have attained their set goals (35, 36). It remains unclear whether intervention decay observed in our study is commensurate with a reduction in coaching intensity. It is likely that more intensive support over a longer duration from a dietitian is needed to promote greater change (37).

An overview of the systematic reviews evaluating intervention components associated with increased effectiveness in dietary generally supports this theory, of a positive relationship between the number of intervention contacts over longer durations and self-reported dietary change (38). Whether or not this is the case specifically in CKD populations requires further investigation.

Furthermore, there were differences in the number of participants who set fruit and vegetable intake goals in each phase of the study. Specifically, in phase 2, less participants chose to continue their goal to improve their fruit and vegetable intake ($n = 16$ in phase 1 compared with $n = 13$ in phase 2). Instead, there appeared to be an increase in the number of participants who set a healthy eating goal in phase 2 compared with that in phase 1 ($n = 15$ and $n = 21$, respectively). This might indicate that participants felt that they had successfully achieved their goal to improve fruit and vegetable intake at 3 months and decided to set a more general healthy eating goal in phase 2. As mentioned above, it is likely that people with CKD still need more coaching over a longer period as mentioned. However, it could also be hypothesized that people with CKD may need more specific goals, such as a fruit or vegetable intake goals, compared with a general healthy eating goal, in order to achieve true change in diet quality. This hypothesis needs testing in future adequately powered randomized controlled trials.

This study has important limitations to consider. This is a *post-hoc* secondary analysis, which only evaluated a single arm of the previous ENTICE-CKD trial with no control group comparison. The findings in this study are therefore exploratory only, hypothesis-generating, and not conclusive. The small sample size is a factor influencing the accuracy of results. Participants showed diverse preferences for dietary goals throughout the previous ENTICE-CKD program, in line with the practical and patient-centered approach the intervention was underpinned by. While most participants determined that “healthy eating” and “fruit and/or vegetable” goals were important to them ($n = 13$ and $n = 21$, respectively), very few participants set goals such as a “weight reduction” goal ($n = 2$), and these goals could not be analyzed in a reliable or meaningful way. This limited our ability to explore the effect of specific goal setting on improving clinical outcomes. Participants showed a “high” diet quality score at baseline indicating healthy volunteer bias; this limitation may underestimate the potential benefits of dietary intervention evaluated in this study. To manage the issue of missing data, the principle of BOCF was applied to impute missing data, which may underestimate the treatment effect, or may ignore the possibility that people could regress or even worsen from baseline. Finally, while this is a patient-centered approach, our study relies almost entirely on self-reported data, which may reduce the confidence in the conclusions inferred.

In conclusion, patient-centered goal setting, which focuses on fruit and vegetable intake appeared to significantly improve diet quality, fruit and vegetable intake, and fiber intake in people with

stages 3–4 CKD. No significant results were found at 6 months, which may be due to the reduction of coaching intensity and suggests that a longer-term telehealth coaching may be required in helping CKD patients to achieve long-term self-management. Further research evaluating longer-term interventions, with an emphasis on goal setting particularly around fruit and vegetable intake, will determine what effect on clinical outcomes are possible for people with CKD.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, upon reasonable request and clear and robust scientific protocol provided, without undue reservation.

ETHICS STATEMENT

The research protocol was approved by Metro South Hospital and Health Service and Bond University Human Research Ethics Committees. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

CC completed the data analysis and wrote the first draft of the manuscript. JK is the supervisor of this study who contributed to the study conception, original research data collection, reviewed the analysis results, and critically reviewed versions of the manuscript. MC participated in the original data collection and intervention design. MR and KC contributed to the study conception and critically reviewed the manuscript. All authors critically reviewed the manuscript and approved the final version of the manuscript.

FUNDING

This study was supported by a research support grant awarded by Kidney Health Australia via Australasian Kidney Trial Network (AKTN) and a Vice-Chancellors Research Award through Bond University.

ACKNOWLEDGMENTS

The authors would like to thank Ms. Evelyne Rathbone for her advice on statistical analysis. JK was supported through a Griffith University Postdoctoral Research Fellowship.

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fnut.2021.627753/full#supplementary-material>

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Ultrasound for Non-invasive Assessment and Monitoring of Quadriceps Muscle Thickness in Critically Ill Patients With Acute Kidney Injury

Alice Sabatino^{1,2*}, Umberto Maggiore^{1,2}, Giuseppe Regolisti^{1,2}, Giovanni Maria Rossi^{1,2}, Francesca Di Mario^{1,2}, Micaela Gentile^{1,2}, Maria Teresa Farina^{1,2} and Enrico Fiaccadori^{1,2}

¹ UO Nefrologia, Azienda Ospedaliera - Universitaria Parma, Parma, Italy, ² Dipartimento di Medicina e Chirurgia, Università di Parma, Parma, Italy

OPEN ACCESS

Edited by:

Alessio Molfino,
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Reviewed by:

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Universidade Federal de Lavras, Brazil

*Correspondence:

Alice Sabatino
alice.sabatino86@gmail.com

Specialty section:

This article was submitted to
Clinical Nutrition,
a section of the journal
Frontiers in Nutrition

Received: 29 October 2020

Accepted: 17 March 2021

Published: 15 April 2021

Citation:

Sabatino A, Maggiore U, Regolisti G,
Rossi GM, Di Mario F, Gentile M,
Farina MT and Fiaccadori E (2021)
Ultrasound for Non-invasive
Assessment and Monitoring of
Quadriceps Muscle Thickness in
Critically Ill Patients With Acute Kidney
Injury. *Front. Nutr.* 8:622823.
doi: 10.3389/fnut.2021.622823

Background and aims: Critically ill patients with acute kidney injury (AKI) undergo major muscle wasting in the first few days of ICU stay. An important concern in this clinical setting is the lack of adequate tools for routine bedside evaluation of the skeletal muscle mass, both for the determination of nutritional status at admission, and for monitoring. In this regard, the present study aims to ascertain if ultrasound (US) is able to detect changes in quadriceps muscle thickness of critically ill patients with acute kidney injury (AKI) over short periods of time.

Methods: This is a prospective observational study with a follow-up at 5 days. All adult patients with AKI hospitalized at the Renal ICU of the Parma University Hospital over 12 months, with a hospital stay before ICU admission no longer than 72 h, and with a planned ICU stay of at least 5 days, were eligible for the study. An experienced investigator assessed quadriceps rectus femoris and vastus intermedius thickness (QRFT and QVIT) at baseline and after 5 days of ICU stay.

Results: We enrolled 30 patients with 74 ± 11 years of age and APACHE II score of 22 ± 5 . Muscle thickness decreased by $15 \pm 13\%$ within the first 5 days of ICU stay ($P < 0.001$ for all sites as compared to ICU admission). Patients with more severe muscle loss had lower probability of being discharged home (OR: 0.04, 95%CI: 0.00–0.74; $P = 0.031$).

Conclusions: In critically ill patients with AKI, bedside muscle US identifies patients with accelerated muscle wasting.

Keywords: acute kidney injury, body composition, critical care, muscle wasting, muscle ultrasound, intensive care unit

INTRODUCTION

Critically ill patients undergo major muscle wasting in the first few days of their ICU stay (1). Clinical consequences are represented by delayed functional recovery, difficult weaning from mechanical ventilation and increased mortality risk (1).

In this clinical setting an important cause of concern is the lack of adequate tools for routine bedside evaluation of the skeletal muscle mass (2). The reference methods considered as the gold

standard for the assessment of skeletal muscle, such as computed tomography (CT), magnetic resonance imaging (MRI), and dual energy X-ray absorptiometry (DEXA) are not feasible for routine evaluation and monitoring of muscle mass and body composition (2). On the other hand, currently used bedside tools, such as bioimpedance analysis (BIA) and anthropometry, are not accurate enough in critically ill patients (2), mainly due to the possible interference of fluid overload, frequently observed especially when Acute Kidney Injury (AKI) coexists. Recently, the use of ultrasound (US) for the assessment of muscle dimensions has aroused considerable interest, and its reliability and validity have been documented also in critically ill patients with AKI (3, 4). US technique seems to be poorly influenced not only by fluid overload, but also by the rapid and relevant fluid shifts typical of patients with AKI undergoing Kidney Replacement Therapy (KRT) (3).

On this premise, in the present study we aimed to assess whether US is able to detect changes in muscle thickness of patients with AKI over a short period of time.

MATERIALS AND METHODS

Patients

This is a prospective, longitudinal (5 days) observational study, conducted in the Renal ICU of the Parma University Hospital. Procedures were held in accordance to the Helsinki declaration and informed consent was obtained from patients or their next of kin. The study was approved by the local ethics committee (Comitato Etico di Area Vasta Emilia Nord, AVEN, Prot n. 43943 –03/12/2015).

All adult patients with AKI hospitalized in the Renal ICU from 15/03/2017 to 15/03/2018, with a hospital stay before ICU admission no longer than 72 hours, and with a predictable ICU stay of at least 5 days were eligible for the study. AKI was diagnosed according to KDIGO guideline criteria (5).

Already available data on quadriceps femoris US evaluation in healthy subjects (body mass index (BMI) > 18.5 Kg/m², Subjective Global Assessment (SGA) class A, absence of chronic or acute illness) (6) were used for comparison with AKI patients, both at ICU admission and after 5 days of ICU stay.

Methods

US Technique

The same experienced investigator (renal dietitian) performed all of the measurements. Quadriceps rectus femoris thickness (QRFT) and quadriceps vastus intermedius thickness (QVIT) were measured by B-mode ultrasonography, wall tracking ultrasound system (Philips hd7xe) with a 7.5 MHz linear array transducer (L12-3 transducer), as previously described in detail (3). All US measurements were performed in duplicate and the average of the scores used in final analyses. The transducer was placed perpendicular to the long axis of the thigh with a large amount of gel and no pressure to avoid compression of the muscle. QRFT and QVIT were measured at the midpoint (RF Prox; VI Prox) and at the border between the lower third and upper two-thirds (RF Dist; Prox Dist) between the anterior superior iliac spine (ASIS) and the upper pole of the patella

(3, 7). The right and left quadriceps values were assessed in both legs with the patient lying in a supine position with both knees extended but relaxed and toes pointing to the ceiling. The assessor was positioned on the side of the patient while performing the measurements, and was allowed to tilt the probe to obtain the best possible image, in which RF and VI would be aligned and centered. Measurements were performed directly on the US machine while obtaining the images. The vertical diameter of the muscles was measured on the inner edge of the muscle fascia. US was performed twice during ICU stay, at baseline (at ICU admission) and after 5 days since the first measurement. Muscle US took <20 min to complete the image acquisition and perform the measurements.

Demographics, clinical data, renal function and outcome: data were collected as per institutional routine at the time of ICU admission and during ICU stay, with special regard to demographic, body weight and height, clinical and laboratory data, renal function, acute and chronic comorbidities, severity of illness (APACHE II score), data on renal replacement therapy (RRT), length of stay and mortality.

Outcomes: muscle loss after 5 days.

Statistical Analysis

Results are expressed as mean and standard deviation for continuous variables with normal distribution, or median and range for non-parametric data, and as frequencies for categorical variables. Group differences were analyzed using Student *t*-test and Mann-Whitney's *U*-test for parametric and non-parametric data, respectively to assess difference between means of the control group and the patient group. ANCOVA was used to adjust the analysis by age and sex.

We examined the difference between muscle thickness at baseline and at 5 days after admission by mixed-effects models with patients fitted as random effects, and the four-way interaction term between time and each of the three sites of measurements (RF vs. VI, Left vs. Right, Proximal vs. Distal) fitted as fixed effects. We examined the relation between baseline comorbidities and change in muscle thickness by mixed-effects ANCOVA models in which muscle thickness at 5 days after admission was included as dependent variate and baseline thickness was included as covariate, in order to adjust for the correlation between change in muscle thickness and random differences in baseline values. We examined the relation between change in muscle thickness and ICU outcome (discharge, transfer to other health care facility, death) in two steps. First, we estimated the individual change over time in muscle thickness by the best linear unbiased predictions (BLUPs) of the random slope from mixed-effects random coefficients models. Then, we fitted a multinomial logistic regression model where outcome (discharge, prolonged stay, death) was the dependent variable and the individual random slope the independent variable. Because of sparse data concerning mortality (five patients only) we did not report the findings on mortality. A two-sided *P* value of less than 0.05 was regarded as statistically significant. Stata Release 16 (StataCorp, College Station, TX, US) was used for all the analyses.

TABLE 1 | Demographic and clinical data.

Variables	Patients (n = 30)	Healthy subjects (n = 35)
Age	74 (10.6)	41 (10.0)*
Male sex (n, %)	21/30 (70)	15/35 (43)*
Body weight (Kg)	82 (13.2)	70.5 (16.6)*
Height (m)	1.67 (0.09)	1.70 (0.09)
BMI (Kg/m ²)	29 (4.6)	24.3 (4.6)*
APACHE II	22 (5)	NA
Main admission diagnosis (n, %)		
— Renal	18/30 (60)	NA
— Sepsis	4/30 (14)	NA
— Respiratory	3/30 (10)	NA
— Vascular	3/30 (10)	NA
— Malignancy	1/30 (3)	NA
— Cardiac	1/30 (3)	NA
Surgical status (n, %)		
— Urgent	2/30 (7)	NA
— Programmed	3/30 (10)	NA
— Non-surgical	25/30 (83)	NA
Chronic comorbidities (n, %)		
— Hypertension	23/30 (77)	NA
— Diabetes mellitus	11/30 (37)	NA
— COPD	7/30 (23)	NA
— Ischemic cardiopathy	7/30 (23)	NA
— Heart failure	8/30 (27)	NA
— Peripheral vascular disease	5/30 (17)	NA
— Immunocompromised	2/30 (7)	NA
— Chronic liver disease	2/30 (7)	NA
— Malignancy	6/30 (20)	NA
— Chronic kidney disease (not on dialysis)	11/30 (37)	NA
Acute complications at first muscle US (n, %)		
— Sepsis	12/30 (40)	NA
— Invasive mechanical ventilation	4/30 (13)	NA
— Non-invasive mechanical ventilation	7/30 (23)	NA
— Oliguria	20/30 (67)	NA
— Vasoactive drug need	7/30 (23)	NA
— Renal replacement therapy	21/30 (70)	NA
ICU outcome (n, %)		
— Death	5/30 (17)	NA
Hospital outcome (n, %)		
— Death	9/30 (30)	NA
— Discharged home	15/30 (50)	NA
— Transferred to long-stay/rehabilitation ward	2/30 (7)	NA
— Transferred to another hospital	4/30 (13)	NA
ICU LOS median, range)	15 (4–72)	NA
Hospital LOS (median, range)	34 (7–138)	NA
Biochemical data		
sCr (mg/dl)	6.3 (4)	NA
BUN (mg/dl)	81.7 (36)	NA
Albumin (g/dl)	2.8 (0.6)	NA
CRP (mg/dl)	109.2 (68.1)	NA

Data expressed as mean (standard deviation), frequencies and median (range). * $P < 0.001$ in comparison to AKI patients. BMI, body mass index; BUN, blood urea nitrogen; COPD, chronic obstructive pulmonary disease; CRP, C-reactive protein; ICU, intensive care unit; IMV, invasive mechanical ventilation; LOS, length of stay; NIMV, non-invasive mechanical ventilation; sCr, serum creatinine.

Sample Size Calculations

No data are currently available in the literature on US evaluation and monitoring of quadriceps muscle mass in patients with AKI. In a recent study on ICU patients (8) 22 patients were enrolled in order to detect a 16% reduction in the quadriceps rectus femoris thickness after 5 days of ICU stay, with a power of 80% and a probability of type I error equal to 0.05. We enrolled 30 patients to account for possible drop-outs.

RESULTS

Table 1 shows the baseline characteristics of the 30 patients studied. Seventy percent (21/30) were male with a mean \pm SD age of 74 ± 11 years, and they represented a severely critically ill cohort (APACHE II was 22 ± 5). A total of 472 images were analyzed across the 30 patients. Eighty-three percent of patients (25/30) were non-surgical patients and the main admission diagnosis in the ICU was renal, followed by sepsis. On average, patients were polymorbid (2.8 ± 1.7 comorbidities per patient), hypertension being the most frequent comorbidity. As to the usual renal function, 37% (11/30) had basal eGFR values <60 ml/min/1.73 m² (CKD stages 2 to 5 non dialysis). At the time of first US evaluation, all of the patients had stage 3 AKI; in 21/30 patients (70%) RRT was started as 10–12 h lasting sustained low-efficiency dialysis. Oliguria was common (67%), as was sepsis (40%). ICU mortality was 17% (5/30); hospital mortality was 30% (9/30). The median (range) length of ICU stay was 15 (4–72) days, while the length of hospital stay was 34 (7–138). C-reactive protein was 109.2 mg/dL (± 68.1).

Patients With AKI in Comparison to Healthy Subjects

Demographic characteristics of control group (35 healthy subjects) are shown in **Table 1**, while US quadriceps muscle data are illustrated in **Figure 1**. In general, the control group was younger and leaner than patients. At univariate analysis (**Figure 1**), muscle thickness of patients differed from that of the control group for all sites, both at T1 and T2. We also performed an adjusted analysis using ANCOVA corrected for age and sex. In the adjusted analysis, no difference was found between T1 values of muscle thickness and control group values; however, the difference between T2 values and the control group values remained statistically significant.

Changes in Muscle Thickness by US

Baseline (T1) and after 5 days (T2) mean \pm SD values for RF and VI thickness are illustrated in **Figure 1**; the difference between means was statistically significant for all sites ($P < 0.001$). On average, there was a mean clinically relevant reduction of 15% ($\pm 12\%$) in every site of measurement within the first 5 days of ICU stay (**Table 2**). Changes in VI Prox and Dist thickness (16–19%) were greater than in RF Prox and Dist thickness (11–13%).

We did not find any association between baseline muscle thickness, inflammatory status (assessed by CRP and serum albumin), as well as chronic and acute comorbidities, with muscle loss (**Supplementary Table 1**). Patients with the more severe muscle loss had a higher probability of having a prolonged

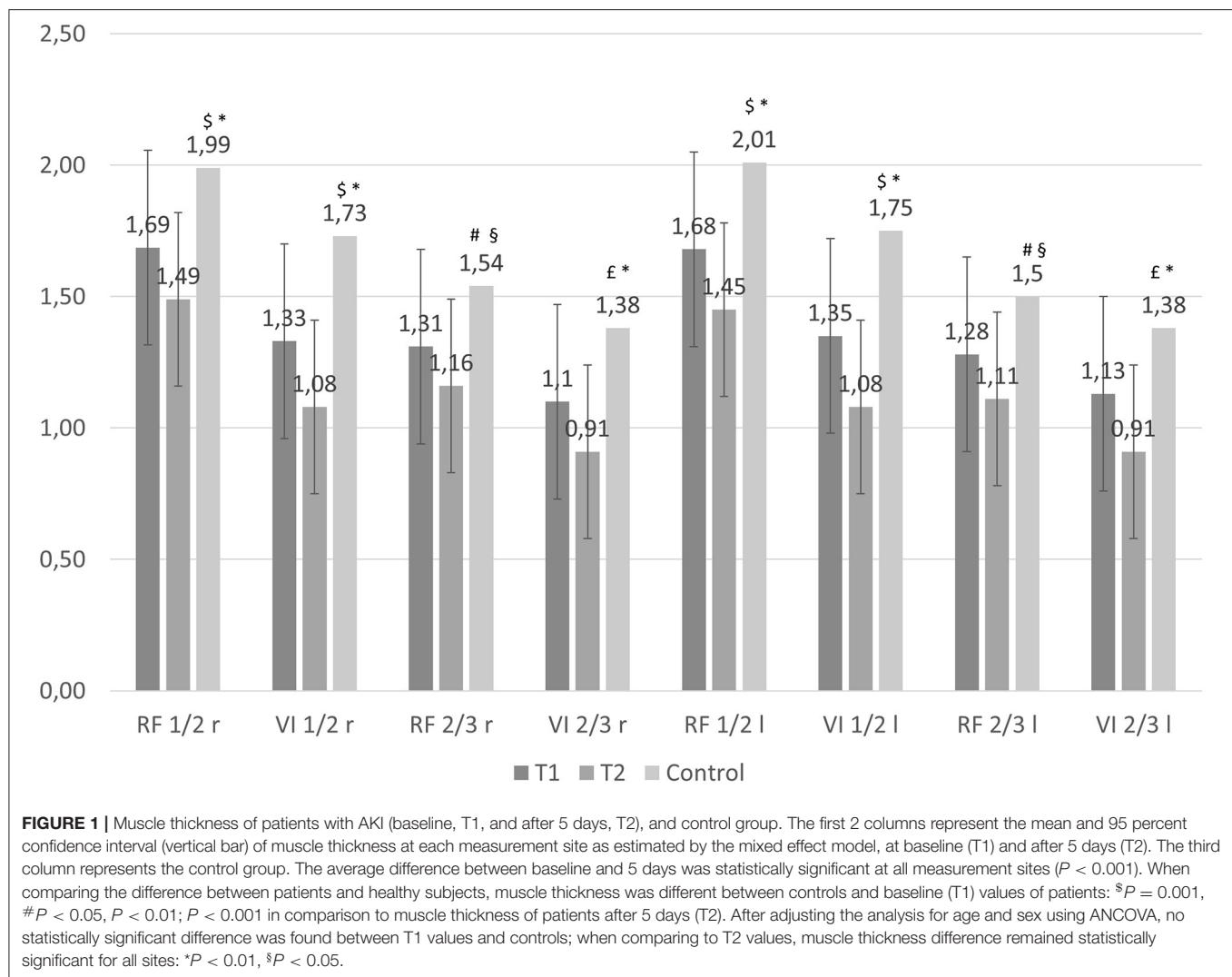


TABLE 2 | Average percent reduction of muscle thickness after 5 days of ICU stay.

Muscle site	Average reduction (SD)
QRFT Prox r ($n = 29$)	11% (8%)
QVIT Prox r ($n = 28$)	18% (14%)
QRFT Dist r ($n = 30$)	11% (10%)
QVIT Dist r ($n = 30$)	17% (12%)
QRFT Prox l ($n = 30$)	13% (10%)
QVIT Prox l ($n = 29$)	19% (16%)
QRFT Dist l ($n = 30$)	12% (10%)
QVIT Dist l ($n = 30$)	16% (15%)
All measurements ($n = 236$)	15% (12%)

Dist, distal; Prox, proximal; QRFT, quadriceps rectus femoris thickness; QVIT, quadriceps vastus intermedius thickness; Values expressed as mean (standard deviation).

hospital stay, which was determined by being transferred to another hospital instead of discharged home (OR: 0.04, 95%CI: 0.00- 0.74; $P = 0.031$) (Figure 2).

DISCUSSION

Our study confirms that muscle loss may occur early and rapidly in the first 5 days of ICU stay, and extends previous observations in general ICU patients also to patients with AKI.

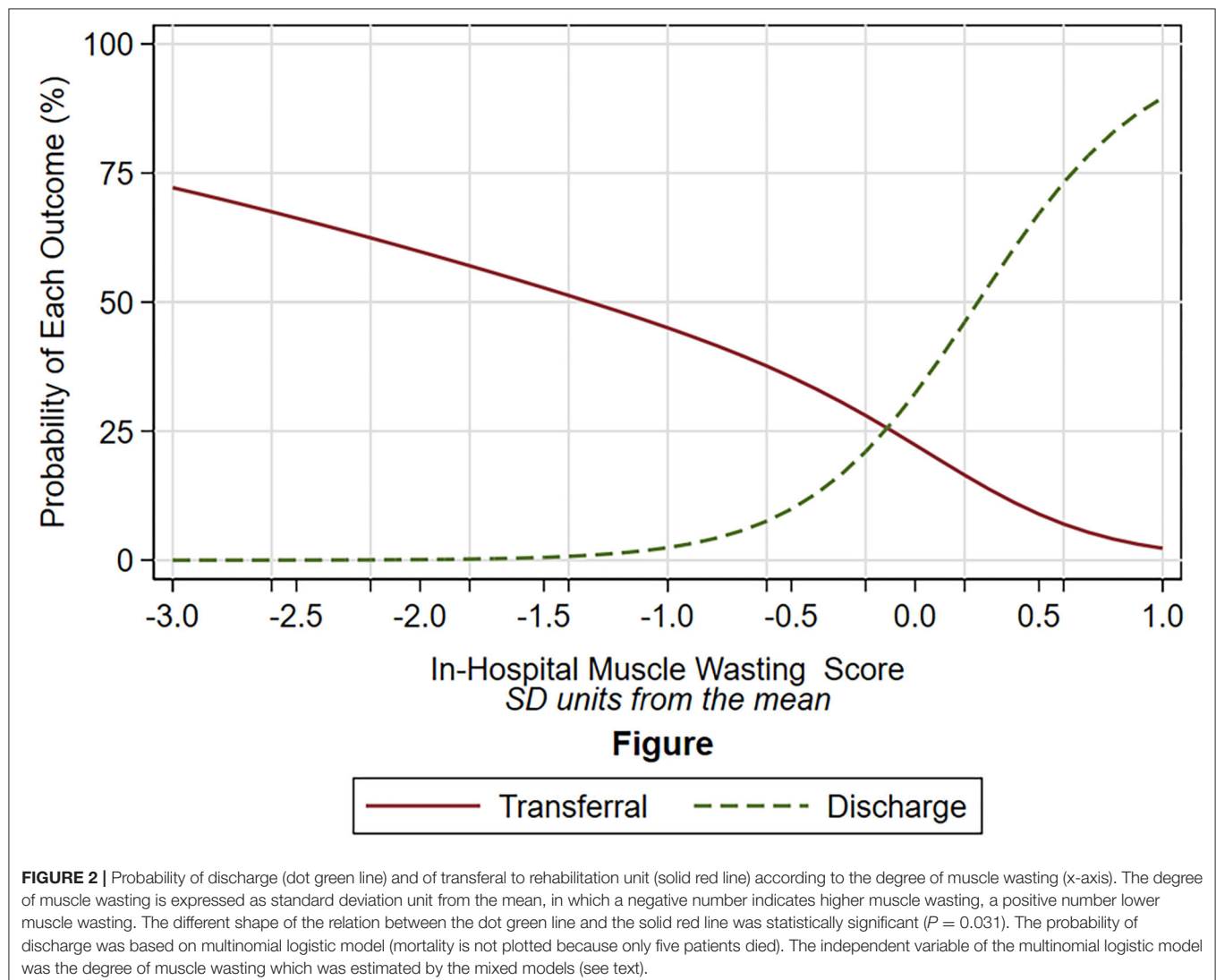
At baseline, values of muscle thickness were not different from those of younger healthy subjects when the analysis were adjusted for age and sex. They were also similar to values previously reported in another study assessing elderly healthy subjects (9). However, at day 5, muscle mass was significantly reduced, both in comparison to baseline values and in comparison to the control group.

In our study, the amount of muscle loss was similar to that observed in critically ill patients (8); in particular, at day 5, the VI muscle had the most important reduction in comparison to the RF muscle. Rectus femoris is often described as a power muscle designed to assist in fast movements, while VI is considered a stabilizing muscle that is important for maintaining posture. The identification of which muscles are more affected by immobilization and critical illness in ICU patients could

provide important indications in order to more precisely target rehabilitation according to the type of muscle predominantly affected (VI-postural or RF-power).

We demonstrated that an inverse relationship exists between the severity of muscle loss and the probability of discharge home. Although we can't ascertain the causal relationship between muscle loss and outcome due to the small sample size, baseline sarcopenia assessed by US predicts adverse discharge disposition (death/transferral to nursing facilities) (10). In addition, muscle loss during ICU stay is a major contributor to functional disability (11). However, despite the rapidly loss faced during the first week of ICU stay, it remains unclear whether it is the change in muscle size from baseline or the total amount of muscle mass at admission that is the most important predictor of functional outcome and mortality (12). In our study, baseline muscle thickness did not correlate to the amount of muscle loss, which is in accordance to previous studies investigating muscle wasting in critically ill patients (11).

Despite its increasingly recognized importance, the assessment of muscle mass is challenging in the ICU setting, especially when inflammation and fluid imbalance are present. Currently available bedside methods (such as anthropometry and bioimpedance spectroscopy [BIS]) have important intrinsic limitations, due for example to fluid shifts typical of critically ill patients with AKI (13). Current recommended reference methods are hardly feasible in this clinical setting. Dual energy X-Ray absorptiometry (DEXA), which has been used and recommended by recent Consensus on sarcopenia to assess appendicular skeletal muscle mass (14), is also influenced by hydration status, because it assumes that lean body mass has a constant hydration of 73%, which is not the case in critically ill patients with AKI (15). In addition, DEXA is not feasible at the bedside and involves patient radiation exposure. Computed tomography (CT) has been recently used to assess muscle mass and its correlation with mortality in critically ill patients (16). Despite its excellent accuracy, CT is expensive,



requires specialized personnel, and is available for muscle mass assessment only when CT is necessary for other diagnostic procedures on the lung or the abdomen. In recent years, muscle US has been increasingly studied in the kidney patient setting. Specifically, the reliability of the muscle US technique applied in the present study has been already reported in critically ill patients with AKI, along with excellent intraclass correlation coefficient (ICC) for inter and intra-operator comparisons (3). This methodology has already been validated in patients with AKI against muscle CT (4). Despite the lack of reference values to be applied at baseline and identify patients with pre-ICU low muscle mass and, therefore, increased nutritional risk, in the present study we showed that quadriceps muscle US is useful to monitor nutritional status of critically ill patients with AKI, and that increased muscle loss reduces the chance of being discharged home.

The main limitation of the present study relies in the small sample size, that does not allow further analysis regarding the effect of muscle loss and baseline muscle assets on mortality or functional outcomes. Further studies will be needed on larger cohorts of patients with AKI to allow for such analyses.

In conclusion, muscle wasting occurs early and rapidly within the first 5 days of ICU stay in critically ill patients with AKI. Muscle US is a sensible and feasible tool for the detection of muscle wasting even in this clinical setting; moreover, it is easy to use, cheap and time-efficient. In the future, studies defining cut-off values for muscularity are needed to allow the early identification of patients with low muscle mass at ICU admission.

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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 Comitato Etico di Area Vasta Emilia Nord, AVEN, Prot n. 43943 –03/12/2015. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

AS: investigation, methodology, validation, and writing—original draft. UM: formal analysis, writing—review and editing. GR, GMR, MG, and MF: writing—review and editing. FD: resources, writing—review and editing. EF: supervision, writing—review and editing. All authors contributed to the article and approved the submitted version.

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fnut.2021.622823/full#supplementary-material>

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Performance of Bioelectrical Impedance and Anthropometric Predictive Equations for Estimation of Muscle Mass in Chronic Kidney Disease Patients

Natália Tomborelli Bellafronte^{1*}, Lorena Vega-Piris², Guillermina Barril Cuadrado³ and Paula Garcia Chiarello⁴

¹ Post-graduate Program in Health Sciences, Ribeirão Preto Faculty of Medicine, University of São Paulo, Ribeirão Preto, Brazil, ² Methodology Unit, Instituto de Investigación Sanitaria del Hospital Universitario de la Princesa, Madrid, Spain, ³ Nephrology Department, Hospital Universitario de la Princesa, Madrid, Spain, ⁴ Department of Health Sciences, Ribeirão Preto Faculty of Medicine, University of São Paulo, Ribeirão Preto, Brazil

OPEN ACCESS

Edited by:

Alice Sabatino,
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Reviewed by:

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São Paulo State University, Brazil

*Correspondence:

Natália Tomborelli Bellafronte
natalia.bellafronte@usp.br

Specialty section:

This article was submitted to
Clinical Nutrition,
a section of the journal
Frontiers in Nutrition

Received: 20 March 2021

Accepted: 26 April 2021

Published: 21 May 2021

Citation:

Bellafronte NT, Vega-Piris L, Cuadrado GB and Chiarello PG (2021) Performance of Bioelectrical Impedance and Anthropometric Predictive Equations for Estimation of Muscle Mass in Chronic Kidney Disease Patients. *Front. Nutr.* 8:683393. doi: 10.3389/fnut.2021.683393

Background: Patients with chronic kidney disease (CKD) are vulnerable to loss of muscle mass due to several metabolic alterations derived from the uremic syndrome. Reference methods for body composition evaluation are usually unfeasible in clinical settings.

Aims: To evaluate the accuracy of predictive equations based on bioelectrical impedance analyses (BIA) and anthropometry parameters for estimating fat free mass (FFM) and appendicular FFM (AFFM), compared to dual energy X-ray absorptiometry (DXA), in CKD patients.

Methods: We performed a longitudinal study with patients in non-dialysis-dependent, hemodialysis, peritoneal dialysis and kidney transplant treatment. FFM and AFFM were evaluated by DXA, BIA (Sergi, Kyle, Janssen and MacDonald equations) and anthropometry (Hume, Lee, Tian, and Noori equations). Low muscle mass was diagnosed by DXA analysis. Intra-class correlation coefficient (ICC), Bland-Altman graphic and multiple regression analysis were used to evaluate equation accuracy, linear regression analysis to evaluate bias, and ROC curve analysis and kappa for reproducibility.

Results: In total sample and in each CKD group, the predictive equation with the best accuracy was AFFM_{Sergi} (men, $n = 137$: ICC = 0.91, 95% CI = 0.79–0.96, bias = 1.11 kg; women, $n = 129$: ICC = 0.94, 95% CI = 0.92–0.96, bias = –0.28 kg). AFFM_{Sergi} also presented the best performance for low muscle mass diagnosis (men, kappa = 0.68, AUC = 0.83; women, kappa = 0.65, AUC = 0.85). Bias between AFFM_{Sergi} and AFFM_{DXA} was mainly affected by total body water and fat mass. None of the predictive equations was able to accurately predict changes in AFFM and FFM, with all ICC lower than 0.5.

Conclusion: The predictive equation with the best performance to assess muscle mass in CKD patients was AFFM_{Sergi}, including evaluation of low muscle mass

diagnosis. However, assessment of changes in body composition was biased, mainly due to variations in fluid status together with adiposity, limiting its applicability for longitudinal evaluations.

Keywords: anthropometry, bioelectrical impedance, body composition, chronic kidney disease, dual energy X-ray absorptiometry, fat free mass, lean mass, sarcopenia

INTRODUCTION

Lean body mass reserve, whose major component is skeletal muscle, is an essential reserve that provides amino acids to support processes such as injury repair and the immune response (1, 2). Therefore, lean body mass plays an important role in clinical outcomes and disease progress, with low lean body mass related to worse prognosis and shorter survival (3). Appendicular lean mass, which encompasses the lean soft tissue in the limbs and is mainly composed of skeletal muscle mass, is the variable of choice for low muscle mass diagnosis (4) and is considered as a key parameter for nutritional status evaluation.

Body composition technologies such as computed tomography, magnetic resonance imaging and dual energy X-ray absorptiometry (DXA) provide objective information about skeletal muscle mass (5). DXA, the most available one of them, has become recognized for its ability to accurately and precisely measure total body composition (5), in a three compartment level (fat mass, lean soft tissue and bone mineral content) (6). However, DXA is not a bedside technique, requires patient transportation to the instrument and has high cost, thus hampering its use in routine practice.

Given the unfeasibility to apply reference methods in clinical settings, there is a growing interest in more suitable techniques for body composition evaluation, such as anthropometry and bioelectrical impedance analysis (BIA). Equations using BIA parameters have been validated to predict fat free mass (FFM, lean mass + bone mineral content) in healthy individuals (7). Methods for prediction of appendicular fat free mass (AFFM, FFM of the limbs) from BIA were also developed in healthy elderly subjects (8), in healthy adults with validation in heart, lung and liver transplant patients (9); and also in non-dialysis-dependent (NDD) chronic kidney disease (CKD) patients (10). Furthermore, methods to predict FFM have been developed from anthropometric measures in healthy (11) and non-obese adults (12), as well as in NDD (13) and hemodialysis (HD) (14) CKD patients.

Patients with CKD are vulnerable to loss of muscle mass due to several metabolic alterations derived from the uremic syndrome (15–18). The metabolic disorders already present in NDD patients (16) become more evident in more advanced stages

of CKD when peritoneal dialysis (PD) and HD treatments are established (3). Even after kidney transplant (KTx) nutritional status is difficult to recover (18). Accordingly, body composition of CKD patients worsens with disease progression (19), with lean tissue loss, sometimes masked by edema and usually preceding weight loss. As body composition is a biomarker for prognosis and helps to monitor clinical interventions, its assessment needs to be part of CKD routine care, although it is often imprecise in clinical settings.

The routine use of simplified methods such as BIA and anthropometry in hospital settings may improve the evaluation of nutritional status allowing clinicians to identify individuals who would benefit most from targeted interventions, as well as to reliably monitor longitudinal changes. Therefore, the primary aim of this study was to assess the accuracy of equations using BIA and anthropometry measures, compared to DXA, to estimate FFM and AFFM for cross-sectional and longitudinal assessment, in NDD, HD, PD, and KTx CKD patients. Secondary, we evaluated the capacity of surrogate methods to diagnose low muscle mass.

MATERIALS AND METHODS

This was a longitudinal study that evaluated clinically stable NDD, HD, PD, and KTx CKD patients from a nephrology outpatient clinic at Ribeirão Preto Medical School University Hospital (University of São Paulo, São Paulo, Brazil) and at a specialized dialysis clinic, the Nephrology Service of Ribeirão Preto. Patients with CKD were enrolled between 2017 and 2019. Inclusion criteria were: age ≥ 18 and ≤ 60 years and under regular treatment for at least 6 months; for NDD patients, estimated glomerular filtration rate (eGFR) ≤ 44 ml/min; for PD patients, absence of peritonitis in the previous 30 days and dialysis for at least 3 months; for HD patients, 4-h dialysis session, three times per week, through an arteriovenous fistula and dialysis for at least 3 months; for KTx patients, transplant for at least 6 months and eGFR ≥ 45 ml/min.

Exclusion criteria were presence of malignant diseases, acute infections and inflammation, human immunodeficiency virus, chronic lung disease, liver and heart failure, pregnancy or lactation, having amputations or an electronic implant, wheelchair user or inpatient, body weight >140 kg or BMI >40 kg/m².

This study was conducted according to the principles of the Declaration of Helsinki, the protocol was approved by the human research ethics committee of Ribeirão Preto Medical School University Hospital. All selected patients were invited by the researcher and those interested in participating the

Abbreviations: AFFM, appendicular fat free mass; ANOVA, analyses of variance; BIA, multifrequency bioelectrical impedance spectroscopy; BMI, body mass index; CKD, chronic kidney disease; DXA, dual energy X-ray absorptiometry; eGFR, estimated glomerular filtration rate; FFM, fat free mass; HD, hemodialysis; ICC, intraclass correlation coefficient; KTx, kidney transplant; NDD, non-dialysis-dependent; OH, over-hydration; PD, peritoneal dialysis; R^2 , coefficient of determination; Δ , body composition changes (prospective measurement—cross-sectional measurement); 95% CI, 95% confidence intervals.

study read and signed the informed consent form before the procedures began.

Protocol and Data Collection

Demographic and clinical data were collected from electronic medical records. Blood samples were taken within 1 d to 1 week before study assessment to determine laboratory parameters. Analyses were performed at the University Hospital's central laboratory. Serum creatinine was determined by kinetic method (creatinine calibrated to IDMS: COBAS 6000 [Roche/Hitachi]). The eGFR was calculated by the Chronic Kidney Disease Epidemiology Collaboration equation (CKD-EPI Creatinine 2009 equation). Dialysis patients had the weekly clearance of urea adjusted for total body water, with Kt/V determination in the week before the assessment.

Anthropometry and BIA were assessed by a single experienced dietician and DXA by a trained technician. All measures were performed consecutively at the same visit, after an 8-h fast (for HD group, 4-h fast), in patients with empty urinary bladder, drainage of the peritoneal dialysate, 30 min after the midweek HD session, advised to avoid strenuous physical activity in the previous day, wearing light clothes, without shoes and on the right side of the body (except if a fistula was present).

The second evaluation was performed 10 ± 2 months later. All measurements were performed exactly with the same protocol as the first evaluation and with the same equipment, dietician and technician for DXA, BIA and anthropometry assessments.

Anthropometric Measurements

Anthropometric measurements included body weight at the nearest 0.1 kg using a balance beam scale (Filizola[®], São Paulo, Brazil); height, waist, calf and mid arm muscle circumferences at the nearest 0.1 cm. BMI was also calculated.

A flexible plastic tape with graduated scale was applied to measure circumferences. Waist circumference was measured at the umbilical scar level (20). Calf circumference was measured with subjects seating down, knees at 90° and at the calf greatest circumference (21). Mid arm circumference and triceps skinfold thickness were measured for mid arm muscle circumference calculation (22). Triceps skin fold thickness was assessed with an adipometer (Lange[®], Cambridge Scientific Industries, Inc), three measures were taken and their mean was applied.

Hand grip strength was evaluated by a handheld pneumatic dynamometer (Charder[®], MG 4800) with subjects seated and asked to grip as hard as possible for three times with 1 min intervals. The highest value was recorded (23).

BIA

Multi-frequency spectroscopy BIA (BCM, Fresenius Medical Care, Bad Homburg, Germany) was performed with a tetra-polar whole-body wrist-to-ankle protocol (24) after 10-min adaptation in a supine position. The variables resistance, reactance, resistance index (resistance [ohm] divided by squared height [cm²]) and phase angle were provided at 50 kHz frequency; total body water, extracellular water and body cell mass (25), were provided by the software manufacturer.

Hydration status was assessed by over-hydration (OH) index provided by the BIA equipment (26, 27). Hydration disturbance was considered if OH > +1.1 L or OH < -1.1 L (28).

DXA

DXA (Hologic Discovery A, USA, Bedford, MA) was performed to evaluate lean mass, appendicular lean mass, AFFM, FFM, fat mass and fat mass of the trunk (6). Daily calibration of the device was performed before each assessment by scanning a spine phantom. A whole body scan was done after 10 min adaptation in supine position.

Predictive Equations of the Study

Supplementary Table 1 presents all the predictive equations herein evaluated, with information about BIA equipment applied, reference method against which the equation was validated and the population used for development of each equation. All BIA equations (7–10) evaluated in this study were developed with a tetra-polar whole-body wrist to ankle protocol.

Low Muscle Mass Definition

As no cutoff points to define low muscle mass are available for CKD patients, we applied definition based on the literature: appendicular lean mass <15 kg for women and <20 kg for men, assessed by DXA, according to the new European consensus on sarcopenia (4).

Statistical Analysis

Sample size was calculated based on the measurement of low muscle mass prevalence in CKD population. We assumed a minimum expected prevalence of 30% for all groups, with estimation of at least 80 patients for each CKD treatment group with a 5% precision and no correction for small population size (29).

Qualitative variables were presented as relative (percentage, %) and absolute (number, n) frequencies. Quantitative variables were expressed as measures of central tendency (mean) and dispersion (standard deviation).

Normality was tested with Shapiro-Wilk test with homoscedasticity evaluated. The differences between groups were performed by independent *T*-test or analysis of variance (ANOVA) adjusted by Bonferroni post-test, as appropriate. Categorical variables were compared by χ^2 -test.

For regression analysis, collinearity of data was observed.

Binary logistic regression analysis was performed to estimate the odds ratio for the potential diagnostic for low muscle mass. A multivariate analysis was carried out to evaluate de confounding effect of sex, age and weight.

DXA was considered the reference method for FFM and AFFM against which the predictive equations were validated for accuracy, for cross-sectional and body composition change data (prospective—cross-sectional data [Δ]).

Intraclass correlation coefficient (ICC) (30) was calculated to analyze agreement in a group level by comparing AFFM and FFM assessments by DXA analysis with predicted values.

The 5% error tolerance between measured and predict value (95% limits of agreement) was calculated as the percentage of

the sample whose predicted value was within 0.95–1.05 fold the measured value.

Multiple regression analysis was also performed between measured and estimated values. The coefficient of determination (R^2) is useful as it reflects the percentage of variation in the measurements by one method that is related to the variation in the other method (31). The standard error of the estimate provides information about the degree of the error (31).

Agreement at individual level between DXA data and predictive equations was evaluated by the Bland-Altman graphic (32) calculating the 95% limits of individual agreement (mean bias between the two methods ± 1.96 SD).

The best performing equation to be considered as surrogate for DXA analysis, in the total sample and in each CKD subgroup, was defined based on the combination of two criteria: (a)

the highest ICC value with the narrowest 95% Confidence interval (95% CI); (b) the highest R^2 and lowest standard error of the estimate. In addition, Bland-Altman bias with limit of agreement and also 5% error tolerance were taken into account.

For the best predictive equations, linear regression analyses were performed to determine the proportional bias between surrogate and reference methods.

Also, the inter-agreement between the DXA predictor of low muscle mass (appendicular lean mass) and the best predictive equation was quantified by Cohen's kappa coefficient. The sensitivity and specificity values were also estimated, and receiving operator characteristics curve analysis was carried out, assigning good reproducibility to area under the curve $>80\%$ (33).

TABLE 1 | Clinical data, anthropometry, BIA and body composition analyze of CKD subgroups stratified by sex.

	NDD		HD		PD		KTx	
	Men	Women	Men	Women	Men	Women	Men	Women
Cross-sectional data								
Sample size	46	37	35	44	8	15	48	33
Age (years)	49 \pm 10 ^a	48 \pm 10 ^a	44 \pm 12 ^a	49 \pm 8 ^a	37 \pm 12 ^b	42 \pm 12 ^b	50 \pm 8 ^a	48 \pm 9 ^a
eGFR (mL/min/1.73 m ²)	19.3 \pm 9.27 ^a	17.8 \pm 7.53 ^a	NA	NA	NA	NA	71.20 \pm 16.78 ^b	69.00 \pm 20.79 ^b
KT/V	NA	NA	1.48 \pm 0.19	1.84 \pm 0.72	2.72 \pm 0.52	2.53 \pm 0.51	NA	NA
Dialysis/KTx time (mo)	NA	NA	66 \pm 55	81 \pm 67	11 \pm 9	18 \pm 20	96 \pm 64	86 \pm 57
Weight (kg)	84 \pm 15.6 ^a	69 \pm 16.3 ^{a*}	69 \pm 13.9 ^b	64 \pm 13 ^b	79 \pm 15 ^b	59 \pm 9 ^{b*}	76 \pm 12 ^b	64 \pm 11 ^{b*}
Hand grip strength (kg)	38.8 \pm 9.57 ^{ab}	21.9 \pm 5.77 ^{ab*}	36.6 \pm 7.21 ^a	20.4 \pm 4.98 ^{a*}	45.5 \pm 6.66 ^{ab}	22.5 \pm 6.85 ^{ab*}	40.6 \pm 7.55 ^b	21.7 \pm 4.90 ^{b*}
Phase angle (°)	6.19 \pm 1.06 ^a	5.6 \pm 0.77 ^{a*}	6.34 \pm 0.94 ^a	5.4 \pm 1.19 ^{a*}	6.2 \pm 0.65 ^a	5.6 \pm 1.01 ^a	6.3 \pm 0.78 ^a	5.5 \pm 0.68 ^{a*}
Resistance index (cm ² /ohm)	67.8 \pm 10.73 ^a	47.5 \pm 9.43 ^a	54.1 \pm 11.8 ^b	39.8 \pm 8.05 ^b	65.5 \pm 10.14 ^{ab}	44.0 \pm 7.87 ^{ab}	59.9 \pm 8.66 ^a	41.2 \pm 6.09 ^a
Total body water (L)	42.9 \pm 5.46 ^a	31.6 \pm 4.93 ^{a*}	36.2 \pm 5.55 ^b	27.7 \pm 5.13 ^{b*}	42.0 \pm 6.08 ^{abc}	29.5 \pm 3.99 ^{abc*}	39.0 \pm 4.96 ^c	28.1 \pm 3.47 ^{c*}
OH (L)	1.05 \pm 2.01 ^a	0.09 \pm 1.38 ^{a*}	−0.18 \pm 1.99 ^b	−0.66 \pm 1.49 ^b	1.30 \pm 0.70 ^a	0.25 \pm 1.51 ^{a*}	0.39 \pm 1.02 ^{ab}	−0.14 \pm 0.87 ^{ab*}
Body cell mass (kg)	28.7 \pm 5.21 ^a	18.6 \pm 3.17 ^{a*}	25.2 \pm 4.50 ^b	15.6 \pm 4.69 ^{b*}	28.6 \pm 5.37 ^{ab}	18.8 \pm 3.22 ^{ab*}	26.5 \pm 5.28 ^{ab}	15.8 \pm 2.78 ^{ab*}
Appendicular lean mass (kg)	23.5 \pm 3.92 ^a	15.9 \pm 3.32 ^{a*}	20.2 \pm 3.54 ^b	13.9 \pm 2.78 ^{b*}	23.9 \pm 5.21 ^{ab}	14.1 \pm 2.30 ^{ab*}	21.4 \pm 3.27 ^{ab}	13.8 \pm 2.15 ^{ab*}
Lean mass (kg)	50.9 \pm 8.03 ^a	36.5 \pm 7.27 ^{a*}	43.9 \pm 7.75 ^b	33.0 \pm 5.89 ^{b*}	48.1 \pm 9.18 ^b	31.7 \pm 4.34 ^{b*}	45.7 \pm 6.49 ^b	32.0 \pm 5.05 ^{b*}
Fat mass (kg)	24.5 \pm 8.62 ^a	26.1 \pm 9.34 ^a	17.3 \pm 8.72 ^b	24.2 \pm 8.50 ^{a*}	22.1 \pm 7.92 ^a	20.8 \pm 5.63 ^a	22.0 \pm 6.67 ^a	25.2 \pm 7.30 ^{a*}
Δ data (prospective analysis—cross-sectional analysis)								
Sample size	16	9	6	12	8	15	20	9
Weight (kg)	−1.35 \pm 3.08	−0.21 \pm 1.97	0.15 \pm 4.64	−0.76 \pm 1.76	−1.84 \pm 2.78	−1.75 \pm 3.35	0.18 \pm 3.17	−2.72 \pm 4.70
Hand grip strength (kg)	−3.71 \pm 5.44	−0.70 \pm 2.28	−2.00 \pm 6.43	−1.07 \pm 4.45	−0.70 \pm 2.89	−1.08 \pm 4.10	−2.24 \pm 5.18	−0.65 \pm 4.36
Phase angle (°)	−0.07 \pm 0.64	−0.22 \pm 0.46	−0.17 \pm 0.89	−0.20 \pm 0.51	0.45 \pm 0.25	0.12 \pm 0.62	0.00 \pm 0.40	0.03 \pm 0.36
Resistance index (cm ² /ohm)	−3.42 \pm 9.61	−0.27 \pm 5.12	4.93 \pm 14.18	1.01 \pm 3.40	−4.42 \pm 5.22	−2.78 \pm 7.64	−0.72 \pm 3.67	−1.90 \pm 4.46
Total body water (L)	−1.68 \pm 3.33	−0.34 \pm 2.25	2.00 \pm 4.98	0.05 \pm 1.66	−1.42 \pm 2.60	−1.34 \pm 2.52	−0.19 \pm 1.91	−1.00 \pm 2.45
OH (L)	−0.32 \pm 2.01	0.20 \pm 0.82	0.80 \pm 2.88	0.30 \pm 0.82	−1.08 \pm 0.47	−0.38 \pm 1.53	−0.27 \pm 0.86	−0.22 \pm 0.46
Body cell mass (kg)	−1.70 \pm 3.66	−0.66 \pm 2.84	1.88 \pm 5.22	−0.04 \pm 2.51	−0.40 \pm 4.22	−0.96 \pm 1.68	−0.07 \pm 2.44	−0.36 \pm 2.78
Appendicular lean mass (kg)	−0.78 \pm 1.78	−0.18 \pm 1.18	−0.98 \pm 1.95	−0.54 \pm 1.32	−1.16 \pm 3.02	−1.23 \pm 1.51	−0.53 \pm 0.98	−1.05 \pm 2.27
Lean mass (kg)	−1.74 \pm 3.61	−1.06 \pm 2.83	−2.23 \pm 3.81	−1.09 \pm 2.83	−1.84 \pm 4.45	−2.27 \pm 2.59	−0.81 \pm 2.27	−2.39 \pm 5.63
Fat mass (kg)	0.18 \pm 3.91	0.78 \pm 2.51	2.26 \pm 6.25	0.19 \pm 2.68	0.25 \pm 6.33	0.57 \pm 2.74	1.05 \pm 2.59	0.20 \pm 4.70

Data presented as mean \pm standard deviation. BIA, bioelectrical impedance analyze; CKD, chronic kidney disease; HD, hemodialysis; KTx, kidney transplant; OH, over-hydration; NA, not applied; NDD, non-dialysis-dependent; PD, peritoneal dialysis. Body cell mass, resistance index, phase angle data by bioelectrical impedance analyze. Appendicular lean mass, lean mass, fat mass by dual energy X-ray absorptiometry analyze. *Independent T-test between sex ($p \leq 0.05$). ^{abc}ANOVA with Bonferroni post-test ($p \leq 0.008$).

Pearson correlation was applied to assess association between variables.

Significance was set at $p \leq 0.05$ except if adjustment for multiple comparisons was necessary. All statistics were performed with SPSS version 23 (IBM, Armonk, NY, USA).

RESULTS

Low Muscle Mass Diagnosis

CKD was secondary to systemic arterial hypertension (29%), glomerulonephritis (25%) and diabetes mellitus (10%). Clinical data, BIA, anthropometric and DXA measurements of CKD subgroups stratified by sex are shown in **Table 1**. PD patients were younger ($p \leq 0.008$). Hand grip strength, appendicular lean mass and lean mass were lower for women than men. HD group had the lowest value of appendicular lean mass and fat mass, and NDD the highest measurement of appendicular lean mass. Total body water and OH were lower for women than men and for HD, and higher for NDD and PD patients. In the prospective analysis, all groups and both sexes lost muscle mass and strength, and gained fat mass.

Low muscle mass affected more women (63%, $n = 81$) than men (37%, $n = 51$) ($p \leq 0.05$), was more prevalent among HD (70%, $n = 55$) and less in NDD (30%, $n = 24$) than the other CKD patients (PD, 52%, $n = 12$; KTx, 52%, $n = 42$) ($p \leq 0.008$).

The evaluation of factors related to the odds ratio for low muscle mass diagnosis is presented in **Table 2**. As NDD patients had the lowest prevalence of low muscle mass and the highest appendicular lean mass they were chosen as reference group. HD patients had more than 5 times and KTx about 3 times the risk for low muscle mass compared to NDD. Adiposity was

a risk factor in the multivariate analysis adjusted for sex, age, and weight.

Accuracy of Predictive Equations to Estimate FFM and AFFM

The agreement of predicted AFFM and FFM in comparison with AFFM and FFM from DXA for the total sample is presented in **Table 3**, and for each CKD group as supplementary material (NDD, **Supplementary Table 2**; HD, **Supplementary Table 3**; KTx, **Supplementary Table 4**; it was not possible to evaluate the PD group because of its small sample size). Agreement analysis was also performed according to the results of R^2 and standard error of the estimate in the total sample and is presented in **Table 4** for cross-sectional and prospective evaluations. Bland-Altman and scatter plot graphics for $\text{AFFM}_{\text{Sergi}}$ and $\text{AFFM}_{\text{Kyle}}$ equations compared with AFFM by DXA are presented in **Figure 1** and **Supplementary Figure 1**, respectively, for cross-sectional and body composition change data.

For cross-sectional data, considering the total sample and each CKD subgroup, FFM prediction equations did not performed well, $\text{AFFM}_{\text{Sergi}}$ and $\text{AFFM}_{\text{Kyle}}$ presented the best performance according to the highest ICC with narrowest 95%CI and the highest R^2 with lowest standard error of the estimate, in addition to the lowest bias and limits of agreement as well as the highest percentage within the 5% tolerance. Regarding body composition changes, none of the predictive equations was able to accurately predict changes in AFFM and FFM, with all ICC lower than 0.5.

Analysis of Interfering Factors for the Best Predictive Equations

We then investigated which variables were associated with bias between the best predictive equations ($\text{AFFM}_{\text{Sergi}}$ and $\text{AFFM}_{\text{Kyle}}$)

TABLE 2 | Odds Ratio of low muscle mass diagnosis in total sample ($n = 266$).

	Unadjusted			Adjusted by sex, age and weight		
	OR	95% CI	p	OR	95% CI	p
Sex (reference: men)	0.351	0.214–0.578	0.000			
CKD treatment (reference: NDD)						
HD	5.978	3.032–11.788	0.000	5.155	1.779–14.940	0.003
PD	2.846	1.102–7.350	0.031	1.236	0.261–5.851	0.790
KTx	2.809	1.468–5.375	0.002	3.154	1.143–8.704	0.027
Weight (kg)	0.837	0.803–0.873	0.000			
BMI (kg/m ²)	0.738	0.683–0.797	0.000	1.079	0.734–1.340	0.679
Hand grip strength (kg)	0.927	0.904–0.951	0.000	0.906	0.849–0.967	0.003
Phase angle (°)	0.649	0.501–0.840	0.0001	0.760	0.482–1.200	0.628
Resistance index (cm ² /ohm)	0.863	0.832–0.894	0.000	0.831	0.772–0.895	0.000
Body cell mass (kg)	0.837	0.786–0.880	0.000	0.746	0.660–0.842	0.000
Fat mass (kg)	0.903	0.872–0.935	0.000	2.035	1.606–2.578	0.000
Fat mass of trunk (kg)	0.831	0.783–0.883	0.000	1.765	1.466–2.125	0.000

BMI, body mass index; CKD, chronic kidney disease; HD, hemodialysis; KTx, kidney transplant; NDD, non-dialysis-dependent; OR, odds ratio; PD, peritoneal dialysis. Body cell mass, resistance index, phase angle data by bioelectrical impedance analyze. Fat mass and fat mass of the trunk by dual energy X-ray absorptiometry analyze. Low muscle mass according to the Revised European consensus (4) applying appendicular lean mass measurement by dual energy X-ray assessment. OR by binary logistic regression. For regression model analysis, the following variables were evaluated: age (years); estimated glomerular filtration ratio (mL/min/1.73 m²); KT/V; creatinine (mg/dL); urea (mg/dL); weight (kg), height (m); overhydration index (L); extracellular water (L); intracellular water (L); total body water (L); waist circumference (cm); calf circumference (cm); mid arm circumference (cm); mid arm muscle circumference (cm); resistance (ohm); reactance (ohm); the variables present in this table.

TABLE 3 | Agreement between DXA and prediction equations for AFFM and FFM in total sample stratified by sex.

Body composition variable	Men												Women															
	DXA or prediction equation		Bland-Altman analysis				ICC analysis				Pearson correlation		5% tolerance		DXA or prediction equation		Bland-Altman analysis				ICC analysis				Pearson correlation		5% tolerance	
			Bias (DXA-prediction)		LOA		ICC	(95% CI)		Bias (DXA-prediction)							LOA		ICC	(95%CI)								
	\bar{x}	SD	\bar{x}	SD	Lower	Upper	r	Lower	Upper	r	p	% (n)	\bar{x}	SD	\bar{x}	SD	Lower	Upper	r	Lower	Upper	r	p	% (n)				
Cross-sectional data $n = 137$												Cross-sectional data $n = 129$																
AFFM _{DXA} (kg)	23.16	4.11											15.32	3.02														
AFFM _{Sergi} (kg)	22.05	3.31	1.11	1.83	-2.47	4.70	0.915	0.795	0.956	0.90	0.000	50 (69)	15.61	2.60	-0.28	1.26	-2.76	2.19	0.945	0.921	0.962	0.91	0.000	42 (54)				
AFFM _{Kyle} (kg)	23.90	3.76	-0.73	1.82	-4.30	2.83	0.935	0.892	0.959	0.90	0.000	44 (61)	16.16	2.93	-0.83	1.24	-3.28	1.61	0.935	0.815	0.969	0.91	0.000	39 (50)				
AFFM _{Macdonald} (kg)	22.10	3.21	1.05	2.11	-3.08	5.18	0.804	0.686	0.880	0.86	0.00	42 (57)	13.86	2.53	1.45	1.82	-2.11	5.01	0.692	0.284	0.846	0.80	0.000	24 (31)				
FFM _{DXA} (kg)	48.99	8.23											34.87	6.42														
FFM _{TianHGS} (kg)	50.32	6.65	-1.29	3.90	-8.93	6.35	0.852	0.782	0.898	0.88	0.000	42 (58)	36.25	5.90	-1.34	2.96	-7.14	4.46	0.865	0.768	0.917	0.89	0.000	41 (53)				
FFM _{TianMAMC} (kg)	53.09	6.82	-4.10	3.77	-11.48	3.28	0.764	0.140	0.909	0.89	0.000	32 (43)	37.56	6.05	-2.69	3.00	-8.57	3.19	0.809	0.366	0.919	0.88	0.000	30 (39)				
FFM _{NooriHGS} (kg)	40.22	7.85	8.80	8.69	-8.23	25.83	0.262	-0.048	0.505	0.42	0.000	22 (30)	23.72	5.12	11.18	6.54	-1.63	23.99	0.129	-0.076	0.360	0.38	0.000	5 (6)				
FFM _{NooriMAMC} (kg)	48.32	6.12	0.66	4.04	-7.96	8.57	0.842	0.785	0.885	0.88	0.000	42 (57)	45.34	5.40	-10.46	3.00	-16.34	-4.58	0.342	-0.036	0.712	0.88	0.000	2 (3)				
FFM _{Hume} (kg)	53.05	6.40	-4.06	4.13	-12.15	4.03	0.732	0.181	0.886	0.87	0.000	30 (41)	41.49	5.64	-6.61	3.16	-12.80	6.19	0.540	-0.078	0.833	0.87	0.000	7 (9)				
FFM _{Janssen} (kg)	30.21	4.53	18.78	5.11	8.76	28.79	0.141	-0.033	0.437	0.83	0.000	0 (0)	18.91	3.44	15.96	4.11	7.90	24.01	0.118	-0.029	0.388	0.81	0.000	0 (0)				
FFM _{Lee} (kg)	33.91	3.84	15.08	5.36	4.57	25.58	0.174	-0.052	0.486	0.85	0.000	0 (0)	23.37	3.39	11.50	3.91	3.83	19.16	0.203	-0.049	0.538	0.86	0.000	0 (0)				
Δ data $n = 47$												Δ data $n = 40$																
AFFM _{DXA} (kg)	-0.78	1.66											-0.78	1.58														
AFFM _{Sergi} (kg)	-2.86	1.50	2.09	1.73	-1.30	5.48	0.361	-0.194	0.671	0.41	0.000	0 (0)	-4.25	0.93	3.47	1.37	0.77	6.17	0.177	-0.115	0.504	0.50	0.000	0 (0)				
AFFM _{Kyle} (kg)	-3.20	1.88	2.43	1.97	-1.43	6.29	0.339	-0.196	0.653	0.39	0.000	0 (0)	-5.08	1.16	2.43	1.97	-1.43	6.29	0.158	-0.085	0.480	0.52	0.000	0 (0)				
AFFM _{Macdonald} (kg)	-0.31	1.84	-0.45	1.94	-4.25	3.35	0.385	0.117	0.603	0.39	0.000	0 (0)	-0.21	1.02	-0.56	1.43	-3.36	2.80	0.391	0.106	0.620	0.46	0.000	0 (0)				
FFM _{DXA} (kg)	-1.48	3.19											-1.71	3.50														
FFM _{TianHGS} (kg)	-0.27	1.33	-1.18	2.95	-6.96	4.60	0.117	-0.144	0.378	0.17	0.263	0 (0)	-0.54	1.31	-1.17	3.00	-7.05	4.71	0.329	0.043	0.572	0.54	0.000	0 (0)				
FFM _{TianMAMC} (kg)	-0.03	1.53	-1.47	3.16	-7.66	4.72	0.173	-0.081	0.419	0.25	0.09	0 (0)	-0.61	1.43	-1.09	3.03	-7.02	4.84	0.338	0.050	0.579	0.51	0.000	0 (0)				
FFM _{NooriHGS} (kg)	-2.27	4.68	0.90	5.58	-10.03	11.83	-0.021	-0.311	0.274	-0.023	0.881	0 (0)	-0.80	3.49	-0.90	3.18	-7.13	5.33	0.472	0.324	0.747	0.58	0.000	0 (0)				
FFM _{NooriMAMC} (kg)	0.04	1.33	-1.54	3.13	-7.67	4.59	0.155	-0.093	0.400	0.25	0.08	0 (0)	-0.51	1.23	-1.19	3.10	-7.26	4.88	0.279	-0.007	0.532	0.48	0.000	0 (0)				
FFM _{Hume} (kg)	-0.18	1.08	-1.32	3.12	-7.43	4.79	0.123	-0.123	0.375	0.22	0.12	0 (0)	-0.39	0.96	-1.32	3.10	-7.39	4.75	0.243	-0.039	0.500	0.52	0.000	0 (0)				
FFM _{Janssen} (kg)	-0.52	3.21	-0.93	3.63	-8.04	6.18	0.352	0.081	0.577	0.36	0.01	0 (0)	-0.35	2.14	-1.36	3.07	-7.37	4.65	0.403	0.115	0.631	0.49	0.000	0 (0)				
FFM _{Lee} (kg)	-0.13	0.80	-1.36	3.10	-7.43	4.71	0.094	-0.152	0.347	0.22	0.12	0 (0)	-0.32	0.79	-1.38	3.16	-7.57	4.81	0.202	-0.075	0.464	0.53	0.000	0 (0)				

AFFM, appendicular fat free mass; DXA, dual energy X-ray absorptiometry; FFM, fat free mass; ICC, intraclass correlation coefficient; LOA, limits of individual agreement; Δ , body composition changes data (second assessment—first assessment). Bias calculated as DXA data—Prediction equation value; 5% tolerance between DXA and prediction equations (Prediction equation/DXA from 0.95 to ≤ 1.05).

TABLE 4 | Performance of prediction equations in total sample ($n = 266$).

Prediction equation	R^2 (coefficient of determination)	Standard error of the estimate (kg)
Cross-sectional data $n = 266$		
AFFM _{Sergi}	0.91	1.58
AFFM _{Kyle}	0.91	1.57
AFFM _{Macdonald}	0.86	1.98
FFM _{TianHGS}	0.88	3.47
FFM _{TianMAMC}	0.88	3.46
FFM _{NooriHGS}	0.52	7.12
FFM _{NooriMAMC}	0.63	6.27
FFM _{Hume}	0.86	3.73
FFM _{Janssen}	0.82	4.25
FFM _{Lee}	0.84	4.05
Δ data $n = 87$		
AFFM _{Sergi}	0.14	1.50
AFFM _{Kyle}	0.13	1.51
AFFM _{Macdonald}	0.16	1.48
FFM _{TianHGS}	0.13	2.97
FFM _{TianMAMC}	0.14	3.09
FFM _{NooriHGS}	0.05	3.11
FFM _{NooriMAMC}	0.13	3.12
FFM _{Hume}	0.13	3.11
FFM _{Janssen}	0.16	3.06
FFM _{Lee}	0.14	3.09

AFFM, appendicular fat free mass; FFM, fat free mass.

and AFFM by DXA (Table 5). The differences between the two methods were affected by sex, resistance index, total body water, AFFM and fat mass for AFFM_{Sergi} (adjusted $r^2 = 0.95$) and AFFM_{Kyle} (adjusted $r^2 = 0.94$), for cross-sectional data and for body composition change data (AFFM_{Sergi} adjusted $r^2 = 0.96$ and AFFM_{Kyle} adjusted $r^2 = 0.97$).

Analysis of Reproducibility for the Best Predictive Equations

The reproducibility of AFFM_{Sergi} and AFFM_{Kyle} for low muscle mass diagnosis, using appendicular lean mass by DXA and according to the cutoffs proposed by the European revised consensus on sarcopenia (4), is presented in Table 6. The inter-agreement was quantified by kappa values, according to sex and CKD subgroups. In total sample and in each CKD subgroup, AFFM_{Sergi} had a better performance than AFFM_{Kyle}, with Kappa from moderate to substantial agreement, and with good performance among NDD and KTx patients and poor performance among HD patients.

Correlations

Given the poor performance of prediction equations in predict changes of body composition, we performed correlation analysis to investigate if there is some surrogate variable with good correlation with cross-sectional data and muscle mass

changes, being as an alternative measurement for longitudinal evaluations (Table 7).

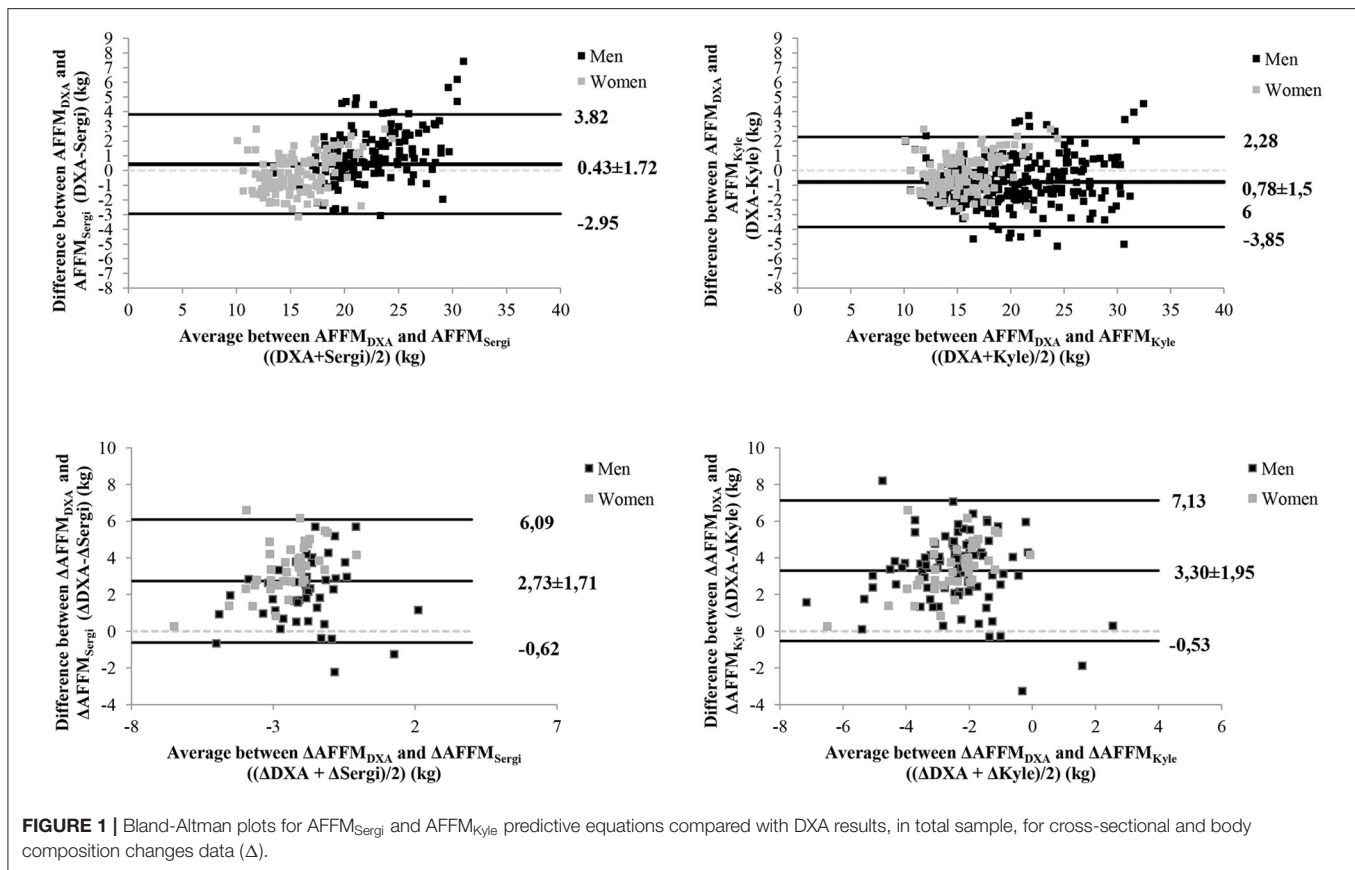
DISCUSSION

Our study showed that AFFM_{Sergi} was the best equation to predict AFFM in CKD, with good performance for low muscle mass diagnosis among NDD and KTx patients, and poor performance among HD patients. FFM equations did not perform well. Fat mass and total body water were important factors that interfered with accuracy of prediction equations, mainly for longitudinal evaluations. None of the predictive equations was accurate for assessment of changes in AFFM and FFM. Resistance index presented the best correlation with appendicular lean mass and lean mass in cross-sectional and longitudinal assessment, highlighting as an alternative measurement for longitudinal evaluations. Multivariate analysis after adjustment for sex, age and weight, revealed HD, KTx and adiposity as risk factors for low muscle mass.

AFFM_{Sergi} (8) equation was recommended by the revised European consensus on sarcopenia (4) as a way to standardize BIA estimation since its assessments vary widely depending on the device applied (34). AFFM_{Kyle} (9) was chosen because it was the first widely evaluated predictive equation for AFFM developed with spectroscopy BIA. Both equations were validated against DXA and presented the best performances. For the others equations, some factors may have interfered with the accuracy of the estimations, such as the protocol applied for body composition assessment. While Noori et al. (14) carried out measurements in a non-HD day, in our study evaluations were conducted after the midweek HD session. Probably, the different time-to-perform lower the accuracy, since fluid status is one of the main factors interfering with accuracy of BIA for body composition evaluations (35).

Macdonald et al. (10), developed the equation with NDD CKD patients with eGFR of 45.9 ± 28.8 ml/min, much higher than eGFR in our NDD group and lower than eGFR in our KTx sample. As AFFM_{Macdonald} is strongly associated with eGFR (10), it is important to take into consideration this parameter for AFFM_{Macdonald} estimations, and probably was the reason for the worse performance in our sample. Tian et al. (13) developed their equations from NDD CKD patients with an eGFR of 27 mL/min/1.73 m². Although this value is higher than the eGFR in our sample, it is closer than eGFR from Macdonald sample. Tian equations presented better performance for FFM than AFFM_{Macdonald}, although lower than AFFM_{Sergi} and AFFM_{Kyle}.

Another important factor is the BIA equipment applied for the development and validation of the predictive equation. All BIA devices involve application of a weak and alternating current through the body. However, they are not interchangeable. The number of frequencies (simple, multiple frequency or spectroscopy), the electronic circuitry, and the mathematical models (linear regression-derived population-specific equations or mathematical biophysical modeling by Cole model and Hanai's mixture theory) (36) applied for each device result in different predictions of water and body composition and also



in raw values that are not interchangeable (35). As could be seen in **Supplementary Table 1**, different BIA was applied for development of the equations herein evaluated.

All methods for evaluation of body composition are indirect, requiring assumptions that may not hold true in illness (5). Failure to account for the precision error in the reference method applied to validate the estimated measurement may contribute to misinterpretation and scaling error (37). This could contribute to decrease the accuracy of $AFFM_{Janssen}$ (7), FFM_{Lee} (12), and FFM_{Hume} (11) as they apply other reference methods than DXA.

It is essential to establish the ability of bedside methods to detect changes over time. In our study, none of the evaluated predictive equations had at least a good ICC for muscle mass changes, including $AFFM_{Sergi}$ and $AFFM_{Kyle}$. We took into account that the expected level of change in the body composition compartment was sufficiently high to be detected by the reference method, and confirmed that it was also detectable by the bedside method (24). Our total sample had a mean percentage of change [(second assessment – first assessment)/first assessment * 100] in $AFFM$ of $-3.71 \pm 8.32\%$, measured by DXA, $-1.37 \pm 6.33\%$ by $AFFM_{Sergi}$ and $-1.46 \pm 7.32\%$ by $AFFM_{Kyle}$. Similarly to DXA, $AFFM_{Sergi}$ and $AFFM_{Kyle}$ were able to detect changes in $AFFM$, although the direction of the change (gain or loss) and the quantity were poorly predicted, as confirmed by the low ICC and correlation and scatter plot graphics with fixed bias. Probably the fluid shift present in CKD and also the variation of adiposity

lowered the accuracy of longitudinal assessment, as total body water and fat mass were important bias for prediction values. Therefore, $AFFM_{Sergi}$ or $AFFM_{Kyle}$ might not be appropriate for follow-up analysis.

The advantages of BIA compared with reference methods are mainly the portability of the device, ease of use, and its affordability. Its use to assess body composition has increased in daily practice, mainly due to validation studies. Close adherence to recommended measurement protocols (38) as well as a standardized time to perform, are important to minimize potential errors, mainly for longitudinal evaluation (38). However, estimation of body composition in CKD is prone to error because underlying assumptions such as hydration of FFM at 73%, stable distribution of extracellular to intracellular water, and predictability of body geometry are not met in this disease, especially in patients with altered hydration or excess adiposity (38). Prediction of $AFFM$ seems to be less affected by the underlying considerations, but, as shown by our results, has limited accuracy in prospective assessment. The use of segmental BIA could improve prediction, despite studies showing controversial results (39, 40).

As pointed out by Mulasi et al. (41) it is unlikely that any algorithm can be relied upon for accurate whole-body estimates in patients with excess adiposity or altered fluid status, both conditions present in CKD. Excess adiposity ($BMI \geq 25 \text{ kg/m}^2$) was present in 60% of our total sample and over-hydration

TABLE 5 | Simple and multiple linear regression models for factors associated with bias in agreement between DXA and prediction equations in total sample.

	AFFM _{DXA} —AFFM _{Sergi} (kg)							AFFM _{DXA} —AFFM _{Kyle} (kg)								
	Simple linear regression model				Multiple linear regression model			Simple linear regression model				Multiple linear regression model				
	Coefficient	95% CI		<i>p</i>	Coefficient	95% CI		<i>p</i>	Coefficient	95% CI		<i>p</i>	Coefficient	95% CI		<i>p</i>
		Lower	Upper			Lower	Upper			Lower	Upper			Lower	Upper	
Cross-sectional data (<i>n</i> = 266)																
Intercept					−0.443	−0.738	−0.148	0.003					0.775	0.491	1.060	0.000
Sex	1.400	1.018	1.783	0.000	−1.404	−1.559	−1.249	0.000	0.099	−0.280	0.478	0.607	−1.885	−2.035	−1.735	0.000
RI (cm ² /ohm)	0.048	0.033	0.062	0.000	−0.038	−0.052	−0.024	0.000	−0.008	−0.022	0.006	0.263	−0.086	−0.099	−0.072	0.000
TBW (L)	0.096	1.018	1.783	0.000	−0.328	−0.363	−0.294	0.000	−0.002	−0.027	0.023	0.856	−0.322	−0.355	−0.288	0.000
AFFM (kg)	0.214	0.185	0.243	0.000	0.865	0.836	0.893	0.000	0.079	0.045	0.114	0.000	0.865	0.837	0.893	0.000
FM (kg)	−0.042	−0.067	−0.018	0.001	−0.077	−0.084	−0.070	0.000	−0.047	−0.069	−0.025	0.000	−0.074	−0.080	−0.068	0.000
Body composition change (Δ) Data (<i>n</i> = 87)																
Intercept					3.958	3.838	4.077	0.000					4.745	4.619	4.870	0.000
Sex	−1.384	−2.062	−0.705	0.000	−1.368	−1.522	−1.215	0.000	−1.874	−2.616	−1.131	0.000	−1.881	−2.042	−1.720	0.000
IR (cm ² /ohm)	−0.073	−0.125	−0.022	0.006	−0.076	−0.107	−0.440	0.000	−0.121	−0.176	−0.065	0.000	−0.125	−0.158	−0.091	0.000
TBW (L)	−0.175	−0.305	−0.045	0.090	−0.224	−0.306	−0.142	0.000	−0.290	−0.431	−0.149	0.000	−0.222	−0.308	−0.136	0.000
AFFM (kg)	0.661	0.481	0.841	0.000	0.846	0.777	0.914	0.000	0.580	0.349	0.811	0.000	0.842	0.770	0.913	0.000
FM (kg)	−0.298	−0.378	−0.219	0.000	−0.083	−0.110	−0.550	0.000	−0.298	−0.396	−0.200	0.000	−0.086	−0.115	−0.057	0.000

AFFM, appendicular fat free mass; DXA, dual energy X-ray absorptiometry; FM, fat mass; RI, resistance index; TBW, total body water. IR and TBW by bioelectrical impedance analysis. AFFM and FM by dual energy X-ray analysis. For regression model analysis, the following variables were evaluated: sex (1 man, 0 woman); age (years); estimated glomerular filtration ratio (mL/min/1.73 m²); KT/V; creatinine (mg/dL); urea (mg/dL); weight (kg), height (m); overhydration index (L); extracellular water (L); intracellular water (L); total body water (L); extracellular to total body water ratio; body cell mass (kg); lean mass, fat-free mass and fat mass by DXA analysis as well as their indexes; the variables present in this table.

TABLE 6 | Reproducibility of AFFM prediction equations to evaluate low muscle mass compared to DXA as reference.

	Inter-agreement		Low muscle mass diagnostic performance					Significance level
	Kappa	<i>p</i>	Sensibility (%)	Specificity (%)	AUC	95% CI		
						Lower	Upper	
DXA vs. AFFM _{Sergi}								
Total sample (<i>n</i> = 266)	0.676	0.000	73.48	94.02	0.838	0.786	0.889	0.000
Men (<i>n</i> = 137)	0.680	0.000	74.50	91.86	0.832	0.753	0.911	0.000
Women (<i>n</i> = 129)	0.650	0.000	72.83	97.91	0.854	0.788	0.920	0.000
NDD-CKD patients (<i>n</i> = 83)	0.809	0.000	78.26	98.33	0.883	0.780	0.986	0.000
Men (<i>n</i> = 46)	0.862	0.000	88.88	97.29	0.931	0.806	1.000	0.000
Women (<i>n</i> = 37)	0.757	0.000	71.42	100.00	0.857	0.708	1.000	0.000
HD-CKD patients (<i>n</i> = 79)	0.464	0.000	72.72	79.16	0.759	0.643	0.876	0.000
Men (<i>n</i> = 35)	0.364	0.031	30.00	66.66	0.683	0.501	0.866	0.050
Women (<i>n</i> = 44)	0.542	0.000	74.28	100.00	0.871	0.769	0.973	0.001
KTx-CKD patients (<i>n</i> = 81)	0.633	0.000	69.04	94.87	0.820	0.723	0.916	0.000
Men (<i>n</i> = 48)	0.689	0.000	70.00	96.42	0.832	0.701	0.964	0.000
Women (<i>n</i> = 33)	0.520	0.001	68.18	90.90	0.795	0.636	0.955	0.006
DXA vs. AFFM _{Kelly}								
Total sample (<i>n</i> = 266)	0.510	0.000	53.03	97.76	0.754	0.694	0.814	0.000
Men (<i>n</i> = 137)	0.585	0.000	33.33	96.51	0.649	0.549	0.750	0.004
Women (<i>n</i> = 129)	0.341	0.000	65.43	100.00	0.827	0.757	0.897	0.000
NDD-CKD patients (<i>n</i> = 83)	0.570	0.000	47.82	100.00	0.739	0.600	0.878	0.001
Men (<i>n</i> = 46)	0.315	0.003	22.00	100.00	0.611	0.383	0.840	0.306
Women (<i>n</i> = 37)	0.691	0.000	64.28	100.00	0.821	0.659	0.984	0.001
HD-CKD patients (<i>n</i> = 79)	0.388	0.000	60.00	87.5	0.738	0.623	0.852	0.001
Men (<i>n</i> = 35)	0.283	0.069	50.00	80.00	0.650	0.466	0.834	0.134
Women (<i>n</i> = 44)	0.439	0.000	65.71	100.00	0.829	0.710	0.947	0.003
KTx-CKD patients (<i>n</i> = 81)	0.443	0.000	45.23	100.00	0.726	0.615	0.838	0.000
Men (<i>n</i> = 48)	0.280	0.005	25.00	100.00	0.625	0.458	0.792	0.143
Women (<i>n</i> = 33)	0.538	0.000	63.63	100.00	0.818	0.677	0.960	0.003

AFFM, appendicular fat free mass; AUC, area under the curve; DXA, dual energy X-ray absorptiometry; HD, hemodialysis; KTx, kidney transplant therapy; NDD, non-dialysis-dependent. Patients were classified with low muscle mass according to cutoffs values for ALM proposed by revised European consensus (4): appendicular lean mass <15 kg for women and <20 kg for men, assessed by DXA.

(OH > + 1.1) in 40%. As BIA provides indirect estimates of body composition from the measurement of resistance of body tissues to an electric current, raw data might be an option to predicted values (42). In our sample, the only parameter with good correlation with longitudinal variation in lean mass and appendicular lean mass was the resistance index. In agreement with other authors (42) we suggest that raw measurements of BIA could provide better objective biomarkers of nutritional status than predicted values. This is an important issue for BIA applicability in clinical settings, mainly in CKD where it is already applied for fluid management. Another point that should not be underestimated is that, more often lately, clinicians have limited time to search the literature to identify the most suitable equation for the patient being evaluated; therefore, the only alternative is to directly apply the prediction provided by the device without knowing its appropriateness or interpreting the raw data.

Low muscle mass was present in more than 30% of patients in all CKD groups, highlighting that the nutritional status was compromised even among non-elderly patients. The NDD group

was less affected by low muscle mass, although these patients also presented an important prevalence. A higher prevalence of low muscle mass was observed in HD patients, in agreement with other studies (43) suggesting that muscle wasting could progress as kidney function declines (19). Likewise, increased inflammation and metabolic acidosis promoted by the dialysis process accelerate protein degradation (44). Similarly to our results, other studies showed an important prevalence of low muscle mass among KTx patients (45), which could be related to previous dialysis therapy (90% of our patients underwent HD before KTx with a median duration of 40 months, 6–65) and with body composition deterioration after transplant (18), demonstrating the challenges of recovering nutritional status in CKD.

CKD patients lost muscle mass and gained fat mass, worsening their nutritional status, which is associated with worse quality of life and higher mortality (46). Thus, screening, prevention, and treatment of muscle loss should have high priority in CKD patients.

TABLE 7 | Correlations between surrogate methods and DXA in total sample ($n = 266$).

Variable	Correlation coefficient	<i>p</i>
Cross-sectional data $n = 266$		
Correlations with appendicular lean mass (DXA)		
Mid arm muscle circumference (cm)	0.73	0.000
Calf circumference (cm)	0.65	0.000
Resistance index (cm^2/ohm)	0.91	0.000
Phase angle ($^\circ$)	0.35	0.000
Body cell mass (kg)	0.82	0.000
AFFM _{Sergi} (kg)	0.95	0.000
AFFM _{Kyle} (kg)	0.95	0.000
Correlation with lean mass (DXA)		
Mid arm muscle circumference (cm)	0.74	0.000
Calf circumference (cm)	0.65	0.000
Resistance index (cm^2/ohm)	0.92	0.000
Phase angle ($^\circ$)	0.29	0.000
Body cell mass (kg)	0.78	0.000
AFFM _{Sergi} (kg)	0.96	0.000
AFFM _{Kyle} (kg)	0.96	0.000
Δ data $n = 87$		
Correlation with Δ appendicular lean mass (DXA)		
Mid arm muscle circumference (cm)		>0.05
Calf circumference (cm)	0.34	0.000
Resistance index (cm^2/ohm)	0.68	0.000
Phase angle ($^\circ$)		>0.05
Body cell mass (kg)	0.37	0.000
AFFM _{Sergi} (kg)	0.38	0.000
AFFM _{Kyle} (kg)	0.37	0.000
Correlation with Δ lean mass (DXA)		
Mid arm muscle circumference (cm)		>0.05
Calf circumference (cm)	0.34	0.000
Resistance index (cm^2/ohm)	0.70	0.000
Phase angle ($^\circ$)		>0.05
Body cell mass (kg)	0.40	0.000
AFFM _{Sergi} (kg)	0.41	0.000
AFFM _{Kyle} (kg)	0.40	0.000

AFFM, appendicular fat free mass; Δ , body composition change (prospective assessment - cross-sectional assessment).

Total fat mass and trunk fat mass are risk factors for low muscle mass. Systemic and muscle oxidative stress, persistent inflammation, insulin resistance and hormonal alterations due to the presence of excess adipose tissue (47) lead to changes in muscle metabolism. These result in reduction of lean mass to fat mass ratio (48). Accordingly, the risk of skeletal muscle loss should also be considered in CKD patients with obesity, the most challenging patients for nutritional status evaluation as not only fluid status may impair their body composition assessment but also excess adiposity could mask loss of muscle mass. As provided by our results, fat mass was an important factor in bias for prediction of AFFM. So, obese patients could benefit most from use of raw BIA data for longitudinal evaluation of muscle mass.

The study's main limitations are the exclusion of patients over 60 years of age, the low sample size of the PD group and the short time of longitudinal assessment, which precludes analysis of exposure/outcome association and limits the evaluation of prediction equations for assessment of changes in body composition. Additionally, body thickness, hydration status, and diseases with water retention can affect DXA results (6). Therefore, in our DXA analysis of body composition, data from CKD patients with excess fat and fluid retention could be prone to errors that may promote a scaling error leading to misinterpretation of predicted values. On the other hand, the present study presents several strengths. First, the evaluation of CKD patients was carried out under treatment conditions such as HD and PD, as well as for NDD patients and patients after KTx therapy, which are more rarely evaluated. Moreover, the study includes DXA as reference method applied to all patients, together with a cross-sectional and longitudinal assessment. Additionally, we evaluated a wide variety of BIA and anthropometry predictive equations through a comprehensive statistical analysis including not only assessment of accuracy and agreement but also sensitivity and specificity analysis with discriminative power to detect low muscle mass.

In conclusion, accuracy and reproducibility analysis in the present study indicates that the predictive equation AFFM_{Sergi} can be an alternative technique to assess muscle mass in CKD patients, and could be useful for low muscle mass diagnosis in NDD and KTx patients, with limited applicability for patients under HD. However, prediction of changes in AFFM may be biased, mainly due to variations in fluid status together with the presence of high adiposity, limiting the applicability of AFFM_{Sergi} equation for longitudinal evaluations. As resistance index from BIA presented the best correlation with cross-sectional and longitudinal data of lean mass evaluated by reference method, we suggest that raw data from BIA could provide better information about progression of muscle mass in CKD, mainly in patients with dehydration or hyperhydration and excess adiposity. More studies are needed to better understand the relationship between nutritional status and also clinical prognosis with raw data from BIA as a way to easily obtain quick and objective measurements suitable to be tracked over time. Body composition evaluations are a priority for CKD patients, given the worsening of their nutrition status over time. Patients in HD, after KTx and with obesity are at higher risk for low muscle mass diagnosis.

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 Human Research Ethics Committee of Ribeirao Preto Medical School University Hospital. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

NB, PC, and GC contributed to the conception of the research. NB contributed to the design of the research, acquisition of the data, and drafted the manuscript. NB and LV-P contributed to the analysis and interpretation of data. All authors critically revised the manuscript, agree to be fully accountable for ensuring the integrity and accuracy of the work, and read and approved the final manuscript.

FUNDING

This research was funded by Coordenação de aperfeiçoamento de pessoal de nível superior (CAPES) (NB, Grant No. 001). The funding agency of the project did not participate in the study

design, data analysis and interpretation, writing of the article, or in the decision to send it to the present journal.

ACKNOWLEDGMENTS

We sincerely appreciate all participants for their kind cooperation in our study. We also wish to acknowledge all of the investigators in our study and the hospital staff members, and also the CAPES for funding the research.

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fnut.2021.683393/full#supplementary-material>

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Association of Phase Angle, but Not Inflammation and Overhydration, With Physical Function in Peritoneal Dialysis Patients

Vanessa Mota Silva¹, Maryanne Zilli Canedo Silva², Barbara Perez Vogt^{3*},
Nayrana Soares Carmo Reis², Fabiana Lourenço Costa², Mariana Souza Dorna²,
Marcos Ferreira Minicucci² and Jacqueline Costa Teixeira Caramori²

¹ Clinical Hospital of Botucatu Medical School, Multiprofessional Specialization in Adult and Elderly Health, São Paulo State University, UNESP, Botucatu, Brazil, ² Internal Medicine Department, Botucatu Medical School, São Paulo State University, UNESP, Botucatu, Brazil, ³ Graduate Program in Health Sciences, Medicine Faculty, Federal University of Uberlândia (UFU), Uberlândia, Brazil

OPEN ACCESS

Edited by:

Carla Maria Avesani,
Karolinska Institutet, Sweden

Reviewed by:

Fabiana Nerbass,
FUNDAÇÃO PRÓ-RIM, Brazil
Nilian Souza,
National Cancer Institute
(INCA), Brazil

*Correspondence:

Barbara Perez Vogt
barbaravogt@ufu.br

Specialty section:

This article was submitted to
Clinical Nutrition,
a section of the journal
Frontiers in Nutrition

Received: 26 March 2021

Accepted: 06 May 2021

Published: 31 May 2021

Citation:

Silva VM, Silva MZC, Vogt BP,
Reis NSC, Costa FL, Dorna MS,
Minicucci MF and Caramori JCT
(2021) Association of Phase Angle,
but Not Inflammation and
Overhydration, With Physical Function
in Peritoneal Dialysis Patients.
Front. Nutr. 8:686245.
doi: 10.3389/fnut.2021.686245

Introduction: Muscle mass depletion, overhydration, and inflammatory state have been related to impaired physical function in chronic kidney disease patients. The relationship between bioelectrical impedance analysis (BIA) parameters, such as hydration status and phase angle (PhA), with physical function in peritoneal dialysis (PD), is still not well-established. Therefore, the objective was to evaluate the association of BIA parameters (overhydration index and PhA) and inflammatory markers with physical function in patients on PD.

Methods: The present cross-sectional study enrolled PD patients. Multifrequency BIA was performed to obtain overhydration index and PhA. The Short Physical Performance Battery (SPPB) test battery was applied to assess physical function. The time to complete the 4-m gait test and sit-to-stand test was also considered for physical function assessment. The inflammatory markers tumor necrosis factor- α and C-reactive protein levels were determined. Multiple linear regression models were performed, with the physical function variables as dependent variables, adjusted for age, diabetes, and sex.

Results: Forty-nine PD patients were enrolled, 53.1% ($n = 26$) women; mean age, 55.5 ± 16.3 years. There were significant correlations between PhA and SPPB ($r = 0.550$, $p < 0.001$), time of 4-m gait test ($r = -0.613$, $p < 0.001$) and sit-to-stand test and ($r = -0.547$, $p < 0.001$). Overhydration index was significantly correlated with SPPB, 4-m gait test ($r = 0.339$, $p = 0.017$), and sit-to-stand test ($r = 0.335$, $p = 0.019$). Inflammatory markers were not significantly correlated with physical function parameters. In the multiple linear regression analysis, PhA was associated with physical function parameters, even after adjustments. Overhydration index was associated with all physical function tests only in the models with no adjustments.

Conclusion: PhA was independently associated with physical function in PD patients. Inflammatory markers and overhydration index were not associated with physical function.

Keywords: bioelectrical impedance, functional capacity, inflammation, phase angle, overhydration, short physical performance battery

INTRODUCTION

Bioelectrical impedance analysis (BIA) has been used in clinical practice to assess patients with chronic kidney disease (CKD) on dialysis. BIA provides raw data regarding electrical current conductivity through the body tissues, such as resistance and reactance. The relation among these parameters may be converted into physiological parameters related to body composition and hydration status (1).

BIA was recommended by the National Kidney Foundation–Kidney Disease Outcomes Quality Initiative in the Nutrition Guideline in CKD, as a method capable of assessing body composition in patients on hemodialysis. However, there is insufficient evidence to suggest using BIA to assess body composition in peritoneal dialysis (PD) patients (2).

Regarding hydration status evaluation, one of the parameters obtained by BIA is the overhydration index (OH), which assesses fluid overload by the difference between measured and expected extracellular water in normal situations (3). OH is considered an important predictor of the evolution of patients on dialysis, as it helps with blood pressure control (4). Moreover, OH has been associated with lower survival in both hemodialysis and PD patients (5, 6).

Phase angle (PhA), a measure of the relationship between resistance and reactance, reflects cell membranes integrity, cellularity, and cell function (7). PhA has been related to hydration status (8), nutritional status (9), and mortality risk (10) in PD patients.

CKD is characterized by metabolic and hormonal disturbances, which culminates in skeletal muscle wasting and disfunction. Decreased physical function has been associated with poor outcomes, such as disability, falls, fracture, hospitalization, and mortality (11). In dialysis and non-dialysis CKD patients, muscle mass depletion, overhydration, and inflammatory state have been related to impaired physical function (12, 13).

However, the relationship between BIA parameters, such as hydration status and PhA, with physical function in PD is still not well-established. Therefore, the objective of the study is to evaluate the association of BIA parameters (OH and PhA) and inflammatory markers with physical function in patients with CKD on PD.

METHODS

This was a secondary analysis of data from a study previously published (14), which enrolled patients aged 18 years or above, with CKD on PD (for at least 3 months) at the Dialysis Unit from the Clinics Hospital of Botucatu Medical School (Botucatu, Brazil). The protocol was approved by the Ethics and Research Committee (CAAE 61634816.4.0000.5411), and the participants gave written informed consent.

Clinical, demographic data, and routine laboratory tests were collected from medical records (sex, age, comorbidities, underlying disease, PD modality, Kt/V, serum hemoglobin, creatinine, urea, albumin, C-reactive protein).

Inflammatory Markers

Blood collection was performed in the morning, after PD. C-reactive protein levels were determined by immunoassay (Vitros, Johnson and Johnson Clinical Diagnostics, Rochester, NY) at the specialized chemistry laboratory of the Hospital of Botucatu Medical School. Blood samples were centrifuged, aliquoted, and kept frozen at -80°C for tumor necrosis factor-alpha (TNF- α) levels determination, using an enzyme-linked immunosorbent assay according to the manufacturer's instructions (R&D System, Inc., Minneapolis, USA).

Bioelectrical Impedance Analysis

All the patients underwent anthropometric and bioimpedance assessment with no dialysate in the peritoneal cavity in the morning, after PD. Measurements of body weight, height, and calculation of body mass index were carried out.

Multifrequency BIA (Body Composition Monitor, BCM—Fresenius Medical Care®, Germany) was performed with the patients in the recumbent position and all metal accessories removed. The BCM measures body resistance and reactance to electrical currents of 50 discrete frequencies, ranging from 5 to 1,000 kHz. Values of OH and PhA in the frequency of 50 kHz were considered. The OH was obtained by the BCM device, considering the difference between the measured extracellular water (ECW) and the predicted values based on fixed hydration on lean and adipose tissue mass.

Physical Function Assessment

To assess physical function, the Short Physical Performance Battery (SPPB) was used. This assessment was performed at the same day of BIA, after PD. SPPB is composed of a set of three tests: balance, a 4-m gait and sit-to-stand tests. These tests are useful to predict the performance of the lower limbs. The scores were assigned to each test, ranging from zero to four (15, 16).

Besides the SPPB total score, the time to complete the 4-m gait test and sit-to-stand test was considered for physical function assessment.

Statistical Analysis

The Kolmogorov-Smirnov test was performed to assess the normality of the data. Data were expressed as mean and standard deviation, median, and interquartile variation, according to the distribution. Frequencies were expressed as percentages.

Comparisons between male and female, and between diabetic and non-diabetic patients were performed using Student's *t* or Mann–Whitney tests.

The patients were grouped according to OH and PhA tertiles. Tertiles were compared, using one-way ANOVA with Tukey *post-hoc* or Kruskal–Wallis with Dunn *post-hoc* test.

Correlations between physical function parameters and other variables were assessed. For this, Pearson and Spearman coefficient correlations were used.

Multiple linear regression models were performed, with the physical function variables as dependent variables, adjusted for age, diabetes, and sex.

TABLE 1 | Clinical and nutritional characteristics of PD patients.

Variables	Total (n = 49)
Dialysis vintage (months)	10.0 (5.0–18.0)
Kt/V total*	2.2 ± 0.5
Kt/V peritoneal**	1.4 (1.1–1.7)
Kt/V renal**	0.6 (0.3–1.1)
Weight (kg)	68.8 ± 15.5
Body Mass Index (Kg/m ²)	25.8 ± 4.3
Phase angle (°)	5.4 ± 1.1
Overhydration (L)	0.8 ± 1.1
Hemoglobin (mg/dL)	11.3 ± 1.5
Urea (mg/dL)	99.3 ± 24.9
Creatinine (mg/dL)	9.1 ± 2.8
Albumin (g/dL)	3.6 ± 0.4
C-reactive protein (mg/L)	0.5 (0.5–1.0)
Tumor necrosisfactor-alpha (ng/L)	16.9 (13.8–21.4)

Data are expressed as numbers only, mean ± standard deviation, number (%), or median (interquartile range).

Kt/V: Fractional clearance of urea; *n = 47; **n = 46.

The analyses were performed, using IBM SPSS Statistics V22.0 (IBM Corp., Armonk, NY, USA). The level of significance was set at $p < 0.05$.

RESULTS

Forty-nine PD patients were enrolled, with mean age 55.5 ± 16.3 years, 53.1% ($n = 26$) female. The patients in different modalities of PD were included, 61.2% ($n = 30$) in continuous cyclic PD, 34.7% ($n = 17$) in night intermittent PD, and 4.1% ($n = 2$) in continuous ambulatory PD. The most prevalent comorbidities were arterial hypertension (79.6%, $n = 39$) and diabetes (26.5%, $n = 13$). The main underlying diseases were glomerulonephritis, 18.4% ($n = 9$); diabetic nephropathy, 16.3% ($n = 8$); hypertensive nephropathy, 16.3% ($n = 8$); undetermined, 24.5% ($n = 12$); and other causes, 24.5% ($n = 12$). Clinical and nutritional characteristics of PD patients are shown in **Table 1**.

Results of physical function assessments of both genders were compared. There were no differences between male and female (SPPB: $p = 0.324$, gait test: $p = 0.109$ and sit-to-stand test: $p = 0.248$). In the comparison between the patients with diabetes or not, there were differences regarding age ($p = 0.003$), SPPB ($p = 0.025$) and sit-to-stand test ($p = 0.009$). OH ($p = 0.184$), PhA ($p = 0.193$), CRP ($p = 0.572$), TNF- α ($p = 0.937$), and gait test ($p = 0.150$) did not differ between diabetic and non-diabetic patients.

Forty-seven patients (96.0%) reached the maximum score in the balance test. The results of physical function assessments and inflammatory markers are shown in **Table 2**, as well as the comparison among OH tertiles. OH ranged from -1.0 to 3.2 L. PhA ranged from 3.2° to 7.8° . The comparison among PhA tertiles is shown in **Table 3**.

In the correlation analysis, there was a positive correlation between PhA and SPPB and negative correlations of PhA with time of a 4-m gait and sit-to-stand tests. OH was

negatively correlated with the total SPPB score, and positively correlated with a 4-m gait and sit-to-stand tests. Inflammatory markers were not significantly correlated with physical function parameters (**Table 4**). Dialysis vintage, renal Kt/V, BMI, urea, and hemoglobin were not significantly correlated with physical function tests. Albumin was correlated only with the gait test ($r = -0.400$, $p = 0.004$). Creatinine was correlated with all the physical function test (SPPB $r = 0.389$, $p = 0.006$; gait test $r = -0.430$, $p = 0.002$; sit-to-stand $r = -0.379$, $p = 0.007$).

Multiple linear regression models were performed with the SPPB, gait test, and sit-to-stand test as dependent variables (**Table 5**). Age and diabetes were chosen as adjustments due to the association with physical function in univariate analysis. Sex was also included as an adjustment. Although serum creatinine was correlated with physical function tests, in multiple linear regression models, creatinine did not improve or change the results. Therefore, as the sample size is small, we did not add creatinine to the final models.

In the multiple linear regression final models, OH was associated with all physical function tests only in the models with no adjustments. PhA was independently associated with all physical function tests after adjustments, except with the gait test when diabetes was included as an adjustment.

DISCUSSION

Although BIA has been used in the last decades for body composition assessment, it is not a direct method for this evaluation (7). BIA is based on electrical properties of tissues, and the body compartments are estimated from equations that include raw impedance parameters and anthropometric measurements (17).

More recently, the use of raw and associated impedance parameters has gained attention. Kidney failure patients are characterized by fluid imbalance, which turn BIA even more valuable. Huang et al. showed that age, diabetes, and fluid overload were independently associated with lower PhA in PD patients. Moreover, in the same study, lower PhA was associated with higher cardiovascular and all-cause mortality (10). Increased OH values were associated with mortality risk, as evidenced in a systematic review, enrolling studies with kidney failure patients (18).

The present study aimed to evaluate the association of BIA parameters (PhA and OH), and inflammation with physical function in PD patients. PhA was positively correlated with the SPPB total score and negatively correlated with the time to complete a 4-m gait and sit-to-stand tests. As expected, these results show the association of increased PhA with better physical function.

In addition to the association of PhA with better physical function, OH was negatively correlated with the SPPB total score and positively correlated with the time to complete the 4-m gait and sit-to-stand tests. The patients in the third tertile had worse physical function compared with the first tertile. However, OH was not independently associated with physical function after adjustments in multiple analyses.

TABLE 2 | Comparison of different parameters among overhydration tertiles in patients on peritoneal dialysis ($n = 49$).

Variables	1st Tertile OH < 0.27 ($n = 17$)	2nd Tertile OH ≥ 0.27 to <1.26 ($n = 16$)	3rd Tertile OH ≥ 1.26 ($n = 16$)	<i>p</i>
Age (years)	52.4 \pm 14.4	52.4 \pm 16.8	61.8 \pm 16.8	0.172
Female sex [n (%)]	10 (58.8)	8 (50.0)	8 (50.0)	0.841
PD modality [n (%)]				0.862
CAPD	1 (5.9)	0	1 (6.2)	
CCPD	10 (58.8)	11 (68.8)	9 (56.3)	
NIPD	6 (35.3)	5 (31.2)	6 (37.5)	
Short physical performance battery (points)	10.0 (9.0–11.0)	9.5 (8.0–10.8)	8.5 (7.0–9.8) ^a	0.035
Gait test (seconds)	4.21 (3.5–5.3)	4.4 (4.1–5.2)	5.0 (4.5–7.1) ^a	0.029
Sit-to-stand test (seconds)	15.2 \pm 3.6	15.8 \pm 3.9	19.5 \pm 4.7 ^{ab}	0.009
C-reactive protein (mg/L)	0.5 (0.5–1.3)	0.5 (0.5–0.6)	0.6 (0.5–1.4)	0.387
Tumor necrosisfactor-alpha (ng/L)	16.94 (14.54–23.23)	17.4 (11.6–21.3)	15.5 (13.7–20.0)	0.527

Data are expressed as numbers only, mean \pm standard deviation, number (%), or median (interquartile range). Comparison using one-way ANOVA with the Tukey post-hoc test or Kruskal–Wallis with Dunn post-hoc. ^a $p < 0.05$ when compared with first tertile; ^b $p < 0.05$ when compared with second tertile.

TABLE 3 | Comparison of different parameters among phase angle tertiles in patients on peritoneal dialysis ($n = 49$).

Variables	1st Tertile PhA <4.92 ($n = 16$)	2nd Tertile PhA ≥ 4.92 to ≤ 5.87 ($n = 17$)	3rd Tertile PhA >5.87 ($n = 16$)	<i>p</i>
Age (years)	67.8 \pm 12.0	54.9 \pm 13.2 ^a	43.7 \pm 14.4 ^{ab}	0.000
Female sex [n (%)]	10 (62.5)	11 (64.7)	5 (31.2)	0.103
PD modality [n (%)]				0.622
CAPD	1 (6.2)	1 (5.9)	0	
CCPD	8 (50.0)	10 (58.8)	12 (75.0)	
NIPD	7 (43.8)	6 (35.3)	4 (25.0)	
Short physical performance battery (points)	8 (7.0–9.0)	10 (9.0–11.0) ^a	10 (9.2–11.0) ^a	0.002
Gait test (seconds)	6.2 (4.8–7.0)	4.5 (4.0–5.1) ^a	4.2 (3.6–4.5) ^a	0.000
Sit-to-stand test (seconds)	20.0 \pm 4.4	16.0 \pm 4.4 ^a	14.5 \pm 2.2 ^a	0.001
C-reactive protein (mg/L)	0.5 (0.5–1.6)	0.5 (0.5–1.3)	0.5 (0.5–0.7)	0.412
Tumor necrosisfactor-alpha (ng/L)	16.7 (14.2–20.9)	14.5 (11.1–17.4)	19.8 (15.6–23.2)	0.058

Data are expressed as numbers only, mean \pm standard deviation, number (%), or median (interquartile range). Comparison using one-way ANOVA with the Tukey post-hoc test or Kruskal–Wallis post-hoc. ^a $p < 0.05$ when compared with first tertile; ^b $p < 0.05$ when compared with second tertile.

Inflammation is considered one of the aspects that lead to decrease of physical function (19, 20). Since overhydration status may increase inflammation in PD patients (21), we expected an association of both inflammatory markers and OH with physical function. However, the inflammatory markers considered in our study were not associated with worse physical function, as well as OH.

In the hemodialysis patients, the presence of fluid overload can slow the gait speed. The fluid overload significantly impacts on the physical performance of the patients over time, as observed in a longitudinal study by Carlos et al. (12). On the other hand, excessive fluid withdrawal in hemodialysis sessions can lead to intradialytic hypotension, which is associated with a reduced gait after a dialysis session (22, 23).

In PD, data about the impact of increased fluid on the physical function assessed by walking are scarce. A walking variable may be reduced in these patients due to conditions such as physical inactivity, peripheral neuropathy, and muscle

strength (24). However, such variables were not evaluated in the present study, which may be an explanation for the lack of association of physical function with OH in the regression model.

In the multiple linear regression analyses, the association of PhA with the sit-to-stand test and SPPB was maintained even after adjusting for age, diabetes, and sex. All the physical function tests used in the present study reflect common daily living activities, i.e., getting up from a chair or walking small distances. Moreover, these tests are easy, fast, portable, and inexpensive procedures to measure physical performance.

PhA has been associated with physical function in many conditions. Recently, Vincenzo et al. (25) have showed in a meta-analysis that PhA is decreased in sarcopenic subjects. Kosoku et al. (26) found a negative correlation between PhA and sarcopenia in kidney transplant recipients. Sarcopenia is defined as the presence of both muscle mass and physical function decrease (27).

TABLE 4 | Correlations between bioimpedance parameters and physical function tests in patients on peritoneal dialysis ($n = 49$).

Variables	Phase angle (°)		OH (L)		CRP (mg/L)		TNF- α (ng/L)	
	<i>r</i>	<i>p</i>	<i>r</i>	<i>p</i>	<i>r</i>	<i>p</i>	<i>r</i>	<i>p</i>
Short physical performance battery (points)	0.550	0.000	−0.316	0.027	−0.013	0.931	0.110	0.450
4-meter gait test (seconds)	−0.613	0.000	0.339	0.017	0.136	0.350	−0.169	0.245
Sit-to-stand test (seconds)	−0.547*	0.000	0.335*	0.019	0.020	0.894	−0.131	0.368

OH, overhydration; CRP, C-reactive protein; TNF- α , Tumor necrosis factor- α .

*Pearson coefficient correlation. Coefficients with no sign refer to Spearman coefficient correlation.

TABLE 5 | Multiple linear regression models with short physical performance battery, the gait test, and the sit-to-stand test as dependent variables in patients on peritoneal dialysis ($n = 49$).

Dependent variable	Independent variable	Model	β	CI 95%	<i>p</i>	Adjusted R ²
Short physical performance battery (points)	Phase angle (°)	1	0.544	0.474–1.261	<0.001	0.281
		2	0.324	0.058–0.977	0.028	0.359
		3	0.340	0.085–1.000	0.021	0.370
		4	0.325	0.008–1.028	0.047	0.356
	Overhydration (L)	1	−0.306	−0.913 to −0.042	0.032	0.075
		2	−0.208	−0.701–0.053	0.090	0.331
		3	−0.194	−0.684–0.079	0.117	0.328
		4	−0.206	−0.701–0.059	0.095	0.339
Gait test (seconds)	Phase angle (°)	1	−0.585	−1.439 to −0.607	<0.001	0.329
		2	−0.356	−1.102 to −0.142	0.012	0.417
		3	−0.369	−1.125 to −0.166	0.009	0.424
		4	−0.293	−1.038–0.014	0.056	0.430
	Overhydration (L)	1	0.289	0.015–0.974	0.044	0.064
		2	0.184	−0.089–0.716	0.124	0.365
		3	0.172	−0.114–0.701	0.154	0.359
		4	0.192	−0.063–0.717	0.098	0.418
Sit-to-stand test (seconds)	Phase angle (°)	1	−0.547	−3.226 to −1.228	<0.001	0.285
		2	−0.372	−2.712 to −0.315	0.014	0.329
		3	−0.391	−2.774 to −0.410	0.009	0.353
		4	−0.373	−2.836 to −0.199	0.025	0.339
	Overhydration (L)	1	0.335	0.231–2.429	0.019	0.093
		2	0.246	−0.007–1.965	0.052	0.296
		3	0.229	−0.084–1.901	0.072	0.300
		4	0.242	−0.019–1.945	0.054	0.319

Model 1: no adjustments; Model 2: adjusted by age; Model 3: adjusted by age and diabetes; Model 4: adjusted by age, diabetes, and sex.

Although the relationship between PhA and physical function in PD is still not well-established, there are studies associating PhA with muscle function in other CKD populations. Muscle function, assessed by handgrip strength, was associated with PhA in maintenance hemodialysis (28, 29) and kidney transplant recipients (30). Moreover, in the hemodialysis patients, PhA was associated with exercise tolerance, which was assessed by a 6-min step test, and peak torque of knee extensors (28).

A possible explanation for the relationship between PhA and physical function is based on cellular structures and its interaction with the electrical current. The structure and the function of cells are dependent on cell membranes, which are composed of electrically non-conducting lipid bilayers with associated proteins (31). The protein–lipid–protein

sandwiched structure provides a reactance to the applied alternating current (32). The reactance arises from cell membranes, while resistance from extra- and intracellular fluid (17). The impedance is the result of two vectors representing resistance and reactance. PhA is the angle between impedance and resistance. Therefore, the higher the reactance, the larger the PhA will be for a given resistance (33). In other words, the higher PhA, the better cell membrane structure, cell mass, cellular integrity, and cell function.

The main limitation of this study is the single-center small sample size, which does not allow the inclusion of many variables in the multiple analyses. However, due to the scarcity of studies assessing the association of PhA and OH with physical function

in PD, the results may be significant for PD patients. Thus, studies with a larger number of patients are still necessary.

In conclusion, bioimpedance parameters (PhA and OH) were correlated with physical function tests, but only PhA was independently associated with physical function tests in the PD patients. Inflammatory markers were not associated with physical function in this sample. In this sense, easy-to-apply tools with good accuracy are important to identify the risk of reduced physical function, allowing interventions to avoid worse outcomes.

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 Research Ethics Committee of the Faculty of

Medicine of Botucatu-Universidade Estadual Paulista (UNESP). The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

VMS, MZCS, BPV, and JCTC were responsible for the research idea and the study design. VMS, MZCS, NSCR, MSD, and FLC performed the data acquisition. VMS, MZCS, and BPV performed the data analysis and the interpretation, were involved in the statistical analysis, and drafted the manuscript. BPV, JCTC, and MFM were responsible for supervision and mentorship. All the authors provided intellectual content to the work and gave final approval of the version to be published.

ACKNOWLEDGMENTS

We acknowledge the patients and health care staff from the Dialysis Unit of the Clinics Hospital of Botucatu Medical School.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Agreement of Single-Frequency Electrical Bioimpedance in the Evaluation of Fat Free Mass and Fat Mass in Peritoneal Dialysis Patients

Nayrana Soares do Carmo Reis, Francieli Cristina Delatim Vaninni, Maryanne Zilli Canedo Silva, Rogério Carvalho de Oliveira, Fabrício Moreira Reis, Fabiana Lourenço Costa, Luis Cuadrado Martin and Pasqual Barretti*

Internal Medicine Department, Botucatu Medical School, São Paulo State University, Botucatu, Brazil

OPEN ACCESS

Edited by:

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*Correspondence:

Pasqual Barretti
pbarretti@uol.com.br

Specialty section:

This article was submitted to
Clinical Nutrition,
a section of the journal
Frontiers in Nutrition

Received: 27 March 2021

Accepted: 06 May 2021

Published: 31 May 2021

Citation:

Reis NSdC, Vaninni FCD, Silva MZC,
de Oliveira RC, Reis FM, Costa FL,
Martin LC and Barretti P (2021)
Agreement of Single-Frequency
Electrical Bioimpedance in the
Evaluation of Fat Free Mass and Fat
Mass in Peritoneal Dialysis Patients.
Front. Nutr. 8:686513.
doi: 10.3389/fnut.2021.686513

Background: Protein-energy wasting is related to impairment of quality of life and lower survival of end-stage kidney disease (ESKD) patients. The evaluation of body composition, especially fat free mass (FFM) and fat mass (FM), is important for the prediction of outcomes in these individuals. The aim of this study was to compare the FFM and FM measurements obtained by single-frequency bioimpedance (SF-BIA) and by a multiple frequency bioimpedance (MF-BIA) device, using dual energy X-ray absorptiometry (DXA) peritoneal dialysis (PD) patients.

Methods: This was a cross-sectional study involving adult patients undergoing regular PD, in which we performed SF-BIA, MF-BIA, and DXA at the same visit. To compare the bioimpedance values with DXA, we used: Person correlation (r), intraclass correlation coefficient (ICC), and Bland-Altman concordance analysis.

Results: The sample consisted of 50 patients in the PD, with mean age of 55.1 ± 16.3 years. Both bioimpedance methods showed a strong correlation ($r > 0.7$) and excellent reproducibility ($ICC > 0.75$) compared to DXA. According to the Bland-Altman diagram, SF-BIA showed agreement in body compartment measurements, with no proportionality bias ($p > 0.05$), without systematic bias for FFM (-0.5 ± 4.9 , 95% CI -1.8 to 0.9 , $p = 0.506$), and for FM (0.3 ± 4.6 , $p = 0.543$). MF-BIA did not present a proportionality bias for the FFM, but it underestimated this body compartment by 2.5 ± 5.4 kg ($p = 0.002$). In addition, MF-BIA presented proportionality bias for FM.

Conclusion: SF-BIA was a more accurate assessing method than MBIA for FFM and FM measurements in PD patients. Because it is a low-cost, non-evaluator-dependent measurement and has less systematic bias, it can also be recommended for fat mass and free-fat mass evaluation in PD patients.

Keywords: nutrition, dialysis, peritoneal dialysis, electrical bioimpedance, dual energy x-ray absorptiometry

INTRODUCTION

Protein-energy wasting (PEW) is a common condition in end-stage kidney disease (ESKD) patients. PEW is reported in 8–54% of dialysis patients and is strongly associated with adverse clinical outcomes, as an increased hospitalization rate and lower survival and has prevention and deceleration of difficult management in these individuals (1, 2). On the other hand, higher body mass index (BMI) values have been associated with better outcomes, in contrast to the association in the general population. This phenomenon has been referred to as the “obesity paradox” (3). However, the causes of this possible protective effect are still unclear since BMI does not provide accurate information about body composition or which compartment has the greatest protective effect, and this measure is likely to perform worse in dialysis patients than in the general population (4–6).

This context highlights the importance of assessing body composition to monitor and predict outcomes in ESKD patients. There are several tools for assessing body composition, with emphasis on dual energy X-ray absorptiometry (DXA), which is considered a reference method capable of more reliably estimating bone, fat, and muscle mass (7, 8). However, few clinics have access to this method, since it requires radiological medicine facilities, generating a high cost for its routine use (9).

Among the most accessible methods are electrical bioimpedance (BIA) analyses. It consists of a non-invasive and relatively low-cost method, which is easy-to-use and portable, and does not require a skilled operator, allowing reproducible results. However, this measurement can be influenced by factors related to ESKD, such as hydration status (8, 10).

BIA methods can be classified into two main categories: single-frequency BIA (SF-BIA) and multiple frequency BIA (MF-BIA). SF-BIA normally operates at a frequency of 50 kHz. This frequency and the impedance are directly proportional the total amount of body water and allows, subsequently, to establish estimates of fat-free mass. In this model, the body is divided into two parts: fat mass (FM) and fat free mass (FFM), with FM defined indirectly as the difference between body weight and FFM. This model assumes that FFM has a constant hydration of 73%. MF-BIA is based on the principle that analyzes of body compartments, using specific frequencies, yield more accurate results. It was found that at low frequencies the current moves around the cells, while at high frequencies the current penetrates the cells. Therefore, proposed that ECW should be estimated at low frequencies (5 kHz), while ICW should be estimated at high frequencies (1 MHz) (11). However, there is no consensus on which of these methods is more reliable for the assessment of body compartments such as fat-free mass (FFM) and fat mass (FM) in dialysis patients, especially peritoneal dialysis (PD), and contradictory results have been reported (12–16).

A recent update to the Clinical Practice Guideline for Nutrition in Chronic Kidney Disease (KDOQI) (8) reports that there is insufficient evidence to suggest the use of bioelectrical impedance to assess body composition of PD patients and recommends conducting future research in this group to determine the validity and reliability of these measurements in PD patients. Therefore, the aim of this study was to compare the

FFM and FM measurements obtained by SF-BIA and a MF-BIA device, using DXA as the reference standard, in PD patients.

MATERIALS AND METHODS

This cross-sectional study was approved by our Institutional Ethics and Research Committee (CAAE 39704314.3.0000.5411) and involved adult end-stage kidney disease (ESKD) patients undergoing PD for at least 90 days at a single Brazilian university center. The sample size calculation was performed based on a pilot study, estimating a minimum correlation coefficient of 0.6 between the tested methods and DXA, with statistical power of 0.9 and an alpha error of 0.05.

Patients' enrollment occurred between January 2017 and May 2018. All eligible patients were invited to participate of the research and those included signed their informed consent term. We did not include patients under 18 years, with cardiac pacemakers, implanted defibrillators, and those with limb amputation, because these factors would make the nutritional assessments unreliable. We performed all body composition in patients without abdominal dialysate. A skilled examiner performed SF-BIA, MF-BIA, and DXA at the same time, with a maximum interval of 2 h between the assessments.

Dual Energy X-Ray Absorptiometry (DXA)

FFM and FM were quantified by DXA using the Hologic® Discovery a device. The integrated software calculated lean mass (kg), FM (kg), bone mineral content (kg), lean mass index (kg/m²), and fat mass index (kg/m²). FFM was considered the sum of lean mass and bone mineral content.

Electrical Bioimpedance

For the SF-BIA assessments, we used a Biodynamics® model 450, 800 μ A, 50 kHz device and evaluated FFM (kg), FM (kg), and phase angle. The equations used to assess these measurements were based on the Kushner & Scholler proposals (16). We performed MF-BIA using the Fresenius Medical Care® Body Composition Monitor, and the body compartment measurements were estimated using a specific software provided by the manufacturer, whose formulas are based on those proposed by Moissl et al. (17). BCM assumes a division of body mass into 3 compartments: lean tissue mass (LTM), adipose tissue mass (ATM), and hyperhydration index (OH). In addition, it provides fat (kg) values consisting of hydration-free fat tissue. For comparison purposes with DXA (which divides the body into 2 body compartments), we initially used: $FFM = LTM \text{ (kg)} + OH \text{ (L)}$ and $FFM = \text{total body weight (kg)} - \text{fat (kg)}$; $FM = ATM \text{ (kg)}$ and $FM = \text{fat (kg)}$. The analyzes using $FFM = LTM \text{ (kg)} + OH \text{ (L)}$ and $FM = ATM \text{ (kg)}$ presented broader agreement limits, with greater need for adjustment. Therefore, our analyzes for BCM were based on the division of body compartments using the fat (kg) measurement provided by the device.

Statistical Analysis

The results were expressed as the mean \pm standard deviation, median (interquartile range) or percentage. Continuous variables with a normal distribution were analyzed using Student's *t*-test,

and those with a non-normal distribution by the non-parametric Mann-Whitney test. To compare categorical variables, we used the Chi square test.

The correlation strength between the measurements was calculated using Pearson's correlation coefficient, while the agreement between the methods was evaluated using the intraclass correlation coefficient (ICC) and the Bland-Altman analysis. The following criteria were considered for ICC: ICC < 0.4, poor reproducibility; $0.4 \leq \text{ICC} < 0.75$, satisfactory reproducibility; and $\text{ICC} \geq 0.75$, excellent reproducibility.

We used Bland-Altman analysis to determine the systematic and proportionality bias between the values obtained by the tested methods and DXA. Systematic bias was assessed using Student's *t*-test for one sample, checking if the average of the differences between the methods would be equal to "0." The proportionality bias was assessed using linear regression to determine if the difference between the methods was biased by the magnitude of the measure, considering the difference between DXA and the tested method values as a dependent variable and the mean of them as an independent variable. The presence of proportionality bias classified the method as not in agreement with DXA.

The total population and subgroups were assessed, in which patients were divided according to sex, median age, and median BMI. The graphical representation of the Bland-Altman diagram is shown by means of dispersion diagrams. Continuous lines correspond to the average of the differences between the tested methods, and the DXA and dashed black lines represent the limits of agreement with 95% reliability (mean difference $\pm 2 \times$ standard deviation).

We used a linear regression analysis to select predictors for the development of new equations aiming to quantify body compartments, in which the dependent variables were the FFM and FM values obtained by DXA. The equations were established using data from the tested methods in combination with demographic variables (age, sex, weight, and height). All analyzes were made using the IBM SPSS STATISTICS version 23 software, and the criterion of statistical significance corresponded to a $p < 0.05$.

RESULTS

Demographic, Clinical, and Nutritional Characteristics

The flowchart of the patients enrolled in the study is described in **Figure 1**. We included 50 PD patients whose demographic and clinical characteristics are shown in **Table 1**. **Table 2** shows the hydration status and nutritional and laboratory parameters of the study group. A strong correlation ($r > 0.7$) and excellent reproducibility ($\text{ICC} \geq 0.75$) of the body composition measurements were observed between the methods tested and DXA in PD patients (**Supplementary Material**).

Agreement Between SF-BIA and DXA

Figure 2 shows the Bland-Altman agreement diagram for FFM and FM measured by SF-BIA and DXA. SF-BIA presented an accurate assessment of the body compartments, with no

systematic or proportionality bias for FFM (-0.5 ± 4.9 , $p = 0.50$) and FM (0.3 ± 4.6 , $p = 0.543$) (**Figure 2** and **Table 3**). This result was maintained in the subgroup assessment, except in the assessment of patients aged <56 years, in which the values obtained were influenced by the magnitude of the measure (proportionality bias). In addition, considering that the excess ECW is not included in the assessment of ATM, unlike DEXA, which can influence the results, we performed two Pearson's simple linear regression analysis between the difference between BCM (FFM and FM) and DEXA measurements with OH. The linear regression coefficient (*r*) regarding FFM and OH and FM and OH was, respectively, -0.2 ($p = 0.115$) and 0.2 ($p = 0.093$) (**Supplementary Material**).

Agreement Between MF-BIA and DXA

Despite not showing proportionality bias, the MF-BIA values underestimated FFM by 2.5 ± 5.4 kg in PD patients (**Table 4**). For this compartment, we found agreement between MF-BIA and DXA only for men and patients with BMI ≥ 25 kg/m². For FM, we observed proportionality bias (**Figure 3** and **Table 4**). In men, MF-BIA values agreed to the reference standard. In those aged ≥ 56 years, FM was overestimated by 2.9 ± 5.4 kg.

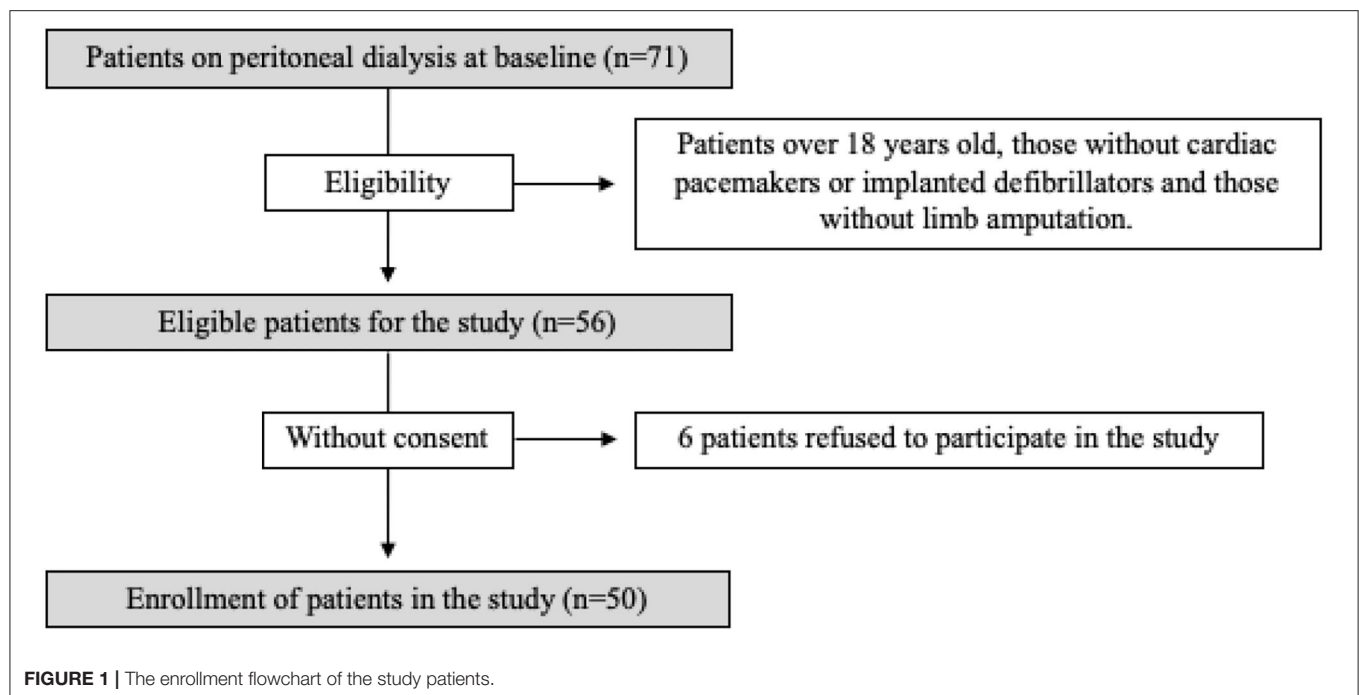
Predictors of FFM and FM

To minimize the systematic errors observed in most evaluations, we developed equations for the prediction of FFM and FM in PD patients (**Chart 1**), based on regression analyses (**Supplementary Material**). Since there was agreement in the PD patients regarding the values of FFM and FM obtained through SF-BIA and DXA, it was not necessary to construct adjustment equations for this subgroup.

DISCUSSION

To our knowledge, this study was the first to test the agreement of FFM and FM measurements performed by SF-BIA and MF-BIA methods against a reference standard, such as DXA, in PD patients. Our results showed that the correlations between these methods and DXA were strong, as described in previous reports (10, 18). However, the correlation coefficient alone is not sufficient to suggest agreement between methods; an adequate concordance analysis is required. Our results showed that SF-BIA can be considered reliable in FFM and FM assessment in PD patients. This finding has great importance, since nutritional status is associated with dialysis outcomes (19). Previous studies by our group have shown that BIA measurements are associated with cardiovascular outcomes (20).

Huang et al. (21) followed up patients for ~ 8 years, reporting that greater FFM was able to predict better patient and technique survival in PD patients. Kang et al. (22) showed an association of low appendicular mass (assessed by DXA) with all-cause mortality in incident PD patients. A recent systematic review and meta-analysis, including four studies and more than 50,000 PD patients, showed a higher mortality risk in patients with lower BMI values (23). However, the BMI *per se* does not discriminate body components such as FFM and FM, which can be estimated by more specific methods.

**TABLE 1 |** Demographic and clinical characteristics of study patients.

	Peritoneal dialysis			<i>p</i>
	Total (<i>n</i> = 50)	Man (<i>n</i> = 23)	Women (<i>n</i> = 27)	
Age (y)	55.1 ± 16.3	54.4 ± 15.4	55.7 ± 17.3	0.787
Race/ethnicity (%)				
Caucasian	90.0	100.0	85.2	0.316
Black	2.0	0.0	3.7	
Brown	2.0	0.0	7.4	
Yellow	6.0	0.0	3.7	
Scholarity (<9 years of study)	64.0	65.2	65.4	0.990
Dialysis vintage (months)	9 (5–17)	7 (5–14)	11 (6–18)	0.163
CKD etiology (%)				
DM	22.0	30.4	14.8	0.658
SAH	16.0	21.7	37.5	
GCN	16.0	8.7	22.2	
Others	46.0	39.2	25.5	
SBP	140 (120–160)	140 (120–170)	140 (120–150)	0.285
DBP	80 (70–100)	80 (70–100)	80 (70–100)	0.889

Values are presented as mean ± standard deviation, median, and interquartile range or percentage. *p* < 0.05. CKD, Chronic kidney disease; DM, Diabetes Mellitus; SAH, Systemic Arterial Hypertension; GCN, Glomerulus Chronic Nephritis; DBP, Diastolic Blood Pressure; SBP, Systolic Blood Pressure.

PD patients have less variation in hydration status due to the continuous nature of their therapy (24), which can contribute significantly to agreement of body compartments using the instruments available in clinical practice, especially when compared to HD patients, who normally retain 1–4 liters over the interdialytic interval (10).

In the evaluation of the FFM, the MF-BIA did not present proportionality bias; however, it underestimated this measure.

As the error occurred systematically, the measurement of FFM by these methods can also be considered possible, provided that adjustments are made to the values obtained. Konings et al. (25) found wide limits of agreement between MF-BIA and DXA, when assessed the body composition of 40 PD patients, with a strong influence on the hydration status. Differently, there was a proportional bias between DEXA and FM, which could be a consequence of the excess ECW not being included in

TABLE 2 | Hydration status and nutritional and laboratory parameters of study patients.

	Peritoneal dialysis			<i>p</i>
	Total (<i>n</i> = 50)	Man (<i>n</i> = 23)	Woman (<i>n</i> = 27)	
BMI (kg/m ²)	25.8 ± 4.3	26.9 ± 4.1	24.8 ± 4.3	0.079
FFM_SF-BIA (kg)	47.8 ± 11.6	57.0 ± 11.1	41.6 ± 6.1	<0.001
FM_SF-BIA (kg)	20.3 ± 7.6	21.1 ± 8.3	19.6 ± 7.0	0.493
LTM_MF-BIA (kg)	37.5 ± 10.8	43.5 ± 10.0	28.8 ± 5.8	<0.001
ATM_MF-BIA (kg)	31.5 ± 11.2	31.7 ± 11.1	31.3 ± 11.4	0.897
Fat_MF-BIA (kg)	23.1 ± 8.2	23.3 ± 8.2	23.0 ± 8.4	0.901
FFM (TBM-Fat)_MF-BIA (kg)	45.8 ± 11.8	54.7 ± 10.7	38.2 ± 5.9	<0.001
FFM (LTM+OH)_MF-BIA (kg)	36.3 ± 10.8	44.3 ± 9.8	29.5 ± 5.6	<0.001
OH	0.8 ± 1.1	0.9 ± 1.1	0.7 ± 1.1	0.517
OH/ECW (%)	5.0 ± 7.2	5.0 ± 6.0	5.0 ± 8.2	0.975
FFM_DXA (kg)	48.2 ± 11.1	56.6 ± 9.2	41.1 ± 6.0	<0.001
FM_DXA (kg)	20.6 ± 6.4	21.1 ± 6.7	20.3 ± 6.3	0.647
FFM_DXA (%)	69.7 ± 6.1	72.9 ± 6.0	67.3 ± 5.0	0.001
FM_DXA (%)	30.1 ± 6.1	27.1 ± 6.0	32.7 ± 5.0	0.001
LTI_DXA (kg/m ²)	18.0 ± 2.7	19.6 ± 2.6	16.7 ± 2.0	<0.001
FTI_DXA (kg/m ²)	7.8 ± 2.5	7.3 ± 2.3	8.3 ± 2.5	0.184
Albumin (g/dl)	3.6 ± 0.4	3.8 ± 0.4	3.5 ± 0.4	0.012
Creatinine (mg/dl)	9.1 ± 2.9	9.9 ± 3.1	8.4 ± 2.6	0.078
Hemoglobin (g/dl)	11.1 ± 1.8	11.8 ± 2.0	10.5 ± 1.3	0.008
TC (mg/dl)	151 ± 31	142 ± 33	160 ± 27.8	0.042
CRP (mg/dl)	0.6 (0.5–0.9)	0.5 (0.5–1.1)	0.6 (0.5–0.9)	0.925

Values are presented as mean ± standard deviation, median and interquartile range. *p* < 0.05. ATM, Adipose tissue mass; ECW, Extracellular water; TC, total cholesterol; DXA, dual energy x-ray densitometry; BMI, body mass index; FTI, Fat tissue index; LTI, Lean tissue index; FM, Fat mass; FFM, Fat-free mass; LTM, Lean tissue mass; OH, hyperhydration index; RCP, C-reactive protein; TG, triglyceride; TBW, total body mass; %, percentage.

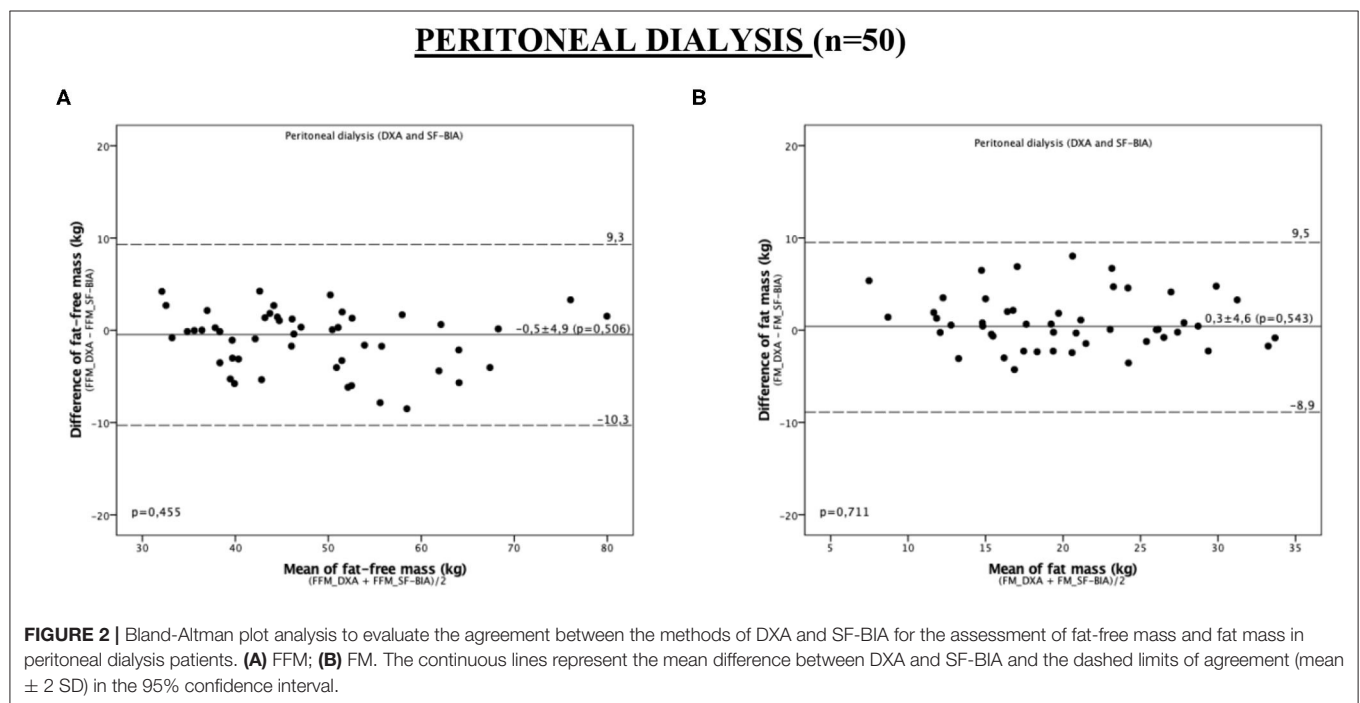


TABLE 3 | Agreement between the measurements obtained by DXA and single-frequency bioimpedance in the assessment of body composition in study patients.

DXA vs. SF-BIA	Peritoneal dialysis		
	Bias	Systematic bias	Proportionality bias
		IC 95%	P
Total (N = 50)			
Fat-free mass (kg)	-0.5 ± 4.9	-1.8 a 0.9	0.506
Fat mass (kg)	0.3 ± 4.6	-0.9 a 1.7	0.543
Man (n = 23)			
Fat-free mass (kg)	-0.4 ± 6.7	-3.2 a 2.5	0.789
Fat mass (kg)	0 ± 6.2	-2.6 a 2.7	0.973
Women (n = 27)			
Fat-free mass (kg)	-0.5 ± 2.6	-1.6 a 0.5	0.298
Fat mass (kg)	0.7 ± 2.6	-0.3 a 1.7	0.178
Age < 56 years (n = 24)			
Fat-free mass (kg)	-0.5 ± 6.3	-3.2 a 2.1	0.681
Fat mass (kg)	0.7 ± 6.0	-1.8 a 3.2	0.564
Age ≥ 56 years (n = 26)			
Fat-free mass (kg)	-0.4 ± 3.2	-1.7 a 0.9	0.535
Fat mass (kg)	0.1 ± 2.8	-1.0 a 1.2	0.854
BMI < 25 kg/m² (n = 24)			
Fat-free mass (kg)	-0.7 ± 3.0	-2.0 a 0.6	0.263
Fat mass (kg)	0.8 ± 3.0	-0.5 a 2.1	0.204
BMI ≥ 25 kg/m² (n = 26)			
Fat-free mass (kg)	-0.2 ± 6.2	-2.7 a 2.3	0.844
Fat mass (kg)	0 ± 5.7	-2.3 a 2.3	0.980

$p < 0.05$. PD, Peritoneal dialysis; DXA, dual energy x-ray densitometry; CI, Confidence interval; BMI, body mass index; SF-BIA, Single-frequency bioimpedance.

TABLE 4 | Agreement between the measurements obtained by DXA and multiple-frequency bioimpedance in the assessment of body composition in study patients.

DXA vs. MF-BIA	Peritoneal dialysis		
	bias	Systematic bias	Proportionality bias
		IC 95%	p
Total (N = 50)			
Fat-free mass (kg)	2.5 ± 5.4	0.9 a 4.0	0.002
Fat mass (kg)	-2.5 ± 4.9	-3.9 a -1.1	0.001
Man (n = 23)			
Fat-free mass (kg)	2.0 ± 6.6	-0.9 a 4.8	0.166
Fat mass (kg)	-2.2 ± 5.8	-4.7 a 0.3	0.087
Women (n = 27)			
Fat-free mass (kg)	2.9 ± 4.1	1.2 a 4.5	0.001
Fat mass (kg)	-2.8 ± 4.1	-4.3 a -1.1	0.002
AGE < 56 years (n = 24)			
Fat-free mass (kg)	2.2 ± 5.0	0.1 a 4.3	0.041
Fat mass (kg)	-2.0 ± 4.4	-3.9 a -0.1	0.036
AGE ≥ 56 years (n = 26)			
Fat-free mass (kg)	2.7 ± 5.8	0.4 a 5.1	0.025
Fat mass (kg)	-2.9 ± 5.4	-5.1 a -0.7	0.011
BMI < 25 kg/m² (n = 24)			
Fat-free mass (kg)	2.2 ± 3.1	0.9 a 3.6	0.002
Fat mass (kg)	-2.0 ± 3.1	-3.3 a -0.7	0.004
BMI ≥ 25 kg/m² (n = 26)			
Fat-free mass (kg)	2.7 ± 6.9	-0.1 a 5.5	0.058
Fat mass (kg)	-2.9 ± 6.2	-5.4 a -0.4	0.024

$p < 0.05$. PD, Peritoneal dialysis; DXA, dual energy x-ray densitometry; CI, Confidence interval; BMI, body mass index; MF-BIA, Multiple frequency bioimpedance.

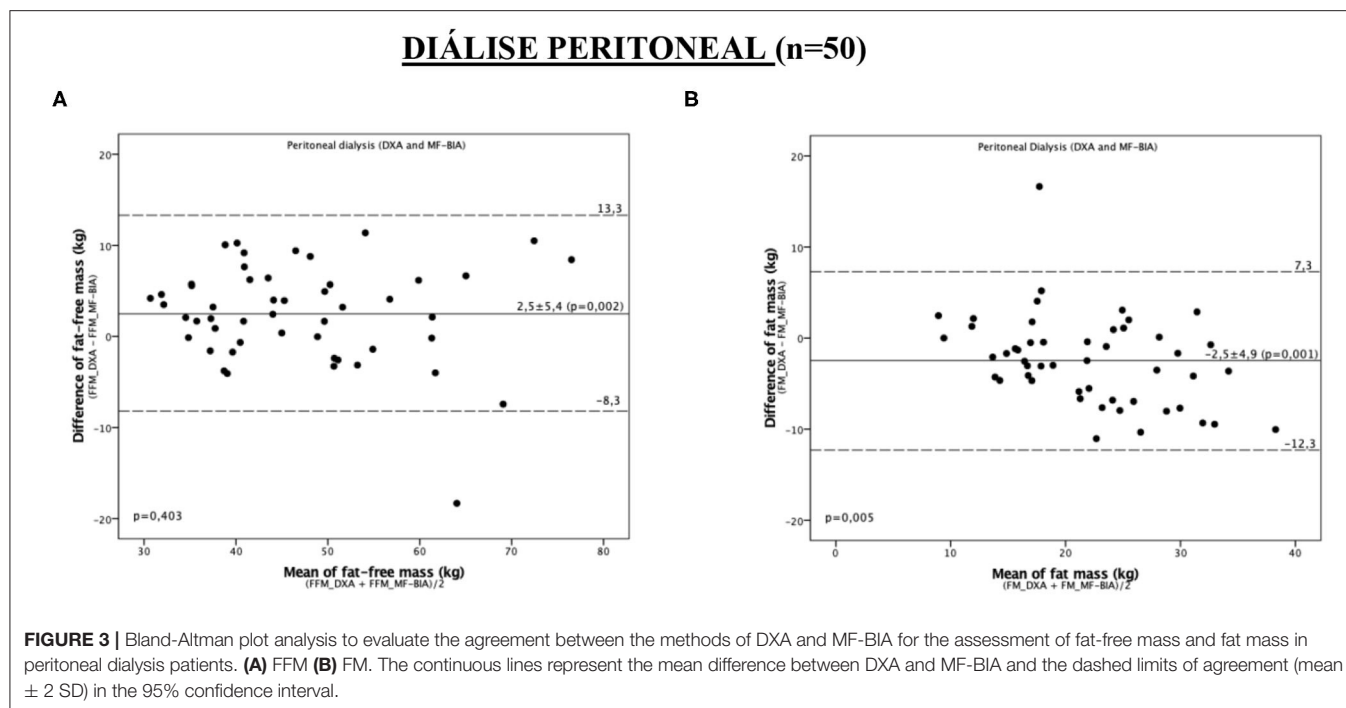


CHART 1 | Prediction equations for FFM and FM in PD patients.

FAT-FREE MASS

$FFM_{MF-BIA} = -22,972 + 0,039 \times \text{age (years)} - 3,077 \times \text{sex (0 if M; 1 if F)} + 0,498 \times \text{weight (kg)} + 15,886 \times \text{height (m)} + 2,244 \times \text{PA (frequency 50)} + 2,011 \times \text{OH}.$

FAT MASS

$FM_{MF-BIA} = \text{weight (kg)} - [-22,972 + 0,039 \times \text{age (years)} - 3,077 \times \text{sex (0 if M; 1 if F)} + 0,498 \times \text{weight (kg)} + 15,886 \times \text{height (m)} + 2,244 \times \text{PA (frequency 50)} + 2,011 \times \text{OH}].$

PA, Phase angle; MF-BIA, Multifrequency Bioimpedance; SF-BIA, Unifrequency Bioimpedance; M, Male; F, Female; FM, Fat mass; FFM, Fat-free mass; OH, hyperhydration index.

the assessment of FM, unlike DEXA. However, the absence of significant correlation between the differences of BCM and DEXA FM measurements and OH contradicts this possibility.

Since our results showed low agreement with MF-BIA in the body composition assessment with DXA, we constructed predictive equations to quantify FFM and FM in PD patients. The main differences in body composition measurements between BIA devices are prediction equations and alternating current frequencies. While SF-BIA analysis depends on the use of regression models, MF-BIA devices often use fit equations for a polynomial curve. In addition, each device is calibrated using its own equation and software. Therefore, despite the theoretical expectation of MF-BIA devices being more promising and reliable from a clinical perspective for the assessment of body composition, we suggest that single frequency devices are not inferior to multifrequency devices, and this has already been shown in previous studies (26). However, some adjustments in mathematical models can be applied in order to improve the agreement of the MF-BIA measures, such as the formulas we propose in our study, despite the need for validation. Predictive equations can have great relevance due to the impossibility of routine evaluation of body compartments using a reference

method such as DXA due to its high complexity and high costs. However, it is important to highlight that the validation of the new equations still needs to be performed in a larger and independent population sample.

A potential limitation of our study is the absence of a gold standard in the assessment of body FFM in dialysis patients, since DXA assumes a constant hydration value. Despite this, DXA is still considered as the reference method by the KDOQI guideline (8) for assessment of body composition in ESKD patients. Its strengths are related to the moment when the evaluations were performed, with all the measurements taken on the same day by a single trained evaluator. In addition, the verification of the agreement was not limited to the evaluation of the systematic bias (as in other existing studies) but also included the objective analysis of the proportionality bias, constituting a more precise analysis.

In conclusion, the current results showed that SF-BIA agreed with DXA in the evaluation of FFM and FM in PD patients, in opposite to MF-BIA measurements. Because its low-cost and being a non-examiner dependent method, SF-BIA can be recommended for the evaluation of free-fat mass and fat mass in PD patients.

DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/**Supplementary Material**, further inquiries can be directed to the corresponding authors.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by Institutional Ethics and Research Committee of the Botucatu Medical School, São Paulo State University (Brazil). The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

NR, FV, and PB were responsible for conceptualization and methodology. NR, MS, FR, and FC performed data acquisition. NR, RO, and LM performed data analysis and interpretation. NR and FV were involved in formal analysis and wrote the original draft. NR, FV, MS, FR, LM, and PB revised and edited the

manuscript. FV, LM, and PB were responsible for supervision and mentorship. All authors provided intellectual content of critical importance to the work and gave final approval of the version to be published.

FUNDING

This study was supported by National Council for Scientific and Technological Development (CNPq).

ACKNOWLEDGMENTS

Our acknowledgments to all patients and the team from the Dialysis Unit of the Clinics Hospital of Botucatu Medical School.

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fnut.2021.686513/full#supplementary-material>

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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The Association Between Dietary Inflammatory Index and Parathyroid Hormone in Adults With/Without Chronic Kidney Disease

Zheng Qin, Qinbo Yang, Ruoxi Liao and Baihai Su*

Department of Nephrology, National Clinical Research Center for Geriatrics, West China Hospital, Sichuan University, Chengdu, China

OPEN ACCESS

Edited by:

Carla Maria Avesani,
Karolinska Institutet, Sweden

Reviewed by:

Alisson Diego Machado,
University of São Paulo, Brazil
Rosa Maria Affonso Moyses,
University of São Paulo, Brazil

*Correspondence:

Baihai Su
imsbh@163.com

Specialty section:

This article was submitted to
Clinical Nutrition,
a section of the journal
Frontiers in Nutrition

Received: 30 March 2021

Accepted: 01 June 2021

Published: 25 June 2021

Citation:

Qin Z, Yang Q, Liao R and Su B (2021)
The Association Between Dietary
Inflammatory Index and Parathyroid
Hormone in Adults With/Without
Chronic Kidney Disease.
Front. Nutr. 8:688369.
doi: 10.3389/fnut.2021.688369

Aims: We aimed to assess the association between dietary inflammation index (DII) with parathyroid hormone (PTH) and hyperparathyroidism (HP) in adults with/without chronic kidney disease (CKD).

Methods: Data were obtained from the 2003–2006 National Health and Nutrition Examination Survey (NHANES). The participants who were <18 years old, pregnant, or missing the data of DII, PTH, and CKD were excluded. DII was calculated based on a 24-h dietary recall interview for each participant. Weighted multivariable regression analysis and subgroup analysis were conducted to estimate the independent relationship between DII with PTH and the HP in the population with CKD/non-CKD.

Results: A total of 7,679 participants were included with the median DII of -0.24 (-2.20 to 1.80) and a mean PTH level of 43.42 ± 23.21 pg/ml. The average PTH was 45.53 ± 26.63 pg/ml for the participants in the highest tertile group compared with 41.42 ± 19.74 pg/ml in the lowest tertile group ($P < 0.0001$). The rate of HP was 11.15% overall, while the rate in the highest DII tertile was 13.28 and 8.60% in the lowest DII tertile ($P < 0.0001$). The participants with CKD tended to have higher PTH levels compared with their counterparts (61.23 ± 45.62 vs. 41.80 ± 19.16 pg/ml, $P < 0.0001$). A positive association between DII scores and PTH was observed ($\beta = 0.46$, 95% CI: 0.25, 0.66, $P \leq 0.0001$), and higher DII was associated with an increased risk of HP (OR = 1.05, 95% CI: 1.02, 1.08, $P = 0.0023$). The results from subgroup analysis indicated that this association was similar in the participants with different renal function, gender, age, BMI, hypertension, and diabetes statuses and could also be appropriate for the population with CKD.

Conclusions: Higher consumption of a pro-inflammatory diet appeared to cause a higher PTH level and an increased risk of HP. Anti-inflammatory dietary management may be beneficial to reduce the risk of HP both in the population with and without CKD.

Keywords: dietary inflammatory index, parathyroid hormone, chronic kidney disease, hyperparathyroidism, National Health and Nutrition Examination Survey

INTRODUCTION

Parathyroid hormone (PTH) is a single-stranded peptide hormone, containing 84 amino acids, which are synthesized and secreted by the chief cells of the parathyroid gland, with the main function of increasing the serum Ca^{2+} and decreasing the serum phosphorus levels (1). The secretion of PTH is also mainly regulated by the concentration of serum Ca^{2+} and phosphorus (2, 3). Serum Ca^{2+} regulates PTH secretion through the interaction with calcium-sensitive receptors (CASR) on the surface of parathyroid cells (4, 5). Serum phosphorus enhances the stability of PTH mRNA and stimulates the proliferation of parathyroid cells to increase the secretion of PTH both directly and indirectly (3, 6). Bone and kidney are the main target organs of PTH (7–9). For a variety of pathological reasons, the parathyroid glands can secrete excessive PTH and cause hyperparathyroidism (HP), which can be classified as primary, secondary, and tertiary (10). HP appeared to be associated with an increased risk of poor clinical outcomes and death, which is often observed in patients with chronic kidney disease (CKD) (11).

Chronic kidney disease refers to a chronic clinical condition of renal, structural, and functional disorders and is characterized by a higher inflammation status (12). Recent studies have revealed the global prevalence of CKD to be about 10%, as well as the increasing disease burdens (13–15). In patients with CKD, abnormal regulation in calcium, phosphorus, vitamin D, and PTH is accompanied by a decline in renal function, which could lead to secondary HP, which is associated with the increased risk of fracture, cardiovascular disease, and death (16–18). Thus, the management of the PTH level in patients with CKD is of great significance.

The association between inflammation and PTH remains unclear. Several animal and human studies indicated that the PTH level may be associated with inflammation (19–21). Previous studies observed a decreased inflammation status after parathyroidectomy in patients with HP (22, 23). However, the inflammation level after parathyroidectomy varies widely among studies; both increased (24) and even no-change results (25) have been reported before. Chen et al. (19) found that inflammatory markers, including C-reactive protein (CRP), red cell distribution width (RDW), and platelet-to-lymphocyte ratio (PLR) levels increased with increasing serum PTH concentration in the US adults, indicating a positive relationship between inflammation and PTH. In *in vitro* studies, several inflammatory cytokines, such as interleukin-8 (IL-8), tumor necrosis factor- α (TNF- α), etc., have been proved to enhance PTH synthesis and secretion through nuclear factor- κ B (NF- κ B) and affect CASR transcription (26–28). It could be inferred that the consumption of an inflammatory diet may also have an impact on the PTH. Dietary inflammatory index (DII) was a literature-derived and population-based scoring system designed to evaluate the inflammatory potential of diets (29). A positive value for DII corresponded to a pro-inflammatory diet, and a negative value for DII corresponded to an anti-inflammatory diet. The higher scores suggested a more pro-inflammatory effect, and the more negative scores suggested a more anti-inflammatory effect. In

fact, previous studies indicated a significant association between DII and the risk of various cancers (30, 31), obesity (32), cardiovascular disease (33), sarcopenia (34), and so on. However, the association between the dietary inflammatory potential and PTH has not been reported before.

In this study, we aimed to assess the impact of DII on PTH and HP, using data from the National Health and Nutrition Examination Survey (NHANES). We hypothesized that the increased consumption of pro-inflammatory diet would be associated with higher PTH levels and increased risk of HP. In addition, with regard to the fact that patients with CKD may have more elevated PTH levels than the population with non-CKD, we further examined this association in subgroups stratified by renal function.

MATERIALS AND METHODS

Study Population

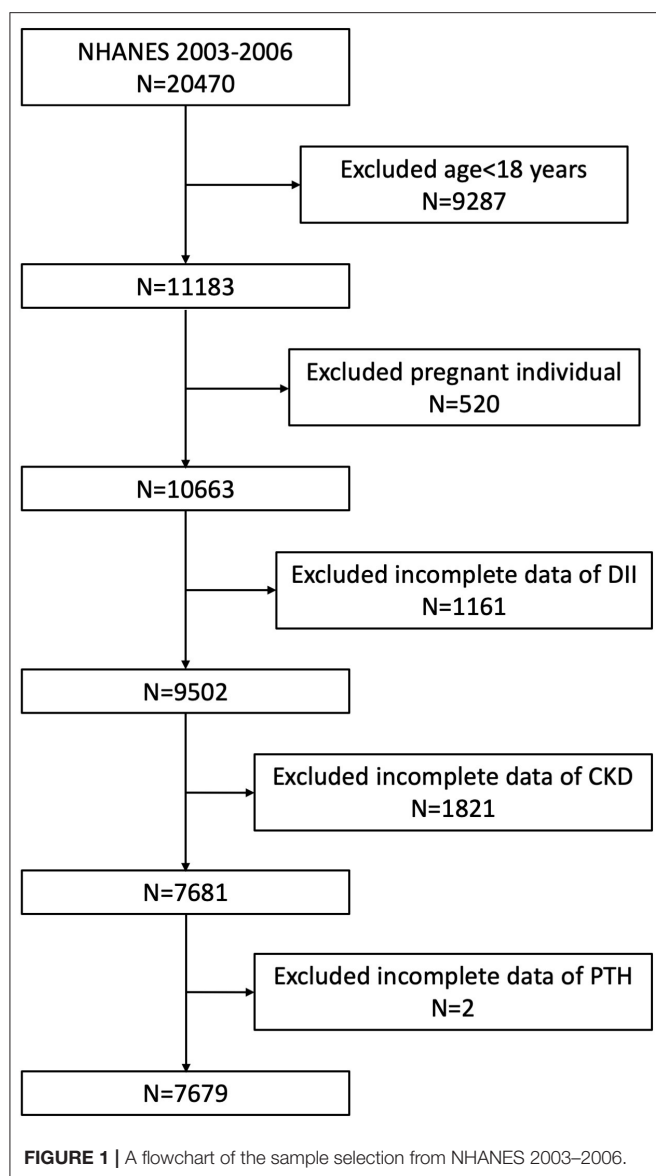
In this study, we obtained data from the NHANES. NHANES is a program of studies administered by the National Center for Health Statistics (NCHS), part of the U.S. Centers for Disease Control and Prevention (CDC), aimed to assess the health and nutrition status of the U.S. population through interviews, physical examinations, and laboratory tests. The NHANES is conducted on a 2-year cycle, and the data are still being updated. Because this study adopted a stratified multistage probability sampling method, the included samples had good representativeness (35). All NHANES data are publicly available at www.cdc.gov/nchs/nhanes/.

This study was based on the data from two 2-year NHANES surveys from 2003 to 2006. A total of 20,470 participants were enrolled at first; after the exclusion of individuals aged <18 years old ($n = 9,287$), who were pregnant ($n = 520$), missing the dietary data relating to DII ($n = 1,161$), missing the data of diagnosing CKD ($n = 1,821$), and missing the data of PTH ($n = 2$), 7,679 participants were included in our final analysis (Figure 1).

The NCHS Ethics Review Board granted the human subject approval for the conduction of NHANES, and written informed consent was obtained from all the participants.

Exposure and Outcome Definitions

Dietary inflammation index was designed as an exposure variable. The dietary data in NHANES were obtained by a 24-h dietary recall interview at the mobile examination center (MEC) (36), which have been validated by the Nutrition Methodology Working Group before (37). The data of 24-h dietary recall interviews were used to calculate the DII scores for each individual, and it could quantify the inflammatory potential of diets. A higher positive DII score indicated a pro-inflammatory diet, and a lower negative DII score indicated an anti-inflammatory diet (29). A total of 27 food parameters were available in NHANES and were used for the calculation of DII, including anti-inflammatory food parameters (alcohol, β -carotene, fibers, folic acid, magnesium, zinc, selenium, vitamin A, vitamin B-6, vitamin C, vitamin E, monounsaturated fatty acid, niacin, riboflavin, n-3 fatty acid, n-6 fatty acid, polyunsaturated fatty acid, caffeine, and thiamin), and pro-inflammatory



food parameters (cholesterol, carbohydrates, energy, fats, iron, vitamin B-12, protein, and saturated fat). Studies have shown that the predictive ability was not affected when <30 dietary parameters were used to calculate DII scores (38–40). DII was analyzed as a continuous variable, and the participants were divided into tertiles from the total sample for further analysis.

The PTH level and HP were designed as outcome variables. The Elecsys 1010 analyzer (Roche Diagnostics) was used to determine the serum intact PTH level. The Elecsys 1010 analyzer was a fully automatic run-oriented analyzer system for the determination of immunological tests, using the ECL/Origen electrochemiluminescent process. All components and reagents for routine analysis were integrated into the analyzer. PTH was measured on the Elecsys 1010, using a sandwich principle. There was no difference in the equipment, lab method, or lab site between the combined two survey cycles in our analysis. HP

was defined as PTH > 65 pg/ml according to previous studies (1, 41). The detailed process of measuring PTH was available at www.cdc.gov/nchs/nhanes/.

Other Study Variables

Baseline variables in this study included age (years), gender, race, educational level, systolic blood pressure (mmHg), diastolic blood pressure (mmHg), body mass index (kg/m²), serum glucose (mg/dl), serum phosphorus (mg/dl), serum iron (ug/dl), serum CRP (mg/dl), serum 25 (OH) D (nmol/L), serum calcium (mg/dl), urinary creatinine (mg/dl), urinary albumin (mg/L), urinary creatinine (mg/dl), eGFR (ml/min/1.73 m²), parathyroid hormone (pg/ml), hypertension, diabetes, albuminuria, low eGFR, and CKD. All detailed measurement processes of study variables were publicly available at www.cdc.gov/nchs/nhanes/.

We calculated urinary albumin: creatinine ratio (ACR) and defined albuminuria as ACR > 30 mg/g. The data about gender, age, and serum creatinine were used to calculate the estimated glomerular filtration rate (eGFR, ml/min/1.73 m²) according to the CKD Epidemiology Collaboration equation for each participant (42), and eGFR lower than 60 ml/min/1.73 m² was used to define low eGFR. We defined CKD as the presence of either albuminuria or low eGFR according to Kidney Disease: Improving Global Outcomes 2012 recommendations (12).

Statistical Analysis

All statistical analyses were conducted according to CDC guidelines (43). All estimates were calculated, accounting for NHANES sample weight. Continuous variables were presented as mean with SD or a median with an interquartile range, and categorical variables were presented as frequency or percentage. Either weighted Student's *t*-test (for continuous variables) or weighted chi-square test (for categorical variables) were conducted to calculate the differences in different DII groups (tertiles). To examine the association between DII and PTH levels, weighted multivariable linear regression explored PTH as a continuous variable, and weighted multivariable logistic regression for HP was used as a categorical variable in three different models. In model 1, no covariates were adjusted. Model 2 was adjusted for gender, age, and race. Model 3 was adjusted for gender, age, race, education level, serum glucose, serum phosphorus, serum iron, serum calcium, serum CRP serum, serum 25 (OH) D, systolic blood pressure, diastolic blood pressure, body mass index, hypertension, diabetes, chronic kidney disease, protein intake, calcium intake, and phosphorus intake. To further explore the association between DII with PTH and HP in different population settings, the subgroup analysis was performed by stratified weighted multivariate regression analysis. In addition, it has been well-recognized that patients with CKD often have elevated parathyroid hormone levels; the CKD status was treated as a prespecified potential effect modifier. An interaction term was added to test the heterogeneity of associations between the subgroups. *P* < 0.05 was considered statistically significant. All analysis was performed using Empower software (www.empowerstats.com; X&Y solutions, Inc., Boston MA) and R version 3.4.3 (<http://www.R-project.org>, The R Foundation).

RESULTS

Baseline Characteristics of Participants

The weighted demographic characteristics and other covariates of included individuals were shown in **Table 1**. A total of 7,679 participants were included in this study, of whom 48.34% were males and 51.66% were females, with an average age of 46.73 ± 18.80 years old. Median DII was -0.24 (-2.20 to 1.80), and the ranges of DII for tertiles 1–3 were -6.60 to -1.50 , -1.50 to 1.08 , and 1.08 to 6.50 , respectively. Among different tertiles of DII, gender, race, education level, systolic blood pressure, diastolic blood pressure, serum iron, serum CRP, serum 25 (OH) D, urinary creatinine, serum creatinine, parathyroid hormone, and HP, whether having hypertension, diabetes, low eGFR and CKD, were significantly different, while no significant difference was observed in BMI, serum glucose, serum phosphorus, serum calcium, urinary albumin, eGFR, and whether having albuminuria between different tertiles. Mean PTH was 43.42 ± 23.21 pg/ml, and the average PTH was 45.53 ± 26.63 pg/ml for the participants in the highest tertile group compared with 41.42 ± 19.74 pg/ml in the lowest tertile group ($P < 0.0001$). The rate of HP was 11.15% overall, while the rate in the highest DII tertile was 13.28 and 8.60% in the lowest DII tertile ($P < 0.0001$).

As for the PTH levels of the participants based on different renal conditions, the participants with albuminuria, low eGFR, and CKD tended to have higher PTH levels compared with their counterparts. In the CKD group, mean PTH was 68.40 ± 55.29 pg/ml for the highest tertile group while 51.12 ± 32.84 pg/ml for the lowest tertile group ($P = 0.0002$). Mean PTH was 68.45 ± 55.36 pg/ml for the highest tertile and 51.42 ± 32.91 pg/ml for the lowest tertile in the low eGFR group ($P = 0.0003$). In the albuminuria group, mean PTH was 131.86 ± 94.31 pg/ml for the highest tertile group while 95.06 ± 110.67 pg/ml for the lowest tertile group; however, there was no significant difference ($P = 0.7248$). Mean eGFR of the patients with CKD was 48.96 ± 10.64 ml/min/1.73 m², and 1.78% of them were <15 ml/min/1.73 m². We also calculated the mean PTH level according to different stages of CKD. The PTH level in the participants with CKD, stages 1 to 5, was 40.50 ± 18.45 , 43.58 ± 19.97 , 56.01 ± 32.16 , 122.30 ± 81.21 , and 239.12 ± 136.89 pg/ml. The participants tended to show a higher PTH level with the CKD progression (**Table 2**).

For the prevalence of HP, participants with reduced renal function tended to have higher rates of HP. In the population with CKD, 28.07% of the participants had HP, while it was 9.61% for those without CKD. Similar results could be observed in the albuminuria and low-eGFR population as well (Albuminuria: 54.86 vs. 11.08%, low eGFR: 28.13 vs. 9.62%) (**Table 2**).

Higher DII Was Associated With Higher PTH Level and Higher Risk of HP

Weighted multivariable regression analysis was conducted to estimate the association of DII with the PTH level and HP in three different models (**Table 3**). The results revealed a positive association between DII scores and PTH with statistical significance (Model 1, $\beta = 0.71$, 95% CI: 0.51, 0.90, $P < 0.0001$;

Model 2, $\beta = 0.58$, 95% CI: 0.38, 0.77, $P < 0.0001$; Model 3, $\beta = 0.46$, 95% CI: 0.25, 0.66, $P \leq 0.0001$). According to the results of the fully adjusted model (Model 3), each unit of the increased DII score was associated with a PTH increase by 46 pg/ml, suggesting that the higher DII scores were associated with a higher PTH level. This association remained statistically significant after DII was grouped as tertiles. The fully adjusted effect size (reference to Tertile 1) was 1.56 (95% CI: 0.28, 2.84, $P = 0.0167$) for Tertile 2 and 2.66 (95% CI: 1.31, 4.01, $P = 0.0001$) for Tertile 3.

In terms of HP, we also observed that increased DII was associated with a higher risk of HP (Model 1, OR = 1.07, 95% CI: 1.05, 1.10, $P < 0.0001$; Model 2, OR = 1.07, 95% CI: 1.04, 1.09, $P < 0.0001$; Model 3, OR = 1.05, 95% CI: 1.02, 1.08, $P = 0.0023$). In Model 3, which adjusted for all covariates, the results indicated that each unit of increased DII score was associated with a 5% increase of risk of HP. In sensitivity analysis, the adjusted OR (reference to Tertile 1) was 1.27 (95% CI: 1.05, 1.55, $P = 0.0157$) for Tertile 2 and 1.32 (95% CI: 1.09, 1.61, $P = 0.0054$) for Tertile 3, suggesting a stable positive relationship between increased DII and higher risk of HP with statistical significance.

Subgroup Analysis

We conducted the subgroup analysis stratified by low eGFR, CKD, gender, age, BMI, hypertension, and diabetes to further explore the association of DII with the PTH level and HP in different population settings by stratified weighted multivariate regression analysis and tested the interactions (**Table 4**). Regarding the correlation between DII scores and the PTH level, the test for interaction was significant for low eGFR (P for interaction = 0.0004) and CKD (P for interaction = 0.0003), indicating significant dependence on renal function. However, the positive association was statistically significant both in subgroups stratified by low eGFR and in subgroups stratified by CKD (all P for trend < 0.05). In subgroups stratified by gender, age, BMI, hypertension, and diabetes, the positive association between DII and PTH was still significant (P for trend < 0.05) and P for interaction > 0.05 , suggesting that the correlation between DII scores and the PTH level was similar in the population with different gender, age, BMI, hypertension status, and diabetes status.

As for the association between DII scores and HP, results of subgroup analysis stratified by CKD or no CKD demonstrated that DII was positively associated with a higher risk of HP (OR = 1.07, P for trend = 0.0023) in the CKD subgroup, and a similar result (OR = 1.04, P for trend = 0.0121) was observed in the non-CKD group. In addition, an interaction test was performed to evaluate if there was any significant dependence of the effect modifier (CKD) on the association. P for interaction > 0.05 means no significant dependence, indicating that the magnitude of the association was the same for the population with/without CKD (P for interaction = 0.4942). Similarly, we did not find any significant dependence on low eGFR, gender, age, BMI, hypertension status, and diabetes status (all P for interaction > 0.05). These results indicated that the positive association between DII scores and HP was similar in the population with different renal function conditions (including CKD and low eGFR), gender, age, BMI, hypertension status, and diabetes status.

TABLE 1 | Weighted baseline characteristics of the participants according to different dietary inflammatory indexes (DIIs).

DII	Overall −0.24 (−2.20~1.80)	Tertile 1 (−6.60~−1.50)	Tertile 2 (−1.50~1.08)	Tertile 3 (1.08~6.50)	P-value
Age (years)	46.73 ± 18.80	45.59 ± 17.88	47.51 ± 19.13	47.26 ± 19.46	0.0003
Gender (%)					
Male	48.34	60.84	46.40	35.06	<0.0001
Female	51.66	39.16	53.60	64.94	
Race (%)					
Mexican American	12.36	12.42	13.10	11.45	0.0003
Other hispanic	3.18	2.92	3.18	3.50	
Non-hispanic white	64.84	66.83	65.22	61.95	
Non-hispanic black	14.77	13.25	13.98	17.53	
Others	4.86	4.58	4.52	5.58	
Education level (%)					
Less than high school	22.34	19.88	22.40	25.31	<0.0001
High school or general educational development	25.39	23.98	25.02	27.56	
Above high school	52.08	55.97	52.43	46.90	
Others	0.18	0.17	0.15	0.23	
Systolic blood pressure (mmHg)	122.59 ± 18.55	121.71 ± 16.48	123.43 ± 19.30	122.76 ± 20.03	0.0043
Diastolic blood pressure (mmHg)	70.18 ± 13.32	71.02 ± 12.86	69.60 ± 13.72	69.78 ± 13.39	0.0002
Body mass index (kg/m ²)	28.20 ± 6.64	28.13 ± 6.47	28.35 ± 6.70	28.11 ± 6.77	0.3685
Glucose, serum (mg/dl)	96.83 ± 30.23	96.63 ± 28.77	96.91 ± 30.54	96.98 ± 31.61	0.9074
Phosphorus, serum (mg/dl)	3.84 ± 0.57	3.85 ± 0.57	3.84 ± 0.57	3.85 ± 0.58	0.7647
Iron, serum (μg/dl)	87.38 ± 36.47	91.77 ± 36.78	85.76 ± 35.00	83.77 ± 37.13	<0.0001
C-reactive protein, serum (mg/dl)	0.41 ± 0.83	0.33 ± 0.63	0.42 ± 0.80	0.48 ± 1.04	<0.0001
25(OH)D, serum (nmol/L)	59.96 ± 22.12	61.73 ± 21.49	60.02 ± 22.31	57.68 ± 22.46	<0.0001
Calcium, serum (mg/dl)	9.52 ± 0.35	9.53 ± 0.35	9.52 ± 0.35	9.51 ± 0.35	0.2978
Hypertension (%)					
Yes	29.58	25.26	32.51	31.71	<0.0001
No	70.42	74.74	67.49	68.29	
Diabetes (%)					
Yes	7.43	5.97	8.69	7.82	<0.0001
No	92.57	94.03	91.31	92.16	
Creatinine, urine (mg/dl)	127.53 ± 79.52	127.38 ± 75.49	123.34 ± 77.40	132.39 ± 86.15	0.0004
Albumin, urine (mg/L)	32.22 ± 225.38	28.30 ± 214.83	33.67 ± 230.78	35.44 ± 231.85	0.4891
Serum creatinine (mg/dl)	0.91 ± 0.29	0.94 ± 0.20	0.91 ± 0.36	0.89 ± 0.30	<0.0001
eGFR (ml/min/1.73 m ²)	91.94 ± 23.47	91.93 ± 21.63	91.79 ± 23.75	92.14 ± 25.27	0.8722
Albuminuria (%)					
Yes	0.16	0.13	0.13	0.22	0.6457
No	99.84	99.87	99.87	99.78	
Low eGFR (eGFR < 60 ml/min/1.73 m ² , %)					
Yes	8.28	6.34	8.64	10.27	<0.0001
No	91.72	93.66	91.36	89.73	
CKD (%)					
Yes	8.33	6.42	8.66	10.30	<0.0001
No	91.67	93.58	91.34	89.70	
Hyperparathyroidism (%)					
Yes	11.15	8.60	12.07	13.28	<0.0001
No	88.85	91.40	87.93	86.72	
Parathyroid hormone (pg/ml)	43.42 ± 23.21	41.42 ± 19.74	43.75 ± 23.29	45.53 ± 26.63	<0.0001

eGFR, estimated glomerular filtration rate; CKD, chronic kidney disease.

TABLE 2 | Parathyroid hormone (PTH) and the prevalence of hyperparathyroidism (HP) based on different renal population settings, weighted.

		DII Tertile 1 (−6.60~−1.50)	DII Tertile 2 (−1.50~1.08)	DII Tertile 3 (1.08~6.50)	P for trend
Parathyroid hormone (pg/ml)					
Albuminuria					
Yes	111.73 ± 89.59	95.06 ± 110.67	99.18 ± 32.35	131.86 ± 94.31	0.7248
No	43.31 ± 22.80	41.35 ± 19.25	43.67 ± 23.19	45.34 ± 25.98	<0.0001
Low-eGFR ^a					
Yes	61.35 ± 45.69	51.42 ± 32.91	61.86 ± 41.57	68.45 ± 55.36	0.0003
No	41.80 ± 19.17	40.74 ± 18.31	42.03 ± 19.91	42.91 ± 19.31	0.0005
CKD ^b					
Yes	61.23 ± 45.62	51.12 ± 32.84	61.91 ± 41.53	68.40 ± 55.29	0.0002
Stage 1	40.50 ± 18.45	39.22 ± 17.38	40.39 ± 18.79	42.19 ± 19.18	<0.0001
Stage 2	43.58 ± 19.97	42.70 ± 19.27	44.29 ± 21.15	43.97 ± 19.44	0.1859
Stage 3	56.01 ± 32.16	48.81 ± 25.91	58.73 ± 36.05	59.15 ± 31.84	0.0011
Stage 4	122.30 ± 81.21	126.26 ± 84.37	93.96 ± 59.24	155.64 ± 89.70	0.0603
Stage 5	239.12 ± 136.89	75.00 ± 0.00	169.02 ± 124.24	265.17 ± 131.93	0.5695
No	41.80 ± 19.16	40.76 ± 18.31	42.02 ± 19.91	42.91 ± 19.31	0.0006
Hyperparathyroidism (%)					
Albuminuria					
Yes	54.86	26.34	67.48	66.91	0.0431
No	11.08	8.57	11.99	13.16	<0.0001
Low-eGFR ^a					
Yes	28.13	19.66	28.24	34.49	0.0018
No	9.62	7.85	10.54	10.85	0.0005
CKD ^b					
Yes	28.07	19.40	28.45	34.39	0.0014
Stage 1	8.88	7.14	8.78	11.08	0.0011
Stage 2	10.63	8.75	12.93	10.52	0.0127
Stage 3	24.87	17.78	25.53	29.86	0.0145
Stage 4	79.47	72.64	66.94	98.94	0.0400
Stage 5	78.57	0.00	42.71	90.54	0.1868
No	9.61	7.86	10.51	10.86	0.0005

eGFR, estimated glomerular filtration rate; CKD, chronic kidney disease.

^aLow eGFR was defined as eGFR < 60 ml/min/1.73 m².

^bCKD stage 1: eGFR ≥ 90 ml/min/1.73 m²; stage 2: 60 ml/min/1.73 m² ≤ eGFR < 90 ml/min/1.73 m²; stage 3: 30 ml/min/1.73 m² ≤ eGFR < 60 ml/min/1.73 m²; stage 4: 15 ml/min/1.73 m² ≤ eGFR < 30 ml/min/1.73 m²; stage 5: eGFR < 15 ml/min/1.73 m².

and could also be appropriate for the participants with reduced renal function.

DISCUSSION

In this cross-sectional study with 7,679 adults, a significant positive association of DII with the PTH level and HP was observed, indicating that higher consumption of pro-inflammatory diet may contribute to the higher PTH level and an increased risk of HP. The association between the exposure variable and outcome variables was still stable after adjustment for covariates. Subgroup analysis stratified by the renal condition (low eGFR and CKD) and other variables showed that this positive association was not affected, suggesting that this association could be appropriate for the population with different

renal function conditions, gender, age, BMI, hypertension status, and diabetes status.

To our knowledge, this is the first study that assesses the association between the dietary inflammatory potential with the PTH level and the risk of HP. Patel et al. (44) investigated whether dietary calcium intake will influence the serum PTH concentrations in apparently healthy Indian adolescents and found that subjects with higher calcium intake had lower PTH. Cheung et al. (45) suggested that low dietary magnesium intake could alter the vitamin D-PTH relationship in adults who were overweight or obese. In another study assessing the effect of the dietary approaches to stop hypertension (DASH) diet, which is rich in fiber and low-fat dairy and is useful for lowering blood pressure on the PTH level, no significant effect was observed (46). Additionally, the cause-and-effect relationship between PTH and inflammation still remains controversial. Cheng et al. (19)

TABLE 3 | Association between DII, PTH, and HP, weighted.

	β /OR ^a (95% CI ^b), P value		
	Model 1 ^c	Model 2 ^d	Model 3 ^e
Parathyroid hormone			
DII (continuous)	0.71 (0.51, 0.90) <0.0001	0.58 (0.38, 0.77) <0.0001	0.46 (0.25, 0.66) <0.0001
DII categories			
Tertile 1	Reference	Reference	Reference
Tertile 2	2.33 (1.09, 3.56) 0.0002	1.77 (0.54, 2.99) 0.0049	1.56 (0.28, 2.84) 0.0167
Tertile 3	4.11 (2.84, 5.39) <0.0001	3.31 (2.03, 4.60) <0.0001	2.66 (1.31, 4.01) 0.0001
Hyperparathyroidism			
DII (continuous)	1.07 (1.05, 1.10) <0.0001	1.07 (1.04, 1.09) <0.0001	1.05 (1.02, 1.08) 0.0023
DII categories			
Tertile 1	Reference	Reference	Reference
Tertile 2	1.48 (1.26, 1.75) <0.0001	1.43 (1.21, 1.68) <0.0001	1.27 (1.05, 1.55) 0.0157
Tertile 3	1.57 (1.33, 1.85) <0.0001	1.49 (1.26, 1.76) <0.0001	1.32 (1.09, 1.61) 0.0054

Insensitivity analysis—dietary inflammatory index was converted from a continuous variable to a categorical variable (tertiles).

^a β : effect sizes; OR: odds ratio.

^b95% CI: 95% confidence interval.

^cModel 1: no covariates were adjusted.

^dModel 2: adjusted for gender, age, and race.

^eModel 3: adjusted for gender, age, race, education level, serum glucose, serum calcium, serum phosphorus, serum iron, serum C-reactive protein serum, serum 25(OH)D, systolic blood pressure, diastolic blood pressure, body mass index, hypertension, diabetes, chronic kidney disease, protein intake, calcium intake, and phosphorus intake.

assessed the association between the PTH level and inflammatory markers among the U.S. adults and reported that higher inflammatory markers were associated with a higher PTH level. Similar results have been reported before in other studies (22, 47–49). Another study observed upregulation of inflammatory genes in the adipose tissue from primary patients with HP when compared with the healthy controls (19). However, conflicting results were found for the effects of inflammatory markers on patients with HP after parathyroidectomy: both increased (24, 50, 51) and decreased (22) and even unchanged levels of inflammatory markers have been reported (52). The inconsistent results indicated an unclear cause-and-effect association between PTH and inflammation. The results of this study indicated that higher pro-inflammatory dietary intake was positively associated with PTH and an increased risk of HP; in another way, a higher inflammatory level may be associated with a higher PTH level and a higher risk of HP. The association was similar in the population with different renal function conditions, gender, age, BMI, hypertension status, and diabetes status according to the subgroup analysis.

Regarding the positive association between DII scores and the PTH level, we observed significant dependence on low eGFR (P for interaction = 0.0004) and CKD (P for interaction = 0.0003), indicating that renal function may be affected by this association.

Serious disorders of calcium and phosphorus metabolism were commonly observed in patients with CKD; we speculated that the effect of the pro-inflammatory diet on PTH may be interfered by the metabolism disorder (53). The interplay between Renin-Angiotensin-Aldosterone System (RAAS) and PTH may also influence this association (54). What is more, patients with end-stage CKD were treated with hemodialysis usually, and the management of hemodialysis can affect the calcium and phosphorus metabolic status (55, 56). However, the data of hemodialysis in NHANES were missing, so we could not exclude the influence of different dialysis modes or conduct further analysis. Additionally, it was worth noting that this positive association was still statistically significant both in subgroups stratified by low eGFR and CKD (all *P*-value < 0.05), suggesting that, although, renal function may affect this association, it was similar in the population with/without low eGFR and CKD.

The exact mechanism of this positive association of DII with PTH and HP remains unclear. A possible explanation to support the results might be the effect of diet on pro-inflammatory markers, such as IL-8, TNF- α , etc.. Previous studies have demonstrated that inflammation was closely associated with the activation of the NF- κ B pathway (57–59). NF- κ B could serve as a direct regulator of genes related to cell proliferation, such as cyclin D1, p21, p27, and p53, thus, involved in cell cycles (60, 61). It also plays a key role in angiogenesis and tumor growth by promoting antiapoptotic mechanisms and the expression of inflammatory cytokines; in turn, these cytokines also contribute to its activation (62, 63). NF- κ B could play a role in PTH regulation as well. Mao et al. (26) found that local NF- κ B activation could promote the synthesis and secretion of PTH and mediated the transcriptional activation of PTH directly. We hypothesized that the elevated inflammatory level may lead to the activation of NF- κ B; thus, similar regulation by NF- κ B may underlie the development of parathyroid hyperplasia and enhanced PTH synthesis and secretion. Angeletti et al. (27) evaluated the effects of IL-8 on the PTH level by incubation of parathyroid cells with recombinant IL-8 *in vitro* and found that IL-8 increased both PTH secretion and PTH mRNA expression. A previous study also reported that TNF- α could affect CASR transcription *via* NF- κ B in human renal tubular cells (64). A significant down-expression of CASR mRNA, thus, upsetting the Ca²⁺ set point and enhancing the sensitivity of parathyroid glands to Ca²⁺ concentration may be another potential explanation of our results.

One of the strengths of our study was that it was based on nationwide, population-based sampling survey data, and sample weights were adopted, which make the study much more representative. It is noteworthy that we performed subgroup analysis stratified by different renal functions and found that this association was similar in different population settings and could also be appropriate for the participants with CKD. However, the limitations of this study cannot be ignored. First, due to the cross-sectional study design, we cannot make a causal inference. Second, the food intake data were based on a 24-h dietary recall; a recall bias was inevitable, and the daily variability of food intake cannot be reflected. With regard to DII scores, a total of 45 food parameters were included according to the design,

TABLE 4 | Subgroup analysis stratified by different variables, weighted.

DII	Parathyroid hormone		Hyperparathyroidism	
	β^a (95% CI) ^b , P for trend	P for interaction	OR ^c (95% CI), P for trend	P for interaction
Low-eGFR ^d				
Yes	1.65 (0.30, 3.01) 0.0171	0.0004	1.10 (1.04, 1.17) 0.0018	0.7108
No	0.33 (0.15, 0.51) 0.0003		1.04 (1.01, 1.08) 0.0121	
CKD				
Yes	1.73 (0.39, 3.08) 0.0116	0.0003	1.07 (1.01, 1.16) 0.0023	0.4942
No	0.33 (0.15, 0.50) 0.0003		1.04 (1.01, 1.08) 0.0121	
Gender				
Male	0.47 (0.20, 0.74) 0.0006	0.8715	1.04 (1.00, 1.09) 0.0469	0.5675
Female	0.43 (0.12, 0.73) 0.0061		1.05 (1.01, 1.10) 0.0206	
Age				
<60 years old	0.32 (0.10, 0.53) 0.0038	0.0588	1.04 (1.00, 1.08) 0.0444	0.5734
≥60 years old	0.89 (0.41, 1.37) 0.0003		1.07 (1.01, 1.12) 0.0137	
BMI				
BMI < 25 kg/m ²	0.48 (0.15, 0.81) 0.0042	0.7475	1.11 (1.05, 1.17) 0.0004	0.1499
BMI ≥ 25 kg/m ²	0.47 (0.21, 0.73) 0.0003		1.02 (1.01, 1.06) 0.0039	
Hypertension				
Yes	0.69 (0.21, 1.17) 0.0053	0.1696	1.04 (0.99, 1.09) 0.0862	0.7086
No	0.34 (0.14, 0.55) 0.0009		1.05 (1.01, 1.09) 0.0071	
Diabetes				
Yes	1.09 (0.12, 2.06) 0.0287	0.1881	1.09 (0.98, 1.21) 0.1276	0.3318
No	0.40 (0.20, 0.61) 0.0001		1.04 (1.01, 1.08) 0.0035	

The results show that the subgroup analysis was adjusted for all presented covariates except the effect modifier. eGFR, estimated glomerular filtration rate; CKD, chronic kidney disease.

^a β : effect sizes.

^b95% CI: 95% confidence interval.

^cOR: odd ratio.

^dLow eGFR was defined as eGFR < 60 ml/min/1.73 m².

but only 27 food parameters were available in the NHANES data. Although, previous studies reported that the predictive ability was not affected when <30 dietary parameters were used with DII calculating (38–40), the impact on accuracy cannot be ignored. Third, HP was defined as intact PTH > 65 pg/ml in our analysis. However, optimal PTH levels in CKD stages 3–5 remain unknown (65). Although most of the participants with CKD in this study were in stage 1 and stage 2, we also used this HP definition in patients with CKD stages 3–5; thus, it may affect the accuracy. In addition, some potential confounders, such as the hemodialysis condition (the patients with non-dialysis dependent or hemodialysis CKD, a hemodialysis type, etc.), drugs use, and fibroblast growth factor 23 (FGF23), may influence this association, but these confounders were not available in NHANES data (56, 66–68). Another limitation was that PTH was only assayed at a single time point; thus, no repeat measurements of PTH were conducted.

CONCLUSION

In this cross-sectional study with 7,679 adults, a significant positive association between DII with PTH and HP was observed, indicating that higher consumption of pro-inflammatory

potential correlates positively with the higher PTH level and an increased risk of HP. This association exists both in the populations with CKD and with no CKD. Our finding suggests that anti-inflammatory dietary management may reduce the risk of HP. Further research and clinical settings are still needed to validate their potential application.

DATA AVAILABILITY STATEMENT

Publicly available datasets were analyzed in this study. This data can be found here: www.cdc.gov/nchs/nhanes/.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by the NCHS Ethic Review Board. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

ZQ: data analysis, software, and writing of the original draft. QY: formal analysis and software. RL: methodology and writing

of the original draft. BS: conceptualization, funding acquisition, writing, reviewing, and editing. All the authors approved the final version.

FUNDING

This work was supported by the State Key Research Programme of China (Grant nos. 2016YFC1103004 and 2016YFC1103003), the National Natural Science Foundation of China (Grant no. 82000702), the Science and Technology Achievement

Transformation Fund of West China Hospital of Sichuan University (Grant no. CGZH19006), National Clinical Research Center for Geriatrics, West China Hospital, Sichuan University (Grant no. Z2018B10), and Med+ Biomaterial Institute of West China Hospital/West China School of Medicine of Sichuan University (Grant no. ZYME20001).

ACKNOWLEDGMENTS

We thank the assistance of Yingfei Xu in data processing.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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How to Overcome Anabolic Resistance in Dialysis-Treated Patients?

Giacomo Garibotto^{1*}, Michela Saio¹, Francesca Aimasso², Elisa Russo^{1,3}, Daniela Picciotto^{1,3}, Francesca Viazzi^{1,3}, Daniela Verzola¹, Alessandro Laudon⁴, Pasquale Esposito^{1,3} and Giuliano Brunori⁴

¹ Department of Internal Medicine, University of Genoa, Genoa, Italy, ² Clinical Nutrition Unit, Istituto di Ricerca a Carattere Scientifico Ospedale Policlinico San Martino, Genoa, Italy, ³ Clinica Nefrologica, Dialisi e Trapianto, IRCCS Ospedale Policlinico San Martino, Genoa, Italy, ⁴ Division of Nephrology and Dialysis, Ospedale Santa Chiara, Trento, Italy

OPEN ACCESS

Edited by:

Alice Sabatino,
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Erasmus Medical Center, Netherlands

*Correspondence:

Giacomo Garibotto
gari@unige.it

Specialty section:

This article was submitted to
Clinical Nutrition,
a section of the journal
Frontiers in Nutrition

Received: 27 April 2021

Accepted: 12 July 2021

Published: 12 August 2021

Citation:

Garibotto G, Saio M, Aimasso F,
Russo E, Picciotto D, Viazzi F,
Verzola D, Laudon A, Esposito P and
Brunori G (2021) How to Overcome
Anabolic Resistance in
Dialysis-Treated Patients?
Front. Nutr. 8:701386.
doi: 10.3389/fnut.2021.701386

A current hypothesis is that dialysis-treated patients are “anabolic resistant” i. e., their muscle protein synthesis (MPS) response to anabolic stimuli is blunted, an effect which leads to muscle wasting and poor physical performance in aging and in several chronic diseases. The importance of maintaining muscle mass and MPS is often neglected in dialysis-treated patients; better than to describe mechanisms leading to energy-protein wasting, the aim of this narrative review is to suggest possible strategies to overcome anabolic resistance in this patient's category. Food intake, in particular dietary protein, and physical activity, are the two major anabolic stimuli. Unfortunately, dialysis patients are often aged and have a sedentary behavior, all conditions which *per se* may induce a state of “anabolic resistance.” In addition, patients on dialysis are exposed to amino acid or protein deprivation during the dialysis sessions. Unfortunately, the optimal amount and formula of protein/amino acid composition in supplements to maximize MPS is still unknown in dialysis patients. In young healthy subjects, 20 g whey protein maximally stimulate MPS. However, recent observations suggest that dialysis patients need greater amounts of proteins than healthy subjects to maximally stimulate MPS. Since unnecessary amounts of amino acids could stimulate ureagenesis, toxins and acid production, it is urgent to obtain information on the optimal dose of proteins or amino acids/ketoacids to maximize MPS in this patients' population. In the meantime, the issue of maintaining muscle mass and function in dialysis-treated CKD patients needs not to be overlooked by the kidney community.

Keywords: nutrition, muscle, protein synthesis, hemodialysis, peritoneal dialysis, CKD

INTRODUCTION

Dialysis-treated patients with end-stage renal disease (ESRD) have a high prevalence of protein-energy wasting (PEW), a condition of muscle and visceral protein stores loss which is not completely accounted for by a low nutrient intake (1). There is a strong association between surrogates of muscle mass and survival in this patients' population (2, 3); therefore, the need for increased medical attention for nutrition in advanced kidney disease is a high clinical priority.

Muscle protein synthesis (MPS), serving either the maintenance of muscle protein during fasting and body anabolism during feeding, is central to the processes of sustaining and shaping body cell mass. Dietary proteins or amino acids (AA), and physical activity are the two major anabolic stimuli that increase MPS (4). If combined, these two factors offer the major available anabolic stimulus in humans (Figure 1). Several other factors, including the dietary protein amount and composition, and the timing of protein ingestion can also influence the anabolic responses.

Recent understanding of the mechanisms which regulate protein metabolism in chronic kidney disease (CKD) has allowed us to understand the role of acidosis, insulin resistance and inflammation on intracellular signals which promote PEW by activating muscle protein degradation (MPD) (5). Even more importantly, Zhang et al. (6) were able to demonstrate that CKD stimulates chromatin-modifying, nucleolar protein 66 (NO66), that downregulates MPS via a demethylase mechanism, giving us a new perspective on the pathophysiology of uremia-related wasting syndrome. Better than to describe mechanisms leading to muscle wasting in dialysis patients, the scope of this narrative review is to analyze current understanding of the dietary and non-dietary modulators of MPS as major determinants of muscle mass in humans, in order to suggest possible strategies to stimulate MPS and, at least in part, overcome anabolic resistance in dialysis-treated patients with CKD.

THE RESPONSE OF MPS TO PROTEIN FEEDING

The MPS response to protein is influenced by both the amount and the type of protein, and the time of feeding.

Protein Amount

The hyperaminoacidemia reached after protein ingestion is “primary anabolic” since it increases MPS rates, and to a minor degree, decreases muscle protein degradation (MPD) (7, 8). Several investigators have studied the optimal amount of protein for the maximal stimulation of MPS in humans (7–14). As demonstrated in early studies (14), the muscle anabolic response to AA/protein ingestion is time-limited in the rested state; after 2–3 h from AA ingestion, MPS undergoes a state of “tachyphylaxis” in spite of ongoing EAA availability and upregulation of mechanistic target of rapamycin complex 1 (mTORC1) signaling (14). This ‘muscle full’ condition is refractory to the intake of additional nutrient signals/substrates regulating MPS, is not induced by muscle blood flow and does not appear to be dependent on the amount of protein ingested (4, 14). Instead, the ‘muscle full’ set-point can be delayed by physical exercise to allow additional use of EAA for MPS (4).

The dose-response of myofibrillar MPS to increasing amounts of whey protein at rest and after exercise has been studied in young subjects by Witard et al. (8). They observed that a 20-g dose of whey protein (containing ~10 g essential AAs) is sufficient for the maximal stimulation of MPS both at rest and during physical exercise, while a greater dose stimulates amino

acid oxidation. However, in elderly subjects, more protein with respect to younger subjects is needed to maximally stimulate MPS in exercised and rested muscle (15).

Protein Type

Also AA composition of the dietary protein influences the MPS response. Lower essential AA (EAA) content, particularly of leucine, lysine, and/or methionine has been shown to be responsible for the lower anabolic capacity of low biological vs. high biological value proteins (16, 17). Apart from their AA content, the nutritional value of proteins ingested with the diet depends also on a series of factors, including their chemical score, net utilization, and digestibility corrected AA score (PDCAAS) (18). Many of these factors are better for animal-based than for plant-based protein sources (19). Both the PDCAAS and the Digestible Indispensable AA Score (DIAAS) are used to assess the ability of dietary protein to satisfy the body's AA requirements (18, 19).

Protein ingestion and physical exercise possess a synergistic interaction on the stimulation of MPS (20, 21). Protein ingestion provides, among the EAA and non-essential AA needed for MPS, also leucine, which stimulates protein translation. In addition, protein and carbohydrate ingestion increase serum levels of insulin, which has a mild stimulatory effect on MPS (20–22) and decreases MPD. Another factor which makes protein differently anabolic is the splanchnic AA removal from circulation (23, 24). As an example, proteins from soy undergo an important splanchnic first-pass, whereas milk proteins are directed to peripheral sites. In comparison to soy or carbohydrate, milk consumption has a greater anabolic effect in young subjects in the post-exercise period (25).

The content of essential AA in plant vs. animal-based proteins is likely the major determinant of their different muscle anabolic effects. Plant proteins have lower essential AA content, in particular of methionine and lysine, than animal-based and muscle protein (17). Of note, the intake of AAs is highest in diets containing prevalently meat protein, followed by fish, and is lowest in diets containing only plant protein (26); accordingly, plasma levels of methionine, tryptophan and tyrosine are lowest in vegans, suggesting dietary deficiency for these AA. In addition, owing to their different structure and/or content in phytic acid and protease inhibitors, plant proteins are less digestible than animal proteins (17). Therefore, it has been suggested to consume different plant proteins with complementary essential AA composition to optimize a full pattern of dietary AA, especially for elderly subjects (17). In long-term studies, at low (~0.8 g/kg/day) protein intake, plant proteins have lower ability to stimulate MPS and to cause muscle anabolism vs. animal proteins. However, the anabolic effect of plant- and animal-based proteins is similar if the protein intake is raised to 1.1–1.2 g/kg (17).

Timing of Protein Ingestion

Protein consumption in western countries exceeds the Recommended Dietary Allowances (RDA), i.e., the protein amount that is consumed daily to meet population needs and to prevent protein deficiencies. However, in many countries the

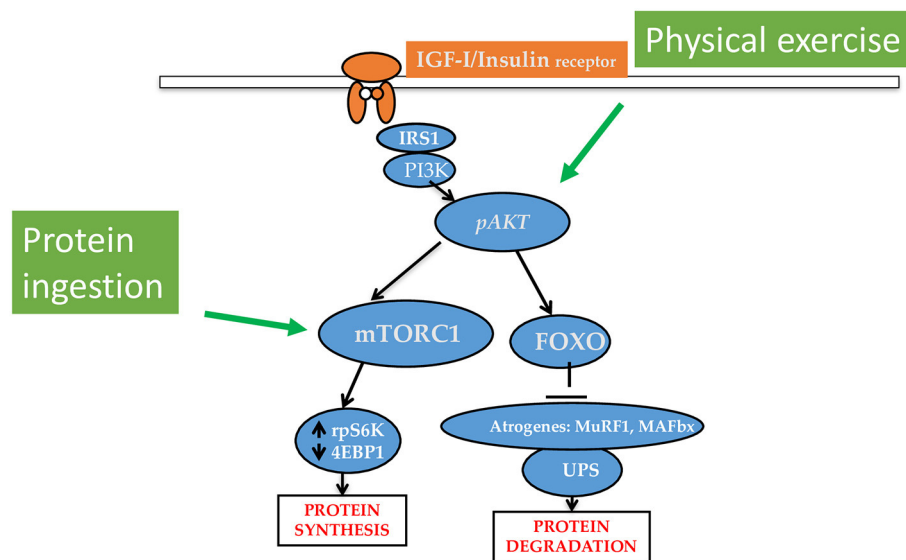


FIGURE 1 | Major intracellular pathways underlying the anabolic effects of protein/amino acid/keto acid intake and physical exercise. The ingestion of protein, amino acids or keto acid of essential amino acids increases muscle protein synthesis (MPS) rates. Physical exercise upregulates phosphorylated Akt to increase MPS and decrease protein degradation. Protein ingestion and physical exercise have synergistic interaction on the stimulation of MPS.

intake is shifted toward the evening meal. In healthy subjects, the rate of MPS is 25% increased when food is distributed during the day, compared to a pattern with protein consumed at the evening meal (27). This finding may have important implications for increasing MPS in subjects who experience a resistance to the stimuli of MPS. Further, immediate post-exercise nutrient ingestion has been shown to increase the accretion of whole body and leg protein more than in a late (3-h) period (28).

Protein ingestion before sleep, if combined with physical activity, is also potentially anabolic (29) and has been suggested to represent a nutritional strategy to increase muscle mass in the elderly. However, there is no study in dialysis-treated patients.

PHYSICAL ACTIVITY

In healthy subjects, resistance exercise augments the MPS rates, resulting in muscle hypertrophy (30) and also increases the expression of contractile, metabolic, and stress response proteins (31). In addition, muscle hypertrophy is associated with improved gait velocity and stair-climbing power even in very old individuals (32, 33). Aerobic fitness increases both markers of mitochondrial biogenesis and MPS (p-P70S6k) (34). In addition aerobic exercise training increases vagal tone and decreases sympathetic tone (35).

Early studies have shown that a single session of exercise increases MPS rates (36, 37). Despite this increase in MPS, if there is no food ingestion, the net muscle protein balance persists to be negative, even if to a lesser extent than in the pre-exercise period (37). During post-exercise recovery period, an increase in AA levels and supply to skeletal muscle from orally administered AA stimulates MPS and, even if to a less extent, inhibits MPD rates, resulting in muscle protein anabolism

(21). Also the ingestion of carbohydrate improves muscle net protein balance (i.e., the difference between MPS and MPD) after resistance exercise; however, this effect is smaller as compared to protein/AA feeding (38). Therefore, post-exercise protein/AA ingestion is more commonly used to maximize anabolism. In addition to protein intake, body composition, age, and/or sex, and usual physical activity may influence the MPS response to feeding (28).

AGING AND COMMON KIDNEY DISEASE COMORBIDITIES WHICH IMPAIR MPS

Several conditions are recognized to impair the MPS response to protein, thus causing “anabolic resistance” and limiting the muscle hypertrophic response.

Aging

The progressive aging-associated muscle atrophy, which frequently presents with muscle weakness and slow motion, is the most common type of muscle atrophy observed in humans (39). With aging, the risk of both CKD and declining capacities such as reduced strength and cognition increases (40); this risk is maximized in subjects developing “accelerated biological aging” (41). Sensory-motor functioning is very common in frail individuals who are at highly vulnerable for loss of independence in activities of day living, fall and mortality (42). Since the population in the western countries aged > 65 is expected to rapidly increase in the next years (43), CKD and age-related losses in skeletal muscle function and mass are considered as an extremely important public health issue (44).

Both quantitative and qualitative modifications of muscle biology and function are observed in elderly subjects. Early

computed tomography studies (45) have shown that aging is associated with a loss of lean mass, with fat and connective tissues replacing muscle. These changes in muscle composition are considered to be consequence of a decreased MPS rates with aging (46).

The age-related muscle atrophy has been described as a slow but unrelenting process, with functional loss varying largely among individuals, which is observed in all elderly subjects, including those healthy and physically active (41, 47). Although the study of sarcopenia is complicated by muscle-specific differences among species and in the response to aging (41), there is consensus that anabolic stimuli do not efficiently activate the IGF1/PI3K/Akt/PKB/mTORC1 pathway (displayed in **Figure 1**) in elderly people, and this anabolic resistance is believed to play a major role in causing sarcopenia and reduced muscle recovery after an injury (41). In accordance with this view, acute activation of the pAkt/FoxO axis has been shown to be of benefit in the elderly (48). In addition, similarly to cancer cachexia, the age-associated low-grade systemic inflammation plays an important role in the pathogenesis of sarcopenia (49). How renal disease abnormalities overlap or potentiate those occurring with aging is still unexplored.

Inactivity

Immobilization or sedentarism induces a status of “anabolic resistance” with reduced fasted and fed-state MPS. The effects of reduced use on skeletal muscle are in part similar to those observed during aging (50), and the increased sedentary behavior occurring in old age (51) has been suggested to be the major cause of the aging-related sarcopenia.

Long-term immobilization in bed causes a more marked decrease in muscle strength than in muscle mass (52). However, changes at cellular and subcellular levels have shown that long-term immobilization causes a significant loss of contractile proteins, that is even greater than the decrease in muscle fiber force generation capacity (53).

Low Energy Intake

A crucial aspect of the nutritional balance is the amount of energy ingested. From studies based on nitrogen (N) balance in the last century, it is known that the protein metabolism is influenced by the amount of energy intake (54–57). Of note, if dietary energy is supplied in amounts sufficient to meet body needs, protein intake strongly influences N balance. In healthy subjects, the administration of excess energy spares the loss of labile proteins, and reduces the time to adapt to a low protein diet. The supply of excess energy results in increased availability of ingested protein, with decrease in protein requirement (58). In keeping with this finding, N balance appears to be more influenced by changes in dietary energy supply at low protein intakes, and the N-sparing effect of energy is blunted when protein intake increases (59). These observations have practical implications, since protein requirements are determined when the requirements for energy are met.

To explain the mechanisms underlying the responses of protein metabolism to changes in protein intake, several years ago Young et al. (60, 61) studied whole body protein metabolism in

12 young men fed a low protein diet (0.6 g egg protein/kg/day) associated with a maintenance energy level or a 25% energy excess. In the fed-state the rate of leucine oxidation tended to be lower, while N balance and net protein balance were higher, when dietary energy was increased. These studies indicate that the oxidation of essential AA is sensitive and varies according to the levels of energy intake, with higher oxidation at low intakes.

HD patients have reduced dietary energy intake (62) and progressively lose muscle mass with time, even if body fat seems to be better preserved than lean mass (63). Studies on the dietary energy requirements of adult HD patients have been assessed by measuring energy expenditure by indirect calorimetry and/or by the use of N balance (63, 64). The results of these studies are not uniform. Some studies reported that dietary energy needs are not different from those of normal adults (65), whereas other studies showed increased energy expenditures (66). Inflammation has been reported to be associated with an increase in energy requirements (67). A few studies in HD patients have evaluated energy expenditure during both at rest and activities daily living. Shah et al. (68) measured dietary energy requirements for 92 days in 13 HD patients residing in a metabolic research ward while receiving a constant energy intake. Their average energy intake was 31 kcal/kg/day calculated from energy intake and change in fat and fat-free calories, which was 28 kcal/day. However, a wide variability in dietary intake among individual patients was observed (range: 26–36 kcal/Kg/day) (68). Dietary intake correlated strongly with their body weight, but was less closely related to their measured resting energy expenditure (REE). Overall, the data from this study show that, on the average, dietary energy requirements of sedentary, clinically stable HD patients are similar to those of age-matched sedentary normal subjects. However, due to the high individual variability in energy requirements, careful individual monitoring of the nutritional status of HD-treated patients is necessary (68).

Despite the importance of energy intake on protein requirements in N balance-based studies, the MPS response to food has been largely focused on the effects of orally administered protein or AA. However, food is usually eaten as a “mixed” meal. One of the main reasons for focusing on isolated protein ingestion is that protein provides essential AA for protein synthesis, while carbohydrates and fats ingested alone are only mildly muscle-anabolic. Protein absorption is delayed when carbohydrate are co-ingested with proteins, without however reducing the MPS response (69). However, protein and carbohydrate ingestion increase systemic insulin, which has a modest stimulatory effect on MPS and decreases MPD (38).

A few studies have focused on the effects of fat co-ingestion on the MPS response. Following physical exercise, the leg removes a greater amount of AA after the ingestion of high-fat milk compared with skim milk (70). In addition, the ingestion of whole eggs (18 g protein, 17 g fat) has been shown to cause a greater increase in post-exercise MPS rates than an isonitrogenous amount of egg whites or egg whites (18 g protein, 0 g fat) (71). This differential response may be due to the greater content of fat and/or micronutrients in whole eggs. Accordingly, fat co-ingestion (237 g of whole milk as compared to fat-free milk) may increase the MPS response to protein in the

post-exercise period (71). However, MPS is blunted by a high fat content (72, 73).

MPS is a process that “*per se*” requires energy (74). HD patients display muscle mitochondrial morphological and dysfunctional changes, which might limit the availability of energy for MPS or impair the efficiency of MPS in terms of energy costs. The energy requirements of whole-body PS have been estimated to account for ~20% of resting energy expenditure (REE) in humans (75). Since muscle protein turnover accounts for ~40% of whole-body protein turnover in humans, one could estimate the cost of MPS to be ~8–10% of REE. The cost of MPS is likely similar to control subjects in wasted HD patients. When both MPS and REE have been concurrently studied in non-inflamed, wasted HD patients before and during Growth Hormone (GH) administration (76) the variations in REE and MPS were significantly correlated suggesting that changes in MPS account for a significant fraction of REE changes. On that ground the energy cost of MPS was estimated to be ~5 kcal/g, which is similar to the normal condition (76).

Diabetes Mellitus

Insulin is an anabolic hormone. Major effects of insulin's action are to suppress protein degradation and up-regulate anabolic pathways (77). While in type 1 diabetic subjects muscle wasting results from overexpression of genes involved in the ubiquitin proteasome pathway, in type 2 diabetes a decrease in insulin sensitivity favors muscle atrophy. A decrease insulin's response in muscle may take place owing to alterations in the insulin signaling pathways secondary to inflammation, glucotoxicity, increased circulating free fatty acids and metabolic acidosis (78, 79). Furthermore, the activation of new anti-anabolic or catabolic pathways, such as the myostatin/activin A system, may occur (79).

Patients with type 2 diabetic kidney disease often show a series of metabolic and nutritional changes linked both to diabetes and kidney failure, including insulin resistance and cardiovascular comorbidities (80), protein-energy wasting and sarcopenia (81). These patients undergo an increase in MPD, an effect primarily mediated by the ubiquitin-proteasome pathway (82, 83). On a kinetic basis, an increase in MPD is often associated with an increase in MPS. However, this seems not to be the case for diabetic kidney disease. In a recent study Zanetti et al. (84) observed that patients with diabetic kidney disease have a catabolic pattern of whole body protein turnover, associated with a 10% decrease in MPS (84). Therefore, multiple changes in protein metabolism, including an increase in protein degradation not associated with an adequately-matched change in protein synthesis account for accelerated wasting in patients with type 2 diabetic nephropathy.

Sepsis

Chronic uremia has for long been recognized as a state of acquired immunodeficiency (85, 86). Sepsis is recognized as the second leading cause of death in ESRD (87, 88). Besides infections from indwelling catheters, numerous other factors, including aging, diabetes, hypoalbuminemia, immunosuppressive therapy,

the dialysis procedure, uremia and increased leptin levels (89, 90) can favor infections in ESRD patients.

Sepsis causes a rapid and significant loss of body protein, in part due to development of an impairment of MPS alone or in association with increases in MPD (91). Septic cachexia has been defined as a life-threatening condition of metabolic inflammatory complex associated with multiple organ dysfunction (91). The metabolic changes induced by the systemic inflammatory response, including mitochondrial dysfunction and metabolic shift (91, 92), are closely connected by a wide array of signaling molecules. In several tissues, including skeletal muscle, the expression of pro-inflammatory cytokines is regulated by Toll-like receptors (TLRs) (92). During sepsis, muscle TLRs monitor for the presence of endotoxin (92, 93) and, when activated by microbial products, induce a local inflammatory response leading to the activation of pro-inflammatory genes. In addition, TLRs can be activated by heat shock proteins and/or endogenous signals of tissue injury (93). Recent observations suggest that TLR4 link the innate immunity activation with the uremic state in skeletal muscle (9). In predialysis CKD patients, an upregulation of TLR4 in muscle is predicted by the residual renal function, suggesting that endotoxins or danger-associated molecular patterns (DAMPs) produced or retained in the pre-uremic state mediate TLR4 activation (94). These findings suggest that in uremic patients muscle inflammation is due to an overexpression of TLR4 (94). The potential for septic cachexia to serve as a novel target disease state to improve clinical outcomes is reviewed elsewhere (92).

Metabolic Acidosis

A normal bicarbonate serum concentration (serum HCO_3^- between 24 and 26 mmol/l) is the recommended target in renal patients. While metabolic acidosis is one of the most important, yet treatable, among factors contributing to accelerated protein catabolism in dialysis patients (79), its effects on MPS are less established. In rats made acidotic (mean pH 7.22) by intragastric administration of NH_4Cl (95), protein synthesis substantially decreased in muscle (96). In another study, induction of metabolic acidosis with NH_4Cl in normal individuals (mean attained HCO_3^- 16 mmol/l) caused a significant reduction in MPS, while albumin synthesis remained unchanged (95); conversely, chronic NH_4Cl -induced metabolic acidosis (mean HCO_3^- 15 mmol/l) was reported to significantly decrease the albumin fractional synthesis, without effects on MPS (97). Lofberg et al. (98) studied the effects of the correction of metabolic acidosis on muscle protein metabolism in 16 HD patients who were randomized to increase or decrease bicarbonate supplementation (blood bicarbonate levels increased from 17.8 to 27.1 mmol/l in the first, and decreased from 26.6 to 18.6 mmol/l in the second group). Muscle protein net balance improved when acidosis was corrected, mainly as a consequence of changes in protein degradation while protein synthesis was unaffected.

In consideration of these discordant findings, it is unclear if mild acidosis may affects MPS. Clearly, metabolic acidosis is potentially anti-anabolic, since it can decrease the release of GH and IGF-1 and induce insulin resistance (79). However, the effects

of acidosis on the anabolic effects of protein feeding in dialysis patients have not been studied so far.

Hemodialysis Procedure

The mechanisms underlying muscle wasting in HD patients involve a large number of factors, including inflammation, anorexia, metabolic acidosis, anabolic hormone resistance, and the loss of AAs and protein (mainly albumin) in the dialysate (99–104). HD acutely decreases MPS and upregulates protein degradation (99–101). Different mechanisms underlie this catabolic response, including losses of AAs and proteins into the dialysis fluid and the upregulation of protein catabolism by the interaction of immunocompetent blood cells with the dialysis membrane. Loss of AAs is very variable (from 4 to 12 g) (102–105), and is directly proportional to the AA plasma levels, dialysate area and blood flow. In addition, protein losses into dialysate (2–3 g) as well as protein adsorption to the dialysis membrane and tubing can occur. Newer dialyzers with medium cut-off membranes, which are designed to improve clearance of middle molecules, are associated with increased albumin losses (106), even if with large variability (107).

A current hypothesis is that the inflammatory state observed in a significant number of HD patients is boosted by the HD procedure, which might increase the risk of cardiovascular complications and cachexia. Boivin et al. (108) showed that HD increases interleukin-6 (IL-6) and the caspase-3-mediated cleavage of actomyosin, the contractile protein made by the actin-myosin complex. In addition, protein degradation was higher in patients during the HD procedure as compared the pre-HD period.

The activation of the complement with its downward inflammatory responses may be an additional mechanism leading to wasting in HD patients (109, 110). By a proteomic approach, Mares et al. (110), by eluting proteins adsorbed to the polysulfone dialyzer membranes, studied the processes that take place on the dialysis membranes. These proteins included ficolin-2 (a component of innate immunity that contributes to the immune recognition of pathogens), and complement C3c fragment. Of note, their data suggest that the polysulfone dialyzer initiates the lectin pathway of complement activation (111). In addition, a current hypothesis is that the anaphylatoxins C3a and C5a, bradykinin, thrombin and factor Xa can induce the activation of endothelial cells in the vascular wall, a process which may lead to inflammation and accelerated atherogenesis (112). However, there is a lack of controlled studies exploring the muscle proteolytic response to different filter materials.

Peritoneal Dialysis

One of the untoward effects of PD is the loss of proteins and AAs into the dialysate effluent. Patients undergoing PD lose both AAs (1–3.5 g/day), and a consistent amount of proteins (5–10 g per day), mainly albumin (113–115). The continuous loss of proteins and AAs is estimated to account for almost one-third of the increase in dietary protein requirements, but poor appetite or anorexia limit intakes in many patients (116, 117). When muscle AA kinetics has been studied during the PD procedure, it was

observed that PD causes an acute decrease in MPS, which has been associated to the combined effect of hyperinsulinemia and AA losses (118).

CKD has been considered for years a state of “anabolic resistance” (119). However, only a few recent studies give support to this hypothesis and show that the muscle anabolic response to protein ingestion is blunted in dialysis-treated patients.

CKD causes a series of gastrointestinal abnormalities, including motility disorders, gastric achlorhydria, pancreopathy and small bowel bacterial overgrowth, which may impair protein absorption as demonstrated by the use of a ^{13}C protein breath test (120). At muscle level, Chen et al. observed that leucine-stimulated mTOR signaling (121) and GH-stimulated insulin-like growth factor-1 (IGF-I) expression (122) are partly attenuated in chronically uremic rats. In addition, the same authors observed that acute metabolic acidosis blunts the leucine-induced increase in MPS in rat muscle (123).

In patients with CKD, after protein ingestion the plasma AA pattern to peripheral tissues is abnormal, with a more marked rise in non-essential AAs, as compared to essential AAs (124–126). Studies performed by whole body leucine kinetics in CKD patients have shown that whole body protein synthesis is resistant to stimulation by hyperaminoacidemia while the response of protein turnover to high insulin levels is normal (127). In addition, in patients with CKD and mild metabolic acidosis insulin sensitivity of muscle protein metabolism is overall preserved at insulin levels in the high (~8–9-fold increased vs basal) physiological range; however, muscle insulin's sensitivity in the low physiological range is impaired (128). This observation suggests that CKD patients may need higher insulin levels to decrease MPD after a low glycemic index meal, such as breakfast. In contrast, the muscle insulin's sensitivity at raised insulin levels, as may take place with larger meals containing refined carbohydrates, appears to be normally maintained.

Recently, van Vliet et al. (129) observed that in HD patients, the ingestion of a mixed-meal containing 20 g protein failed to increase MPS. Unfortunately, basal rates of MPS were ~2-fold increased in HD patients as compared to control subjects, a finding that might have impaired the ability of researchers to demonstrate any meal-induced MPS effect. In a recent study, Draicchio et al. (130) observed that whole body protein turnover rate decreased in HD patients after the ingestion of a mixed-meal compared with controls, suggesting that AA were poorly absorbed from the meal or dysregulated AA metabolism.

HD patients may also display “anabolic resistance” to physical exercise. In the “PEAK” study Cheema et al. (131) randomized 49 patients to a 12-week resistance training during routine HD treatment or usual care. Physical exercise improved muscle strength, mid-thigh and mid-arm circumference, as compared to controls, but was not followed by statistically significant difference in muscle cross sectional area (an index of muscle hypertrophy). Mulsted et al. (132) studied 29 HD patients before and after completing 16 weeks of strength training. Before the training period, the participants were randomly assigned to receive a protein or a non-protein drink after every training session. Strength training did not result in muscle hypertrophy,

while it was associated with increases in muscle strength and power (132).

Taken together, the available data in experimental models and in humans suggest that HD patients are resistant to the most important anabolic stimuli, i.e., protein and physical exercise. If so, HD patients, in addition to well-tailored exercise programs, need to eat more than the 20 g high-quality protein which is necessary to obtain an optimal MPS response in healthy subjects (8). As an example, elderly subjects, who are also considered to be “anabolic resistant”, need 35 grams of protein i.e. ~18 g essential AAs for the maximal stimulation of MPS in exercised and rested muscle (132, 133). However, this large amount of AAs, if not efficiently used for MPS, could boost ureagenesis, toxins and acid production. As an alternative, the use of keto acids (KA) of essential AA might offer an anabolic stimulus with less N. It appears therefore urgent to obtain information on the optimal dose of protein and/or oral AA/KA to maximize MPS in HD patients.

DISCUSSION: HOW TO OVERCOME ANABOLIC RESISTANCE IN DIALYSIS-TREATED PATIENTS?

In this narrative review, we have briefly summarized our current evidence on factors which regulate MPS in humans, as a background information for effective treatment of patients with advanced CKD, mainly those on maintenance HD.

Consider Amount, Quality, and Timing of Ingested Proteins

According to the current NKF KDOQI/AND guidelines, “nutritional counseling should aim to achieve a dietary intake of 1.2 g protein and 30–35 kcal/kg body weight/day in dialysis patients at risk of malnutrition” (134). This can be obtained through increased intake of nutrients, food fortification, and/or the provision of oral nutritional supplementation. However, despite strong attention to nutritional requirements, many dialysis-treated patients do not meet the current guidelines (1, 103, 104).

Quality of dietary proteins needs also to be checked. Plant proteins contribute to a large part of daily protein intake in the world population (135) and plant-based low protein diets are encountering increasing success for the prevention and treatment of several comorbidities associated with CKD (136). In addition, plant protein-based diets may correct many of the uremia-associated abnormalities (137–140). However, both in young and elderly subjects, plant proteins have a lower ability to increase MPS than animal proteins, when protein intake is low (≈ 0.8 g/kg/day) (17, 141). Of note, in elderly subjects, plant- and animal-based proteins are similarly anabolic when plant protein intake is adequate (1.1–1.2 g/kg) (17, 141), although no study is available in HD patients. Therefore, increasing the intake of plant protein to 1.2 g/kg may be the first option to offer when dietary plant-based protein content is low. Other approaches have been suggested to increase the anabolism induced by plant protein (141). These may include: 1) considering different plant

protein sources; 2) combining animal and plant proteins; 3) supplementing plant-based diets with essential AA or KA of essential AA (141).

One of the current debates on nutritional treatment of dialysis patients is how to increase dietary protein intake without causing hyperphosphatemia, which is associated with all-cause and cardiovascular mortality (142). However, HD patients may have both poor nutritional status and hyperphosphatemia. Often nephrologists handle with hyperphosphatemia by prescribing both phosphate binders and restriction of protein-rich foods, an effect which may, however, boost wasting. Recently, Yamamoto et al. (143) observed that the practice of recommending an increase in dietary protein intake for HD patients with concurrent low serum albumin and high phosphate levels was associated with higher serum creatinine and potentially lower all-cause mortality in the Dialysis Outcomes and Practice Patterns Study (DOPPS) study-phase 4 (2009–2011, follow-up 1.6 years). According to this observation recommending the liberalization of protein intake together with hyperphosphatemia management may be a critical practice for better nutritional status and outcomes in HD patients. However, prospective RCTs are needed to determine the impact of protein intake liberalization on protein status and hard outcomes in malnourished, hyperphosphatemic HD patients.

Allow Eating During HD

In many dialysis centers HD patients are fasting throughout the entire dialysis session, for fear of hypotension or vomiting (144). In addition, patients are often suggested not eating before the session, so it is very common that those on an afternoon HD session miss their noon meal (or they miss their breakfast if they have a planned morning HD session). For these reasons, nutrient intakes are often lower than recommended on dialysis-treatment days as compared with non-dialysis days.

In acute studies, feeding protein during HD treatment has been shown to cause a positive whole body protein balance (145). Providing meals during the HD treatment can increase overall nutrient intake and as well as the percentage of patients who reach nutritional targets (146). To replete AA pools, protein-rich foods have been recommended during and after the HD session; additionally, it is worth recognizing that wanting of protein-rich foods increases immediately after dialysis (147). Allowing patients to eat regularly during HD has been associated with better essential AA kinetics (148), nutritional status, quality of life, and possibly clinical outcomes [see for review ref. (149)].

In patients with gastrointestinal symptoms or who do not tolerate eating during HD treatment, intradialytic parenteral nutrition (IDPN) is used to prevent HD-related catabolism (150, 151), but the nutritional effects of this treatment may require long-time. Although a prospective study did not show an advantage of adding IDPN to oral nutritional supplementation on in malnourished HD patients, the improvement in serum prealbumin was associated with a decrease in morbidity and mortality (152).

Deleaval et al. (153) recently showed that the addition of branched-chain AAs (BCAAs) in the dialysis fluid may help to

maintain their physiological plasma levels and MPS. Clearly, the clinical usefulness of this approach needs to be tested.

Intraperitoneal AA + Glucose in PD Patients

PD shifts muscle protein metabolism to a status of reduced anabolism, a condition potentially harmful if nutrient intake is diminished or during superimposed diseases (154). Results from clinical trials on the use of intraperitoneal AAs (IPAA), including both RCTs and non-RCTs, were inconclusive [see ref. (121) for review] because of insufficient sample size and/or the occurrence of confounding conditions (121). Currently, IPAA are recommended when protein and energy intake are unattainable (121).

In an acute study (118), MPS was stimulated when AAs were supplemented intraperitoneally in combination with dextrose. The combined use of dextrose and AAs resulted in a cumulative anabolic and anticatabolic effect, because of the decrease in MPD (induced by insulin) and the stimulation of MPS (induced by AA availability) as compared to dextrose. Canepa et al. (155) in a study using an automatic PD cycler (APD) to simultaneously infuse glucose and AA (proportion of 3:1) in pediatric patients observed that the combined treatment was associated with a better metabolic profile, with maintenance of adequate non-protein calorie/nitrogen ratio. Therefore, these findings support the use of combined glucose-AA solutions, to improve utilization of AA for MPS while controlling urea levels (155). More recently, Tjong et al. (156, 157) in an open label, cross-over study, compared the effects of a mixture of AA and glucose vs. glucose only containing dialysate, in 2 periods of 7 days each in eight APD-treated patients. They concluded that APD with dialysate composed of a mixture of AA and glucose acutely improves protein metabolism, and that this gain made during the night persists to a considerable extent for 24 h. However, no long term RCT is available on the effects of combined use of IPAA + glucose on nutritional status and harder outcomes.

Oral Nutrition Supplements

The administration of oral nutritional supplements (ONS) is used to maintain nutritional status (156), when dietary counseling alone is not sufficient to fill the gap between actual intakes and target requirements. Patients are advised to take ONS preferably 1 h after meals rather than as a meal replacement (156). Essential AA supplements offer the possibility to increase MPS; many observational studies have shown that in the short term, ONS can improve albumin and blood EAA profile in HD patients (156). A recent meta-analysis in HD patients shows that if short-term (3 months) ONS use increases BMI and serum albumin levels, prolonged (> 3 months) supplementation does not cause such effects (158). Recently, Pokkrong et al. (159) observed that ONS administered to 80 HD patients was associated with an improvement in energy, protein, fat, fiber and magnesium intake and 29–24% decrease in malnutrition-inflammation score (MIS), while the improvement in serum albumin was slight (5.3–3.3%). A Cochrane meta-analysis (160) included 22 studies (1,278 participants, 79% on HD and 21%

on PD); however, the authors pointed out that many studies were at unclear risk of selection, performance, and reporting bias. Overall, it is likely that protein-based ONS increase serum albumin and prealbumin (mainly in malnourished patients), as well as anthropometric measures. However, it is unclear if these results may translate into clinically relevant outcomes. In a recent meta-analysis Liu et al. (161) pointed out that large well-designed RCTs in this population are required. There is also little information on the effects of different AA formulas prescription.

Ketoacids and Hydroxyacid Analogs

Even less information on the anabolic effect for ONS containing AA and KA is available in the HD or PD setting. Currently, AA and KAs containing supplements are associated to low protein diets (LPDs) or very-low protein diets (VLPDs) in patients with CKD4-5 (162, 163); these supplements have phosphate-chelating properties, are likely anabolic, can substantially postpone time to renal replacement therapy and decrease uremic toxicity (162); very recently, they have also been shown to protect patients with diabetic CKD from CV mortality (164).

Ketoacids (KA) of essential AA in general and ketoisocaproate (Kic) in particular, have been shown to reduce muscle protein degradation. In particular, in experimental models of CKD, a LPD supplemented with KAs compared to an LPD alone was able to downregulate the activity of the ubiquitin-proteasome system and to protect skeletal muscle from atrophy and oxidative damage (165). KAs can also maintain the activity of the mitochondrial electron transport chain complex and increase mitochondrial respiration (166). In the perfused heart muscle, branched-chain α -ketoacids (BCKA) increase the phosphorylation of the translational repressor 4E-BP1 as well as multiple proteins in the MEK-ERK pathway, leading to an increase in PS (167).

The Branched-chain amino acid transaminase (BCAT) enzyme reaction is reversible and maintains the equilibrium between the BCAA and their respective KA. Skeletal muscle contributes by the largest fraction of leucine deamination, reamination, and oxidation in the body in humans (168, 169). During fasting leucine is preferentially deaminated to Kic in muscle (168, 169). However, because of the reversibility of the BCAT reaction, the administration of KA can increase intracellular their respective AA level, mostly important the concentration of leucine. In an earlier study, Escobar et al. (170) observed that the infusion of Kic in the neonatal pig, stimulated MPS similarly to infusion of leucine. Both treatments increased intracellular leucine concentrations within the range 2–4-fold, a level considered to be essential for mTORC1 activation. More, recently it has been shown that BCKA are preferentially reaminated and activate protein synthesis in the heart (167) an effect that on a long term, could favor heart hypertrophy in conditions associated with elevated BCAA levels, such as obesity.

Recently, Fuchs et al. (171) were able to show that feeding old male subjects with 6 g BCAA, 6 g BCKA and 30 g milk proteins equally increased MPS; however, following the ingestion of milk protein MPS rates persisted to be high, while the postprandial MPS increase was short-lived following the ingestion of both

BCAA and BCKA. On the one hand these findings suggest that a complete essential AAs pattern needs to be provided to allow a prolonged postprandial increase in MPS; on the other hand, it leads to the speculation that KA are truly anabolic and may be considered to treat PEW in HD patients. However, Li et al. (172) in a small prospective, randomized, controlled, single-center study observed that the KA supplementation did not improve neither plasma AA nor body composition. The ability to use a low-volume/small-quantity BCKA supplement to efficiently stimulate MPS, while avoiding excess N intake, warrants additional trials to define formulas, dosing and examine long-term clinically relevant outcomes.

Endurance and Resistance Training

Early studies in HD patients have suggested that physical exercise prevents cardiovascular diseases, by ameliorating cardiovascular risk factors as well as cardiac autonomic control and left ventricular systolic function (173–175). In the HD-associated catabolic outline, it is not surprising that great interest has been given to physical exercise as a means to improve muscle strength and function. Kopple et al. (176), to study potential mechanisms by which exercise training ameliorates exercise capacity, randomized 80 HD patients into endurance training, strength training, endurance plus strength training or no training. At the end of the intervention, there was no whole body neither regional lean and fat mass in any groups. However, the investigators were able to demonstrate several muscle transcriptional changes that would favor muscle anabolism, including increases in muscle IGF-I and decrease in myostatin (176).

Among the treatment options to prevent loss of muscle mass and function in ESRD patients, endurance or resistance exercise appears to be the most useful. Several cohort studies have investigated the effects of endurance exercise training on physical function, aerobic capacity and muscle strength in HD patients. Results from these trials have showed benefits in terms of physical function, cardiovascular disease (CVD) risk, and quality of life (QOL) [see for review refs. (177, 178)]. Also expert opinion reports (179, 180), position statements (181) and guidelines (182) have suggested that physical exercise needs to be considered as standard of care treatment in patients with CKD, including HD patients. These indications are similar to recommendations for the general population (183).

Either “intradialytic cycling,” with patients cycling on ergometers during the HD session and out-of-center (“interdialytic”) exercise appear to have similar results (184). However, some apparently discrepant results have also been reported (185). Two major RCTs have been recently published on the effect of physical exercise in HD patients. Koh et al. (174) randomized 70 HD patients to intradialytic or home-based exercise training or usual care for 6 months. In the intradialytic arm, patients underwent three training sessions per week on a cycle ergometer, while in the home-based exercise arm patients were provided with a walking program to achieve the same weekly physical activity; the primary outcome was assessed as a change in performance on the 6-min walk test. Unfortunately, the authors found no significant differences among groups in

the 6-month study period (174). More recently, Manfredini et al. (186), in a large 6-month randomized multicenter trial in HD patients, showed that personalized walking exercise program at home was associated with significant benefits on physical function, including the 6-min walking test, five times sit-to-stand test, cognitive function score and quality of social interaction score (186). Importantly, patients with the highest adherence to the exercise protocol had the largest performance improvement, suggesting a dose-response effect from the exercise. More recently, Exel et al. (187) studied 107 HD patients randomly divided into two groups: stretching and resistance exercise. Intervention programs were performed for 8 weeks, three times a week. Resistance exercise caused an increase in muscle strength and distance walked, as compared to stretching (187).

It has been outlined (185) that the reasons for the apparently discrepant results from training studies are possibly related to the need of enrolling a large number of patients to detect significant results. Another problem is the low volume and intensity of the exercise prescription, which may have accounted for the relatively modest functional improvements (185). So a current hypothesis is that exercise interventions might fail to produce clinically significant improvements in HD patients, primarily because the volume and intensity of the exercise prescribed is insufficient (185).

In summary, several meta-analyses (174, 177, 178, 184, 188, 189) show a statistically significant improvement in physical function and quality of life after 3–6 months of endurance exercise training in HD patients. However, no conclusion can be reached for elderly HD patients, since a few studies are existing focusing on them (190). In addition, a major problem stems from the low percentage of HD patients who are able to perform physical exercise. ESRD patient are more commonly sedentary and tend to avoid exercise (191); additionally, the dialysis population is more often aged, frail, depressive and with many comorbid conditions which are major obstacles to exercising. Finally, often the HD personnel has lack of expertise or resource to start an exercise program (192).

Combining Nutrition and Physical Exercise

The combination of physical exercise and nutrition provides the strongest muscle anabolic stimulus in humans. In the elderly, who are “anabolic resistant” combining exercise with protein supplementation provides an enhanced anabolic response (193). Several studies have addressed the issue of the efficacy of this combined treatment in HD patients [see ref. (192) for review]. Likewise, early studies in HD patients has showed that a single bout of either endurance or resistance exercise enhances the anabolic response to nutritional supplements (194). However, the results from long-term interventions have been less favorable (132, 195–197).

Many factors are likely to contribute to the modest advantages of exercise in HD patients, including limited exercise prescriptions and intradialytic training (192). A greater exercise dose or enhanced nutritional support may be needed to demonstrate the potentiated additive benefits of these treatments (192). Clearly, defining the role of exercise in CKD remains a top research priority.

Other Treatments to Increase MPS and Muscle Mass

Beta-Hydroxy-Beta-Methyl Butyrate

HMB is a downstream metabolite of leucine. HMB is present at low concentrations in muscle, and its turnover is about 0.7% of that of leucine (198). Better than being used alone, HMB is often consumed with other AAs or as part of a multiple ONS. Initial studies on the use of HMB in healthy volunteers demonstrated a high anabolic effect (199). Despite some RCTs have not detected any positive effects on muscle mass, strength and function from HMB supplementation, HMB is still considered a nutritional compound that may possess the potential to attenuate the rate of muscle loss in conditions of anabolic resistance (198, 200), probably as a long-term treatment (201).

There is very little information on HMB metabolism in CKD. A large percentage of plasma HMB is excreted into urine; however, from the observation that leucine and Kic pools are reduced in CKD4-5 (94), one would imply that also HMB is reduced in muscle. We also have no information on the effects of HD on HMB plasma levels. Since HMB molecular weight is low (m.w.118) it is likely removed by HD. HMB is available in many ONS on the market, but whether it can promote anabolism in dialysis-treated patients is not known. Fitschen et al. (202) observed no significant effect of HMB supplementation on body composition, bone density, strength, fall risk and quality of life in a double-blind, placebo-controlled, randomized trial in 35 HD patients. However, on analysis of plasma HMB concentrations, 5 of 16 patients (31%) in the HMB arm were found to be non-compliant at 3 or 6 months (202). Therefore, the role of HMB supplementation in patients who are on dialysis is still not completely explored.

Omega 3 (n-3) Fatty Acids

Omega 3 (n-3) Fatty Acids (n-3 FA) have been shown in both young and older adults to have favorable effects on muscle insulin sensitivity, inflammation and anabolism (203–205). McGlory et al. (206), by utilizing intravenous hyperinsulinemic and hyperaminoacidemic clamps have shown that a moderate dose (~4 g) of n-3 FA supplementation augments MPS rates both in healthy young and older adults (206).

Deger et al. (207) studied the effect of the administration of n-3 FA (2.9 g/d) over 12 weeks on muscle protein turnover in HD patients with systemic inflammation. N-3 FA supplementation was associated with decrease of forearm MPD but did neither influence MPS nor muscle net protein balance. In a recent meta-analysis Rondanelli et al. (208) studied the effect of n-3 FA and docosahexaenoic acid (DHA) supplementation on fat free mass and physical performance in patients with various chronic diseases. Daily n-3 FA + DHA supplementation (from 0.7 to 3.36 g) decreased the time of Time Up and Go (TUG) test, and the fat free mass had an improvement trend which was however not statistically significant. Overall, these data suggest that n-3 FA + DHA might have a positive effect on physical performance, and that their use may improve some sarcopenia component.

Testosterone

Hypogonadism is commonly observed in men with CKD, which occurs as an effect of both depressed hypothalamic-pituitary-gonadal axis functionality and androgen synthesis (209, 210). Hypogonadism may contribute to several common adult complications in males with CKD (209, 210). Testosterone's downward signal acts to offset some of the catabolic pathways which are activated by uremia. Testosterone promotes satellite cell recruitment and increases MPS through stimulation of androgen receptors and activation of the insulin-like growth factor-1 (IGF-1) pathway (211). In humans, the anabolic action of testosterone is mediated via an enhancement of the efficiency of MPS, i.e., the MPS rate relative to the availability of AA (211). Long-term testosterone users develop hypertrophy of both Type I and Type II muscle fibers (211).

Low testosterone levels are associated with reduced fat free mass and muscle strength both in CKD and in HD patients (209, 210). Furthermore, Chiang et al. (212) recently were able to show that low plasma testosterone is strictly related to declining physical function, frailty, and muscle wasting in HD patients, which suggests that the CKD-associated decrease in testosterone levels may contribute to the procatabolic environment. However, current guidelines do not recommend routinely prescribing testosterone to all elderly men with low testosterone concentrations, but suggest that testosterone therapy is offered on an individualized basis, after explicit discussion of its potential risks and benefits, to elderly men who have symptoms suggestive of testosterone deficiency and consistently low morning testosterone concentrations (211).

Growth Hormone

Growth Hormone (GH) is anabolic both through the IGF-I downward pathway and a direct effect on skeletal muscle (213). GH deficiency is associated with muscle atrophy and GH administration causes muscle hypertrophy (214). GH treatment has been reported to have beneficial effects in elderly subjects (215). Although GH and IGF-I replacement in GH deficient adults has proven anabolic, this is not an unequivocal finding, possibly due to differences in dosage, and time on treatment (216).

Short stature is commonly observed in children with CKD, even after renal transplantation. The CKD-related GH insensitivity is characterized by deficiency of functional IGF-1, and can be overcome by the administration of supraphysiological doses of recombinant human GH (rhGH); long-term rhGH treatment stimulates IGF1 synthesis, increases longitudinal growth and likely improves adult height (216).

RhGH has been licensed for the treatment of CKD-related growth failure in Europe, North America and many other high-income countries (217). In children, CKD is associated to significant muscle wasting (217). Concurrent rhGH therapy causes higher leg lean body mass Z-scores, compared to untreated-children (218). Similarly, small randomized trials of rhGH in malnourished adult dialysis patients showed benefits including weight gain, improved MPS (219) and performance (220, 221). The OPPORTUNITY™ Trial (222) examined whether rhGH reduces mortality in hypoalbuminemic HD

patients. Secondary end points were effects on number of hospitalizations, cardiovascular events, lean body mass (LBM), serum proteins, exercise capacity, QoL and adverse events. Although the OPPORTUNITY™ Trial was terminated early owing to slow recruitment, treatment with rhGH, compared to placebo, improved certain factors associated with cardiovascular disease risk factors, without adverse outcomes.

Denosumab

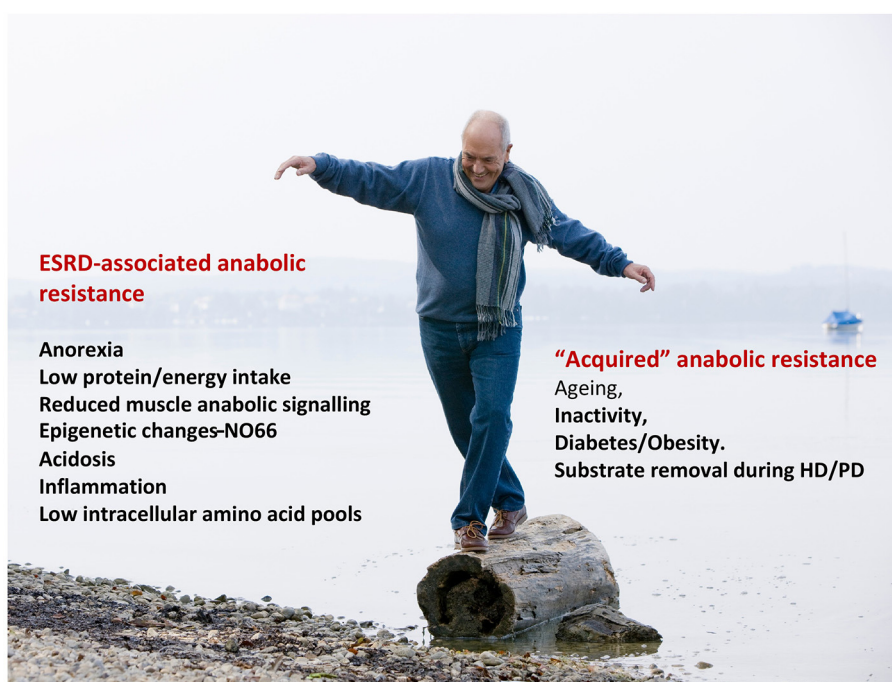
In postmenopausal women, a major role in the development of osteoporosis is played by the increased activity of receptor activator of nuclear factor kappa-B ligand (RANKL) (223). The binding of RANKL to its cognate receptor RANK drives a series of events triggering differentiation, activity, and survival of osteoclasts. Denosumab is a monoclonal antibody which binds RANKL, thus interfering with osteoclast differentiation, activation and survival. Denosumab is currently approved for the treatment of osteoporosis in postmenopausal women at risk of fracture. It is interesting that in elderly subjects osteoporosis and sarcopenia have similar risk factors, underscoring muscle-bone interactions, which may result in wasting, falls, and fractures. The combined existence of osteoporosis and sarcopenia (osteosarcopenia) is an emerging definition, which is cause of a significant health burden (224). Kirk et al. (224) observed in a large cohort of community-dwelling older adults that risk factors associated with osteosarcopenia include older age, female gender, physical inactivity, low body mass index, higher fat mass and the coexistence of chronic diseases, including CKD. Osteosarcopenia is suggested to be caused by reduced

mechanical loading, and altered crosstalk between muscle, bone, and fat cells. It is interesting that, in addition to nutritional treatment and physical exercise, Denosumab, may be offered to osteosarcopenic patients (225). By inhibiting RANKL, Denosumab decreases bone resorption, increases bone mineral density (BMD), and reduces new fractures in postmenopausal women with osteoporosis (225). In a non-randomized study of community-dwelling older adults denosumab treatment improved balance, fear of falling, and physical function (226), suggesting that denosumab may increase muscle strength and mass.

Studies have examined the denosumab-activated pathways in skeletal muscle. RANK is also expressed in skeletal muscle and can activate the NF- κ B pathway, which mainly inhibits AKT/mTORC1 signal and myogenic differentiation, leading to muscle loss (227, 228). Mice overexpressing RANKL (HuRANKL-Tg+) undergo muscle atrophy, lower limb force and maximal speed (228). Interestingly, the administration of denosumab to HuRANKL-Tg+ mice increases limb force and muscle mass, increases muscle insulin sensitivity and downregulates myostatin gene expression (228). These findings suggest that RANKL decreases, while its inhibition improves, muscle strength and insulin sensitivity both in osteoporotic mice and humans. In dialysis patients one might expect a similar effect, although no study is available.

Vitamin D

Vitamin D deficiency is associated with muscle functional impairment and falls (229). Even if the vitamin D receptor



ESRD-associated anabolic resistance

Anorexia
Low protein/energy intake
Reduced muscle anabolic signalling
Epigenetic changes-NO66
Acidosis
Inflammation
Low intracellular amino acid pools

“Acquired” anabolic resistance

Ageing,
Inactivity,
Diabetes/Obesity.
Substrate removal during HD/PD

FIGURE 2 | The dialysis patient balancing between the anabolic resistance associated to End Stage Renal Disease (ESRD) and the “acquired” anabolic resistance led by advancing age, inactivity, diabetes and substrate depletion during dialysis treatments (Shutterstock source).

(VDR) is expressed at low levels in skeletal muscle, the deletion of VDR in the myocyte decreases muscle size and strength (230). Vitamin D has in muscle both genomic and non-genomic effects, which regulate cellular differentiation and proliferation. The non-genomic effects include the regulation of membrane calcium channels (which suggest a role for vitamin D in muscle contraction) mitochondrial function, insulin signaling and muscle substrate metabolism. In addition, studies conducted in cell culture systems and animals suggest that both vitamin D and conjugated linoleic acids (CLAs) stimulate MPS. Although some studies have shown that supplementation with vitamin D in the general population has a positive effect on muscle function, including athletic performance, falls and strength (231, 232), not all results are univocal. In a recent trial in 32 sedentary, older adults both Vitamin D and/or CLA supplementation, did not have effects on MPS (233). Similarly, there is not enough evidence to understand the role of vitamin D on musculoskeletal outcomes in the CKD population (231). However, given that this lack of evidence does not necessarily indicate no effect on musculoskeletal health (231), vitamin D might be considered to improve muscle strength and physical performance in renal patients, especially those who have low 25(OH)D plasma levels (<20 ng/mL–50 nmol/L).

Other Micronutrients

There is some evidence that the MPS response to feeding may be augmented by the use of micronutrients, mainly those contained in the yolk (vitamin D, vitamin E, vitamin A, zinc, selenium, and cholesterol) (234, 235). However, if and to what extent micronutrients potentiate the anabolic effects of protein feeding or physical exercise in dialysis-treated patients still needs to be established.

Other Treatments

Several other targets to increase MPS, such as targeting pro-inflammatory cytokines, manipulation of transforming growth

factor (TGF) family members, satellite cells and stimulation of mitochondrial biogenesis, which are currently under study, are reviewed elsewhere (3, 236).

In conclusion, the importance of maintaining muscle mass and MPS is often overlooked in patients with kidney diseases. HD patients are often exposed to AA or protein deprivation, which causes low circulating and tissue levels of essential AAs. Anorexia or fasting prescribed during the dialytic treatments can potentially decrease MPS. These settings associate with several abnormalities occurring in CKD that stimulate protein degradation and/or decrease MPS. Recent observations suggest that skeletal muscle is “anabolic resistant” in HD patients and that greater amounts of proteins than in healthy subjects are needed to maximally stimulate MPS. Accordingly, the dialysis patient needs to cope with the CKD-associated intrinsic anabolic resistance, and the “acquired” anabolic resistance led by advancing aging, inactivity, diabetes and substrate depletion (Figure 2). The combination of physical exercise and protein/AA feeding provides the strongest muscle anabolic stimulus in humans; however the major gaps in our current knowledge of nutritional treatment of dialysis-treated patients with CKD include optimal formulas and amount of protein, ONS and exercise paradigms, and research on how to incorporate effective management approaches into clinical care.

AUTHOR CONTRIBUTIONS

GG and DV conceived the review. MS and FA performed literature research. GG, FV, GB, and DV wrote the manuscript. ER, DP, AL, and PE reviewed and edited the manuscript. All authors have read and agreed to the published version of the manuscript.

FUNDING

This study was supported by grants from the University of Genoa (Finanziamento di Ateneo).

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Estimation of Muscle Mass in the Integrated Assessment of Patients on Hemodialysis

Alice Sabatino¹, Natascha J. H. Broers^{2,3}, Frank M. van der Sande^{2,3},
Marc H. Hemmelder^{2,3}, Enrico Fiaccadori¹ and Jeroen P. Kooman^{2,3*}

¹ Nephrology Unit, Department of Medicine and Surgery, Parma University Hospital, University of Parma, Parma, Italy,

² Division on Nephrology, Department of Internal Medicine, Maastricht University Medical Centre, Maastricht, Netherlands,

³ NUTRIM School of Nutrition and Translational Research in Metabolism, CARIM School for Cardiovascular Diseases, Maastricht University, Maastricht, Netherlands

OPEN ACCESS

Edited by:

Michele Barone,
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Reviewed by:

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Queensland, Australia
Calvin Lloyd Cole,
University of Rochester, United States

*Correspondence:

Jeroen P. Kooman
Jeroen.kooman@mumc.nl

Specialty section:

This article was submitted to
Clinical Nutrition,
a section of the journal
Frontiers in Nutrition

Received: 19 April 2021

Accepted: 16 July 2021

Published: 16 August 2021

Citation:

Sabatino A, Broers NJH, van der
Sande FM, Hemmelder MH,
Fiaccadori E and Kooman JP (2021)
Estimation of Muscle Mass in the
Integrated Assessment of Patients on
Hemodialysis. *Front. Nutr.* 8:697523.
doi: 10.3389/fnut.2021.697523

Assessment of muscle mass (MM) or its proxies, lean tissue mass (LTM) or fat-free mass (FFM), is an integral part of the diagnosis of protein-energy wasting (PEW) and sarcopenia in patients on hemodialysis (HD). Both sarcopenia and PEW are related to a loss of functionality and also increased morbidity and mortality in this patient population. However, loss of MM is a part of a wider spectrum, including inflammation and fluid overload. As both sarcopenia and PEW are amendable to treatment, estimation of MM regularly is therefore of major clinical relevance. Whereas, computer-assisted tomography (CT) or dual-energy X-ray absorptiometry (DXA) is considered a reference method, it is unsuitable as a method for routine clinical monitoring. In this review, different bedside methods to estimate MM or its proxies in patients on HD will be discussed, with emphasis on biochemical methods, simplified creatinine index (SCI), bioimpedance spectroscopy (BIS), and muscle ultrasound (US). Body composition parameters of all methods are related to the outcome and appear relevant in clinical practice. The US is the only parameter by which muscle dimensions are measured. BIS and SCI are also dependent on either theoretical assumptions or the use of population-specific regression equations. Potential caveats of the methods are that SCI can be influenced by residual renal function, BIS can be influenced by fluid overload, although the latter may be circumvented by the use of a three-compartment model, and that muscle US reflects regional and not whole body MM. In conclusion, both SCI and BIS as well as muscle US are all valuable methods that can be applied for bedside nutritional assessment in patients on HD and appear suitable for routine follow-up. The choice for either method depends on local preferences. However, estimation of MM or its proxies should always be part of a multidimensional assessment of the patient followed by a personalized treatment strategy.

Keywords: ultrasound, bioimpedance, sarcopenia, hemodialysis, creatinine index

INTRODUCTION

Assessment of nutritional state is of high relevance in patients on hemodialysis (HD). This given the relation between protein-energy wasting (PEW), a condition characterized by reduced body stores of protein and energy fuels characteristic of patients with chronic kidney disease (CKD) and end-stage kidney disease (ESKD), and mortality (1), and since abnormalities in the nutritional state may be amenable to therapeutic intervention (2). Muscle wasting in patients on HD can be due to

multiple factors including insufficient dietary intake and a loss of nutrients through the dialysate, or an increased muscle breakdown due to inflammation or metabolic acidosis (3). Measurement of muscle mass (MM), or its proxies lean tissue mass (LTM) or fat-free mass (FFM), is an integral part of the assessment of the nutritional state, as well as in the diagnosis of sarcopenia. Sarcopenia is characterized by reduced MM and strength and is frequently observed in the elderly, but can also happen earlier as a consequence of chronic conditions, such as CKD/ESKD and patients on HD (3–5). Malnutrition as well as sarcopenia are part of a spectrum including impaired functional status, low physical activity, low quality of life, and frailty (6–8), and they are the important components of the premature aging phenomenon in this patient population (9). Next to this, inflammation and fluid overload were also found to be related to a decrease in LTM and intracellular water (ICW) (8, 10). Thus, loss of MM is a central part of the multimorbid spectrum of patients on HD, and should be interpreted in view of both its consequences as well as in the context of potentially amendable underlying factors. The aim of this short review is to give a concise overview of instrumental methods that can be used on a daily clinical basis in patients on HD, and their use in the context of a multidimensional assessment in these patients will be discussed.

CLINICAL SYNDROMES ASSOCIATED WITH A LOSS OF MM

Loss of MM or its proxies is included in various syndromes related to the nutritional and functional status of the patient on HD, as summarized in **Table 1**. Except frailty, in which only a reduction in muscle strength is a parameter, a reduction of MM or FFM is included in the diagnostic criteria of other syndromes, such as PEW, cachexia, and sarcopenia. These partly, but not entirely, overlapping syndromes (3, 11) are part of a wide spectrum of nutritional and functional abnormalities in patients on HD, although an important common denominator appears to be tissue loss (11). Importantly, one of the criteria for the definition of PEW, also referred to as kidney cachexia, is an increase in inflammatory parameters (4, 12). In contrast, inflammation is not included in the diagnostic criteria of sarcopenia (13). This division is relevant, as the pathophysiology and also possibly the clinical approach to a patient on HD with a pure “sarcopenic” phenotype may differ from that of a patient with a “cachectic” phenotype (14). Furthermore, patients can have both an increase in fat mass and a decline in MM (15), an entity also known as sarcopenic obesity, which is prevalent in patients on HD, although its relation with the outcome is yet uncertain (16, 17). The development of sarcopenic obesity is not captured by the estimation of changes in body weight or body mass index (BMI) (18).

The assessment of body composition is complicated by the fact that various parameters are used to express (loss) of MM or LBM. For instance, LBM, FFM, and MM are not equivalent, although they are often used as interchangeable surrogates. FFM, as the name suggests, is the total body mass except for the

body fat, and it includes LBM and bone mineral tissue. The LBM, in turn, is composed of the total body water, appendicular skeletal muscle mass (ASMM), and the fat-free mass of organs. Since different techniques measure different compartments, the identification of the body compartment of interest, along with the availability of the method, must precede the choice of the method of assessment. As an example, some available techniques, such as bioelectrical impedance analysis (BIA), assess body composition by dividing the body into two compartments (2-C), the FFM, which conceptually includes all non-fat tissue and the fat mass (FM) (19). While other methods, such as dual-energy X-ray absorptiometry (DXA) and one application of bioelectrical impedance spectroscopy (BIS), divides the body into three compartments (3C). DXA assesses LBM, which includes total body protein and total body water (TBW), but excludes bone and fat mass (20, 21) (**Figure 1**). On the other hand, 3C-BIS assesses a “normohydrated” LTM, which reflects a compartment that is separate from adipose tissue mass (ATM; fat mass and adipose water) and a virtual “overhydration” compartment, but that includes bone mineral tissue (8) (**Figure 1**), as will be discussed later in more detail.

These specific examples show that although all entities reflect comparable physiological dimensions, they are not interchangeable. FFM assessed by a 2-C model can be different from LTM assessed by a 3-C model (22). Thus, parameters obtained by a specific method cannot be used interchangeably with comparable parameters assessed by different methodologies. In addition, ideally, reference values should also be developed for specific techniques, devices, and populations. As this may not always be feasible, it is important to have these caveats in mind when assessing literature on body composition in patients on kidney replacement therapy (KRT).

BEDSIDE TECHNOLOGY FOR THE ASSESSMENT OF MM

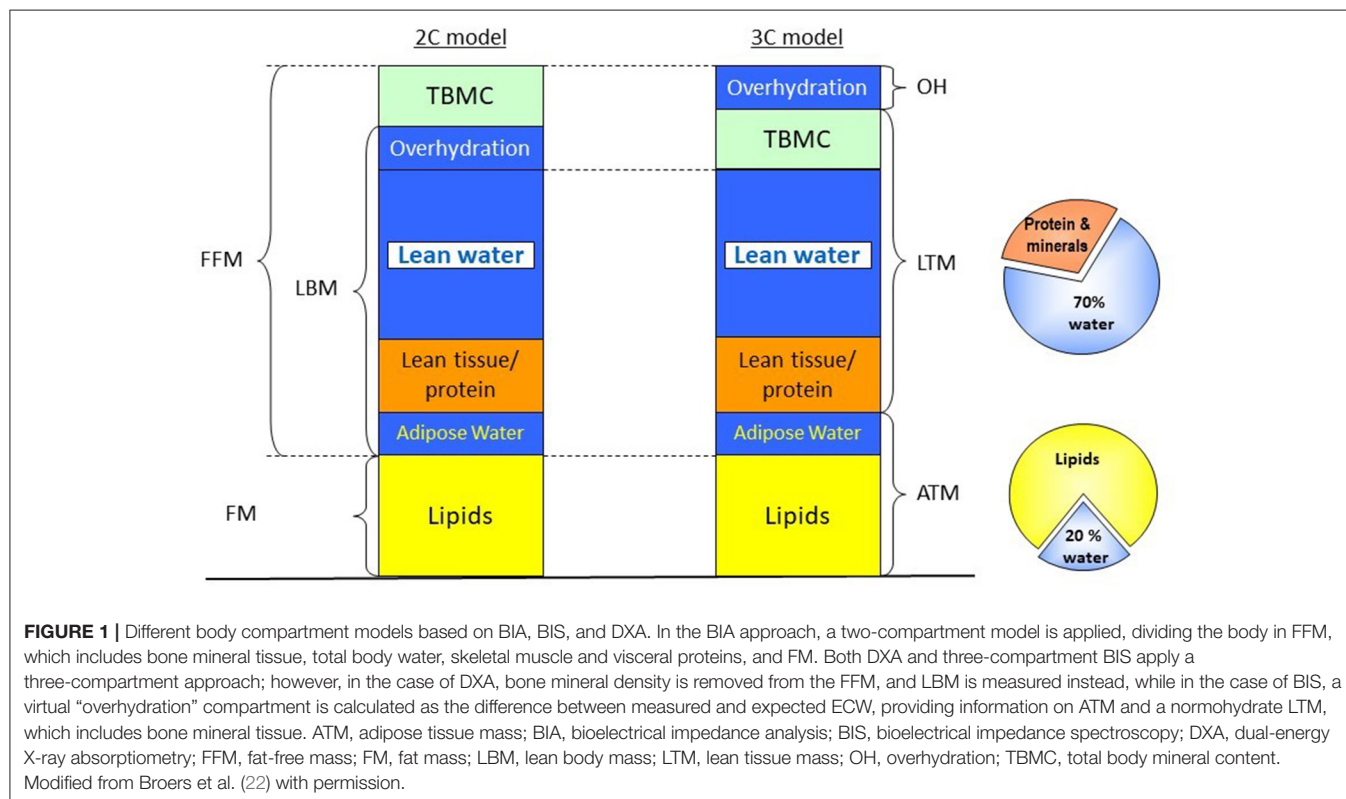
For the assessment of MM or its proxies, various options are available. Computer-assisted tomography (CT) or MRI are considered gold standard methods but are impractical to be used on a routine basis (13, 23). DXA is generally considered a reference method to estimate LBM as well as ASMM in guidelines (13) but may be difficult to perform frequently in clinical practice. Furthermore, because DXA assumes a hydration ratio with LBM of 0.73 (24), results can be influenced by severe fluid overload (25). Still, DXA, when routinely available, provides important information on changes in body composition on HD and might also serve as a calibration for bedside methods.

Various methods are available to assess MM or LBM in patients which can be used on a routine basis in patients on KRT. These can be conceptually divided into methods that indirectly estimate body composition (such as bioimpedance or anthropometry), biochemical methods (based on creatinine kinetics), and methods that measure muscle dimensions at an anatomical level [MRI, CT, or ultrasound (US)]. Anthropometry is a time-honored method that is also included in the original diagnostic criteria for PEW (4). When performed by a skilled

TABLE 1 | Categories of assessment for the diagnosis of different syndromes related to the nutritional and functional status of patients.

Criteria	Malnutrition ESPEN	Malnutrition GLIM	PEW	Cachexia	Sarcopenia	Frailty
Weight loss/BMI	+	+	+	+		+
Muscle mass/FFM/LTM	+	+	+	+	+	
Muscle strength					+	+
Functional performance					+	+
Fatigue/exhaustion						+
Biochemical markers			+	+		
Etiology		+				

BMI, body mass index; FFM, fat free mass; LTM, lean tissue mass; PEW, protein energy wasting.



investigator it was able to predict a reduction in MM with an accuracy comparable with instrumental methods (16). However, the emphasis of the present article is on biochemical and technological tools in the assessment of body composition.

Bioelectrical Impedance

Although the theoretical backgrounds of BIA are complex and discussed in excellent reviews (26, 27), in general, this method measures the opposition (impedance) of the body or a body segment to an alternating current. The impedance (Z) is a composite of the resistance to this flow, which is related to TBW, and reactance (X_c) is related to the capacitance of the cellular membrane. With the single-frequency (SF) approach, FFM and ASMM are estimated using population-derived regression equations, including Z or R , measured at 50 kHz as the resistance index $H^2/R50$ along with anthropometric parameters, sex, and

age (27). With SF-BIA, also ASMM can be estimated using regression equations (27, 28).

The multifrequency (MF) approach delivers different frequencies that vary from 5 to 1,000 kHz, and depending on the method it can use only several frequencies or, in the case of BIS, broadband of frequencies within this range. The lower the frequency, the more the difficulty with which it has to pass across the cell, passing only at the extracellular water (ECW) at frequencies < than 1 kHz and not through the ICW, with higher frequencies it passes through both, with TBW being measured at frequencies > 5,000 kHz. For technical reasons, measurements at very low and very high frequencies are not possible; however, with BIS, the resistance at zero (R_0) and infinity (R_∞) are extrapolated by applying a Cole-Cole plot to predict ECW and TBW (27). In classic 2C-models of SF-BIA and MF-BIA, FFM is subsequently calculated from TBW, assuming fractional

hydration of 0.73 (29). A drawback of this method, which divides the body into two compartments, is that the excess of ECW due to fluid overload is added to the TBW which can subsequently result in an overestimation of FFM (25) (**Figure 1**). In the case of BIS, Moissl et al. developed a model for the assessment of ECW, ICW, and TBW using Hanai mixture theory adjusted for BMI, which showed good agreement with dilution methods (30). Chamney et al. further developed the so-called 3C-model, which assumes fixed hydration of LTM and ATM, and divides the body in normohydrate LTM and ATM, and a virtual “overhydration” (OH) compartment (31).

With regard to the estimation of body composition, a definite superiority of either approach has not yet been proven. Donadio et al. found a slightly lower prediction error for FFM with SF-BIA as compared with an MF-BIA using a two-compartment approach, with a highly significant correlation of both methods with DXA (32). Another study observed a stronger relation between FFM estimated by MF-BIA and creatinine kinetics as compared with SF-BIA (33). Raimann et al. found a slightly improved estimation of ICW with the SF method, and conversely, improved estimation of ECW with BIS. With regard to the detection of changes in ECW, MF-BIA was found to have a higher precision (34, 35). On the other hand, ASMM predicted by the Sergi equation using SF-BIA showed high accuracy in predicting sarcopenia with DXA as the reference method (16); however, this equation with the form: $ASMM\ (kg) = -3.964 + (0.227 \times [height\ (cm)]^2/R) + (0.095 \times weight) + (1.384 \times sex) + (0.064 \times Xc)$, was primarily validated in an elderly Caucasian population (36).

With the 3C BIS method, estimation of body composition is based on a theoretical approach without the use of population-specific regression equations. Lean tissue index (LTI), which corresponds to LTM, divided by $height^2$, below the 10th percentile of a healthy age-matched reference population was independently related to outcome in most, but not all studies (8, 15, 37, 38). Still, in a meta-analysis including over 15,000 patients, a low LTI was associated with increased mortality [Hazard ratio 1.53 (95% CI: 1.41–1.64)] (39). Especially, the combination of a low LTI and fat tissue index (FTI, the $height^2$ -normalized ATM) appears to be associated with increased mortality risk (15).

However, in several studies, despite reasonable agreement at a population level, relatively wide limits of agreement were observed between body compartments assessed by 3C-BIS and reference techniques, such as DXA (40, 41). To some degree, these differences may be explained by the fact that even the reference method is not free of errors. Indeed, the excess ECW with overhydration is added to the LBM compartment with DXA, but not with the 3C-BIS approach (8). Also, it should be taken into account that ATM assessed by 3C-BIS includes intra adipose water, unlike FM measured by DXA. Using ASMM measured by DXA as the reference method, LTM measured by whole-body BIS was able to predict sarcopenia with acceptable accuracy (mean AUC 0.79 for females and 0.77 for males) (16); however, it should be noted that in the current EWGSOP2 guidelines, cut-off values based on the 3C-BIS model were not yet included in the definition, while the Sergi equation based on SF BIA was advocated for standardization (13). However, in the case of tissue edema, these estimations may be less reliable in patients on HD.

To standardize measurements and avoid this kind of problem, the recent KDOQI guidelines on nutrition in CKD recommends that BIA/BIS should be performed at least 30 min after the HD session to allow for the distribution of body fluids (42). Still, in the case of 3C-BIS, measurements of ATM and normohydrated LTM were slightly (0.77 and 0.40 kg, respectively) affected by the timing of measurements (43), whereas predialytic fluid status was more consistently related to the outcome as compared with postdialytic measurements (44).

An alternative approach is the construction of a vector plotting R and Xc at 50 kHz within tolerance ellipses of a healthy population. The advantage of this method is that results are displayed without the need for underlying theoretical assumptions or population-based equations (28). A potential disadvantage is that the direct translation of the findings into constructs such as sarcopenia, fluid overload, and PEW may be more difficult as compared with a numerical approach.

To summarize, whereas various BIA approaches can be used to assess body composition in dialysis patients, it must be kept in mind that estimations are dependent on population-specific regression equations or theoretical assumptions regarding the conversion from bioelectrical signals to estimations of body water compartments. Measurements obtained with a specific device or method can therefore not be used interchangeably (45), even with regard to raw parameters such as Z, Xc, and R (46). It is important to acknowledge that as long as a device is correctly calibrated, the magnitude of the differences are small and generally within the precision of the specifications of the manufacturer. Whereas, a definite superiority of a specific BIA methodology for estimations of body composition has not been proven, in our view, the 3-C model holds the advantage that it also provides a separate estimation of fluid status in a single measurement, whereas it has shown high-construct validity in predicting outcome in large datasets.

Biochemical Methods: The Creatinine Index

Serum creatinine is a breakdown product of creatine phosphate in muscle tissue that was found to be strongly related to LBM assessed by DXA in patients on HD (47). Serum creatinine is inversely related to mortality in patients on renal replacement therapy (48); however, serum creatinine in patients on HD is also dependent on dialysis adequacy. Therefore, the concept of creatinine index (CI) was developed (49). CI was found to be an independent predictor of outcome in patients with HD (50). However, as creatinine kinetics may be complicated to use in routine clinical practice, a simplified form [simplified creatinine index (SCI)] was developed, which was also found to be related to the outcome (51). SCI (mg/kg/d) is calculated according to the formula $16.21 + 1.12 \times [1\ \text{if male}; 0\ \text{if female}] - 0.06 \times \text{age (years)} - 0.08 \times \text{spKt/V urea} + 0.009 \times \text{predialytic serum creatinine } (\mu\text{mol/l})$. Also, LTI derived from SCI was strongly related to LTI assessed by BIS, although the mean BIS-derived value was 4.7 kg lower than the SCI-estimated value (51). SCI is easy to apply in clinical practice as only routinely gathered data that are already present in electronic health records (EHR) are needed,

with the advantage that longitudinal trends can be tracked easily. Indeed, SCI declined 6 months before death, potentially serving as an early warning sign (51). However, a potential pitfall in the follow-up of the SCI is changes in residual renal function, which may independently affect serum creatinine values apart from muscle mass, as well as changes in dietary intake (42). Serum creatinine is also a parameter included in the nutritional component score (NCS), an aggregate score that also consists of routinely captured parameters interdialytic weight gain, serum phosphate, serum albumin, and normalized protein catabolic rate (nPCR). The use of this score, which was shown to decline 1–2 months before hospitalization and also up to 6 months before death (52, 53), allows for the interpretation of changes in parameters like interdialytic weight gain (IDWG) and serum phosphate, which have a bidirectional relationship with the outcome (54). Whereas, a decrease in these parameters is usually regarded as a positive sign given their detrimental effect on the cardiovascular status of the patient, a sharp decline in IDWG and serum phosphate can also be a sign of impending malnutrition and adverse outcomes when accompanied by a decrease in the other nutritional parameters (52).

To Summarize, following trends in biochemical indices derived from EHR can aid in the early detection of changes in nutritional state and MM, whereas changes in serum creatinine or SCI cannot replace validated questionnaires to establish the risk for sarcopenia (55), they can aid in case finding given the fact that they can be performed on a frequent and routine basis with the potential for automated processing of the findings.

Muscle Ultrasound

Recently, regional muscle US has been applied in patients with kidney disease for the assessment and monitoring of skeletal muscle. Its major advantages, compared to other imaging techniques, are represented by lower cost, portability, lack of radiation exposure, and the possibility to be applied by non-specialized staff (56–58). In comparison to other bedside techniques, such as anthropometry, US allows real-time visualization of the target structure, allowing for the assessment of muscle size (thickness and area) and/or quality, through echogenicity, which provides information about the presence of inflammation, fibrosis, and adipose infiltration (59). Its portability is of particular interest in the CKD research setting and clinical practice since patients can be evaluated during HD session or outpatient visits.

Quadriceps muscle US has been studied extensively in patients with renal disease. In the available research, two muscles were most frequently studied, the quadriceps rectus femoris (RF) and vastus intermedius (VI), in two different points, the midpoint, and at the border of the lower third and upper two-thirds between the anterior superior iliac spine and the upper pole of the patella (**Figure 2**) (56, 57, 60–62). Abundant contact gel to avoid any pressure is needed to prevent muscle deformation. Regarding the accuracy and reproducibility of the method, its reliability has been tested in critically ill patients with acute kidney injury (AKI), showing excellent intraclass correlation coefficients (ICC) for inter- and intra-operator comparisons, as well as for measurements performed before and after HD (56).



FIGURE 2 | Quadriceps muscle ultrasound (US) methodology. The points of interest correspond to the midpoint and the lower third between the anterior superior iliac spine (ASIS) and the upper pole of the patella. Using a B-mode ultrasound with a linear transducer, we obtain the image on the right. Rectus femoris (RF) and vastus intermedius (VI) thickness, a measure on the inner edge of the muscle.

In the same clinical setting, US assessment of quadriceps muscle has also been validated against CT (60), showing small and non-significant differential and proportional bias in comparison with CT (60). Also in patients with AKI, US was successfully used to monitor the quadriceps muscle in the first 5 days of stay in the intensive care unit, being able to identify early muscle loss (63). In non-acutely ill patients, US was performed in patients on HD before and after the dialysis session to assess whether the presence of fluid overload or the rapid fluid shifts caused by the treatment could influence measurements (57). No differences were found between measurements performed before and after dialysis, and the correlation between measurements was very high, ranging from 0.91 to 0.95 (57), showing that muscle US is not influenced by fluid overload. Also in the outpatient setting, the assessment of muscle cross-sectional area (CSA) was validated using CT in patients with CKD and not on HD (61). In another study, RF-CSA was assessed before and after 12 weeks of resistance exercise in patients with CKD not on HD, showing a high correlation with MRI at baseline and follow-up, and moderate positive association observed between changes in RF-CSA by US and quadriceps volume by MRI following exercise training (64).

More recently, studies investigating the role of quadriceps muscle US in identifying patients with PEW or sarcopenia have been published. Sabatino et al. (57) used the PEW cutoffs for BMI and albumin to stratify patients on chronic HD in two groups and found that in the multivariable analysis patients with lower BMI had lower muscle thickness, whereas no difference was found between patients with serum albumin

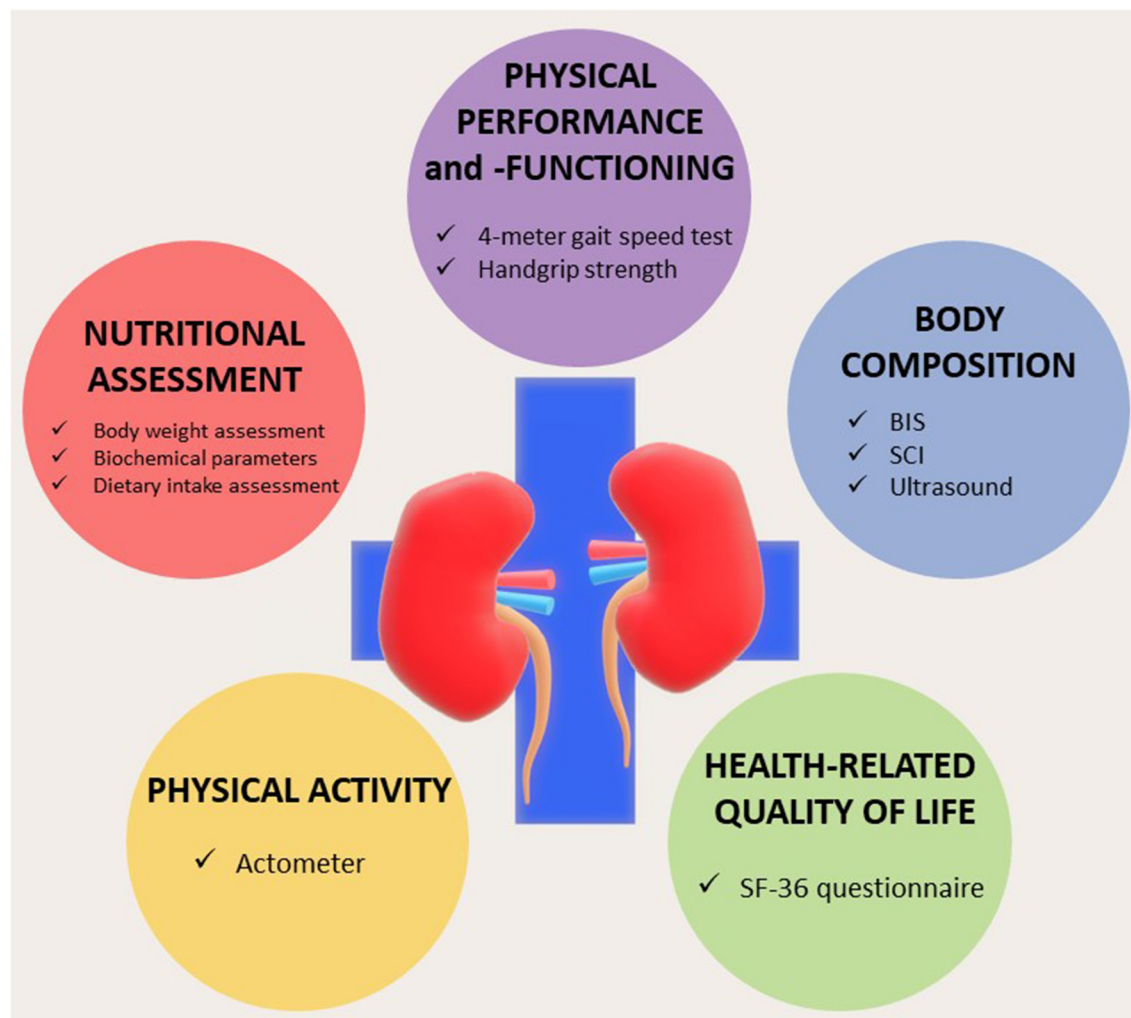


FIGURE 3 | Dashboard summarizing the integrated nutritional assessment of patients with chronic kidney disease (CKD)/end-stage kidney disease (ESKD). BIS, bioelectrical impedance spectroscopy; SCI, simplified creatinine index. Modified from Broers et al. (8) under Creative Commons.

below or above the reference value (57). A similar analysis was performed using the malnutrition inflammation score (MIS), an internationally recognized tool to assess malnutrition and predict outcomes in patients on HD, and found that patients with worse scores (≥ 6) had lower RF and VI thickness (57). In another study, quadriceps CSA cut-offs have been derived using receiver-operation characteristic curves based on the presence or absence of PEW in a Malaysian population (62). In that study, patients diagnosed with PEW had significantly lower RF and VI thickness and RF CSA in comparison with well-nourished patients, and the area under the curve (AUC) for RF CSA was high (men = 0.74, 95% CI: 0.66–0.82 and women = 0.82, 95% CI: 0.73–0.91, both $p < 0.001$). In addition, the correlation between US and LTI by BIS ranged from moderate to high (0.28–0.52) depending on the measurement site, with a higher correlation for RF thickness and CSA in comparison to VI.

Despite such encouraging results, more work is needed before assuming muscle US as a reference method for the diagnosis of sarcopenia. Studies defining reference values derived from healthy subjects from populations with different ethnic backgrounds should be performed to allow the early identification of patients with low MM. However, considering its validity, reliability, and sensitivity in detecting changes in skeletal muscle, its use as a tool for the monitoring of the regional muscularity of a patient could be recommended.

The Role of Estimation of MM in the Integrated Functional Assessment of Patients on KRT

Assessment of MM or its derivatives achieves its full potential in combination with other parameters. As shown in **Table 1**, with a relatively limited battery of measurements, various clinical

syndromes such as PEW, sarcopenia, and frailty can be easily diagnosed. For the diagnosis of sarcopenia, it should be combined with an assessment of muscle strength, for instance, handgrip strength (HGS), and in case of a positive diagnosis, with a measure of physical function such as the 4 m gait speed test (13). HGS and the gait speed tests are easy to perform even in a routine clinical setting. Muscle quantity and strength, though interrelated (22), are not equivalent. Muscle strength appears to be a more powerful predictor for the outcome as compared to MM (65). Furthermore, following renal transplantation, we observed a profound increase in HGS without a significant increase in LTM (66).

Which bedside test should be used for the assessment of MM/FFM/LBM depends on local preferences and availability. In the case of 3C-BIS, information on body composition as well as the fluid overload is combined in a single measurement. Muscle ultrasound may be superior to BIS in the diagnosis of skeletal muscle depletion, but reference values need to be defined in larger populations, whereas a trained investigator is necessary. A possibility is to use SCI or BIS for case finding, followed by the US for a more precise estimate of MM depletion.

A diagnosis of PEW or sarcopenia should be combined with an estimate of potentially modifiable factors such as dietary intake and physical activity, for example, by performing actimetry regularly. Also, impaired physical activity, nutritional status, and physical performance should be interpreted given its relation with a low-health-related QoL (7, 67), which is especially important as these factors are often amenable to therapeutic intervention. Lastly, complications that are frequently associated with loss of MM, such as inflammation and fluid overload, should be assessed. A proposal for an integrated assessment, preferably summarized in an easily interpretable dashboard, is illustrated in **Figure 3**. Such a dashboard could be the basis for a personalized

approach. For instance, a patient with a low LTI or MM and reduced muscle strength with adequate nutritional intake and absence of inflammation (a “sarcopenic” phenotype), but with low physical activity may primarily benefit from both aerobic as well as resistance training (68). In addition, activity trackers or smartphone applications may provide the patient with feedback on his/her physical activity patterns.

On the other side of the spectrum, a patient with a “cachectic” phenotype, with inflammation, and reduced protein intake, will primarily benefit from a search into the cause of inflammation and targeted nutritional intervention. Participation in an active rehabilitation program will be much more difficult for this patient, although interventions such as neuromuscular electrical stimulation may be beneficial (69). In addition, the risk of fluid overload may be increased in this patient, which may only be resolved by an increase in dialysis time due to the altered distribution between the interstitial and intravascular compartment (70).

In conclusion, a reduction in MM is an important determinant of various clinical syndromes in patients on dialysis, which are related to increased morbidity and mortality but also potentially amenable to therapeutic intervention. Whereas, different bedside methods can be used to assess MM or its proxies in patients on dialysis, it is important to maintain a critical view of their relative advantages and potential pitfalls. Assessment of MM should be part of a multidimensional approach and a personalized treatment strategy.

AUTHOR CONTRIBUTIONS

AS and JK conceived the paper. NB, FS, MH, and EF contributed to and reviewed the paper. All authors contributed to the article and approved the submitted version.

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Malnutrition and Sarcopenia Combined Increases the Risk for Mortality in Older Adults on Hemodialysis

Catarina Macedo¹, Teresa F. Amaral¹, Juliana Rodrigues², Fernanda Santin² and Carla Maria Avesani^{2,3*}

¹ Faculty of Nutrition and Food Science, University of Porto, Porto, Portugal, ² Graduation Program in Food, Nutrition and Health, Nutrition Institute, Rio de Janeiro State University, Rio de Janeiro, Brazil, ³ Department of Clinical Science, Intervention and Technology, Karolinska Institutet, Stockholm, Sweden

OPEN ACCESS

Edited by:

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*Correspondence:

Carla Maria Avesani
carla.avesani@ki.se

Specialty section:

This article was submitted to
Clinical Nutrition,
a section of the journal
Frontiers in Nutrition

Received: 07 June 2021

Accepted: 20 August 2021

Published: 17 September 2021

Citation:

Macedo C, Amaral TF, Rodrigues J,
Santin F and Avesani CM (2021)
Malnutrition and Sarcopenia
Combined Increases the Risk for
Mortality in Older Adults on
Hemodialysis. *Front. Nutr.* 8:721941.
doi: 10.3389/fnut.2021.721941

Aim: Sarcopenia and malnutrition are highly prevalent in older adults undergoing hemodialysis (HD) and are associated with negative outcomes. This study aimed to evaluate the role of sarcopenia and malnutrition combined on the nutritional markers, quality of life, and survival in a cohort of older adults on chronic HD.

Methods: This was an observational, longitudinal, and multicenter study including 170 patients on HD aged >60 years. Nutritional status was assessed by 7-point-subjective global assessment (7p-SGA), body composition (anthropometry and bioelectrical impedance), and appendicular skeletal muscle mass (Baumgartner's prediction equation). Quality of life was assessed by KDQoL-SF. The cutoffs for low muscle mass and low muscle strength established by the 2019 European Working group on sarcopenia for Older People (EWGSOP) were used for the diagnosis of sarcopenia. Individuals with a 7p-SGA score ≤ 5 were considered malnourished, individuals with low strength or low muscle mass were pre-sarcopenic, and those with low muscle mass and low muscle strength combined as sarcopenic. The sample was divided into four groups: sarcopenia and malnutrition; sarcopenia and no-malnutrition; no-sarcopenia with malnutrition; and no-sarcopenia and no-malnutrition. Follow-up for survival lasted 23.5 (12.2; 34.4) months.

Results: Pre-sarcopenia, sarcopenia, and malnutrition were present in 35.3, 14.1, and 58.8% of the patients, respectively. The frequency of malnutrition in the group of patients with sarcopenia was not significantly higher than in the patients without sarcopenia (66.7 vs. 51.2%; $p = 0.12$). When comparing groups according to the occurrence of sarcopenia and malnutrition, the sarcopenia and malnutrition group were older and presented significantly lower BMI, calf circumference, body fat, phase angle, body cell mass, and mid-arm muscle circumference. In the survival analysis, the group with sarcopenia and malnutrition showed a higher hazard ratio 2.99 (95% CI: 1.23: 7.25) for mortality when compared to a group with no-sarcopenia and no-malnutrition.

Conclusion: Older adults on HD with sarcopenia and malnutrition combined showed worse nutritional parameters, quality of life, and higher mortality risk. In addition, malnutrition can be present even in patients without sarcopenia. These findings highlight the importance of complete nutritional assessment in patients on dialysis.

Keywords: chronic kidney disease, hemodialysis, malnutrition, older adults, mortality

INTRODUCTION

Chronic kidney disease (CKD) has been recognized as one of the main and most prevalent public health problems worldwide. In fact, in 2017 it was estimated that 1.2 million of people died from CKD with an increase of 41% in the global mortality rate between 1990 and 2017 (1). The global prevalence of CKD is of 9.1% (1), which is similar to the CKD prevalence reported of 8.9% in Brazil in 2015 (2). The main factors justifying this increase in CKD are related to the population aging and to the increase in the prevalence of hypertension, diabetes mellitus, and obesity, which are known as the main risk factors for the development of CKD (1).

The stages of CKD evolve as the glomerular filtration rate decreases and the clinical condition worsens, requiring renal replacement therapy in the later stages of the disease, with hemodialysis (HD) being one of the therapeutic options. If, on the one hand, HD enables patients with CKD to live, on the other hand, it can contribute to the development of nutritional disturbances malnutrition, sarcopenia, and frailty. Among these patients on HD, malnutrition and sarcopenia stand out due to their high prevalence in dialyzed patients (3).

According to the sarcopenia consensus from the European Working Group on Sarcopenia for Older People (EWGSOP2), sarcopenia is defined as a progressive and generalized skeletal muscle disorder characterized by the occurrence of low muscle strength and low quality or quantity of muscle mass (4). The coexistence of both conditions constitutes the confirmatory criteria for the diagnosis of sarcopenia. Particularly in CKD, sarcopenia has multicausal etiology, with factors that overlap with traditional factors of sarcopenia in the elderly (5). According to a group of experts from the International Society of Renal Nutrition and Metabolism (ISRNM), malnutrition is characterized by multiple changes caused by a set of factors that lead to an increase in protein catabolism, thus leading to a negative protein balance (6). Sarcopenia and malnutrition are highly prevalent in patients on HD, with the former varying from 51 to 68% and the latter varying from 3.9 to 63.3%, depending on the detection method, disease stage, type of treatment, age, and cutoff points used (7). Both conditions are associated with adverse outcomes, including not only decreased quality of life and functionality, but also increased susceptibility to infection, high hospitalization rates, healthcare costs, morbidity, and mortality (8, 9).

Moreover, data from the United States shows that 40% of end-stage patients were over 65 years in 2013, and projections for 2030 indicate that this proportion will increase to 55–61% (10). Given the fast increase in the prevalence of older adults

on dialysis in the recent decades, and the effect of senescence on decreasing skeletal muscle mass, the assessment of the outcomes of malnutrition and sarcopenia in patients undergoing chronic HD is of major relevance. Importantly, research has shown that early intervention in these patients increases the quality of life and reduces mortality (11). However, the diagnosis and monitoring of malnutrition and sarcopenia in dialyzed patients is not yet carried routinely out in many dialysis clinics, hindering the early intervention for these two conditions. Although it is well-known that malnutrition and sarcopenia are related to higher mortality risk, we now investigate the effect of these two conditions combined on markers of nutritional status, clinical condition, quality of life, and on mortality.

METHODS

Study Protocol

This is an observational, longitudinal, and multicenter study including 170 patients under HD treated in six dialysis units in Brazil. A detailed description of the methodology can be found elsewhere (8). All participants were included from March 2010 to February 2014 and were followed for mortality events up to 36 months. Patients who changed dialysis modality or were transferred to other dialysis units or had kidney transplantation were censored.

Patients

Patients were eligible for inclusion if aged over 60 years, undergoing HD for at least 3 months, three times per week, with each session lasting 3.5–4 h. The exclusion criteria comprised patients using a wheelchair, with amputated limbs and with HIV, cancer, and Alzheimer's and Parkinson's diseases. The study was approved by the Ethics and Research Committee of Rio de Janeiro State University, Brazil registered with protocol number 039.3.2011, and a written informed consent was obtained from all patients before their admission in the study.

Methods

At baseline, all participants had the nutritional status assessed by the 7-point-subjective global assessment (7p-SGA) translated to Portuguese (12), by anthropometric measurements [body weight, height, midarm circumference, triceps skinfold (SKF) thickness, and hip and calf circumference], bioelectrical impedance (BIA), and handgrip strength (HGS) after 30–60 min the dialysis session in a midweek dialysis day to minimize the influence of fluids overload on body composition (13). SKFs were measured using an SKF caliper (Lange, Cambridge Scientific Industries, Cambridge, MD, USA), following the Lohman's Protocol (14).

Body fat was estimated by BIA (Biodynamics® 450; Biodynamics Corporation, Seattle, WA, USA) with the patient in a supine position after 5 min of rest. The measurements of body weight, height, resistance, and reactance were entered in the software Fluid & Nutrition version 3.0 (RJL body composition analyzer) to obtain body fat and phase angle. The study from Heo et al. (15) that evaluated body composition by BIA in non-CKD individuals was used to classify obesity, by using body fat percentage above 32.3% for men and above 44.1% for women. The arm contrary to the arteriovenous fistula was used for the assessment of arm circumference, triceps SKF, and HGS. Muscle strength was measured by a mechanical handgrip dynamometer (Baseline, Fabrication Enterprises, Inc, Elmsford, NY, USA). The highest value of three measurements was taken, with arms along the body after a voice command asking to use the maximal force in the dynamometer. The midarm muscle circumference (MAMC) was calculated using the Frisancho equation (16). The standard values of MAMC and triceps SKF were calculated by the equation: (measured value/value on P50 from NHANES III) $\times 100$ (9).

The 7p-SGA was applied by experienced renal dietitians. The nutritional status of a patient was classified as well-nourished when the 7p-SGA score was equal to 7 and 6 and, as malnourished when the 7p-SGA score ≤ 5 (12).

The Baumgartner's prediction equation (17) was used to estimate appendicular skeletal muscle mass (ASM):

$$\text{ASM (kg)} = 0.2487 (\text{weight, kg}) + 0.0483 (\text{height, cm}) - 0.1584 (\text{hip circumference, cm}) + 0.0732 [\text{HGS, kgf (kilogram force)}] + 2.5843 (\text{male}) + 5.8828 \quad (17)$$

A previous study, conducted by our research group and which included patients on HD, showed that this equation had good agreement with the ASM assessed by dual-energy X-ray absorptiometry (DXA) with an intraclass coefficient correlation (ICC) of 0.92 (95% CI: 0.86–0.95) (9). The ASM was divided by the square height (m) for calculation of the ASM index (ASMI).

The laboratorial measurements were performed before the dialysis session and included assessment of serum albumin (bromocresol green method), high-sensitive C-reactive protein (hs-CRP; by nephelometry), and 25 hydroxyvitamin D [(25 (OH) D); by chemiluminescence immunoassay]. Serum urea was assessed before and after dialysis session for calculation of the urea Kt/V according to the formula of Daugirdas (18) from a midweek dialysis session.

Quality of life was assessed using the Short Form 1.3 questionnaire (KDQoL-SF) (19), which was applied during the dialysis session in 154 patients from the total sample (170 patients). The reason for a smaller sample having data on quality of life is that this assessment did not start at the beginning of the data collection.

For sarcopenia diagnosis, the cutoffs for low muscle mass and low muscle strength established by the 2019 EWGSOP (4) were used. Low muscle strength was considered when HGS was <27 kilogram force (kgf) for men and <16 kgf for women and low

muscle mass was considered when the ASMI was $<7.0 \text{ kg/m}^2$ for male and $<5.5 \text{ kg/m}^2$ for female (4). Patients were considered with pre-sarcopenia when presenting only one of the muscle abnormalities, that is, low muscle mass or low muscle strength and with sarcopenia when both conditions were present.

The patients were classified into four groups considering the presence of malnutrition, pre-sarcopenia, and sarcopenia:

- Group sarcopenia and malnutrition ($n = 56$): Comprised by patients with positive criteria for sarcopenia/pre-sarcopenia and for malnutrition (7p-SGA score ≤ 5).
- Group sarcopenia and no-malnutrition ($n = 28$): Comprised by patients with positive criteria for sarcopenia/pre-sarcopenia, but without criteria for malnutrition (7p-SGA score = 6 and 7).
- Group no-sarcopenia with malnutrition ($n = 44$): Comprised by patients without criteria for sarcopenia/pre-sarcopenia, but with positive criteria for malnutrition (7p-SGA score ≤ 5).
- Group no-sarcopenia and no-malnutrition ($n = 42$): Comprised by patients without criteria for sarcopenia/pre-sarcopenia and for malnutrition (7p-SGA score = 6 and 7).

Statistical Analysis

The Shapiro–Wilk test was applied to test normality. Categorical variables are described as absolute number and percentage and continuous variables as mean and SD or as median and interquartile range, as appropriate. The comparisons of the variables among the groups of sarcopenia and malnutrition were performed using the chi-square test for categorical variables, and one-way ANOVA or Kruskal–Wallis tests for continuous variables, as appropriate. The Bonferroni test was used to verify the differences among the groups for the variables presenting normal distribution.

The comparisons between the survival and deceased groups were done by chi-square test, independent *t*-test, or Mann–Whitney test, as appropriate. The survival analyses were performed by the Kaplan–Meier graphic using the log-rank test to compare the survival curves among the sarcopenia and malnutrition groups. The Cox's proportional risk model adjusted for gender, age, and hs-CRP was used to assess the hazard ratio for mortality, using the no-sarcopenia and well-nourished group as reference. The value of $p < 0.05$ will be used for statistical significance. All analyses will be performed using the SPSS software version 27 (IBM Corp. Released 2015, Armonk, NY, USA).

RESULTS

Table 1 shows the main characteristic of the studied sample comprised of older adults on chronic HD. In general, the mean age was around 70.6 years, the majority of the sample was comprised of males, and the urea Kt/V was indicative of adequate dialysis. The mean BMI, calf circumference, standard triceps SKF, and MAMC indicated adequate nutritional status according to cutoffs established for non-CKD individuals, which are well-accepted for use in patients with CKD (20). However, when the

TABLE 1 | Main demographic, nutritional, and clinical characteristics of older adults on hemodialysis.

	Results (n = 170)
Age (years)	70.6 ± 7.2
Male (n; %)	111 (65.3)
Dialysis length (years)	2.9 (1.3; 5.6)
Urea Kt/V	1.5 (1.3; 1.6)
Diabetes (n; %)	44 (37.7)
BMI (kg/m ²)	25.4 ± 4.5
Pre-sarcopenia (n; %)	60 (35.3)
Sarcopenia (n; %)	24 (14.1)
No sarcopenia (n; %)	86 (50.6)
Standard triceps skinfold thickness (%)	102.5 (72.7; 142.1)
Standard Midarm muscle circumference (%)	98.1 ± 14.7
Calf circumference (cm)	
Male	34.5 ± 3.9
Female	33.3 ± 3.5
Malnutrition (n; %)	100 (58.8)
Body fat (%)	
Male	27.5 ± 7.0
Female	37.7 ± 5.3
Phase angle (°)	
Male	5.5 ± 1.3
Female	5.3 ± 1.2
Body cellular mass (kg)	
Male	22.1 ± 4.8
Female	17.0 ± 3.6
Appendicular skeletal muscle mass index (kg/m ²)	
Male	7.48 ± 0.77
Female	4.60 ± 0.81
Pre-sarcopenia (n; %)	
Low HGS	38 (22.4)
Low ASMI	22 (12.9)
Total	60 (35.3)
Albumin (g/dl)	3.9 ± 0.4
Hemoglobin (mg/dl)	11.3 ± 1.6
Hematocrit (%)	34.4 ± 5.0
S Creatinine (mg/dl)	8.7 ± 2.8
S Urea (mg/dl)	138 ± 39.5
PTH (mg/dl)	223 (101; 402)
25(OH)D (ng/ml)	19.2 (12.7; 27.1)
hs-CRP (mg/dl)	0.42 (0.2; 1.1)

BMI, body mass index; hs-CRP, high-sensitive C-reactive protein; 25(OH)D, 25-hydroxyvitamin D; PTH, parathormone.

Data described as absolute values and its percentage for categorical variables; as mean ± SD as or as median and interquartile range for continuous variables according to the variable's distribution. Malnutrition defined as 7p-SGA ≤ 5.

Pre-sarcopenia defined as either low muscle mass (appendicular skeletal muscle mass index below 7 kg/m² for male and below 5.5 kg/m² for female) or low muscle strength (handgrip strength below 27 kgf for male and below 16 kgf for female). Sarcopenia was defined by the concomitant condition of low muscle mass and low muscle strength.

nutritional status was assessed by 7p-SGA, 58.8% of the sample had a score ≤ 5, indicating malnutrition. As for body composition assessed by BIA, the mean ± SD values for body fat percentage

showed that 26.7% of male and 7% of female were obese when applying the cutoffs suggested by Heo et al. (15). When assessing the presence of sarcopenia, about one-third of the sample had either low muscle mass or low muscle strength, here defined as pre-sarcopenia, whereas 14.1% had both conditions combined, defined as sarcopenia, and 50.6% had no signs of low muscle mass or low muscle strength. The laboratory exams were compatible to that observed for patients on dialysis treatment, and the mean serum albumin was within the acceptable values to patients with CKD (>3.8 mg/dl) (21).

Considering that the presence of two nutritional disturbances—malnutrition and sarcopenia were investigated, we evaluated whether the frequency of malnutrition differed among the sarcopenia groups. **Figure 1** shows the frequency of patients with malnutrition (assessed as 7p-SGA ≤ 5) in the groups stratified as sarcopenia, pre-sarcopenia, and no-sarcopenia. As can be observed, the prevalence of patients with malnutrition did not differ among the sarcopenia groups, indicating that malnutrition was present even in the group of no-sarcopenia. We then expanded our analysis by exploring the role that these two conditions combined (sarcopenia and malnutrition) have on other nutritional markers, clinical condition, quality of life, and mortality events.

Table 2 shows the comparison of demographic, nutritional, clinical characteristics and quality of life among the groups classified by the presence of sarcopenia and malnutrition. Age and the percentage of males differed significantly among the groups, with the age higher in the group sarcopenia and malnutrition while male gender had higher prevalence in the group sarcopenia and no-malnutrition. Except for serum albumin that did not differ among the groups, the other nutritional markers differed significantly, indicating worse nutritional status in the group sarcopenia and malnutrition as compared to the group no-sarcopenia and no-malnutrition. Regarding clinical condition, the urea Kt/V and 25(OH)D differed among the groups, being the group sarcopenia and malnutrition presenting higher Kt/V and lower 25(OH)D as compared to the group no-sarcopenia and no-malnutrition. Regarding the domains related to the quality of life, most of them did not differ among the groups. Those that showed significant differences were quality of social interaction, role physical, social function, and SF12 mental composite with the group no-sarcopenia and no-malnutrition showing better scores when compared to the remaining groups. After 23.5 (12.2; 34.4) months of follow-up (median and interquartile ranges), there were 62 events of death. The group of deceased patients was older, with higher Kt/V and hs-CRP as compared to the patients who survived (**Table 3**).

The survival analysis showed that there was a significant difference in the survival curves among the groups, being the group combining both conditions (sarcopenia and malnutrition) the one with lower survival rate (**Figure 2**, Kaplan–Meier; log-rank test, $p = 0.019$). This finding was confirmed in the Cox regression analysis adjusted for age, gender, and hs-CRP, where the group with sarcopenia and malnutrition had a hazard ratio of 2.99 (95% CI: 1.23; 7.25) as compared to the reference group no-sarcopenia and no-malnutrition (**Table 4**).

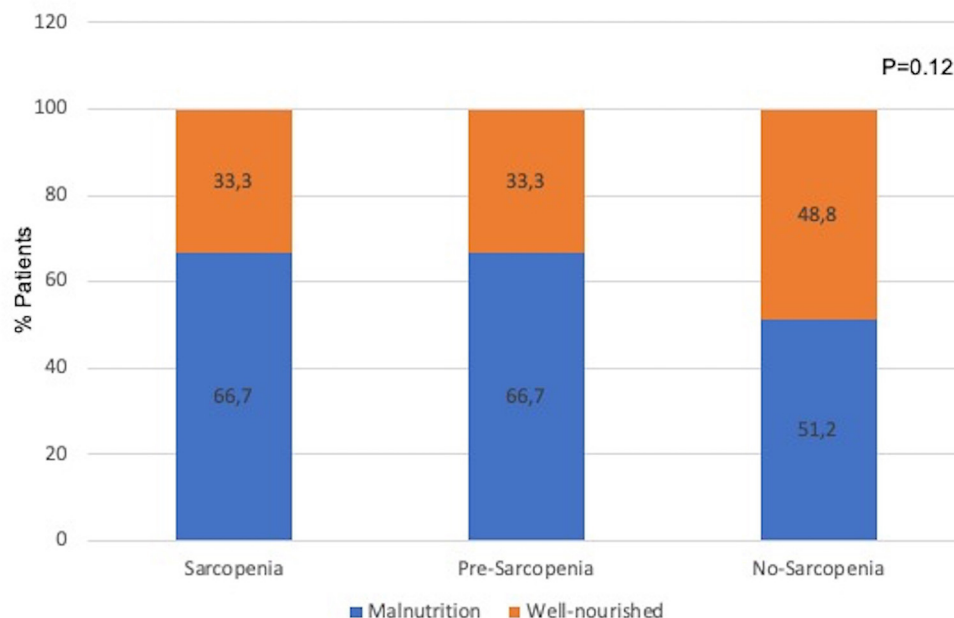


FIGURE 1 | Prevalence of malnutrition (assessed by 7p-SGA) in groups classified as sarcopenia, pre-sarcopenia, and no-sarcopenia. Differences among groups were tested by chi-square test.

DISCUSSION

In this study, we aimed to evaluate the role of sarcopenia and malnutrition on the nutritional markers and survival in a cohort of older adults on chronic HD. Malnutrition (diagnosed by 7p-SGA) was present in 58.8%, which was similar to that found by Cianciaruso et al. (22) where 51% of older adults (>65 years) in HD and peritoneal dialysis had malnutrition assessed by SGA. Moreover, in a meta-analysis aiming to describe the prevalence of malnutrition in patients with CKD (assessed by SGA or malnutrition inflammation score), it was shown that the 25–75th percentile ranges of malnutrition among studies on dialyzed patients was 28–54% (3). Therefore, our findings on the presence of malnutrition are somehow higher than that from previous studies, most likely due to the inclusion of only older adults on HD.

Markers of muscle abnormality, such as low muscle strength or low muscle mass, named as pre-sarcopenia in the current study, were present in 35.3% of the patients, a percentage similar to that found by Isoyama et al. in incident dialyzed patients (39% of the patients with either low muscle strength or low muscle mass) (23). Moreover, we found that sarcopenia (diagnosed by the concomitance of low muscle mass and low muscle strength) was present in 14.1% of the patients, a frequency lower to that from previous studies in dialysis patients (20–40%) (23–25). This discrepancy is most likely due to the diagnostic methods used and the cutoffs applied for the diagnose of low muscle mass and low muscle strength in patients on HD, as previously shown by Lamarca et al. (7). In addition, these divergent

results, when comparing with findings from other studies, can also be explained by different characteristics from the studied sample, such as the CKD stage, dialysis modality, presence of comorbidities, and the mean age of the sample (5).

Furthermore, we identified that among the groups stratified by sarcopenia status, 66.7% of the patients from the group sarcopenia had also malnutrition (**Figure 2**). In another study including older adults (>65 years) with CKD stages 3b to 5 not on dialysis, 52% of the patients with sarcopenia were diagnosed with protein energy wasting (PEW) using the diagnostic criteria from the ISRNM (26). In incident dialysis patients, 64.7% of the patients with sarcopenia had malnutrition diagnosed by SGA (23). The similar frequency of sarcopenia and malnutrition combined found in our study and in the aforementioned ones underlines that these nutritional disturbance can coexist, and a careful assessment for both conditions should be performed in patients on HD.

Surprisingly, when comparing the frequency of malnutrition among the sarcopenia groups (**Figure 1**), 51.2% of the patients in the group no-sarcopenia had malnutrition, a proportion not different from that observed in the groups sarcopenia and pre-sarcopenia. Similarly, in two previous studies including either patients with CKD stages 3b–5 or before the start of dialysis therapy, 14–20% of the patients with no-sarcopenia had malnutrition (23, 26). In other words, the absence of sarcopenia does not exclude the existence of malnutrition. This finding highlights that although sarcopenia and malnutrition share some common criteria, these are different nutritional abnormalities, and the investigation of both is crucial. In the current study, malnutrition was diagnosed by 7p-SGA, which

TABLE 2 | Comparisons of demographic, nutritional, clinical characteristics, and quality of life of older adults on hemodialysis according to the groups sarcopenia and malnutrition.

	Sarcopenia and malnutrition (n = 56; 33%)	Sarcopenia and no-malnutrition (n = 28; 16.5%)	No-sarcopenia and malnutrition (n = 44; 25.8%)	No-sarcopenia and no-malnutrition (n = 42; 24.7%)	p*
Age (years)	73.2 ± 8.0 ^a	72.4 ± 7.9 ^{a,c}	69.3 ± 6.0 ^{b,c}	67.4 ± 5.4 ^{b,d}	<0.001
Male (n; %)	34 (60.7)	24 (85.7)	23 (52.3)	30 (71.4)	0.02
Dialysis length (years)	3.01 (1.66; 5.60)	2.96 (1.37; 6.32)	2.08 (0.93; 5.87)	3.15 (1.24; 5.60)	0.67
Urea Kt/V	1.51 (1.40; 1.70)	1.34 (1.22; 1.71)	1.47 (1.30; 1.63)	1.41 (1.28; 1.53)	0.04
Diabetes (n; %)	14 (25)	12 (42.8)	24 (54.5)	14 (33.3)	0.06
BMI (kg/m ²)	23.6 ± 4.7 ^a	25.0 ± 3.3 ^{a,c}	25.9 ± 4.6 ^{b,c}	27.7 ± 3.9 ^{b,d}	<0.001
Standard triceps skinfold thickness (%)	89.5 (63.6; 125.7)	110.1 (91.4; 175.7)	96.15 (72.1; 121.2)	134.4 (96.5; 169.1)	<0.001
Standard midarm muscle circumference (%)	93.6 ± 13.4 ^a	94.0 ± 13.2 ^{a,c}	100.9 ± 16.7 ^{a,b,c}	103.9 ± 12.7 ^b	0.001
Calf circumference (cm)	32.6 ± 3.4 ^a	33.7 ± 2.3 ^a	33.6 ± 4.6 ^a	36.6 ± 2.9 ^b	<0.001
Body fat %	28.5 ± 7.8 ^a	30.0 ± 8.5 ^{a,b}	32.3 ± 8.2 ^{a,b}	33.9 ± 7.3 ^b	0.007
Phase angle (°)	4.9 ± 1.2 ^a	5.2 ± 0.9 ^{a,c}	5.4 ± 0.9 ^{a,b,c}	6.2 ± 1.4 ^b	<0.001
Body cellular mass (kg)	17.7 ± 4.3 ^a	20.0 ± 3.8 ^b	19.2 ± 3.8 ^{a,b}	24.2 ± 5.0 ^c	<0.001
Appendicular skeletal muscle mass index (kg/m ²)	6.42 ± 1.1 ^a	7.03 ± 0.83 ^{a,b,c}	6.99 ± 0.96 ^b	7.56 ± 0.94 ^c	<0.001
Albumin (g/dl)	3.83 ± 0.41	3.98 ± 0.40	3.92 ± 0.41	3.91 ± 0.41	0.44
Hemoglobin (mg/dl)	11.3 ± 1.9	11.2 ± 1.7	11.2 ± 1.3	11.4 ± 1.5	0.96
Hematocrit (%)	34.6 ± 6.0	34.3 ± 5.2	34.0 ± 4.1	34.4 ± 4.5	0.94
S Creatinine (mg/dl)	8.5 ± 2.4	8.9 ± 2.9	8.2 ± 2.8	9.4 ± 3.1	0.24
S Urea (mg/dl)	130.6 ± 41.7	150.8 ± 40.9	137.2 ± 37.2	140.4 ± 36.7	0.16
PTH (mg/dl)	165.3 (59.6; 331.6)	218.3 (111.6; 454.1)	262.3 (106.6; 442.8)	256 (171.2; 402.6)	0.23
25(OH)D (ng/ml)	17.5 (11.7; 30.1)	14.7 (10.5; 21.4)	18.0 (13.5; 25.4)	25.7 (19.0; 34.0)	0.001
hs-CRP (mg/dl)	0.37 (0.20; 1.23)	0.26 (0.09; 0.69)	0.52 (0.26; 1.18)	0.52 (0.22; 1.15)	0.19
Quality-of-life domains					
Symptom problem list	75.1 ± 16.8	74.5 ± 21.6	67.7 ± 24.0	75.6 ± 18.4	0.24
Effects of kidney disease	68.7 (50.0; 84.4)	71.9 (40.6; 90.6)	59.4 (42.2; 78.1)	75.0 (66.7; 91.7)	0.38
Burden of kidney disease	37.5 (18.8; 56.3)	50.0 (18.8; 75.0)	25.0 (12.5; 50.0)	50.0 (25.0; 68.8)	0.10
Work status	50.0 (0.0; 50.0)	50.0 (0.0; 50.0)	50.0 (0.0; 50.0)	50.0 (0.0; 100.0)	0.83
Cognitive function	80.4 ± 21.0	78.3 ± 26.4	75.9 ± 25.8	83.5 ± 19.4	0.50
Quality of social interaction	81.5 ± 17.9 ^a	77.1 ± 20.4 ^a	70.7 ± 24.0 ^{a,b}	82.8 ± 19.2 ^{a,c}	0.03
Sexual function	95.8 ± 10.2	85.9 ± 14.5	77.3 ± 26.8	84.2 ± 19.7	0.50
Sleep	65.0 (47.5; 83.8)	57.5 (35.0; 75.0)	57.5 (46.3; 70.0)	72.5 (52.5; 81.3)	0.08
Social support	100.0 (66.7; 100.0)	83.3 (66.7; 100.0)	100.0 (66.7; 100.0)	100.0 (66.7; 100.0)	0.78
Dialysis staff encouragement	75.0 (56.3; 93.8)	87.5 (75.0; 100.0)	100.0 (75.0; 100.0)	87.5 (75.0; 100.0)	0.07
Overall health	60.0 (50.0; 100.0)	50.0 (50.0; 100.0)	60.0 (50.0; 85.0)	60.0 (50.0; 80.0)	0.61
Patient satisfaction	68.4 ± 19.3	75.4 ± 21.2	72.4 ± 21.6	71.9 ± 20.2	0.56
Physical functioning	45.0 (22.5; 80.0)	50.0 (25.0; 70.0)	40.0 (25.0; 70.0)	55.0 (40.0; 75.0)	0.21
Role physical	50.0 (0.0; 75.0)	50.0 (0.0; 100.0)	0.0 (0.0; 62.5)	50.0 (25.0; 100.0)	0.009
Pain	62.5 (45.0; 90.0)	70.0 (45.0; 90.0)	55.0 (22.5; 95.0)	67.5 (45.0; 100.0)	0.43
General health	60.0 (30.0; 70.0)	50.0 (35.0; 75.0)	50.0 (32.5; 62.5)	65.0 (40.0; 82.5)	0.15
Emotional well-being	76.0 (52.0; 92.0)	84.0 (60.0; 96.0)	68.0 (42.0; 90.0)	80.0 (64.0; 92.0)	0.18
Role emotional	33.3 (0.0; 100.0)	66.7 (0.0; 100.0)	33.3 (0.0; 66.7)	66.7 (33.3; 100.0)	0.09
Social function	62.5 (37.5; 100.0)	87.5 (62.5; 100.0)	62.5 (25.0; 87.5)	75.0 (62.5; 100.0)	0.04
Energy fatigue	55.0 (32.5; 75.0)	60.0 (40.0; 80.0)	45.0 (32.5; 75.0)	65.0 (42.5; 80.0)	0.27
SF12 Physical composite	37.9 (31.1; 45.8)	36.4 (32.3; 45.3)	36.5 (26.2; 46.5)	40.6 (35.1; 47.2)	0.22
SF12 Mental Composite	44.9 ± 11.1 ^a	50.3 ± 13.5 ^{a,b}	44.3 ± 12.3 ^a	50.5 ± 10.1 ^b	0.02

NA, non-applicable; hs-CRP, high-sensitive C-reactive protein; 25(OH)D, 25(OH)D, 25-hydroxyvitamin D; PTH, parathormone. Quality of life was evaluated in a subgroup of 154 patients (n = 49, n = 23, n = 41, and n = 41, respectively in the four groups). Data described as absolute values and its percentage for categorical variables; as mean ± SD as or as median and interquartile range for continuous variables according to the variable's distribution.

*Chi-square or one-way ANOVA or Kruskal–Wallis test, as appropriate. The Bonferroni post-hoc test for ANOVA $p \leq 0.05$: Significant differences among the groups are signed by the different superscript letters.

TABLE 3 | Comparison of older adult patients on hemodialysis according to the group alive and deceased ($n = 170$).

	Alive ($n = 108$)	Deceased ($n = 62$)	p^*
Male	71 (65.7)	40 (64.5)	0.87
Age (years)	69.6 ± 6.7	72.5 ± 7.8	0.013
Kt/V	1.42 ± 0.3	1.60 ± 0.5	0.002
hs-CRP	0.34 (0.18; 0.82)	0.58 (0.27; 1.48)	0.004
Dialysis length (years)	2.9 (1.2; 5.4)	2.9 (1.3; 6.0)	0.68

hs-CRP, high-sensitive C-reactive protein.

*The *t*-test; chi-square test, or Mann–Whitney test, as appropriate.

evaluate several domains of nutritional status (involuntary loss of body weight, food intake, gastrointestinal symptoms, poor appetite, functional status, comorbidities, and physical exam for subcutaneous fat and muscle loss) (27). Therefore, it provides a broad assessment of nutritional status including aspects not included in the criteria for sarcopenia diagnosis. This likely explains the reason why individuals in the group no-sarcopenia had malnutrition when diagnosed by 7p-SGA. Adding to these findings, we also demonstrated, as expected, that when sarcopenia and malnutrition occurred concomitantly (group sarcopenia and malnutrition), all parameters of nutritional status, except for albumin, were worse when compared with the group no-sarcopenia and no-malnutrition. The non-difference in serum albumin among the groups of sarcopenia and malnutrition corroborates the findings from Gama-Axelsson et al. (28) The authors reported that in prevalent dialyzed patients, serum albumin correlated poorly with markers of nutritional status, including SGA score and body composition parameters, but it was significantly correlated with hs-CRP (28). Altogether, this is aligned with the statement from the updated guidelines in nutrition and CKD from the NKF-KDOQI that albumin is a predictor of hospitalization and mortality, and not a marker of nutritional status (29).

It was interesting to note that within the sarcopenia and malnutrition group, the mean values of body fat markers, such as BMI, percentage of body fat, and standard triceps SKFs were within the normal range for non-CKD individuals (15, 30, 31). Similarly, Lee et al. (32) also observed in a group of older adults on HD that patients with low gait speed and low HGS combined had BMI within the normal range. Additionally, in the study from Ren et al. (24), no significant differences were found between no-sarcopenia and sarcopenia in relation to anthropometric indexes, namely TSE, BMI, MAC, and MAMC. The remaining markers of nutritional status differed among the groups stratified as malnutrition and sarcopenia status, being this difference more marked between the group sarcopenia and malnutrition and group no-sarcopenia and no-malnutrition. Among those, the phase angle, which is not much explored in patients on HD, could discriminate adequately the nutritional status in the four studied groups. In our study, the phase angle differed mainly between the sarcopenic and malnourished group and the group no-sarcopenia and no-malnutrition. This finding is in agreement with studies in non-elderly adults on HD, where the

phase angle also differed significantly between malnourished and non-malnourished groups (24, 33, 34). Therefore, one marker of nutritional status should not be used alone to evaluate nutritional status, but rather a combination of markers can be used, as in fact stated in the guideline for nutrition and CKD from the NKF-KDOQI (29).

Regarding quality of life, the domains most affected were social interaction, role physical, social function, and SF12 mental composite, which had worse scores in the group sarcopenia and malnutrition. We are not aware of studies in patients with CKD evaluating the role of sarcopenia and malnutrition combined on quality-of-life domains, but in a previous study from our group, we showed that patients on HD with low muscle strength had worse quality-of-life domains than that of the group with low muscle mass (9). Moreover, in another study including patients on dialysis, malnutrition was associated with worse quality of life (35–37) and with the presence of depression and sleep disorders (35).

Finally, when evaluating survival, we found that the mortality risk of the groups with sarcopenia and malnutrition was close to three times higher than the group without any of these abnormalities. As far as we are concerned, there are no previous studies assessing the role of malnutrition and sarcopenia combined in older adults on HD, but studies in older adults hospitalized without CKD, showed that older adults with combined sarcopenia and malnutrition had a risk for mortality of close to five times higher when compared to the group with none of these nutritional disturbances (38). In patients on dialysis, previous studies have consistently shown that malnutrition (36, 39), sarcopenia (25), and low muscle strength (23, 32, 40) were associated with increased mortality.

Some limitations and strengths of this study can be listed. As a limitation, the observational study design can impair the identification of a causality–effect association. Second, the relatively small sample size can underpower the comparison among the sarcopenia and malnutrition groups, though statistical differences were already listed with this sample size. Third, the lack of robust methods to estimate muscle mass can hinder muscle abnormalities related to muscle mass. As positive aspects, we consider the originality of evaluating the concomitance of malnutrition and sarcopenia in elderly patients on HD, and the relationship of these conditions with quality of life and survival. In addition, although the methods used to evaluate muscle mass could be influenced by the variation in the hydration status, all measurements were performed after the dialysis session to minimize the influence of fluid retention. In addition, since these are the methods used in the routine care of dialysis clinics and also recommended by the updated guidelines in nutrition and CKD from NKF/KDOQI (29), our findings can be used to support a nutritional assessment with methods that are suitable for the routine use.

In conclusion, patients on HD aged 60 years and older who have sarcopenia and malnutrition showed worse nutritional parameters, quality-of-life domains, and higher mortality risk. In addition, we reported that malnutrition can occur in patients without sarcopenia, and that the body fat markers within the normality range can occur concomitantly with

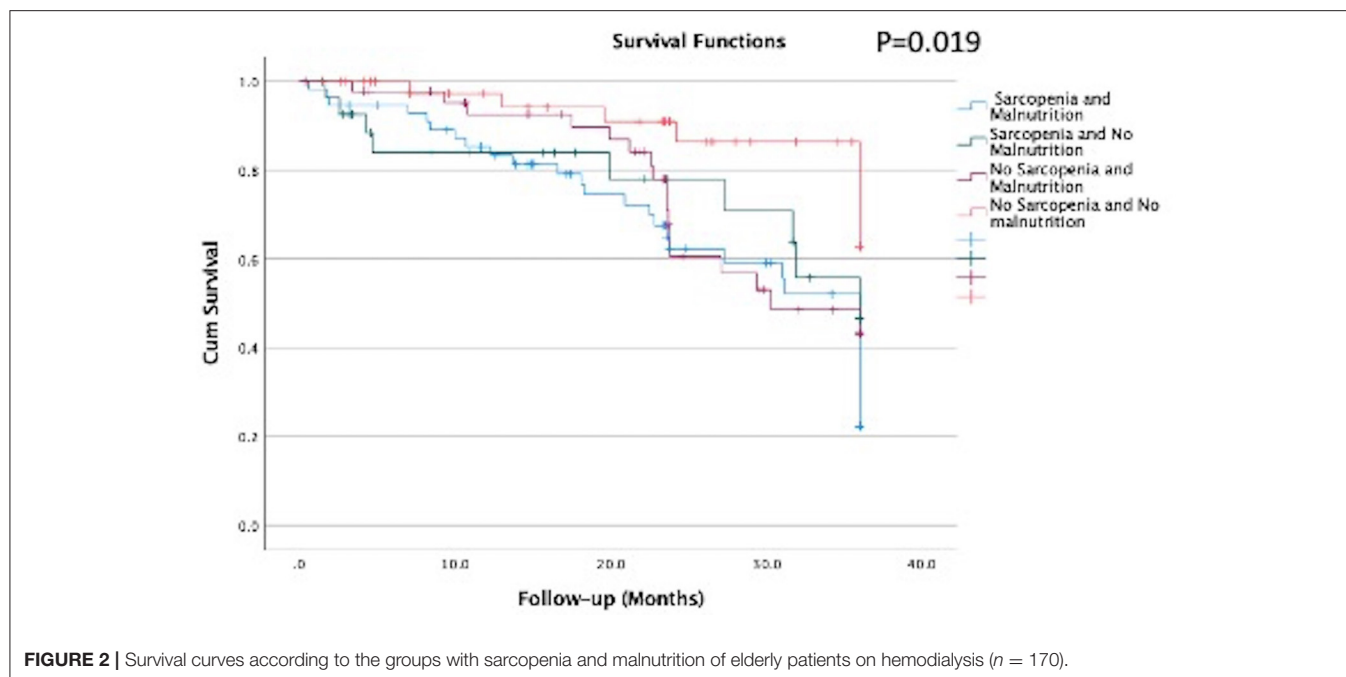


TABLE 4 | Risk for mortality events, expressed as hazard ratio* according to the combination of group with sarcopenia and malnutrition in older adults undergoing maintenance hemodialysis ($n = 170$).

	Hazard ratio*	95% CI		p-value
		Lower	Upper	
Male	0.44	0.24	0.83	0.01
Age (years)	1.02	0.96	1.06	0.27
hs-CRP (mg/dl)	1.35	2.00	1.58	0.03
No-sarcopenia + no-malnutrition (reference group) ($n = 42$)				
Sarcopenia + no-malnutrition ($n = 28$)	2.65	0.86	7.05	0.09
No-Sarcopenia + malnutrition ($n = 44$)	2.43	0.97	6.05	0.06
Sarcopenia + malnutrition ($n = 56$)	2.99	1.23	7.25	0.03

hs-CRP, high-sensitive C-reactive protein.

*Cox's proportional risk model.

malnutrition and sarcopenia. Altogether, these findings highlight the importance of complete nutritional assessment in older patients on dialysis. Further studies to better understand the role of these abnormalities in the health of older adults undergoing maintenance HD are needed.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will not be made available by the authors.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by Ethics and Research Committee of Rio

de Janeiro State University, Brazil. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

CM and TA analyzed and interpreted the data, drafted the article, and gave final approval of the submitted version. JR and FS acquired the data, critically revised the important intellectual content, and gave final approval of the submitted version. CA conceived and designed the study, acquired, analyzed, interpreted the data, drafted the article, critically revised the important intellectual content, and gave final approval of the submitted version. All authors contributed to the article and approved the submitted version.

FUNDING

This research received two grants from Fundação Carlos Chagas Filho de Amparo à Pesquisa do Estado do Rio de Janeiro (FAPERJ); Grant Numbers: E-26/111.653/2010 and E-26/103.209/2011).

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ACKNOWLEDGMENTS

The authors would like to thank Renata Lemos Fetter, Fernanda Guedes Bigogno, Fernando Lamarca, Juliana Giglio, and Ana Lucia Mendes Pereira who helped during the data collection and also thank Sergio Franco Laboratory for the laboratorial analysis.

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