



CKD Stages, Bone Metabolism Markers, and Cortical Porosity Index: Associations and Mediation Effects Analysis

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Xiong Y, He T, Wang Y, Liu WV, Hu S, Zhang Y, Wen D, Hou B, Li Y, Zhang P, Liu J, He F and Li X (2021) CKD Stages, Bone Metabolism Markers, and Cortical Porosity Index: Associations and Mediation Effects Analysis. Front. Endocrinol. 12:775066. doi: 10.3389/fendo.2021.775066 Chronic kidney disease (CKD) has a significant negative impact on bone health. However, the mechanisms of cortical bone deterioration and cortical porosity enlargement caused by CKD have not been fully described. We therefore examined the association of CKD stages with cortical porosity index (PI), and explored potential mediators of this association. Double-echo ultrashort echo-time magnetic resonance imaging (UTE MRI) provides the possibility of quantifying cortical porosity in vivo. A total of 95 patients with CKD stages 2-5 underwent 3D double-echo UTE-Cones MRI (3.0T) of the midshaft tibia to obtain the PI. PI was defined as the ratio of the image signal intensity of a sufficiently long echo time (TE) to the shortest achievable TE. Parathyroid hormone (PTH), β -CrossLaps (β -CTX), total procollagen type I amino-terminal propeptide (T-P1NP), osteocalcin (OC), 25-hydroxyvitamin D (25OHD), and lumbar bone mineral density (BMD) were measured within one week of the MRI. Partial correlation analysis was performed to address associations between PI, eGFR and potential mediators (PTH, β -CTX, T-P1NP, OC, 25OHD, BMD, and T-score). Multiple linear regression models were used to assess the association between CKD stages and PI value. Then, a separate exploratory mediation analysis was carried out to explore the impact of CKD stages and mediators on the PI value. The increasing CKD stages were associated with a higher PI value (P_{trend} < 0.001). The association of CKD stages and PI mediated 34.4% and 30.8% of the total effect by increased PTH and β-CTX, respectively. Our study provides a new idea to monitor bone health in patients with CKD, and reveals the internal mechanism of bone deterioration caused by CKD to some extent.

Keywords: chronic kidney disease, CKD-MBD, Ultrashort TE, porosity index, cortical porosity, mediation analysis, bone metabolism markers

INTRODUCTION

Chronic kidney disease (CKD) has become a recognized global public health problem, which affects approximately 0.75 billion human beings worldwide (1). CKD has a significant negative impact on bone health (2, 3), which results in a two-fourfold higher risk of fracture in CKD patients compared with general population, and the occurrence of fractures is associated with increased mortality (4, 5). Cortical bone is the main factor that determines the overall bone strength, accounting for more than 80% of the total bone (6). Cortical porosity is the main factor that determines the strength of cortical bone, and the development of cortical pores leads to an increase in bone fragility, which increases the fracture (7). A study by Nickolas et al. (8) found that hemodialysis has an independent effect on the increase in cortical porosity, which suggests that cortical porosity may increase as CKD progresses.

CKD-mineral and bone disorder (CKD-MBD), a disease of bone metabolism that manifests as systemic mineral and bone metabolism disorders, is the main complication of CKD and affects most patients with moderate to severe CKD (9). Elevated parathyroid hormone (PTH) is a potential mediator for patients with MBD, and it is also the most widely studied and most recognized bone metabolism marker for predicting bone turnover in CKD patients (10). In terms of the mechanism, with the progression of CKD and calcium and phosphorus metabolism disorders, patients develop secondary hyperparathyroidism to varying degrees, and increased PTH has a catabolic effect on cortical bone (11). Cortical porosity in patients with CKD showed highly positive correlation with PTH (r = 0.62; P = 0.021) (3). In addition, tartrate-resistant acid phosphatase 5b (TRAP5b) and bone-specific alkaline phosphatase (BSAP) are less affected by renal clearance and can more accurately assess MBD (12), but these two indicators are not as widely used in clinical practice as PTH. The application of other bone metabolism markers (e.g., β -CrossLaps (β-CTX), total procollagen type I amino-terminal propeptide (T-P1NP), osteocalcin (OC), and 25-hydroxyvitamin D (25OHD), etc.) in CKD is controversial, and there are few related studies. These bone metabolism markers are relatively common in the study of osteoporosis. In this study, we wanted to explore which metabolic markers influenced the relationship between CKD and cortical porosity, and therefore some of these metabolic markers were included.

Early imaging studies on CKD-MBD mainly focused on trabecular bone (13). However, recent research has shown that cortical bone plays a more crucial part than trabecular bone in the bone health of CKD-MBD patients (14). In patients with CKD, the loss of cortical bone is associated with increased cortical porosity and decreased cortical thickness (8). Nevertheless, cortical bone is not taken as a key parameter in the "turnover, mineralization, volume" classification of CKD-MBD (3).

At present, information on the porosity of human cortical bone *in vivo* is mainly obtained through high-resolution peripheral quantitative computed tomography (HR-pQCT) and bone biopsy (15). However, HR-pQCT involves radiation, and its limited resolution underestimates porosity (16). And bone biopsy is expensive and invasive and requires professional knowledge to perform, process, and interpret; thus, only a few laboratories worldwide carry out this test (15). With the development of magnetic resonance technology, the adoption of ultrashort echo time (UTE) in vivo has proven to be feasible for quantitative assessment of cortical pore information (17). Magnetic resonance quantification of bone tissue has always been a difficult problem in the field of magnetic resonance research. UTE magnetic resonance imaging (MRI) is an advanced and promising method. Its echo time is very short, which solves the difficult problem of the inability of conventional MRI sequences to obtain bone signals. At present, this sequence has been used for the quantitative and qualitative assessment of cortical bone pore water (e.g., bicomponent analysis, tricomponent analysis, single/double adiabatic inversion of UTE technology, etc.) (18), and increased pore water content has been confirmed to be related to increase in cortical porosity and decrease in mechanical properties (19). However, these methods are complicated, the scanning time is very long, and the postprocessing technology requirements are very strict, so they are hardly used in clinical research. Rajapakse et al. (20) evaluated cortical bone porosity using a novel double-echo UTE MR sequence recently. The porosity index (PI) was defined as the ratio of the image signal intensity of a sufficiently long echo time (TE) (mainly from the pore water signal) to the shortest achievable TE (from all water signals), representing the fraction of pore-water. The selection of TE value needs to consider the T2* value of bound water (0.3-0.4 ms) and pore water (> 1 ms) (20, 21). The researchers found that the PI measured by this method exhibits highly positive correlation with cortical porosity measured by micro-CT ($R^2 = 0.79$). Furthermore, this means showed high repeatability when measuring vivo tibial PI (with a variation of 2.2% in betweenday coefficient). Thus, it is reasonable for us to conclude that PI can reflect porosity to a large extent. This method, which is noninvasive and does not involve radiation, has a short scanning time and a simple processing protocol, which results in a high potential for application in clinical practice. Chen et al. (22) applied this method to healthy volunteers. However, there are almost no relevant studies on applying this method to various metabolic bone diseases, including CKD-MBD.

The mediators selected in this study are currently the most recognized mediators in bone metabolism, and their application in CKD-MBD needs to be further studied. Therefore, the aim of this research is to (1) explore the association between the increasing stages of CKD and the development of cortical PI; (2) judge the mediation effect of several bone metabolism markers (i.e., PTH, β -CTX, T-P1NP, OC, and 25OHD) and dual-energy X-ray absorptiometry (DXA) lumbar spine bone mineral density (BMD) on this association.

MATERIALS AND METHODS

Subjects

This cross-sectional cohort study was prospective, which was approved by Medical Ethics Committee of Tongji Hospital of Huazhong University of Science and Technology, TJ-IRB20210108. We obtained written informed consent from every participant before the research. This trial was registered with ClinicalTrials.gov as NCT04564924. The subjects were recruited from September 1, 2020, to April 11, 2021. We recruited one hundred patients from the Department of Nephrology of Tongji Hospital. Three initially collected patients were excluded as a result of the replacement of coils, one patient was excluded because the tibia cortex was too thin to draw the ROI, and one patient was excluded due to poor image quality. The remaining 95 patients were included. All participants were over 18 years old and ambulatory. The specific diagnostic criteria for CKD are: (1) any evidence of kidney damage (pathological abnormalities, abnormal blood or urine components, or abnormal imaging tests.) or (2) eGFR < 60 mL/min/1.73m² longer than 3 months. The inclusion criteria were hospitalized patients diagnosed with CKD stage 2-5. The reason why we did not include CKD stage 1 (eGFR > 90 mL/min/ $1.73m^2$) is (1) The onset of CKD is insidious in the ultra-early stage, and the clinical symptoms of CKD stage 1 patients are atypical and cannot be firmly diagnosed by eGFR alone. The evidence of kidney damage should be acquired. Fewer than 5% of early-stage CKD patients are aware of their disease (23). Therefore, there are few such patients in clinical practice; (2) In the process of recruiting patients, we hardly recruited patients with CKD stage 1; (3) Almost none studies on CKD-MBD in last decades have included CKD stage 1 patients (8, 15, 24, 25). Key exclusion criteria included any disease known to affect bone metabolism (e.g., diabetes, hyperthyroidism, rickets, osteomalacia, Paget's disease, acromegaly, scurvy (vitamin C deficiency), history of malignant tumors, treatment with radiotherapy or chemotherapy, fractures within 6 months, lumbar or calf trauma surgery, rheumatic immunity disease, scoliosis, anorexia nervosa, and motor neuron disease); use of medications known to influence bone metabolism (e.g., bisphosphonates, oral glucocorticoids, steroid hormones, and salmon calcitonin); and general contraindications to MRI exam (e.g., IUD, pacemaker, cochlear implant, and claustrophobia).

MRI Scanning

Studies were conducted on a 3 Tesla (T) clinical scanner (Signa Pioneer, GE Healthcare, USA) using a medium soft coil, and the dominant leg was scanned. A 3D double-echo UTE-Cones sequence (TR/TE₁/TE₂, 11.5/0.032/4.4 ms; FOV, 16 cm; slice thickness, 3 mm; flip angle, 9°; bandwidth, 62.5 kHz; in-plane spatial resolution, 0.89mm*0.89mm; 12 axial images; and scan time, 54 s) was adopted to get tibial images. To locate the thickest part of the tibial cortex, in the positioning phase, the spine coil was used to image the full length of the tibia and was then positioned at the thickest plane (approximately in a 38% place of the tibial length close to the lateral malleolus) of the tibia in the coronal and sagittal planes.

The reason for tibial cortex used for our study is mainly due to two reasons: (1) In the study of bone pore water, the tibial cortex and the femoral neck cortex are often selected parts (8, 20, 22, 26, 27). (2) According to other research and our preexperiment, compared with the femoral neck cortex, the tibial cortex is very thick, and the boundary with the bone marrow is very clear, which is more beneficial to accurately draw the ROI of bone cortex.

Image Processing

All images were processed with ImageJ (National Institutes of Health). Two experienced musculoskeletal radiologists (one had three years of experience and one had five years of experience), who were blinded to the subjects' clinical data, independently drew the ROI. The ROI was drawn on TE₁ and then copied to TE₂. PI = TE_{4.4}intensity/TE_{0.032}intensity. The ROI was limited to a compacting cortex in appearance, and the trabecularized transition zone was not included, because most UTE signals in these parts were derived from adipose tissue, interfering with water signal of the cortical, thereby interfering with the measurement of PI. The tibial cortex was analyzed on the middle 5 layers of the 12 axial images. Subsequently, the PI value of each layer was calculated separately, and then the average value of PI was obtained. **Figure 1** shows examples of the ROI.

Kidney Function

All subjects completed the kidney function examination within one week of the MRI examination. We used serum creatinine level to obtain the estimated glomerular filtration rate (eGFR) through the CKD-EPI formula (28). The stages of CKD were based on the eGFR.

Bone Metabolism Markers

Bone metabolism markers, namely, PTH, $\beta\text{-}CTX$, T-P1NP, OC, and 25OHD, were assessed within one week of the MRI examination.

Dual-Energy X-Ray Absorptiometry

A Prodigy Lunar scanner (GE Healthcare, Waukesha, WI, USA) was used to evaluate the areal BMD and T-score of the lumbar spine (from L1 to L4).

Statistical Analysis

We summarized continuous variables as the mean \pm SD, and categorical variables as frequency and percentage. Based on eGFR, patients were divided into four stages. The baseline characteristics between the four stages were compared for linear trends using one-way analysis of variance (ANOVA) for continuous variables and Chi-squared statistics for categorical variables. Partial correlation analysis was performed to address associations between PI, eGFR and potential mediators, controlling for age and BMI. Multiple linear regression models were used to assess the association between CKD stages and PI value. Crude effects were tested in an unadjusted model. Next model, we adjusted for age, sex, and BMI. In the final model, we introduced the potential mediators that were correlated with PI value in partial correlation analysis as well as the mediators' interactions with CKD stages. We calculated P for the linear trend by analyzing CKD stages as a continuous instead of an ordinal variable. Then, a separate exploratory mediation analysis was carried out for each metabolic indicator to explore the



impact of CKD stages and mediators on the PI value. The mediation R package (29) was used for mediation analysis. In this analysis, statistical models were simplified by treating CKD stages as interval variables. P-values and nonparametric bootstrap confidence intervals were estimated from one thousand simulations. Figure 2 shows the tested hypothesized pathways in this study, which is the underlying model of this analysis. The model assumes that there is a causal relationship between the selected bone metabolism markers and the progression of CKD and cortical bone PI. The model includes age, sex, and BMI to adjust for potential confounding factors. Direct and indirect effects are obtained from multiple linear regression. The interobserver variability of PI was assessed between two independent raters using intraclass correlation coefficients (ICCs). The ICCs were computed by using the two-way mixed model for absolute agreement and single

measures. An ICC value greater than 0.90 was considered excellent, and a value between 0.75 and 0.90 was considered good. All statistical analyses were performed in R v4.0.3. We set the statistical significance at P < 0.05.

RESULTS

Baseline Characteristics

Table 1 reports the baseline characteristics for the four CKD stages (12 people in stage 2, 23 people in stage 3, 20 people in stage 4, and 40 people in stage 5). The mean age, sex and BMI distributions were comparable across all four groups. Patients with more severe CKD had significantly higher intact PTH, β -CTX, T-P1NP, OC and PI values. No difference could be observed in 25OHD or BMD.



outcome (PI). Adjusting for confounders: age, sex, and BMI. Direct and indirect effects are obtained from multiple linear regression. PTH, parathyroid hormone; β CTX, β-CrossLaps; T-P1NP, total procollagen type I amino-terminal propeptide; OC, osteocalcin; 25OHD, 25-hydroxyvitamin D; BMD, bone mineral density; PI, porosity index.

CKD Cortical Bone Mediation Analysis

TABLE 1 | Baseline characteristics by the stages of CKD.

	Total	Stages of CKD				P_{trend}
		2	3	4	5	
n	95	12 (12.6)	23 (24.2)	20 (21.1)	40 (42.1)	na
Males (%)	48 (50.5)	9 (75)	10 (43.5)	12 (60)	17 (42.5)	0.158
Age (years)	49.7 (12.5)	52.9 (8.1)	49.8 (12.8)	49.5 (13.8)	48.7 (13.0)	0.360
BMI (kg/m ²)	23.2 (3.4)	24.3 (4.2)	22.5 (3.3)	23.8 (3.7)	22.9 (2.9)	0.542
eGFR (mL/min/1.73m ²)	27.2 (22.0)	71.0 (5.7)	41.1 (9.0)	23.3 (4.1)	8.1 (3.5)	< 0.001
intact PTH (pg/mL)	150.2 (124.7)	64.9 (51.2)	84.9 (54.5)	139.9 (81.5)	218.4 (149.2)	< 0.001
β-CTX (ng/mL)	1.23 (1.00)	0.58 (0.39)	0.75 (0.47)	0.91 (0.54)	1.86 (1.18)	< 0.001
T-P1NP (ng/mL)	122.7 (140.2)	29.8 (19.9)	40.6 (49.9)	88.6 (74.9)	214.9 (164.8)	< 0.001
OC (ng/mL)	73.9 (66.3)	27.2 (19.2)	30.4 (24.2)	58.6 (39.6)	120.5 (72.5)	< 0.001
250HD (ng/mL)	18.5 (8.5)	21.9 (8.2)	19.8 (8.5)	16.8 (7.7)	17.6 (8.8)	0.102
L1-L4 BMD (g/cm ²)	1.11 (0.16)	1.17 (0.14)	1.14 (0.17)	1.07 (0.15)	1.10 (0.16)	0.120
L1-L4 T-score	-0.83 (1.23)	-0.45 (1.07)	-0.59 (1.40)	-1.11 (1.02)	-0.93 (1.25)	0.151
PI (%)	20.2 (2.4)	18.2 (1.8)	19.8 (2.0)	20.2 (2.1)	21.1 (2.4)	< 0.001

Data are presented as means (standard deviation) and numbers (%). In Appropriate time, one-way ANOVA and Chi-square are used. Bold P_{trand} indicates statistical significance.

Partial Correlation Between PI, eGFR and Potential Mediators

Significant correlations were revealed between the PI value and different bone metabolism markers (PTH: r = 0.381, P < 0.001; β -CTX: r = 0.375, P < 0.001; T-P1NP: r = 0.327, P = 0.001; OC: r = 0.357, P < 0.001). Similarly, significant correlations were revealed between the eGFR and different bone metabolism markers (PTH: r = -0.499, P < 0.001; β -CTX: r = -0.470, P < 0.001; T-P1NP: r = -0.512, P < 0.001; OC: r = -0.538, P < 0.001) (**Table 2**).

Multiple Linear Regression Analysis

In the crude model (model 1), we observed a positive association of CKD stages with PI, again with a significant gradient [stage 3 versus 2: β (95% CI) 1.58 (0.02, 3.14), stage 4 versus 2: β (95% CI) 2.01 (0.41, 3.61), stage 5 versus 2: β (95% CI) 2.93 (1.48, 4.37), $P_{\text{trend}} < 0.001$; In model 2, after adjusting for age, sex and BMI, the association was attenuated, but significance remained [stage 4 versus 2: β (95% CI) 1.96 (0.32, 3.59), stage 5 versus 2: β (95% CI) 2.83 (1.32, 4.33), $P_{\text{trend}} < 0.001$]. In model 3, after introducing all potential mediators (PTH, β -CTX, T-P1NP, and OC) that were correlated with PI value in partial correlation analysis as well as the mediators' interactions with CKD stages, we found that the association between CKD stages and PI were no longer significant. (**Table 3**).

Mediation Analysis

According to model 3, we found that the potential mediators were in the four bone metabolism markers (PTH, β -CTX, T-P1NP, and

OC). At the same time, due to the multicollinearity between these four markers and their correlation with eGFR, in order to explore the impact of CKD stages and these four markers on PI value, we conducted simple causal mediation analysis for these four markers respectively. The main effects of PTH and β -CTX were significant (Table S2). We tested interactions of the stages of CKD with each potential mediator respectively (adjusted for age, sex and BMI); none of them was significant (P > 0.05). The association of CKD stages and PI mediated 34.4% (P = 0.012) and 30.8% (P = 0.024) of the total effect by increased PTH and β -CTX, respectively. No significant mediation was observed by T-P1NP and OC, 25(OH) vitamin D or BMD (Table 4). The results of mediation analysis of PTH and β -CTX were shown in Figure 3. After adjusting for confounding factors, one increase in CKD stage was associated with an increase of 58.87 pg/mL (95% CI: 38.00, 79.73) in PTH, and associated with an increase of 0.545% (95% CI: 0.096, 1.027) in PI; for every 1 pg/mL increase in PTH was associated with an increase of 0.005% (95% CI: 0.001, 0.009) in PI; the increased PTH mediated 34.4% of the total effect of CKD stages on PI. Similarly, one increase in CKD stage was associated with an increase of 0.47 ng/mL (95% CI: 0.30, 0.63) in β-CTX, and associated with an increase of 0.575% (95% CI: 0.099, 1.058) in PI; for every 1 ng/mL increase in β -CTX was associated with an increase of 0.549% (95% CI: 0.027, 1.071) in PI; the increased β -CTX mediated 30.8% of the total effect of CKD stages on PI. The association of CKD stages with potential mediators was shown in Table S1. The main effect of potential mediators was shown in Table S2.

 TABLE 2 | Adjusted correlation coefficients between PI, eGFR and potential mediators.

	PI	eGFR
PTH	0.381 (<0.001)	-0.499 (<0.001)
β-CTX	0.375 (<0.001)	-0.470 (<0.001)
T-P1NP	0.327 (0.001)	-0.512 (<0.001)
OC	0.357 (<0.001)	-0.538 (<0.001)
250HD	0.058 (0.578)	0.171 (0.101)
L1-L4 BMD	-0.139 (0.183)	0.227 (0.029)
L1-L4 T-score	-0.135 (0.198)	0.215 (0.039)

Partial correlation coefficients adjusted by age, and eGFR. Data are presented as means (P-value). The bold part indicates significant meaning.

CKD Cortical Bone Mediation Analysis

Outcome		Stages of CKD				
	2	3	4	5		
Model 1	(ref.)	1.58 (0.02, 3.14)	2.01 (0.41, 3.61)	2.93 (1.48, 4.37)	<0.001	
Model 2	(ref.)	1.49 (-0.13, 3.11)	1.96 (0.32, 3.59)	2.83 (1.32, 4.33)	<0.001	
Model 3	(ref.)	1.35 (-0.25, 2.96)	1.53 (-0.12, 3.18)	1.73 (-0.02, 3.49)	0.084	

TABLE 3 | Association of the stages of CKD with Pl.

Data are presented as regression coefficient (95% Cl). The bold part indicates significant meaning.

Model 1: crude stages of CKD (Stage 2 is reference category).

Model 2: Model 1 + age, sex, and BMI.

Model 3: Model 2 + potential mediators (PTH, β-CTX, T-P1NP, and OC) (and their interactions with CKD stages).

Interobserver Variability

The ICC was 0.85 (95% CI: 0.430, 0.942).

DISCUSSION

We examined the associations of CKD stages with the PI of the tibial cortex. More severe CKD is associated with higher PI values, independent of age, sex and BMI, while dependent of potential mediators. It was suggested that the association between CKD stages and tibial PI was partly due to PTH and β -CTX through exploratory mediation analysis.

Although the importance of cortical porosity for bone biomechanics has been increasingly appreciated (7, 30, 31), almost no one has assessed it noninvasively *in vivo*. Through PI measurements, we evaluated cortical porosity of the tibia by using double-echo UTE MRI sequences. The mean PI value of all subjects measured in our study was $20.2\% \pm 2.4\%$, which is consistent with the tibial cortical PI value of $20\% \pm 3.8\%$ (range, 15%-31%) of 34 old menopausal women aged 55-80 years (20). This method can quantify the changes in cortical PI in CKD patients, providing a new choice to monitor and detect the deterioration of cortical bone microarchitecture in CKD patients.

The mechanisms underlying the association between CKD stages and cortical porosity are not completely clear, and few related studies have been conducted. Nickolas et al. (8) performed HR-pQCT scans on the distal tibia and distal radius. They found that dialysis was an independent factor in the increase in cortical porosity. On the basis of this research, our study further covered more complete CKD stages, supplemented the current evidence, and further showed that the progression of CKD is closely associated with increased cortical porosity. Nevertheless, we should not assume that more severe CKD

necessarily corresponds to greater PI values, because the actual mechanism by which CKD drives cortical pore enlargement is still largely unknown (32). The factors leading to bone deterioration in patients with CKD are very complex and include the severity of hyperparathyroidism, the level of various bone metabolism markers, the degree of bone turnover and the impact of CKD itself on the bones, and these factors are interrelated (33–35). Consequently, we further conducted an exploratory mediation analysis. We tested each mediator separately with corrected age, sex and BMI to prevent overadjustment.

Among the indicators included in this study, high PTH was the strongest mediator. PTH can directly stimulate osteocytes to release receptor activator of nuclear factor kB ligand (RANKL), which results in increased osteoclast generation (36). In a study of CKD mice, Metzger et al. (37) found that PTH and cortical porosity increased with time, meanwhile RANKL increased and osteocyte apoptosis decreased. Therefore, they speculated that high PTH could prolong the life of osteocytes and lead to sustained RANKL and thus osteoclastic bone resorption in the cortex, promoting the cortical porosity formation. At present, PTH is still the most commonly used diagnostic aid for MBD. On the basis of the 2017 Kidney Disease: Improving Global Outcomes (KDIGO) guidelines, the level of bone turnover can be classified according to the level of PTH (2). Recently, some studies have shown that increased porosity is associated with increased PTH (3, 8, 26, 37, 38), which is consistent with our results. At the same time, our study also found that the increase in cortical porosity in people at different CKD stages is partly due to higher PTH.

Second, our research found that β -CTX was also a mediator. β -CTX is a marker of bone resorption, which can reflect osteocyte activity (39). At present, there is no consensus on the application of β -CTX in MBD. Some studies indicate that β -CTX

TABLE 4 Mediators of the association between CKD stages and Pl.							
Mediators	IDE	Р	DE	Р	Proportion mediated	Р	
PTH	0.286 (0.051, 0.518)	0.012	0.545 (0.096, 1.027)	0.012	0.344 (0.051, 0.824)	0.012	
β-CTX	0.256 (0.040, 0.499)	0.024	0.575 (0.099, 1.058)	0.020	0.308 (0.038, 0.795)	0.024	
T-P1NP	0.198 (-0.064, 0.477)	0.140	0.633 (0.096, 1.191)	0.022	0.238 (-0.079, 0.794)	0.140	
OC	0.242 (-0.081, 0.529)	0.114	0.589 (0.067, 1.168)	0.024	0.292 (-0.083, 0.862)	0.114	

Data are presented as means (95% Cl). Nonparametric bootstrap Confidence Intervals are estimated from 1000 simulations adjusted for age, sex, and BMI. Bold P-values indicate statistical significance. IDE, indirect effect; DE, direct effect.



FIGURE 3 | Estimates of the mediation effect of (A) PTH and (B) β -CTX about the association between the stages of CKD and PI value, after adjusting for age, sex and BMI. Note: (A) the effect of grouping on mediator; (B) the main effect of mediator on PI; IDE, indirect effect (Stages of CKD \rightarrow PTH/ β -CTX \rightarrow PI); DE, direct effect (Stages of CKD \rightarrow PTH, parathyroid hormone; β -CTX, β -CrossLaps; PI, porosity index. *P < 0.05.

is a bone turnover biomarker which exists direct release during bone resorption and could be used to more accurately assess the bone turnover in CKD (12). At the same time, Nishiyama et al. (24) found that higher levels of β -CTX and worsening cortical porosity have a linear dose-response relationship, which suggested that β -CTX could predict cortical deterioration. However, some studies suggested that β -CTX was not suitable as a diagnostic aid for MBD due to the influence of renal function (40). The role of this indicator in CKD-MBD needs further study. T-P1NP and OC did not show mediation effects in this study, possibly because these two indicators were greatly affected by renal clearance and could not accurately reflect bone deterioration in patients with CKD. There was also no mediation effect of 25OHD, which may due to supplement with vitamin D for the most patients with end-stage CKD, so that the 25OHD level in end-stage patients was higher than that in patients at earlier stages in our study. Due to the analytical and biological variability of these serum markers, there are few studies on bone metabolism markers other than PTH in CKD, and the clinical applications of these indicators are limited. The results of this study could provide some hints for further research in the future.

Interestingly, we did not observe a mediation effect of lumbar BMD measured by DXA, and we found that as CKD progressed, BMD did not continue to decrease. It might be that the BMD measured by DXA was a real BMD, which is not distinguishable between trabecular and cortical bone (11, 41), or that BMD itself has a small contribution to porosity. Because cortical and trabecular bone behave differently in answer to increased parathyroid activity (decrease and increase, respectively) and larger trabecular BMD can mask the loss of cortical bone, BMD measured by DXA in CKD patients may be higher, equal to or lower than that in healthy controls (13). Some studies were consistent with our study. Rajapakse et al. (20) showed that the femur and spine BMD measured by DXA had no significant correlation with PI. They also found that the cortical BMD obtained using pQCT imaging in bone specimens can merely partially explain the change in porosity, highlighting the need to estimate porosity without the

dependence of BMD. Similarly, Carvalho et al. (42) found that there was no significant correlation between DXA parameters and cortical porosity for either internal cortical bone or external cortical bone.

The advantages of this study include: (1) The assessment of relatively complete indicators related to bone deterioration. (2) There is currently little evidence for indicators other than PTH in CKD, and this study can be used as a supplement to some extent. (3) We prospectively included CKD stages 2-5 patients in strict accordance with the inclusion criteria and exclusion criteria that may affect bone metabolism. (4) We also explored the mediation effect of a series of potential mediators, which can reflect the inherent causal relationship between CKD progress and the changes in PI values, which is not seen in the current CKD-MBD research. At the same time, this study has some limitations: (1) Many drugs and diseases can affect bone metabolism, so we excluded diseases and drugs that are known to affect bone metabolism as much as possible based on relevant research. Dialysis has been proved to be an independent factor of PI increase (8), but this study did not include dialysis patients. Our follow-up study will take dialysis patients as a separate group for further study. (2) Due to the limited detection level, we did not include TRAP5b or BSAP, two markers of bone metabolism that are considered to be independent of glomerular filtration rate.

In conclusion, the progression of CKD is associated with increased cortical PI values, and this association may be partly mediated by increased PTH and β -CTX. Our study provides a new idea to monitor bone health in patients with CKD, and reveal the internal mechanism of bone deterioration caused by CKD to some extent. Further work needs to be done to study other deep possible mechanisms about the association.

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 Medical Ethics Committee of Tongji Hospital, Tongji Medical College, Huazhong University of Science and Technology. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

FH and XL concept and design. DW, BH, and YL devised the outline of the manuscript. YX and TH designed the study and evaluated the data. YW, SH, and YZ collected the information and analyzed the data. YX wrote the manuscript. PZ and JL created the figures for the manuscript. WL and XL critically

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fendo.2021. 775066/full#supplementary-material

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