

# Evaluation of prolidase activity and oxidative status in patients with knee osteoarthritis: relationships with radiographic severity and clinical parameters

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**Abstract** We investigated serum prolidase activity and oxidative/antioxidative status in patients with knee osteoarthritis (OA) and evaluated its relationships with radiographic severity and clinical parameters. The study population consisted of 137 patients with knee OA and 134 healthy volunteers. The severity of knee OA was classified according to the Kellgren–Lawrence criteria. Each patient was also evaluated clinically according to the Western Ontario and McMaster University Osteoarthritis Index (WOMAC). Serum prolidase activity was measured spectrophotometrically. Oxidative status was assessed by measuring serum lipid hydroperoxide (LOOH) and total oxidative status (TOS). Antioxidative status was assessed by measuring serum-free sulfhydryl groups (–SH = total thiol) and total antioxidant capacity (TAC). Oxidative stress index (OSI) was calculated. Serum prolidase activity was significantly lower in the knee OA group than in the control group ( $p < 0.001$ ). The serum prolidase activities decreased with the severity of knee OA. Furthermore, serum LOOH, TOS, and OSI levels of the knee OA group were significantly higher than those of the controls ( $p < 0.001$  for all), whereas TAC and –SH levels did not differ between the two groups ( $p > 0.05$ ). In a multiple regression analysis, WOMAC score was independently associated with serum prolidase activity ( $\beta = -0.340$ ,  $p < 0.001$ ). Decreased serum prolidase activity and elevated LOOH, TOS, and OSI levels may be associated

with knee OA, and serum prolidase activity may be a useful adjunctive indicator of the progression of knee OA in follow-up.

**Keywords** Prolidase · Oxidative status · Knee osteoarthritis · WOMAC score · X-ray grading

## Introduction

Osteoarthritis (OA) is a common degenerative joint disease, and knee is the most frequently involved joint. Radiographic appearance and clinical features are still often used for diagnosis of the disease. However, the etiology of OA is not fully understood, although mechanical, biochemical, and genetic factors are accepted to play roles [1, 2]. Intraarticular and periarticular structures are affected in knee OA, and histopathological changes occur in the joint cartilage, subchondral bone, ligaments, periarticular muscle, nerve, and synovial membrane [3]. The general consensus is that these changes help to explain the persistence of knee OA and progression of radiographic severity and clinical parameters. Understanding the histopathological changes that characterize OA and investigating adjunctive tools for evaluation of radiographic severity and clinical parameters of the disease are the most important areas of current research. The major component of the structures involved in knee OA is collagen. Serum prolidase enzyme activity, which is required for collagen biosynthesis, plays a key role in the breakdown of collagen, and prolidase enzyme activity has been shown to be related to the collagen turnover rate [4, 5]. Prolidase enzyme activity and its pathophysiological role have been investigated in the context of different musculoskeletal system diseases [6, 7]. In a recent study, serum

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prolidase activity was evaluated in patients with knee OA and found to be significantly reduced compared to a control group [8]. These studies showed that serum prolidase enzyme activity reflects the abnormal collagen turnover in affected joints. Nothing on the correlation between serum prolidase enzyme activity and radiographic severity of knee OA and clinical parameters has yet appeared in the literature.

One possible cause of OA is oxidative stress. The levels of pro-inflammatory mediators, such as reactive oxygen species (ROS), are elevated in OA [9, 10]. Thus, the increased levels of these reactive species with oxidative activity mediate the effects of many pro-inflammatory cytokines, such as interleukin (IL)-1 and tumor necrosis factor (TNF)- $\alpha$  [9–11]. IL-1 and TNF- $\alpha$  may play a crucial role in cartilage matrix degradation by stimulating matrix metalloproteinase (MMP) expression in patients with OA [10, 11]. Additionally, recent studies have demonstrated that mitochondrial DNA damage can be induced by these inflammatory cytokines or ROS, and this damage may promote chondrocyte death [12–14]. Thus, ROS contributes to cartilage degradation by reducing cartilage repair capacity and cell death in extracellular cartilage matrix.

Thus, we investigated serum prolidase enzyme activity, oxidative status [lipid hydroperoxide (LOOH), total oxidative status (TOS), and the oxidative stress index (OSI)], antioxidative status (free sulfhydryl groups,  $-SH =$  total thiol), and total antioxidative capacity (TAC) in knee OA and evaluated their relationship with radiographic severity of knee OA and clinical parameters.

## Materials and methods

### Study population

This study included 137 patients with knee OA (90 females, 47 males; mean age  $58.5 \pm 9.8$  years, range 40–79) diagnosed according to the American College of Rheumatology criteria [15] and 134 healthy controls (85 females, 49 males; mean age  $59.3 \pm 8.7$  years, range 43–80) who were free of symptoms. Exclusion criteria included the use of supplemental vitamins, secondary posttraumatic OA, previous knee injury, inflammatory rheumatic disease, infectious- or endocrine-related arthropathy, clinically unstable medical illness, or the use of any medication within 4 weeks prior to initiation of the study. The controls were matched in terms of age, body mass index (BMI), and gender with the patients.

Written informed consent was obtained from all subjects. The use of their medical records was approved by the local research ethics committee.

### Radiological assessment

All radiographs of the patients (weight-bearing anteroposterior and weight-bearing lateral and Merchant's X-rays of both knees) were evaluated by a single observer (CE) who was not involved in the clinical care of the patients. The severity of OA was assessed and classified according to the Kellgren–Lawrence (KL) grading scale [16]. Grade 0 was accepted as normal, grade 1 as possible osteophytes only, grade 2 as absolute osteophytes and possible joint space narrowing, grade 3 as moderate osteophytes and/or absolute joint space narrowing, and grade 4 as large osteophytes, severe joint space narrowing, and/or bony sclerosis. The grade used was the highest of the right or left knee. Patients with knee OA were divided into four subgroups according to the KL grading scale.

### Clinical assessment

Clinical assessments included the Western Ontario and McMaster University Osteoarthritis Index (WOMAC). The WOMAC questionnaire consists of three items: pain, stiffness, and physical function. The WOMAC scores were recorded on a Likert scale of 0–4 (0 = no pain/limitation, 1 = mild pain/limitation, 2 = moderate pain/limitation, 3 = severe pain/limitation, and 4 = very severe pain/limitation). Maximum scores for pain, stiffness, and physical function were 20, 8, and 68, respectively, for a total score of 96 (0–96 scale with 0 as the best and 96 as the worst) [17]. Additionally, clinical baseline parameters such as weight, height, BMI, age, gender, side of involvement (left or right or both knees) were recorded.

### Blood sample collection

Blood samples were collected into Vacutainer serum clot activator tubes (BD<sup>®</sup>, USA) and immediately stored at 4 °C. Serum samples were then separated from cells by centrifugation (3000 rpm, 10 min). The remaining serum portions were stored at  $-80$  °C until analysis.

### Measurement of serum prolidase activity

Prolidase activity was measured by a photometric method, based on the measurement of proline levels produced by prolidase [18]. The samples (100  $\mu$ L) were mixed with 100  $\mu$ L of physiological saline (NaCl, 0.9 %). A volume of 25  $\mu$ L of the mixture was preincubated with 75  $\mu$ L of the preincubation solution (50 mmol/L Tris–HCl buffer pH 7.0 containing 1 mmol/L GSH, 50 mmol/L  $MnCl_2$ ) at 37 °C for 30 min. The reaction mixture, which contained 144 mmol/L gly-pro, pH 7.8 (100  $\mu$ L), was incubated with 100  $\mu$ L of preincubated sample at 37 °C for 5 min. To stop the reaction, 1 mL glacial acetic acid was added. After

adding 300  $\mu\text{L}$  Tris–HCl buffer (pH 7.8) and 1 mL ninhydrin solution (3 g/dL ninhydrin was dissolved in 0.5 mol/L orthophosphoric acid), the mixture was incubated at 90 °C for 20 min and then cooled on ice. Absorbance was then measured at 515 nm to determine proline levels, according to the method of Myara et al. [19], as optimized by Ozcan et al. [18]. Intra- and interassay coefficients of variation (CV) were lower than 10 % for the assay.

### Measurement of lipid hydroperoxide levels

Serum LOOH levels were measured by a ferrous ion oxidation-xylenol orange ('FOX-2') assay, which involves the oxidation of ferrous ions to ferric ions via the effects of various oxidants. The ferric ion level is then measured using xylenol orange. LOOH levels are reduced by the application of triphenyl phosphine (TPP), a reductant specific for lipids. LOOH levels were then calculated as the difference between the absence and presence of TPP [20]. The CV for the determination of the LOOH level was 5 %.

### Measurement of total free sulfhydryl groups

Free sulfhydryl groups (–SH) in serum samples were assayed according to the method of Ellman [21] as modified by Hu et al. [22]. Briefly, 1 mL of buffer containing 0.1 M Tris, 10 mM EDTA (pH 8.2), and 50  $\mu\text{L}$  serum was added to cuvettes, followed by 50  $\mu\text{L}$  of 10 mM DTNB in methanol. Blanks were run for each sample with no DTNB. Following incubation for 15 min at room temperature (RT), sample absorbance was read at 412 nm on a Cecil 3000 spectrophotometer (Cecil Instruments, Cambridge, UK). The concentration of sulfhydryl groups was calculated using reduced glutathione as the free sulfhydryl group standard, and the results are expressed in millimoles. The CV for the measurement of serum –SH levels was 3.6 %.

### Measurement of total oxidative status (TOS)

The TOS of serum was determined using an automated measurement method (Rel assay diagnostics kits) developed by Erel [23]. Oxidants present in the sample oxidized the ferrous ion-*o*-dianisidine complex to ferric ions. The oxidation reaction is enhanced by glycerol molecules, which are abundant in the reaction medium. The ferric ion produces a colored complex with xylenol orange in an acidic medium. The color intensity, which can be measured spectrophotometrically, is related to the total amount of oxidant molecules present in the sample. The assay was calibrated with hydrogen peroxide ( $\text{H}_2\text{O}_2$ ), and the results are expressed as micromoles of  $\text{H}_2\text{O}_2$  equivalents per liter ( $\mu\text{mol H}_2\text{O}_2$  equiv/L). The CV for the measurement of the serum TOS level was <3 %.

### Measurement of total antioxidative capacity (TAC)

TAC of serum was determined using an automated measurement method (Rel assay diagnostic kits) of Erel [24], in which the hydroxyl radical, the most potent biological radical, is produced. Ferrous ion solution in reagent-1 is mixed with  $\text{H}_2\text{O}_2$ , which is present in reagent-2. Sequentially produced radicals include the brownish-colored dianisidiny radical cation, produced by the hydroxyl radical. Using this method, the antioxidative effect of the sample against the potent free radical reactions, which was initiated by the hydroxyl radicals produced, was measured. The data are expressed as mmol Trolox equiv/L. The CV for the measurement of the serum TAC level was <3 %.

### Oxidative stress index (OSI)

The ratio of TOS to TAC yielded the OSI, an indicator of the percentage degree of oxidative stress [23, 24]. For calculations, the unit of TAC was changed to  $\mu\text{mol/L}$ , and the OSI value was calculated according to the following formula: OSI (arbitrary units):  $\text{TOS } (\mu\text{mol H}_2\text{O}_2 \text{ equiv/L}) / \text{TAC } (\mu\text{mol Trolox equiv/L}) \times 10$ .

### Statistical analysis

All analyses were conducted using the SPSS software (version 16; SPSS Inc., Chicago, IL, USA). Continuous variables are expressed as mean  $\pm$  SD. Comparisons of categorical and continuous variables between the knee OA and control groups were performed using the Chi-square test and independent-samples *t* test, respectively. The paired-samples *t* test was used for comparison of continuous variables categorized according to age within both groups. Comparison of laboratory variables between groups categorized according to the grade of knee OA was performed using one-way analysis of variance (ANOVA) with the least significant difference post hoc test. The correlation between serum prolidase activity and clinical and laboratory parameters was evaluated using Pearson's correlation test. Standardized  $\beta$ -regression coefficients and their significance in a multiple linear regression analysis were reported. A two-tailed  $p < 0.05$  was considered to indicate statistical significance.

### Results

Demographic characteristics of patients with knee OA and the control group are shown in Table 1. No significant difference was observed in age, BMI, or female/male ratio between patients and controls.

Serum prolidase activity was significantly lower in patients with knee OA compared with controls ( $p < 0.001$ ).

**Table 1** Demographic characteristics of the knee osteoarthritis and control groups

	Patients <sup>a</sup> (n = 137)	Controls <sup>a</sup> (n = 134)	p value
Age (years)	58.5 ± 9.8	59.3 ± 8.7	>0.05
Gender (M/F)	47/90	49/85	>0.05
BMI (kg/m <sup>2</sup> )	30.0 ± 4.4	30.5 ± 4.5	>0.05
Prolidase activity (U/L)	754.6 ± 19.1	772.7 ± 34.6	<0.001
LOOH (μmol/L)	7.78 ± 0.66	5.93 ± 0.39	<0.001
TOS (μmol H <sub>2</sub> O <sub>2</sub> equiv/L)	15.8 ± 2.31	10.6 ± 1.25	<0.001
OSI (arbitrary unit)	1.71 ± 0.53	1.12 ± 0.29	<0.001
Total -SH (mmol/L)	0.38 ± 0.11	0.39 ± 0.09	>0.05
TAC (mmol Trolox equiv/L)	0.99 ± 0.23	0.99 ± 0.23	>0.05
Side of involvement [left (L), right (R), or both (B)]	64 L, 48 R, 25 B	52 L, 63 R, 19 B	>0.05

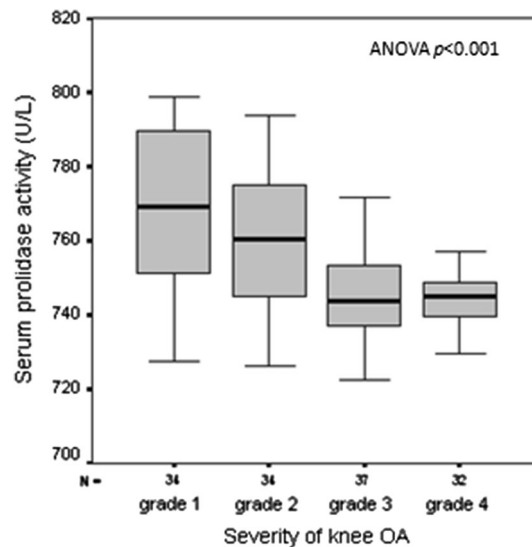
BMI body mass index, LOOH lipid hydroperoxide, TOS total oxidant status, OSI oxidative stress index, SH free sulfhydryl groups (total thiol), TAC total antioxidant capacity

<sup>a</sup> Measurements are given as the mean ± SD

The comparison included 137 patients with OA in four subgroups according to the KL grading scale—grade 1 ( $n = 34$ ), grade 2 ( $n = 34$ ), grade 3 ( $n = 37$ ), and grade 4 ( $n = 32$ )—along with 134 controls. Figure 1 demonstrates levels of serum prolidase activity in the patient group divided by X-ray grades of the knee joints. The ANOVA test indicated significant differences between the groups. Serum prolidase activities of the patients with grades 3 and 4 (late knee OA) were significantly lower than those of knee grades 1 and 2 (early knee OA; ANOVA,  $p < 0.001$ ). Although prolidase activity was significantly lower in the patient group than in the controls ( $p < 0.001$ ), when all participants were stratified into two groups (40–60 and 61–80 years) according to age, there was no statistically significant difference between the two layers within each group (Table 2). The relationship between serum prolidase activity and clinical characteristics is presented in Table 3. Serum prolidase activity was inversely correlated with WOMAC score in a bivariate analysis ( $r = -0.516$ ,  $p < 0.001$ ; Table 3). The correlation between serum prolidase activity and WOMAC score is shown in Fig. 2. In a multiple regression analysis, we observed that WOMAC score ( $\beta = -0.340$ ,  $p < 0.001$ ) was independently associated with serum prolidase activity (Table 3).

Furthermore, oxidant parameters (LOOH, TOS, and OSI) were significantly higher in knee OA patients ( $p < 0.001$  for all). However, antioxidant parameters (TAC and -SH) did not differ between the groups ( $p > 0.05$ ). In a bivariate analysis, LOOH level was positively correlated with TOS ( $r = 0.609$ ,  $p < 0.001$ ) in patients with OA of the knee.

The relationship between WOMAC score and clinical characteristics and laboratory parameters is presented in Table 4. The WOMAC score was positively correlated with severity of knee OA ( $r = 0.619$ ,  $p < 0.001$ ), but inversely



**Fig. 1** Serum prolidase activities in patients with grades 3 and 4 were significantly lower than those with knee grades 1 and 2 (ANOVA,  $p < 0.001$ )

correlated with serum prolidase activity ( $r = -0.516$ ,  $p < 0.001$ ) in a bivariate analysis. In a multiple regression analysis, we found that severity of knee OA ( $\beta = 0.429$ ,  $p < 0.001$ ) and serum prolidase activity ( $\beta = -0.298$ ,  $p < 0.001$ ) were independently associated with the WOMAC score (Table 4).

## Discussion

The main results of this study were as follows. First, prolidase activity was lower in the knee OA group than in the control group. Second, after the participants were stratified

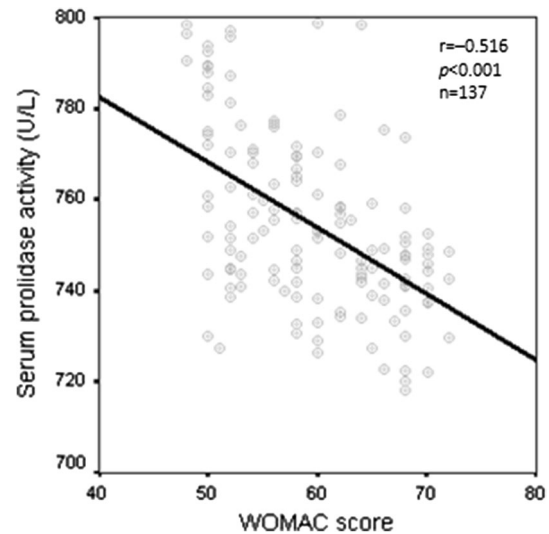
**Table 2** Serum prolidase activity of the knee osteoarthritis and control groups according to age

Age (years)	Patients <sup>a</sup> (n = 137)	Controls <sup>a</sup> (n = 134)	p value
40–60	755.8 ± 20.5 (n = 86)	774.6 ± 35.4 (n = 75)	<0.001
61–80	753.2 ± 16.6 (n = 51)	769.2 ± 34.2 (n = 59)	<0.001
p value	0.435	0.358	

<sup>a</sup> Measurements are given as the mean ± SD

into two groups according to age, serum prolidase activity was significantly lower in the patients group than in the controls in both age groups. Third, prolidase activity was significantly decreased in late-grade versus early-grade knee OA. Fourth, the oxidative stress parameters (LOOH, TOS, and OSI) of the OA group were significantly greater than those of the control group, whereas the antioxidative stress parameters (–SH groups and TAC) were unchanged. Fifth, the WOMAC score was independently correlated with prolidase activity and knee OA grading.

We observed that serum prolidase activity was significantly lower in patients with knee OA than in the controls. The pathogenic mechanisms underlying this reduction in serum prolidase activity in patients with OA are unclear. However, three major causes have been suggested. First, it may be related to decreased collagen resynthesis in patients with knee OA. Prolidase (E.C. 3.4.13.9) is a cytosolic Mn(II)-activated metalloproteinase that hydrolyzes imidodipeptides and imidotriptides with a C-terminal proline or hydroxyproline. Then, it releases these two amino acids for collagen resynthesis and cell growth, and its activity determines the rate of collagen turnover [4, 5, 25]. Although prolidase activity is found in a wide variety of diseases, the reports have been inconsistent. Several studies have reported that prolidase activity decreases in diseases such as chronic uremia [26] and cardiomyopathy [27], while increased prolidase activity has been described in Legg–Calve–Perthes disease [6] and idiopathic clubfoot [7]. Under normal circumstances, chondrocytes stimulate synthesis of the new extracellular matrix, such as aggrecan and type II collagen. However, in patients with OA, synovial cells and



**Fig. 2** Correlation between serum prolidase activity and severity of knee osteoarthritis

chondrocytes serve to increase the levels of inflammatory cytokines. Thus, this anabolic phase of matrix remodeling is considered insufficient or defective. Second, the reason for this imbalance in cartilage remodeling in OA may be related to aging [12, 28]. Previous studies have demonstrated that the accumulation of advanced glycation end products (AGEs) is associated with age in OA. The accumulation of AGEs causes damage to matrix synthesis in articular cartilage due to decreased collagen turnover [28]. In this study, to evaluate the effect of age, we stratified all participants into two groups according to age. We found lower prolidase enzyme activity in the patient group in each age group than in the controls. Although a partial decrease was seen in the >60-year-old group, we did not find a statistically significant difference between the two layers within each group. These results may be related to the severity of the knee OA in terms of prolidase activity rather than age. Another possible reason for decreased prolidase activity in knee OA may be related to the low physical activity levels of the elderly population. Physical activity levels decreased in patients with knee OA, and collagen turnover seems to be positively correlated with the degree of exercise [27, 29].

**Table 3** Relationship between serum prolidase activity and clinical characteristics

	Pearson’s correlation coefficient	p value	β regression coefficient <sup>a</sup>	p value
BMI (kg/m <sup>2</sup> )	0.003	0.085		
Severity of knee OA	–0.495	<0.001	–0.285	0.002
WOMAC score	–0.516	<0.001	–0.340	<0.001

OA osteoarthritis, WOMAC Western Ontario and McMaster University Osteoarthritis Index

<sup>a</sup> From multiple linear regression

**Table 4** Relationship between the WOMAC score and clinical characteristics and laboratory parameters

	Pearson's correlation coefficient	<i>p</i> value	$\beta$ regression coefficient <sup>a</sup>	<i>p</i> value
Age (years)	0.216	<0.05	0.103	0.10
BMI (kg/m <sup>2</sup> )	0.262	<0.01	0.231	<0.001
Severity of knee OA	0.619	<0.001	0.429	<0.001
Prolidase activity (U/L)	-0.516	<0.001	-0.298	<0.001

*BMI* body mass index, *OA* osteoarthritis

<sup>a</sup> From multiple linear regression

Furthermore, we found significant decreases in the levels of prolidase activity in the late stages of knee OA. The cause of the decreased serum prolidase activity with advancing OA stage is unclear. However, we believe that these results may be due to grades 3 and 4 being the late stages of OA, in which most of the cartilage has already degenerated and morphological changes are manifest.

In our study, we observed significantly elevated oxidant parameters in subjects with knee OA compared to the controls. These results confirmed the presence of oxidative stress. Under healthy conditions, when ROS production decreases, LOOH and TOS are inhibited by various antioxidants present in plasma. However, in the case of excessive ROS production, as demonstrated in our study by the rise in LOOH and TOS, this protection may be inadequate as a defense mechanism against continuing oxidative stress. Oxidative stress is associated with OA, but the precise mechanism is unclear. However, oxidative stress causes telomere shortening and aging in chondrocytes. Senescent cells have decreased tolerance to oxidative stress [14, 30]. Furthermore, ROS promote cellular senescence and apoptosis. Thus, ROS may play important roles in the development of OA. We used serum -SH groups and TOS levels as markers of antioxidant status. In a previous study in our laboratory, Altindag et al. [8] showed that serum -SH groups, TAC levels, and prolidase activity were lower, whereas LOOH levels were higher, in patients with knee OA compared with healthy individuals. However, we detected no difference in the serum levels of -SH groups or TAC in patients with knee OA compared with controls. As a result, the cause of the unchanged antioxidative stress parameters (-SH groups and TAC) may be attributable to increased oxidative stress compensatory mechanisms in patients with knee OA.

In the present study, the WOMAC score was positively correlated with the severity of knee OA, but inversely correlated with serum prolidase activity. The WOMAC scale allows detailed analysis of pain and dysfunction [31]. The articular cartilage is aneural and avascular and thus does not directly generate pain. However, the changes in articulation caused by structural and associated degenerative changes in articular cartilage may result in the manifestation of pain in joint tissues [32].

BMI is routinely used as a screening tool for obesity, and higher BMI has been associated consistently with an elevated risk of knee OA [33]. However, in our study, there was no relationship between the BMI and serum prolidase activity. This may be due to our patient's BMI values not being very high, and as mentioned recently, BMI does not describe the pattern of fat distribution or body composition and does not distinguish adipose from non-adipose body mass due to the changes in fat-free mass and fat mass with height, weight, and age [34].

Several limitations of this study should be noted. First, it had a cross-sectional design and was a single-center study. Second, radiographically visualized degeneration is only part of knee OA. Thus, these findings should be confirmed by magnetic resonance imaging or arthroscopy, which we did not perform. Third, prolidase enzyme activity was evaluated only in serum samples. We were unable to obtain synovial fluid samples due to ethical restrictions. Further prospective longitudinal studies are needed to evaluate whether the results in serum are comparable to those in synovial fluid or cartilage and to determine the molecular mechanisms of the association between prolidase activity and the severity of knee OA.

## Conclusions

In conclusion, decreased serum prolidase activity, correlated inversely with WOMAC scores, and elevated LOOH, TOS, and OSI levels may be associated with knee OA. Additionally, according to the significant decreases in the levels of prolidase activity in the late stages of knee OA, prolidase enzyme activity may be a useful adjunctive indicator to assess the progression of knee OA.

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**Conflict of interest** The authors have no conflicts of interest to declare.

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