ANIMAL STUDY

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Received: 2020.03.30 Accepted: 2020.08.11 Available online: 2020.09.28 Published: 2020.11.24		-	Exploring the Mechanism of Icariin in Osteoporosis Based on a Network Pharmacology Strategy		
Authors' Contribution:ABCDE 1,2Study Design AABCDE 3,4Data Collection BABCDE 5Statistical Analysis CABCDE 6Data Interpretation DABCDE 6Manuscript Preparation EABCDE 7Literature Search FABCD 7		ABCDE 1,2 ABCDE 3,4 ABCDE 5 ABCDE 6 ABCDE 7 ABCD 7	Zhiyong Long* Jiamin Wu* Wang Xiang* Zhican Zeng* Ganpeng Yu* Jun Li	 Shantou University Medical College, Shantou University, Shantou, Guangdong P.R. China Department of Rehabilitation Medicine, Institute of Geriatric Medicine, Guangdong Academy of Medical Sciences, Guangdong General Hospital, Guangzhou, Guangdong, P.R. China Shanghai University of Traditional Chinese Medicine, Shanghai, P.R. China Shuguang Hospital Affiliated to Shanghai University of Traditional Chinese Medicine, Shanghai, P.R. China Affiliated Hospital of Guilin Medical University, Guilin, Guangxi, P.R. China Affiliated Hospital of Guilin Medical University, Guilin, Guangxi, P.R. China Tianjin Medical University, Tianjin, P.R. China Department of Orthopaedics, People's Hospital of Ningxiang City, Ningxiang, Hunan, P.R. China 	
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Background: Material/Methods:		ground: ethods:	With the aging of the world's population, the incidence of osteoporosis (OP) has become a public health prob- lem of worldwide concern. Research shows that icariin may have a therapeutic effect on OP. PharmMapper was utilized to predict the potential targets of icariin. GeneCards and Online Mendelian Inheritance in Man (OMIM) were used for the collection of OP genes. The STRING database was utilized to obtain the pro- tein-protein interaction (PPI) data. We used Cytoscape 3.7.2 to construct and analyze the networks. The genes		
Results:		Results:	(DAVID) to undergo Gene Ontology (GO) and pathway enrichment analysis. Finally, animal experiments were performed to verify the prediction results of this study. A total of 297 icariin potential targets and 262 OP genes were obtained, and an icariin-OP PPI network was con- structed and analyzed. The results of the GO enrichment analysis showed that icariin can regulate the steroid hormone-mediated signaling pathway, skeletal system development, extracellular space, cytosol, and steroid hormone receptor activity. The results of the pathway enrichment analysis showed that icariin can regulate os- teoclast differentiation, FoxO, estrogen, and PPAR signaling pathways. The results of the experiments showed that icariin can increase estradiol, β -catenin, and Receptor Activator of Nuclear Factor- κ B Ligand (RANKL)/os- teoprotegerin (OPG) ratio in postmenopausal OP rats (<i>P</i> <0.05).		
Conclusions:			This research found that the icariin can regulate OP-related biological processes, cell components, molecular functions, and signaling pathways.		
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Background

With the aging of the world's population, the incidence of osteoporosis (OP) has been increasing year by year, and it has become a public health problem of worldwide concern [1,2]. Primary OP is divided into type I and type II; type I OP is postmenopausal osteoporosis (PMOP) and type II OP is senile osteoporosis [3]. PMOP is caused by endocrine disorders due to hypoovarian function after menopause, especially the decrease in estrogen level. Because bone resorption is greater than bone formation, osteoblasts required for bone reconstruction are relatively reduced. The relative increase in osteoclasts leads to a decrease in systemic bone mass and destruction of bone tissue microstructure, resulting in increased bone fragility and metabolic bone disease, predisposing bones to fracture [3-5]. At present, the drugs recognized for the treatment of OP mainly include bone resorption inhibitors, bone formation promoters, and bone minerals [6]. However, from the perspective of drug treatment effects, the current drugs only improve clinical symptoms and delay the progression of the disease, and they cannot restore the patient's bone remodeling process to a balanced state. Therefore, it is urgent to find new ideas to solve this problem [7,8].

Traditional Chinese medicine (TCM) has a role in treating OP [9]. Modern pharmacological studies have found that herbal formulas for treating OP have the combined effects of improving intestinal calcium absorption, promoting osteoblast mineralization, enhancing cell osteogenic activity, and inhibiting the formation of osteoclasts [9,10]. Hence, the active ingredients of herbal medicines have unique advantages and broad development prospects in the prevention and treatment of OP. Icariin is a compound derived from Epimrdii Herba (TCM believes it can strengthen bones). Current studies have found that icariin can regulate osteoblast proliferation and differentiation through bone-specific matrix proteins, transcription factors, cytokines, and Receptor Activator of Nuclear Factor-κ B Ligand (RANKL)/osteoprotegerin (OPG) signaling pathways [11,12]. However, the specific mechanism underlying the effects of OP treatment is still unknown. Network pharmacology, as a new discipline developed based on the intersection and fusion of multidisciplinary technologies such as classic pharmacology, computer technology, and bioinformatics, can systematically study the interactions between drugs and the human body, as well as their laws and essences, from multiple levels such as molecules, cells, and organs [13–15]. This strategy has been widely used in TCM multi-component-multi-targetmulti-path-multifunctional research, analysis of herbal formulas, new drug development, and interpretation of TCM basic theory. Hence, this study used a network pharmacological strategy to explore the specific mechanism involved in icariin treatment of OP. The research process is shown in Figure 1.



Figure 1. The research process.

Material and Methods

Potential targets prediction of icariin and OP genes collection

The molecular structure of icariin was queried at PubChem (*https://pubchem.ncbi.nlm.nih.gov/*), and its SDF file can be downloaded from this website. Then, the SDF file was input into the PharmMapper server (*http://lilab-ecust.cn/pharmmapper/*) to predict the potential targets of icariin [16]. Finally, a total of 297 of icariin's potential targets were predicted by PharmMapper (Supplementary Table 1, see supplementary materials).

The GeneCards (*http://www.genecards.org*) and Online Mendelian Inheritance in Man (OMIM) (*http://omim.org/*) databases were utilized to collect the OP genes [17,18]. After the search, a total of 2983 OP-related genes were obtained. The genes with relevance score \geq 8.0 were selected for sequence research (Supplementary Table 2).

Network construction and analysis methods

The protein-protein interaction (PPI) data of icariin targets and OP genes were collected from STRING 11.0 (*http://string-db.org/*) [19]. After entering the website, select "Multiple Proteins", we imported icariin targets and OP genes into "List of Names", set Organism to "Homo sapiens" for PPI, and then downloaded the file in "tsv" format. Cytoscape 3.7.2 software (*https://cytoscape.org/*) was utilized to draw the networks, such as icariin-OP PPI network, and to perform network analysis [20].

The targets and genes in the icariin-OP PPI network were input into the Database for Annotation, Visualization and Integrated Discovery (DAVID) 6.8 (*https://david.ncifcrf.gov/*) to undergo Gene Ontology (GO) enrichment analysis and Kyoto Encyclopedia of Genes and Genomes (KEGG) pathway enrichment analysis [21].

Animals

Forty-eight (48) specific pathogen-free (SPF)-grade female SD rats, weighing 220–250 g, 10–12 months old, were purchased from Hunan Slake Jingda Experimental Animal Co., Ltd. [Certificate No. 43004700005517, License No. SCXK (Xiang) 2013–0004]. The animals were housed in an air-conditioned room with constant temperature $(23\pm2^{\circ}C)$ and humidity (45–50%), and a 12h/12h light/dark cycle. These animals were housed at the Experimental Animal Management Center of People's Hospital of Ningxiang (Ningxiang, China), Hunan University of Chinese Medicine. This study was approved by the Ethics Committee of People's Hospital of Ningxiang (Ningxiang, China),

China), Hunan University of Chinese Medicine (HUCM-0018). All animal care and experimental procedures were in accordance with the National Institute of Health's Guide for the Care and Use of Laboratory Animals.

Instruments and reagents

We used the following instruments and reagents: Estradiol (E2) enzyme-linked immunosorbent assay (ELISA) Kit (Shanghai Bangyi Biotechnology Co., Ltd.), rat osteoprotectin (OPG) ELISA test kit, rat β -catenin ELISA kit, rat nuclear factor κB receptor activating factor ligand (RANKL) ELISA test kit (Suzhou Calvin Biotechnology Co., Ltd), TRIZOL reagent (Invitrogen, Inc.), Reverse Transcription System, Go Tag Green Master Mix (Promega, Inc.), liquid nitrogen (Changsha Rizhen Gas Co., Ltd.), Estradiol valerate (Bayer Healthcare Co., Ltd. Guangzhou Branch), chloroform, isopropanol (Sinopharm Group Chemical Reagent Co., Ltd.), icariin (content greater than 98%) (Shanghe Biotechnology Co., Ltd.), Eppendorf BioPhotometer Plus Nucleic Acid Protein Analyzer (Eppendorf Inc.); InfiniteF50 microplate reader (Tecan Inc.), gel imaging analysis system (Bio-Rad, Inc.), Horizontal electrophoresis apparatus (Beijing Liuyi Biotechnology Co., Ltd.), Electronic balance (Ohaus Instrument Co., Ltd.), Enzyme-labeled instrument (MB53Q) (Shenzhen Huisong Technology Development Co., Ltd.), and low-temperature high-speed centrifuge (1-14VK) (Sigma Inc.).

Animal modeling, grouping, and intervention

The PMOP pathological model of adult female rats with ovariectomy for 3 months was used in this study.

Modeling method

Rats were anesthetized with 2% sodium pentobarbital (0.2 mL/100 g body weight), and under sterile conditions, the skin and both sides of the muscles were longitudinally excised at a distance of 1 cm from the outside of the thoracolumbar vertebrae of the rats, and bilateral ovaries were removed. The wound was sutured in 2 layers and the blood stains were scrubbed with saline. Penicillin sodium (40 000 units each) was intramuscularly injected for 3 consecutive days after the operation, and the sutures were removed 5 days after the operation. The animals were kept in a clean environment at room temperature of 23-25°C and relative humidity of 40~60%. In the sham operation group, we only removed the corresponding volume of fat around the ovary.

Animal grouping

The rats were randomly divided into 4 groups: the sham operation group, the model group, the icariin group [150 mg/kg/d)], and the E2 group [0.167 mg/kg/d] according to body weight, with 12 rats in each group. All groups except the sham operation group were resected with bilateral ovaries to construct PMOP pathological models. Interventions started 1 week after surgery.

Intervention methods

Rats in the icariin group were orally administered icariin 150 mg/kg daily, and rats in the E2 group were orally administered E2 0.167 mg/kg daily. Rats in the sham operation group and the model group were intragastrically administered an equal volume of 0.9% sodium chloride daily. The intervention lasted 12 weeks.

Specimen collection

After the intervention, 8 female rats were randomly selected from each group for testing. All rats were anesthetized by intraperitoneal injection of 2% pentobarbital (0.2 mL/100g body weight), and blood was collected through the abdominal aorta. The blood sample was naturally coagulated at room temperature, and then the serum was separated by centrifugation (1000 g), and stored at -80° C for later use. The left femur of the rat was stripped, the muscles on the bone were removed, and the periosteum was retained for pathomorphological observation.

Bone mineral density (BMD) measurement

After taking blood from the abdominal aorta, the 2nd to 4th lumbar vertebrae of the rats were collected and the muscles on the bones were removed; the samples were wrapped with 0.9% sodium chloride gauze without paraformaldehyde and stored at -4°C. The samples were later sent to the Bone Density Test Room of People's Hospital of Ningxiang, Hunan University of Chinese Medicine to measure the BMD of the lumbar spine.

Determination of serum E2, $\beta\mbox{-}catenin,$ RANKL, and OPG levels

The serum E2, β -catenin, RANKL, and OPG levels were detected by enzyme-linked immunosorbent assay (ELISA). The serum samples were coagulated at room temperature for 20 min and centrifuged at 3000 rpm for 10 min. The supernatant was collected and stored at -20°C. Then, the standard was diluted and loaded. After that, blank wells and experimental wells were set, in which 40 μ L of sample dilution and 10 μ L of samples were added to the experimental wells. Subsequently, the enzyme-labeled coated plate was incubated at 37°C for 30 min. After washing the plate, 50 μ L of enzyme-labeled reagent was added to each well (except the blank well) and incubated at 37°C for 30 min. After washing the plate again, 50 μ L of A and B color reagents were added to each well and incubated at



Figure 2. Venn diagram of icariin potential targets and OP genes.

37°C in the dark for 15 min. Finally, a stop solution was added to each well to stop the reaction, and the optical density (OD) value was measured at 450 nm.

Statistical analysis

Statistical analysis was performed using SPSS 22.0 software. The experimental data were expressed as mean \pm SD (x \pm SD) and the data that conformed to normal distribution were analyzed by one-way ANOVA (for comparison of differences between groups) or *t* test (for comparison between 2 groups). *P* 0.05 was considered statistically significant.

Results

Icariin potential targets and OP genes

A total of 297 icariin potential targets and 262 OP genes were obtained. There are 22 identical targets between the 2 target sets (Figure 2). The PPI data of icariin targets and OP genes were collected from STRING 11.0 for sequence research.

Icariin-OP PPI network

The icariin targets, OP genes, and PPI data were input into Cytoscape 3.7.2 to construct an icariin-OP PPI network. This network is composed of 213 OP gene nodes, 272 icariin target nodes, 22 icariin-OP target nodes and 7823 edges (Figure 3). The nodes were sorted according to their degree, and the top 20 were: (1) Icariin targets: AKT1 (188 edges), EGFR (157 edges), MAPK1 (138 edges), MMP9 (137 edges), MAPK8 (123 edges), CASP3 (119 edges), HSP90AA1 (100 edges); (2) OP genes: INS (235 edges), IL6 (200 edges), TNF (170 edges), MAPK3 (150 edges), CTNNB1 (131 edges), JUN (131 edges), IL18 (117 edges), LEP (110 edges), SPP1 (106 edges); (3) Icariin-OP targets: ALB (227 edges), IGF1 (151 edges), SRC (136 edges), and ESR1 (129 edges).



Figure 3. Icariin-OP PPI Network (Pink circles stand for icariin targets. Blue circles stand for OP genes; Purple circles stand for Icariin-OP targets).

GO enrichment analysis results of icariin-OP PPI network

The targets and genes in icariin-OP PPI network were input into DAVID to undergo GO enrichment analysis and returned many biological processes, cell components, and molecular functions (Figure 4). The results of the GO enrichment analysis were further screened to obtain OP-related biological processes, cell components, and molecular functions. They were sorted according to their degree of enrichment (negative correlation with P value) and counts.

The top 10 OP-related biological processes were: steroid hormone-mediated signaling pathway, skeletal system development, positive regulation of cell proliferation, negative regulation of apoptotic process, ossification, extracellular matrix disassembly, collagen catabolic process, peptidyl-tyrosine autophosphorylation, response to glucocorticoid, and bone resorption (Figure 5, Supplementary Table 3)

The top 10 OP-related cell components were: extracellular space, extracellular region, extracellular exosome, cytosol, extracellular matrix, proteinaceous extracellular matrix, extrinsic component of the cytoplasmic side of plasma membrane, receptor complex, membrane raft, and endoplasmic reticulum lumen (Figure 6, Supplementary Table 4).

The top 10 OP-related molecular function were: steroid hormone receptor activity, RNA polymerase II transcription factor activity, ligand-activated sequence-specific DNA binding, protein homodimerization activity, growth factor activity, identical protein binding, serine-type endopeptidase activity, protein tyrosine kinase activity, receptor binding, cytokine activity, and steroid binding (Figure 7, Supplementary Table 5).



Figure 4. The results of GO enrichment analysis.



Figure 5. Bubble chart of biological processes (X-axis stand for fold enrichment).

KEGG pathways enrichment analysis results of icariin-OP PPI network

The targets and genes in the icariin-OP PPI network were input into DAVID to undergo KEGG pathway enrichment analysis and returned many signaling pathways (Figure 8). The results of KEGG pathway enrichment analysis were further screened to obtain OP-related signaling pathways. They were sorted according to their degree of enrichment (negative correlation with P value) and counts. The top 10 signaling pathways were: Osteoclast differentiation, FoxO signaling pathway, Estrogen signaling pathway, PPAR signaling pathway, Ovarian steroidogenesis, PI3K-Akt signaling pathway, T cell receptor signaling pathway, TNF signaling pathway, Insulin resistance, and MAPK signaling pathway (Figure 9, Supplementary Table 6). The role of each target in the Wnt and NF- κ B signaling pathway is shown in Figure 10. Based on the prediction results of network pharmacology, we found that icariin can regulate multiple signaling pathways through core targets, thereby directly and indirectly regulating bone metabolism to exert an anti-OP effect. Next, the prediction results were further verified through animal experiments.

Effect of icariin on BMD

The difference in lumbar BMD between the sham operation group and the model group was statistically significant (*P*<0.05), indicating that the animal pathological model of PMOP was successful. Compared with the model group, the BMD of the icariin group increased (*P*<0.05), but compared with the estrogen group, there was no statistically significant difference (*P*> 0.05) (Figure 11).



Figure 6. Bubble chart of cell components (X-axis stand for fold enrichment).



Figure 7. Bubble chart of molecular function (X-axis stand for fold enrichment).





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Figure 9. Bubble chart of signaling pathway (X-axis stand for fold enrichment).

Effect of icariin on morphology of left femur bone in rats

In the sham operation group, the trabecular bones were evenly distributed, and the tibial trabecular bone density and trabecular bone area/medullary cavity area ratio were normal. In the model group, the trabecular bone was sparse and damaged, the bone marrow cavity was enlarged, and the trabecular bone density and the trabecular bone area/medullary cavity area ratio were significantly decreased. In the icariin group, the trabecular bone distribution was relatively uniform, and the trabecular bone density was roughly restored (Figure 12).

Effect of icariin on serum E2 and $\beta\mbox{-}catenin$ level and RANKL/OPG ratio

Compared with the model group, the serum E2 levels of the rats in the sham operation group, icariin group, and E2 group were all increased, and the differences were statistically significant (P<0.05).

Compared with the sham operation group, the serum β -catenin level in the model group was significantly decreased (*P*<0.01). Compared with the model group, the serum β -catenin level in the E2 group was significantly increased (*P*<0.01), and the serum β -catenin level in the icariin group was also increased (*P*<0.05) (Figures 13, 14).

Compared with the sham operation group, the serum RANKL/OPG ratio in the model group was significantly increased (P<0.01). Compared with the model group, the serum RANKL/OPG ratio of the icariin group and the E2 group was significantly decreased (P<0.01) (Figures 15, 16).

Discussion

This research constructed and analyzed the icariin-OP PPI network. The results showed that the icariin potential targets can interact with many OP genes in the OP disease target network. Targets such as ALB, IGF1, SRC, and ESR1 are at the center of the network; these targets play a crucial role in the OP's pathological process. For example, IGF-1 plays a key role in the balance between bone resorption and bone formation. During bone resorption, osteoclasts produce IGF-1 and stimulate new bone formation [22]. In addition, IGF-1 also inhibits osteoblast apoptosis [23]. During bone formation, IGF-1 and IGFBP synthesis and secretion increase. IGF-1 binds to osteoblast surface receptors to increase collagen synthesis and inhibit collagenase production in osteoclasts. IGF-1 can also acidify osteoblasts and bone matrix, which is beneficial to bone mineralization [24,25]. ESR1, a type of ligand-activated nuclear transcription factor, mediates most of the estrogen response and is a nuclear macromolecule that mediates the



Figure 10. Signaling pathways [A: Wnt signaling pathways adapted from KEGG (hsa04310); B: NF-κB signaling pathways adapted from KEGG (hsa04064). The icariin targets were marked in red; the OP genes were marked in yellow; the Icariin-OP target was marked in orange].



Figure 11. Effect of Icariin on BMD (After 12 weeks of drug intervention. n=8, * Compared with model group, P<0.05).</p>

biological role of estrogen [26]. Estrogen is an important factor in increasing bone density and preventing bone loss after menopause. ESR was found on the surface of both human osteoblasts and osteoclasts, which indicates that estrogen plays a direct regulatory role on bone cells [27,28]. From the results of GO enrichment, we found that icariin mainly regulates the production of estrogen, steroid hormones and their signaling pathways, bone development, proliferation and apoptosis, ossification, and bone resorption. The above biological processes play an important role in the physiological processes of ossification and bone resorption. The core signaling pathways are FoxO, estrogen, PPAR, PI3K-Akt, Wnt, NF-κB, T cell receptor, TNF signaling pathway, and ovarian steroid production. These pathways directly and indirectly affect bone metabolism. The FoxO signaling pathway mainly regulates oxidative stress during osteoporosis. Inhibiting the transcription of FoxO activated by oxidative stress through anti-oxidative stress and positively regulating the Wnt signaling pathway can improve the osteogenic differentiation and bone formation of bone tissue [29,30]. It can also reduce high conversion levels of bone metabolism and reduce bone resorption [29,30]. Estrogen is a steroid hormone produced by the endocrine system. It is involved in the physiological and pathological processes of human bones and plays this important role in bone reconstruction [31]. The estrogen drugs currently used for OP include exogenous estrogen, phytoestrogens (soy isoflavones), and selective estrogen



Figure 12. Effect of icariin on morphology of left femur bone in rats (×400, HE staining. After 12 weeks of drug intervention. A: Sham operation group; B: model group; C: E2 group; D: Icariin group. Black arrow indicates trabecular bone; the red arrow indicates the marrow cavity.)



Figure 13. Effect of icariin on serum E2 level (After 12 weeks of drug intervention. n=8, * Compared with model group, *P*<0.05).



Figure 14. Effect of icariin on serum β-catenin level (After 12 weeks of drug intervention. n=8, * Compared with model group, P<0.05).</p>



Figure 15. Effect of Icariin on RANKL and OPG level (A: the level of RANK; B: the level of OPG. After 12 weeks of drug intervention. n=8, * Compared with model group, P<0.05).



Figure 16. Effect of Icariin on RANKL/OPG ratio (After 12 weeks of drug intervention. n=8, * Compared with model group, *P*<0.05).

receptor modulators [32,33]. Recent research indicates that the PI3K/Akt signaling pathway inhibits the progression of OP by promoting osteoblast proliferation, differentiation, and bone formation [34]. Wu et al. found that the mechanism of Si-Wu-Tang, a TCM formula used in treating OP, is to activate the PI3K/ Akt and NF-KB signaling pathways, increases the expression of ALP, BMP-2, and OPN genes, promote osteoblast differentiation, and increases bone formation [35]. In addition, PI3K and downstream signals RANKL, c-Fms, and αvβ3 integrin play important roles in the processes of osteoclast survival, differentiation, and bone resorption [36]. The Wnt/ β -catenin signaling pathway, as an important regulator of bone health and bone disease, can change bone mass through multiple pathways and is closely involved in OP [37,38]. The RANKL/RANK/OPG/NF-κB signaling pathway mainly regulates the differentiation and absorption activity of osteoclasts and participates in physiological and pathological bone reconstruction [39]. RANKL binds to the Receptor Activator of Nuclear Factor-κ B (RANK) receptor on the surface of osteoclasts, allowing TRAF-6 to aggregate in osteoclasts and OCPs, activating IKK/NF-κB, JNK, AP-1, c-Myc, c-Fos, and NFATc1, thereby promoting the differentiation and function of osteoclasts

[39,40]. In addition, TGF-beta, BMP, Notch, Hedgehog, MAPK, and other signaling pathways also play important regulatory roles in the occurrence and development of OP [41–43]. These signaling pathways are intertwined with each other, and together influence the occurrence and development of OP.

Icariin, a flavonoid, is an effective pharmacological component extracted from Epimrdii herba; it has estrogen-like activity, can enhance the activity of osteoblasts, and inhibits the proliferation and differentiation of osteoclasts [44,45]. Recent studies have shown that estrogen receptor (ER) plays an important role in regulate of bone metabolism by estrogen. ER is a glycoprotein that can specifically bind to estrogen. It is located in the cytoplasm and nucleus, and has the characteristics of strong specificity and high affinity [46,47]. Estrogen directly stimulates osteoblasts and inhibits the function of osteoclasts by activating $ER\beta$, thereby regulating bone formation, bone resorption, and bone rebuilding rate during the bone rebuilding cycle [46-49]. The present experimental study found that after ovariectomy, the serum E2 level in the model group was significantly lower than in the sham operation group. Then, the serum E2 levels of each drug treatment group increased, suggesting that estrogen levels increased after icariin treatment. Icariin may play a role in regulating bone metabolism by inhibiting osteoclast differentiation through this effect, thereby preventing the occurrence of PMOP. In addition, recent pharmacokinetic studies have shown that icariin's prototypes and metabolites in animals and humans are safe [50,51].

The classic signaling pathway Wnt/ β -catenin also plays an important role in bone. It increases bone mass in a variety of ways, including renewing stem cells, stimulating pre-osteoblastic replication, inducing osteoblastogenesis, and inhibiting osteoblasts and osteoblast apoptosis [52,53]. The key molecule of the Wnt/ β -catenin signaling pathway is β -catenin, which is closely involved in the development of bones and is a molecule necessary for osteoblasts to complete the differentiation

process [53]. The present study showed that, compared with the sham operation group, the serum β -catenin level in the model group was significantly lower. After treatment with icariin, rat serum β -catenin levels were higher than in the model group, suggesting that icariin activates the Wnt/ β -catenin signaling pathway through this effect and plays a role in treating PMOP.

The RANKL/RANK/OPG system is an important signaling pathway that was recently discovered; it plays a role in bone reconstruction and osteoclast differentiation [54]. RANK is an agonist that can bind to RANKL on the surface of osteoclast precursor cells. When RANK is activated, osteoclasts begin to differentiate and mature, and an abundance of activated osteoclasts causes osteolytic bone resorption [55]. OPG is a soluble inhibitory protein and a competitive inhibitor of RANKL; it inhibits the differentiation and maturation of osteoclast precursor cells by competing with RANKL. The expression levels of RANKL and OPG reflect the degree of bone resorption [56]. Generally speaking, when the ratio of RANKL/OPG increases, the number and activity of osteoclasts will increase; when the RANKL/OPG ratio decreases, the number and activity of osteoclasts will decrease. The RANKL/OPG ratio is the decisive factor for the final effect of the RANKL/RANK/OPG system [57]. The present study shows that, compared with the sham operation group, the RANKL/OPG ratio of the model group rats increased significantly. After icariin treatment, the RANKL/OPG ratio decreased more obviously, indicating that icariin can affect the RANKL/RANK/OPG system, and thus can play a role in the treatment of PMOP. In addition to RANKL and OPG, there are other signaling molecules related to OP, such as CTX-1, ALP, TRACP 5b, and PINP [58-60]. In the future, our group plans to further study these OP-related signaling molecules and pathways.

The strength of the present study is that it elucidates the regulatory effect of icariin on the OP biomolecular network, and discovered OP-related targets, biological modules (i.e., clusters), and signaling pathways. Among them, most of the biological

Supplementary Data

Supplementary Table 1. Compound targets of icariin. Supplementary Table 2. OP genes. Supplementary Table 3. Biological processes of icariin-OP PPI network. Supplementary Table 4. Cell components of icariin-OP PPI network. Supplementary Table 5. Molecular functions of icariin-OP PPI network. Supplementary Table 6. Pathway of icariin-OP PPI network.

Supplementary/raw data available from the corresponding author on request.

modules and signal pathways (such as the regulation of osteoblasts and osteoclasts) have been verified by previous studies; therefore, we chose a new direction that has not yet been studied to verify the prediction results, assessing Wnt pathway and RNKL pathway expression in serum. The limitation of this study is that the prediction results were mainly verified by animal experiments, and relevant clinical trials have not yet been carried out, so there is a lack of support from clinical evidence.

We constructed a molecular network model of icariin intervening in OP through network pharmacology strategies, and found that icariin may play a therapeutic role through osteoclast differentiation, FoxO, Wnt, Estrogen, PPAR, PI3K-Akt, NF- κ B, and T cell receptor signaling pathway. Then, we selected the RANKL/RANK/OPG system and Wnt signaling pathway for experimental verification. The experiment revealed that icariin can treat PMOP by regulating serum E2 levels, Wnt/ β -catenin signaling pathway, and RANKL/RANK/OPG system at the level of bone metabolism. Recent research also shows the expression of these 2 pathways in bone tissue after icariin intervention [61]. This discovery provides a more in-depth interpretation of the pathogenesis of PMOP and provides a new theoretical basis for the clinical application of icariin.

Conclusions

This research constructed and analyzed an icariin-OP PPI network and found that icariin can regulate OP-related biological processes, cell components, molecular functions, and signaling pathways. Experiments also showed that it can affect PMOP rats by regulating estrogen, Wnt/ β -catenin, and RANKL/ RANK/OPG signaling pathways.

Conflict of interest

None.

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