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ORIGINAL RESEARCH

Network Pharmacology and Experimental Validation of the Therapeutic Effect of Baji Capsule on LPS-Induced Osteoporosis

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Purpose: Osteoporosis is a common skeletal disease characterized by impaired bone microarchitecture, decreased bone mineral density and increased bone fragility, leading to a heavy physical and economic burden due to its greatly increased risk of fracture. Baji capsule is a proprietary medicine that can treat menstrual disorders and postmenopausal osteoporosis. However, the efficacy of Baji capsule has not been reported for osteoporosis caused by oxidative stress and inflammation. Therefore, the aim of this study was to evaluate whether Baji capsule has a therapeutic effect on lipopolysaccharide (LPS)-induced inflammatory osteoporosis and to explore the underlying mechanisms through network pharmacology.

Patients and Methods: Osteoporosis model in ICR mice induced with LPS. Mice were treated with vitamin E (100mg/kg), PBS, high-dose Baji capsule (810mg/kg) and low-dose Baji capsule (90mg/kg), respectively. The therapeutic effect of Baji capsule was evaluated by high-resolution micro-computed tomography (Micro-CT) and tissue section staining, serum inflammatory factor levels were assessed by ELISA, serum oxidative stress-related marker levels were determined by kits, and finally the mechanism was explored by network pharmacology and then verified by immunohistochemistry.

Results: Micro-CT results showed that Baji capsule attenuated LPS-induced inflammatory bone loss.Baji capsule also reduced serum inflammatory factor levels and oxygen free radical production. Target screening by network pharmacology yielded a total of 236 active ingredients of Baji capsule, as well as 278 common targets after taking the intersection of Baji capsule active ingredient targets and osteoporosis disease-related targets.

Conclusion: Baji capsule can treat osteoporosis by reducing inflammation and oxidative stress. The therapeutic effects of Baji capsule were shown to be multi-targeted and multi-pathway through network pharmacology. In the future, its anti-inflammatory and antioxidant properties can be utilized to further explore its therapeutic effects on inflammatory diseases, as well as a prospective study for the clinical treatment of osteoporosis.

Keywords: inflammatory, oxidative stress, osteoporosis, traditional Chinese medicine, network pharmacology

Introduction

Osteoporosis is a chronic metabolic disease characterized by impaired bone microarchitecture, decreased bone mineral density and increased bone fragility.¹ The prevalence of osteoporosis is high, affecting approximately 75 million people worldwide, including 10 million women.² Each year, the US healthcare system spends roughly \$7 billion on osteoporotic fractures, and by 2040, the annual cost is projected to approach \$50 billion. And this economic burden is similar across the globe.³

Currently, the clinically drugs used to treat and prevent osteoporosis are bisphosphonates, parathyroid hormone analogs, and selective estrogen receptor modulators. However, in addition to different degrees of side effects,^{4–6} these drugs are more effective in osteoporosis caused by estrogen aging, deficiency, and reduced mechanical weight-bearing, and lack specificity for inflammation and oxidative stress-induced osteoporosis. LPS is a component of the outer

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Traditional Chinese medicine (TCM) has been widely used because of its long history, small side effects, most of the raw materials from natural plants and other wonderful characteristics, and now there are a number of Chinese medicines with Chinese medicine extracts have been proved to be able to treat osteoporosis, and has shown good efficacy.^{14,15} Baji capsule is a proprietary Chinese medicine containing 16 natural herbs: Radix Morindae Officinalis, Polygonum multiflora, Cortex Eucommiae, Herba Cistanches,Dipsaci Radix, Curculiginis Rhizome, Fructus Rosae Laevigatae, Herba Epimedii, Rubus idaeus Linn, Radix Angelicae Sinensis,Codonopsis Radix, Radix Rehmanniae Praeparatae, Lycii Fructus, Radix Glycyrrhizae, Radix Astragali, Rhizoma Cibotii. Among them, the Radix Morindae Officinalis, Cortex Eucommiae, Herba Epimedii,Curculiginis Rhizome, etc. have been confirmed in the literature to be able to reduce the level of serum inflammatory factors and regulate the balance between osteogenesis and osteoblastogenesis thereby treating osteoporosis, ^{16–20} and there are also reports on the therapeutic effect of Baji capsule on the rats model of ovariectomized osteoporosis, but the therapeutic mechanism, active ingredients and molecular targets have not been elucidated.²¹

Network pharmacology can exhaustively elucidate the molecular mechanisms of TCM in the treatment of various diseases through its key concept of "network target, multi-component" model.^{22,23} So far, the therapeutic effects of various Chinese medicinal preparations on osteoporosis have been demonstrated by means of network pharmacology, such as Zhuanggu Busui Formula,²⁴ Liuwei Dihuang pill²⁵ and Zhuangguguanjie Formulation.²⁶ Therefore, we also exploited network pharmacology to investigate the route, target, and active ingredient of Baji capsule for its anti-osteoporotic effects.

Materials and Methods

Animals

Animal experiments were supervised and approved by the Animal Ethics Committee of Tongji Medical College, Huazhong University of Science and Technology. All the methods were performed in accordance with relevant guidelines and regulations. All the methods were performed in accordance with NC3Rs ARRIVE guidelines. Twenty-five ICR mice (females, 8 weeks) were purchased from Wuhan Hualianke Biotechnology Co. Ltd (Hubei, China) and housed in the Animal Center of Liyuan Hospital, affiliated with Tongji Medical College, Huazhong University of Science and Technology, all of them were maintained in an environment with a 22°C and a 12-hour circadian rhythm.

Animal Models of Osteoporosis

LPS (derived from E. coli 055:B5) was purchased from Sigma-Aldrich, Baji capsule was purchased from Chongqing Xieran Pharmaceutical Co. Ltd, vitamin E,Tween 80 and 4% paraformaldehyde solution were purchased from Wuhan Hualianke Biotechnology Co. Ltd. The powder inside the Baji capsule was removed and dissolved in PBS to make Baji capsule solutions (BJT) of different concentrations. Twenty-five 8-week-old female ICR mice were randomly divided into three treatment groups, namely, the high-concentration BJT group (BJTH), the low-concentration BJT (BJTL) group and the vitamin E group (vitE); as well as the LPS-induced osteoporosis model (LPS) group and the untreated blank control group, with five mice in each group. LPS powder was dissolved in PBS solution and then injected intraperitoneally at a dose of 5 mg/kg into mice in the BJTH, BJTL, vitE and LPS groups on days 0 and 4, whereas the blank control group was injected intraperitoneally with PBS. High-concentration BJT (810 mg/kg), low-concentration BJT (90 mg/kg), and vitamin E (100 mg/kg, dissolved in 0.1% Tween 80) were administered by gavage to the BJTH, BJTL, and vitE groups, respectively, starting on day 0, and continued for 8 days; PBS was gavaged to both LPS and control groups. Gavage was performed at 0.1 mL /10 g body weight. At the end of modeling, all mice in this study were

anaesthetised with the inhalational anaesthetic isoflurane, and the mice were anaesthetised and died after ocular blood sampling and autopsy operations.

The collected femurs were fixed in 4% paraformaldehyde for three days and then were scanned and analyzed by micro-CT (SkyScan 1176, Medical sub-Center of Analytical and Testing Center, Huazhong University of Science and Technology, China) to evaluate the bone-trabecular microstructure. Bone volume as a percentage of single bone tissue (BV/TV), trabecular number (Tb.N)., trabecular thickness (Tb.Th)., and trabecular separation (Tb.Sp)., were obtained.

Assay for Serum Chemistry

Serum collected by centrifugation was measured by standard colorimetric methods using commercial kits for alkaline phosphatase (ALP), malondialdehyde (MDA) and catalase (CAT) (Nanjing Jiancheng Bioengineering Institute, China). Serum tumor necrosis factor (TNF- α) and interleukin 6 (IL-6) were measured by enzyme immunoassay using commercial kits (Wuhan Bioswamp Biotechnology Co., Ltd.) according to the manufacturer's instructions.

Histomorphometric Analysis

Mouse femurs were fixed with 4% paraformaldehyde in for three days at room temperature and decalcified using EDTA decalcification solution (Wuhan servicebio Biotechnology Co., Ltd.) for one month at room temperature, then the tissue samples were embedded in paraffin wax and sectioned in 5µm thick slides. The slides were processed for hematoxylin and eosin (H&E) and tartrate-resistant acid phosphatase (TRAP) staining. Under the microscope, with the femoral head in the centre, select areas in the subchondral region and near the centre point, and the number of bone traps and empty bone traps were counted in each region to obtain the empty bone trap rate. In addition, the number of osteoclasts stained red was counted in each region.

Network Pharmacology

BJT active compounds and targets were screened using the Traditional Chinese Medicine System Pharmacology Database and Analysis Platform (TCMSP, <u>https://tcmspw.com/tcmsp.php</u>). Oral bioavailability (OB \geq 20%) and drug-likeness (DL \geq 0.1) were used as the criteria for screening the active compounds. The chemical information repository PubChem (<u>https://pubchem.ncbi.nlm.nih.gov</u>) was used to find the Canonical SMILES of the screened active compounds,²⁷ and then the obtained Canonical SMILES were analyzed using SwissTargetPrediction (STP) (<u>http://www.swisstargetprediction.ch</u>) to screen the potential target genes of the active compounds.²⁸ Osteoporosis disease targets were obtained by searching the GeneCards (<u>https://www.genecards.org/</u>) and OMIM (<u>https://omim.org/</u>) databases with the keyword "osteoporosis".²⁹

The STRING database (<u>https://www.string-db.org/</u>) was utilized to screen the active ingredients of Baji capsule and the targets corresponding to osteoporosis diseases according to the criterion of combined score >0.90, and the chosen target proteins were restricted to Homo sapiens. Cytoscape is a platform for integrating molecular interaction network data.³⁰ The intersection of the screened targets was then input into Cytoscape (<u>https://cytoscape.org/</u>) to build a PPI network of drug and disease targets.

DAVID (<u>https://david.ncifcrf.gov/</u>) is a Web-based online database available as a bioinformatics resource for interpreting a large number of gene/protein functions.³¹ DAVID was used to implement gene ontology (GO) enrichment analysis and KEGG enrichment analysis of intersection targets. GO enrichment analysis was used to explain three biological processes including molecular function (MF), biological process (BP), and cellular component (CC).³² While KEGG enrichment analysis was used to systematically analyze gene functions.³³

Molecular Docking

The targets related to inflammation and oxidative stress were found in DAVID, and then the cross-targets with the top 20 CytoNCA scores in the PPI network of drug and disease targets were cross-targeted with the targets related to inflammation and oxidative stress to obtain the key targets, ie, STAT3 and AKT1. Prior to molecular docking, we validated the expression of key targets in bone tissue based on the Human Protein Atlas (HPA, <u>https://www.proteinatlas.org/</u>) to determine whether they were expressed in bone tissue. Next, using PyMoL-2.6.0 and AutoDockTools-1.5.7 software, classical molecular dynamics

was used to analyze the binding sites, binding affinities, and interactions between STAT3 and AKT1 and the active ingredients of Baji capsule with CytoNCA scores ranked among the top 35 active ingredients. Crystal structures of key intersecting proteins were obtained from the RCSB Protein Data Bank (PDB, <u>http://www.pdb.org/</u>).³⁴ Mol2 format files of Baji cansule active compounds were downloaded through the TCMSP database. The co-crystallized ligands isolated from the receptors and the active pockets of each key intersecting proteins were probed using AutoDockTools-1.5.6. The molecular docking of active compounds to key targets and the determination of their free binding energies were realized by aotodock. PyMOL and was used for the interaction and binding of active compounds mode visualization and analysis.

Immunohistochemical Analysis

Femurs were fixed, decalcified, wax-packed, and sectioned and the slides were incubated at 37°C for 1 hour with anti-STAT3 and anti-AKT1 (proteintech. Wuhan Three Eagles Biotechnology Co., Ltd). The slides were incubated with HRPcoupled secondary antibody for 30 min and under microscopic observation, ImageJ software was employed to analyze the average optical density (AOD) at 100x magnification.

Statistical Analysis

All data were analyzed using SPSS software (version 23.0), and the results are expressed as mean \pm standard deviation. We used t-tests to analyze differences between two groups, and ANOVA to analyze differences between multiple groups. *P*-value <0.05 was considered statistically significant.

Results

Effect of Baji Capsule on Morphology and Parameters of Bone Tissue in Mice

Figure 1A and B show typical 3D images of the femur Micro-CT scans of rats in each experimental group. It can be seen that mice in the LPS group had a large number of cavities in the femoral diaphysis and poor continuity of trabecular. Compared with the mice in the control group, the mice in the LPS group had lower relative bone volume, BV/TV and the Tb.N decreased. In additional, the Tb.Th decreased, Tb.Sp increased. The aqueous solution of Baji capsule improve the bone microstructure changes of bone trabecular in osteoporotic mice and reversed BV/TV, Tb.N, Tb.Th and Tb.Sp. But the trends of Tb.Th and Tb.Sp was not significant and not statistically significant (Figure 1C-F, p<0.05; p>0.05; p>0.05). The BJTH group even had a more significant intervention effect than the vitE group, which was the standard medication group. The BJTH group showed a more significant improvement in bone volume, increasing the area covered by trabecular and the Tb.N.

Chemical Indicators for Serum Testing

Serum levels of the inflammatory factors TNF- α and IL-6 were significantly increased in the LPS group of mice compared with the control group (both p<0.001; Figure 2A–B). In contrast, BJTH, BJTL and vitE treatments all significantly inhibited the LPS-induced increase in serum TNF- α and IL-6 levels (both p<0.001; Figure 2A–B). When the endogenous system fails to provide sufficient antioxidant capacity, oxygen free radicals will lead to lipid peroxidation and MDA accumulation.³⁵ Therefore, MDA level can reflect the level of oxidation in the body.³⁶ Figure 2C shows that MDA levels in serum were higher in the LPS group than in the control group (p<0.01), whereas BJTH treatment significantly reduced MDA levels (p<0.05).Catalase (CAT) is an antioxidant enzyme whose activity reflects the body's antioxidant capacity.³⁷ LPS intraperitoneal injection decreased the level of CAT activity in serum (Figure 2D, p<0.001). BJTH, BJTL and vitE treatment all significantly increased the level of CAT activity in serum (Figure 2D, p<0.001). p<0.01, p<0.01). ALP is an incipient marker of osteoblast differentiation, and intraperitoneal injection of LPS significantly reduced serum ALP levels in mice (Figure 2E, p<0.01), while BJTH and BJTL treatments significantly reversed serum ALP levels (p<0.05, Figure 2E). Curiously, mouse serum ALP levels were not significantly different between the vitE-treated group and the LPS modeling group.

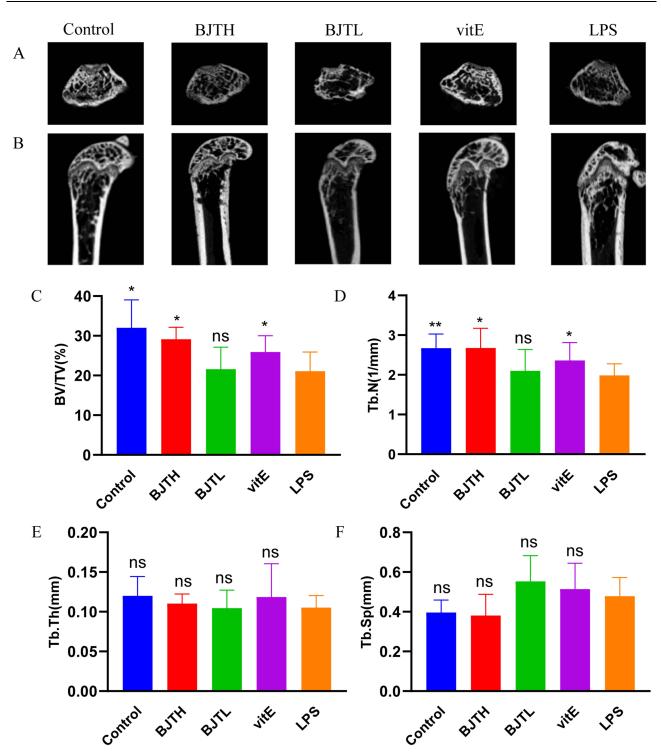


Figure I Baji capsule prevents LPS-induced microstructural destruction of femur in mice. (**A**). Micro-CT horizontal plane image of the distal femur. (**B**). Micro-CT coronal image of the distal femur. (**C**), (**D**), (**E**), and (**F**) graphs represent histogram depictions of the four trabecular parameters of BV/TV, and Tb.N, Tb.Th and Tb.Sp, respectively. Bar graphs are presented as mean \pm standard deviation (n=5), p<0.05*, p<0.01**, ns: not significant, compared with LPS group.

Morphologic Changes in Bone Tissue and Immunohistochemical Verification of Key Genes Expression in Bone Tissue

Histologic evidence of H&E staining demonstrated intact trabecular structure and morphology, clearly visible osteocytes within the trabecular, and fewer bone traps in the control group. In the LPS group, the number of trabecular in the femur

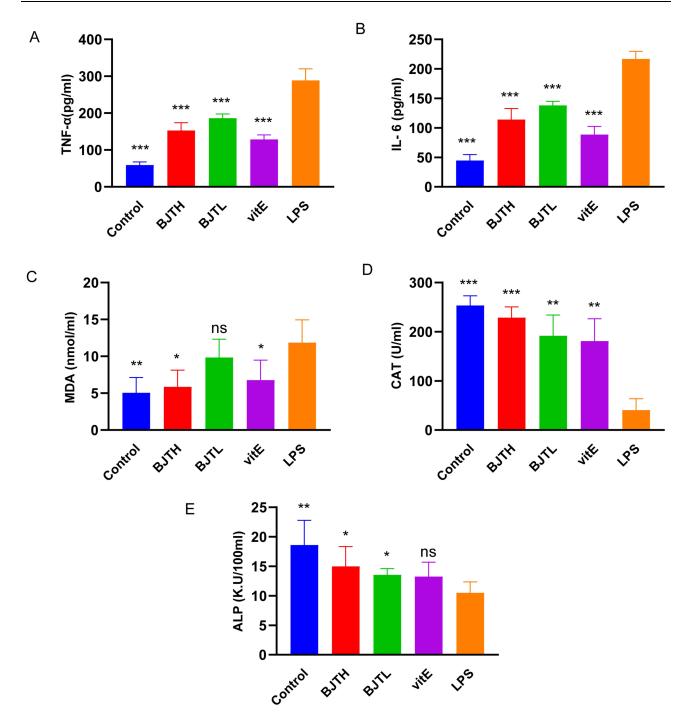


Figure 2 Serum levels of inflammatory cytokines, oxidative stress factors and bone formation markers in different groups. (**A**). TNF- α levels in serum. (**B**). IL-6 levels in serum. (**C**). MDA levels in serum. (**D**). CAT levels in serum. (**E**). ALP levels in serum. Bar graphs are presented as mean ± standard deviation (n=5), p<0.05*, p<0.01***, p<0.001****, ns: not significant, compared with LPS group.

was reduced and structurally disorganized, with a significant increase in empty bone trabecular. In the BJTH group, the pathologic changes in the trabecular structure were reduced, most osteocytes were normal, and there were not many empty bone sockets. The BJTL and vitE groups also reduced the pathologic changes in the bone tissue to different degrees, but to a lesser extent than the BJTH group (Figure 3A). Moreover, quantitative analysis of TRAP staining showed a significant increase in the number of osteoclasts in the LPS group compared to the control group (p<0.05) (Figure 3B). Then, we used immunohistochemical staining to verify the expression of STAT3 and AKT1. The staining sites of STAT3 and AKT1 (Figure 3C–D) were concentrated in the femoral neck and diaphysis. Immunohistochemical

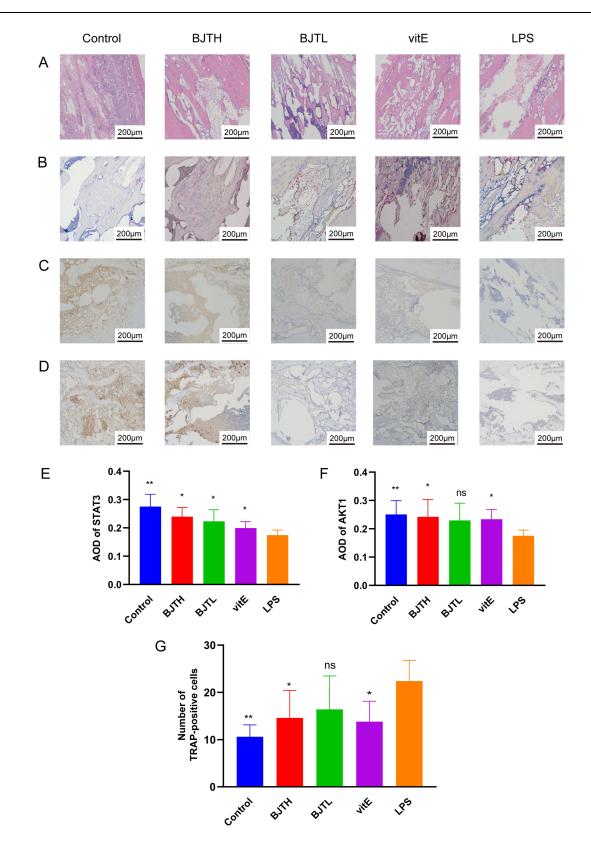


Figure 3 Morphologic changes and immunohistochemical in bone tissue. (A). H&E staining of mouse femur tissue sections (Images are at x100 magnification. Scale bars are 200 μ m). (B). Representative images of TRAP staining of mouse femur sections (Images are at x100 magnification. Scale bars are 200 μ m). (C) and (D) represent immunohistochemical staining of STAT3 and AKT1 in femoral tissue, respectively (Images are at x100 magnification. Scale bars are 200 μ m). (E) and (F) represent the mean optical density values for immunohistochemical analysis of STAT3 and AKT1, respectively. Data are shown as mean ± standard deviation (N=5, compared with LPS group, *p<0.05, **p<0.01, ns: not significant, independent samples t-test).(G). Quantitative statistics of osteoclasts per magnified 100x section. (Results are expressed as mean ± standard deviation (N=5, compared with LPS group, *p<0.05, **p<0.01, ns: not significant, independent samples t-test).

stained areas were quantified by comparing optical density values. As shown in Figure 3E and F, LPS intraperitoneal injection significantly inhibited the expression of AKT1 and STAT3 in mice, whereas vitE, BJTH, and BJTL all improved the inhibitory effect of LPS on the expression of AKT1 and STAT3 to varying degrees, of which the improvement effect of BJTH treatment was the most pronounced (p<0.05). However, BJTL treatment did not seem to prevent the inhibitory effect of LPS intraperitoneal injection on AKT1 expression. Treatment with BJTH significantly reduced the number of osteoclasts (p<0.05) (Figure 3 G).

Screening, Collecting Active Compounds and Building Compound-Disease-Target Networks

The obtained interpretation relationships and classifications of 16 herbal medicines, 145 active ingredients and 278 crosstargeting annotations were loaded into Cytoscape software, and the active ingredient target network of herbal medicines of Baji capsule for the treatment of osteoporosis was mapped (Figure 4). The targets corresponding to Baji capsule and osteoporosis respectively were obtained by STRING by setting the interaction score to the highest confidence level (0.9). Figure 4 shows the targets corresponding to the 145 active ingredients of Baji capsule and Figure 5 shows the targets corresponding to osteoporosis disease, and the Venn diagram was constructed by taking the intersection of the two quantities after counting them (Figure 6A). Then, the intersection targets were transferred into Cytoscape software to draw the PPI interaction diagram (Figure 6B). Finally, two PPI diagrams were plotted in order to clearly show the

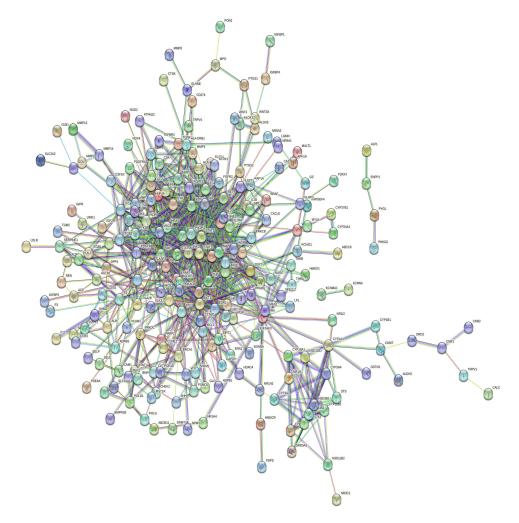


Figure 4 Targets corresponding to the active ingredients screened for Baji capsule. The 3D conformation of each target, target name, and interconnections are given in the figure, with different colors of the connecting lines representing different types of interactions.

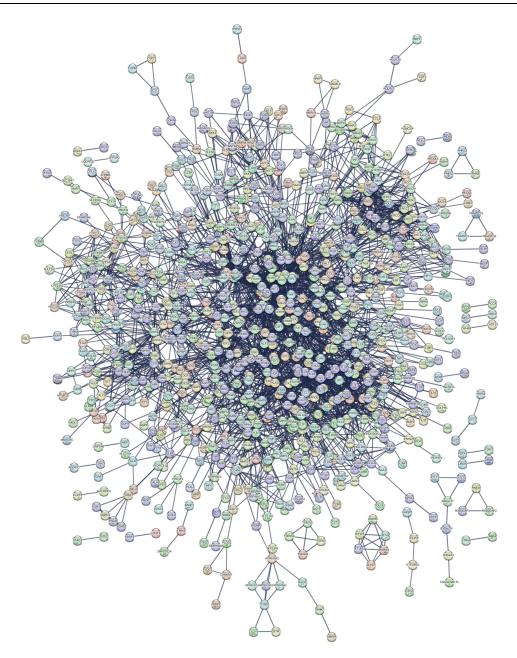


Figure 5 Screened targets corresponding to osteoporosis. Line color indicates the type of interaction evidence. The 3D conformation of each target, target name, and interconnections are given in the figure, with different colors of the connecting lines representing different types of interactions.

relationship between drug composition, active ingredients and target interactions. Figure 7 represents the active ingredients corresponding to the 16 constituent drugs of Baji capsule. The correlation between the 145 active ingredients and the protein targets is shown in Figure 8.

GO Enrichment Analysis and KEGG Pathway Analysis of Baji Capsule With Osteoporosis Intersection Targets

This partial intersection of targets with targets related to inflammation and oxidative stress was first taken one more time to obtain the core targets. We then performed GO enrichment analysis (p<0.05) on the core targets using the DAVID database and its extension package, and visualized the first 8 items enriched for BP, CC and MF (Figure 9A-C). The results indicate that BP is mainly related to the response to chemicals, organic substrates and oxygen-containing

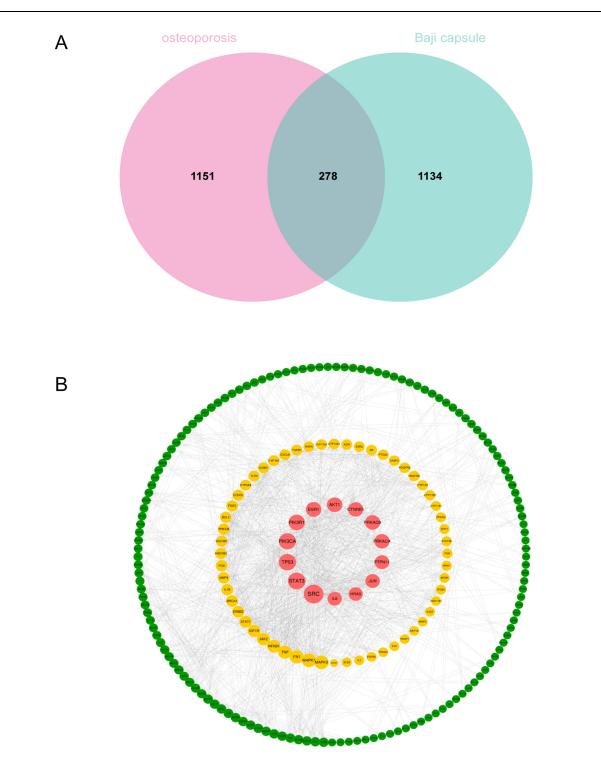


Figure 6 (A) Venn diagram showing cross-targeting of Baji capsule active ingredient targets and osteoporosis targets. (B) PPI network of drug and disease targets. Different colors and sizes indicate cytoNCA scores, and node size is proportional to cytoNCA score, the more inwardly aligned the node the higher the cytoNCA score.

compound, etc. CC mainly includes the plasma membrane, extracellular space and cell surface, etc. MF mainly includes enzyme binding, signaling receptor binding, identical protein binding, and molecular transducer activity, etc.

To further understand the potential pathways of Baji capsule for osteoporosis, we performed KEGG pathway enrichment analysis (p<0.05). And the top 8 pathways with the most statistically significant were visualized, including the pathway in cancer, proteoglycans in cancer, PI3K-Akt signaling pathway, and MAPK signaling pathway (Figure 9D).

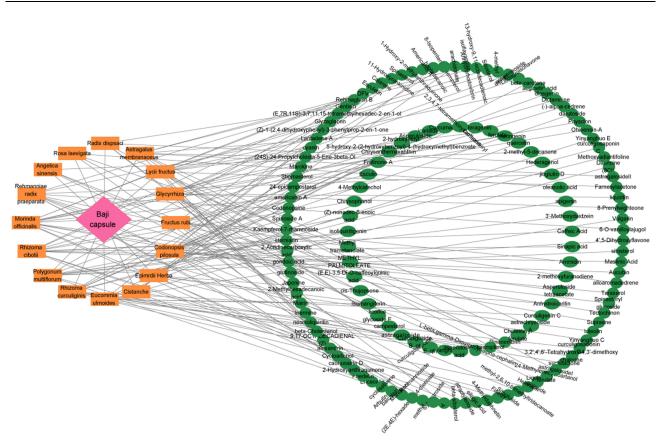


Figure 7 PPI network of Baji capsule drug composition and active ingredients. The Orange squares on the left indicate the 16 Chinese herbs contained in Baji capsule, and each green circle on the right indicates the 145 drug active ingredients corresponding to the Chinese herbs. Each type of Chinese medicine contains the active ingredients of the Baji capsule are connected by straight lines, the more straight lines are connected means that the Chinese medicine contains more active ingredients.Larger circles indicate a higher degree of association between the constituent drugs and the active ingredients.

The connectivity between the core targets and the active ingredients of Baji capsule was calculated and analyzed using cytoNCA in the Cytoscape software, and the two most highly correlated targets, STAT3 and AKT1, were identified.

Molecular Docking Validation

Two key targets (STAT3 and AKT1) screened in the PPI network analysis were molecularly docked with the top 35 active ingredients in the cytoNCA "degree" of the active ingredient target network. In order to analyse whether the key targets could play a role in osteoporosis, we first examined the expression of the key targets recorded in the HPA database before molecular docking and presented them as histograms. The results showed that STAT3 and AKT1 were moderately expressed in bone marrow (Figure 10A and B). The binding energy of \leq -5 kcal/mol was a criterion that an active ingredient could be docked to key targets.³⁸ Therefore, molecular docking was performed several times to screen out the active ingredients that could be docked to key targets. We describe the names and binding energies of the active ingredients screened for their alignment to key targets uniformly in Table 1. Finally, we visualized the seven pairs of active compounds that could bind to the key targets by PyMOL software. Among them, Chrysanthemaxanthin was docked to residue GLU-341 of AKT1 through hydrogen bonding; Helenalin was docked to residue GLN-43 of AKT1 through hydrogen bonding; Helenalin was docked to residue GLN-43 of AKT1 through hydrogen bonding; Helenalin was docked to residue GLN-43 of AKT1 through hydrogen bonding; Marckine was docked to residue ASP-323 and GLU-355 of AKT1 through hydrogen bonding (Figure 11A–D). In addition, Friedelin was docked to residue ASN-647 of STAT3 via hydrogen bonding; Frutinone A was docked to residue ASP-369, LYS-370 and THY-440 of STAT3 via hydrogen bonding; 2-Hydroxyanthraquinone was docked to residue TYR-640 of STAT3 via hydrogen bonding. (Figure 12A–C).

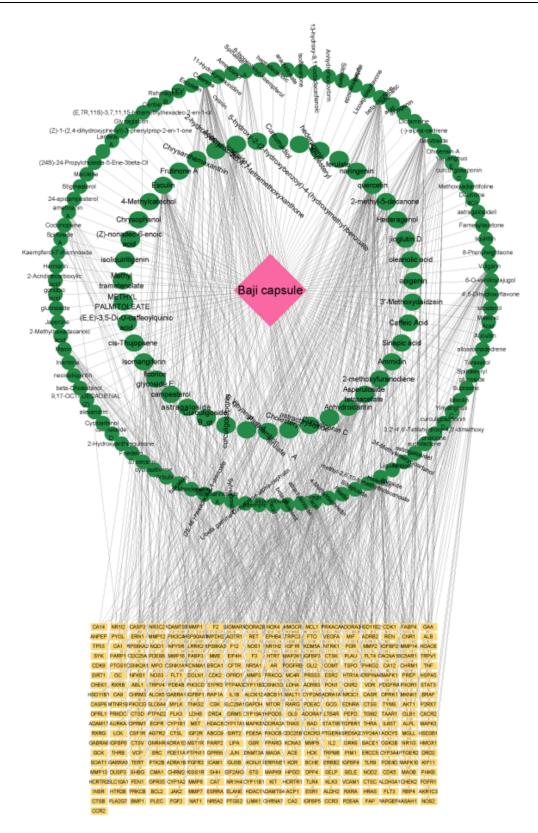


Figure 8 PPI network of Baji capsule active ingredients and corresponding targets. Larger nodes indicate stronger correspondence between the active ingredient and the target site. The green circle at the top of the figure indicates the drug active ingredient contained in Baji capsule, and the yellow box indicates the protein target corresponding to the drug active ingredient, and the protein target corresponding to each active ingredient is connected by a straight line.

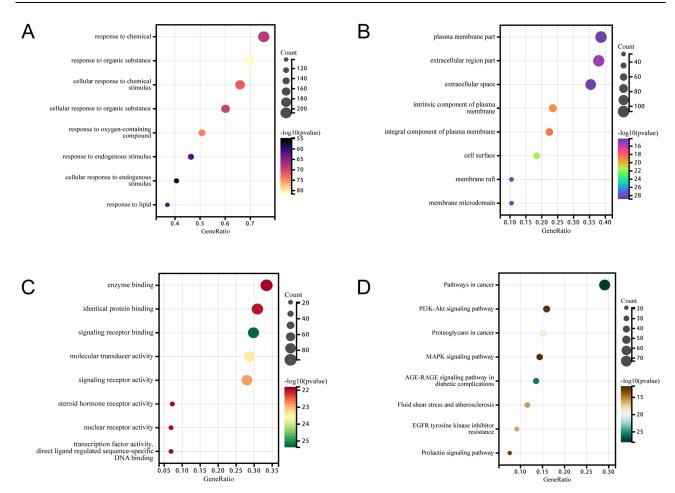


Figure 9 GO enrichment analysis and KEGG pathway enrichment analysis of core targets. (A). Biological Processes, (B). Cellular Components. (C). Molecular Function. (D). KEGG pathway enrichment result.

Discussion

LPS-induced inflammation and oxidative stress promote bone resorption and lead to bone loss in animals.^{39,40} Effective treatment for bacterial-induced bone destruction is limited to the use of antibiotics and surgery[18], but the resulting antibiotic resistance and surgical intolerance as well as the poor efficacy of traditional anti-osteoporotic medications have prompted the search for safer and more effective treatment modalities. In our experiments, we inferred that LPS-induced osteoporosis was associated with increased serum levels of pro-inflammatory factors and oxidative stress; we also demonstrated that Baji capsule counteracted LPS-induced bone loss by promoting STAT3 and AKT1 expression.

Bone tissue structure is an instrumental criterion for assessing bone health.⁴¹ In this research, bone micro-CT and bone histopathologic techniques were applied to assess bone structure. Our experimental results showed that both vitE and BJTH treatments improved the bone microstructure changes of bone trabecular in osteoporotic mice and reversed the alterations in parameters such as BV/TV, Tb.N. The results imply that Baji capsule improved LPS-induced bone microstructural damage and was effective in restoring bone morphologic parameters, but it must be instituted with a BJTH treatment, and BJTL treatment cannot achieve the desired results.

Oxidative stress and inflammation are drivers of bone loss.⁴² Many studies have illustrated that inflammatory cytokines such as TNF- α and IL-6 promote osteoclastogenesis and activation while inhibiting osteoblast proliferation.⁴³⁻⁴⁶ Excessive accumulation of reactive oxygen species (ROS) hinders osteoblast proliferation and differentiation; enhances osteoclast differentiation and ultimately leads to more bone resorption.^{47,48} Moreover, osteoblast death is induced in the presence of dramatically increased ROS levels, leading to structural bone damage and decreased bone density.⁴⁹ In this study, we found that compared to the LPS group, CAT and ALP activity levels were up-



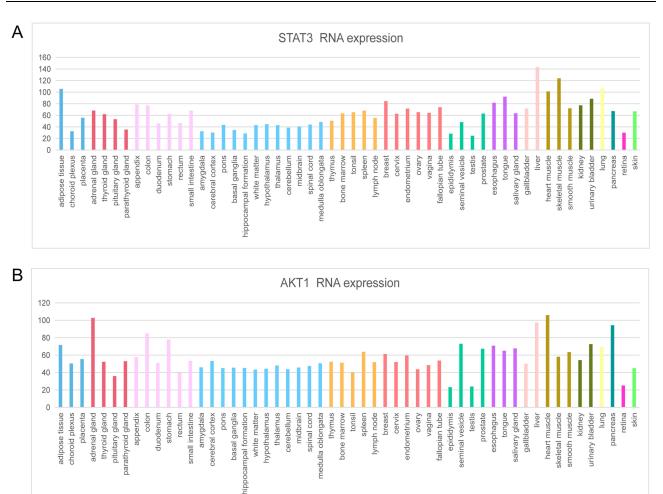


Figure 10 Exploring the expression of key targets. (A) Histogram of STAT3 expression in different tissues based on the HPA database. (B) Histogram of AKT1 expression in different tissues based on HPA database.

regulated in Baji capsule aqueous solution administration groups; MDA, IL-6 and TNF- α activity levels were decreased. This suggests that Baji capsule can inhibit inflammation and oxidative stress induced by intraperitoneal injection of LPS. Curiously, mouse serum ALP levels were not significantly different between the vitE-treated group and the LPS

NO	Molecule PubChem CID	Molecule Name	Key protein Modeling	cytoNCA (Betweenness)	Bind Energy (kcal/mol)
I	21160900	Chrysanthemaxanthin	AKTI 4EKL	42000.0	-6.07
2	5,318,332	II-Hydroxyrankinidine	AKTI 4EKL	44608.0	-6.09
3	23205	Helenalin	AKTI 3096	31,548.0	-5.16
4	72341	Marckine	AKTI 4EKL	21064.0	-5.89
5	91472	Friedelin	STAT3 6NJS	28930.0	-6.2
6	441965	Frutinone A	STAT3 6NJS	52420.0	-5.62
7	11,796	2-Hydroxyanthraquinone	STAT3 6NJS	96348.0	-5.18

Abbreviations: AOD, average optical density; ALP, alkaline phosphatase; AGE, Advanced glycation endproduct; BJT, Baji capsule solutions; BJTH, high concentration BJT; BJTL, low concentration BJT; BV/TV, bone volume as a percentage of single bone tissue; BP, biological process; BMSCs, bone marrow mesenchymal stem cells; CC, cellular component; CAT, catalase; COL-10, collagen type X; DL, drug-likeness; GO, gene ontology; HE, hematoxylin and eosin; HPA, Human Protein Atlas; IL-6, interleukin 6; iNOS, inducible nitric oxide synthase; LPS, Lipopolysaccharide; MDA, malondialdehyde; MF, molecular function; MSC, mesenchymal stromal stem cell; NF-κB, Nuclear factor-kappa B; Nrf2, nuclear factor erythroid type 2-associated factor; OB, Oral bioavailability; PDB, RCSB Protein Data Bank; ROS, reactive oxygen species; RANKL, nuclear factor kappa-B ligand; RAGE, AGE receptors RAGE; STP, SwissTargetPrediction; TCM, Traditional Chinese medicine; Tb.N, trabecular number; Tb.Th, trabecular thickness; Tb.Sp, trabecular separation; TNF- α , tumor necrosis factor; TRAP, tartrate-resistant acid phosphatase; TCMSP, Traditional Chinese Medicine System Pharmacology Database and Analysis Platform; vitE, vitamin E.

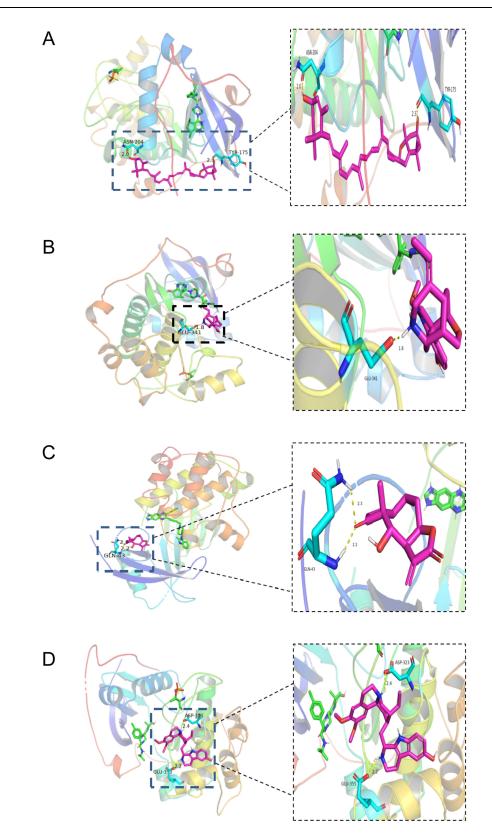


Figure 11 Molecular docking modeling diagram. (A) Molecular docking of AKT1 with Chrysanthemaxanthin. (B) Molecular docking of AKT1 with 11-Hydroxyrankinidine. (C) Molecular docking of AKT1 with Helenalin. (D) Molecular docking of AKT1 with Marckine. Purple molecules represent active ingredients that can be docked to key targets, blue molecules represent key target residues that dock to the active ingredient, and hydrogen bonds are indicated by yellow dashed lines.

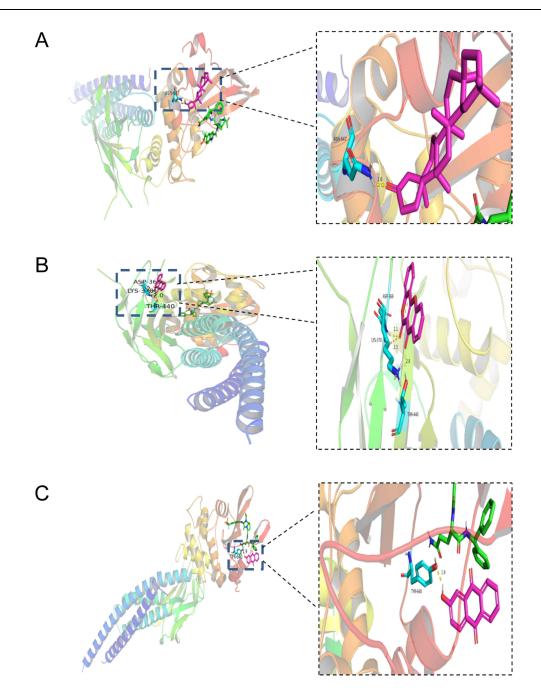


Figure 12 Molecular docking modeling diagram. (A) Molecular docking of STAT3 with Friedelin. (B) Molecular docking of STAT3 with Frutinone A. (C) Molecular docking of STAT3 with 2-Hydroxyanthraquinone. Purple molecules represent active ingredients that can dock to key targets, blue molecules stand for key target residues that dock to the active ingredient, and hydrogen bonds are indicated by yellow dashed lines.

modeling group, which may be caused by the toxicity of Tween 80, the organic solvent used to dissolve vitE. Furthermore, Intraperitoneal injection of LPS significantly increased the number of osteoclasts in bone tissue, whereas BJTH treatment inhibited the pro-osteoclastogenic effect of LPS. It suggests that at high concentrations of Baji capsule may treat osteoporosis by inhibiting osteoclast formation and promoting osteoclastogenesis.

The study demonstrated the anti-inflammatory, anti-oxidative stress and thus anti-osteoporotic effects of Baji capsule and explored the underlying mechanisms through network pharmacology. The active compounds that can interface with the key targets include: Friedelin, Frutinone A, 2-Hydroxyanthraquinone, Chrysanthemaxanthin, 11-Hydroxyrankinidine, Helenalin and Marckine. Friedelin, a triterpenoid derived from Codonopsis Radix, has demonstrated promising free radical scavenging, anti-inflammatory, and antioxidant properties in animal models.⁵⁰ In addition, friedelin induces the

expression of collagen type X (COL-10), a homotrimeric collagen associated with chondrocyte hypertrophy that promotes cartilage calcification, in mesenchymal stromal stem cell (MSC)[51]. Chondrogenic hypertrophy is a process by which chondrocytes in the central areas increase in size and become hypervascularized during calcification, a process that is accompanied by the differentiation of bone progenitor cells into osteoblasts and the replacement of cartilage by mineralized bone tissue.⁵¹ This means that friedelin has the ability to indirectly promote osteogenesis by promoting chondrocyte calcification. Frutinone A is a coumarin derived from Codonopsis Radix. Studies have demonstrated that coumarins exhibit many pharmacological characteristics such as antioxidant, anti-inflammatory and anti-tumor properties.⁵² Furthermore, some studies have shown that coumarin has an inhibitory effect on osteoclast activity.⁵³ and also promotes osteogenic differentiation of stem cells.⁵⁴ 2-hydroxyanthraquinone is an anthraquinone compound derived from Radix Morindae Officinalis. Although 2-hydroxyanthraquinone has not been successfully isolated from Radix Morindae Officinalis, anthraquinones are thought to have anti-inflammatory, anti-osteoporotic and other biological activities.⁵⁵ Chrysanthemaxanthin and 11-Hydroxyrankinidineis are also active constituents of Codonopsis Radix. Although its pharmacological properties have not been reported, Codonopsis Radix extracts have been demonstrated to have anti-inflammatory, antioxidant, and anti-osteoporotic effects in various studies.⁵⁶⁻⁵⁸ Nuclear factor-kappa B (NF- κB) is a transcription factor that controls cell proliferation and differentiation, and when NF- κB is activated and translocated to the nucleus, it can up-regulate the transcription of relevant genes involved in inducing osteoclast differentiation.^{59,60} Activation of nuclear factor erythroid type 2-associated factor (Nrf2) binds to and promotes the expression of promoters of certain antioxidant genes, thereby enhancing the body's antioxidant defense.⁶¹ Helenalin is a natural active ingredient derived from Cortex Eucommiae. Helenalin analogs have been reported to activate the Nrf2 pathway and specifically inhibit NF-KB activation.^{62,63} Marckine (Tubulosine) is an active component of the natural plant Herba Cistanches, belonging to the isoquinoline alkaloids group.⁶⁴ We know that revitalisation of the inducible nitric oxide synthase (iNOS) pathway facilitates inflammatory induced osteoporosis through repressing bone formation and contributing to osteoblast apoptosis.⁶⁵ And isoquinoline alkaloids can suppress iNOS expression by inhibiting NF-KB activation and thus may help to combat inflammation-induced osteoporosis.⁶⁵

Runt-related transcription factor 2 (Runx2) is a key transcription factor in the differentiation of bone marrow mesenchymal stem cells (BMSCs) into osteoblasts.⁶⁶ STAT3 is a key signaling protein that integrates a variety of cytokines, growth factors, and oncoproteins.⁶⁷ STAT3 is activated by phosphorylation and transported to the nucleus where it combines with the Runx2 promoter and augments its transcriptional capacity, thereby stimulating the differentiation of BMSCs into osteoblasts.⁶⁸ In addition, it has been reported that STAT3 deficiency attributes to skeletal defects including craniofacial deformities, osteoporosis and spontaneous fractures, and inhibits osteoblast differentiation.⁶⁹ In contrast, high expression promotes osteoblast formation and thus increases bone mass.^{70–72} Akt is an serine/threonine kinase that is a central transducer of the PI3K-Akt signaling pathway,⁷³ and its isoform, Akt1, is expressed in a variety of tissues.⁷⁴ Activation of AKT1 by phosphorylation activates a variety of downstream substrates, which inhibits apoptosis and promotes cell survival.⁷⁵ It has been shown that stimulating AKT1 expression can inhibit glucocorticoid-induced apoptosis of osteoblasts, thereby attenuating glucocorticoid-induced osteoporosis.⁷⁶ It is well known that macrophages are heavily polarised towards M1 during the early inflammatory phase, whereas macrophage M2 polarisation.⁷⁹ This can be deduced that AKT1 has a certain anti-inflammatory capacity by promoting macrophage M2 polarisation, thus inhibiting osteoporosis.

The results of KEGG enrichment analysis indicated that the active ingredients of Baji capsule may exert their curative effects on osteoporosis by modulating multiple pathways. The pathways were predominantly enriched in the pathway in cancer, proteoglycans in cancer, PI3K-Akt signaling pathway, MAPK signaling pathway and AGE-RAGE signaling pathway in diabetic complications, etc. It has been demonstrated that binding of the receptor for nuclear factor kappa-B (RANK) on the surface of osteoclast precursor cells to the receptor activator of nuclear factor kappa-B ligand (RANKL) present in stromal cells stimulates osteoclast differentiation and activation through activation of the downstream P13K/AKT pathway.^{80–82} Furthermore, activated PI3K/AKT signaling also promotes RANKL expression in osteoblasts and regulates the differentiation of mesenchymal stromal cells to osteoblasts.^{83,84} RANK binding to RANKL also activates the MAPK pathway, which regulates osteoclast differentiation and osteoclast precursor proliferation and apoptosis.⁸⁵

Moreover, activation of the MAPK pathway promotes osteogenic differentiation and matrix mineralization.^{86,87} Advanced glycation endproduct (AGE) and AGE receptors (RAGE) bind to each other to stimulate IL-6 production by osteoblasts, inhibit osteoblast phenotypic expression and bone matrix mineralization, and ultimately increase osteoclast resorption.^{88,89} Type I collagen promotes the progression of an osteoblast phenotype, but it has been shown that collagen osteoblasts modified by AGE are inhibited in function and bone nodule formation, ultimately affecting osteoblast differentiation.⁹⁰ Collectively, each of these pathways regulates bone homeostasis by modulating the balance between osteoclasts and osteoblasts, and when these pathways are inhibited or activated, an imbalance in bone homeostasis causes the development of osteoporosis.

As we expected, Baji capsule had a therapeutic effect on LPS-induced osteoporosis in mice and could improve the bone microstructural disorders caused by LPS. We found that Baji capsule declined serum IL-6, TNF- α , and MDA levels, implying that Baji capsule can modulate oxidative stress and inflammation-related pathways. In animal experiments, LPS mice were intraperitoneally injected to establish an osteoporotic mouse model, micro-CT was performed to detect bone microstructural changes in mice, serum inflammation, oxidative stress biomarkers and osteogenesis-related markers were detected, and the number of TRAP-positively stained cells in each mouse femur section was counted. Meanwhile, the underlying mechanism of Baji capsule against osteoporosis was explored by network pharmacology. Finally, the key targets predicted by network pharmacology were validated by immunohistochemistry. However, the therapeutic effects and extraction process of some active ingredients screened by network pharmacology were rarely reported and lacked some background support. There was also no correlation study along with in vitro experiments, which are essential to validate and explore more accurate pharmacological effects and mechanisms of Baji capsule.

Conclusion

Baji capsule had a therapeutic effect on LPS-induced osteoporosis in mice and could improve the bone microstructural disorders caused by LPS. Network pharmacological analysis indicated that Baji capsule exerts its therapeutic effects through multi-targets, multi-components, and multi-pathways. The key targets for osteoporosis treatment with Baji capsule may be STAT3 and AKT1, and the potential mechanism may be linked to cancer pathway the regulation of pathway in cancer, proteoglycans in cancer, PI3K-Akt signaling pathway, MAPK signaling pathway, the AGE-RAGE signaling pathway in diabetic complications, and so on. Our study provides basic evidence for osteoporosis treatment with Baji capsule and provides a new drug candidate for the treatment of osteoporosis.

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Disclosure

The authors report no conflicts of interest in this work.

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