

Kaempferol promotes bone formation in part via the mTOR signaling pathway

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Abstract. Previous research indicates that kaempferol (Kae) promotes osteogenesis, but its underlying mechanism of action remains unclear. The present study hypothesized that the osteogenic effects of Kae were mediated through mammalian target of rapamycin (mTOR). To validate this hypothesis, bone marrow mesenchymal stem cells (BMSCs) from ovariectomized (OVX) rats were differentiated into osteoblasts. The bone mineral density and bone microarchitecture of the OVX rats was measured *in vivo*, while osteogenesis was evaluated *in vitro* via Alizarin Red S staining and alkaline phosphatase activity measurements in cultured BMSCs. The levels of phosphorylated eukaryotic translation initiation factor 4E-binding protein 1 (p-4E/BP1) and phosphorylated ribosomal protein S6 kinase B1 (p-S6K), and the expression of Runt-related transcription factor 2 and Osterix, were concurrently quantified by western blot analysis. The data suggested that Kae prevented OVX-induced osteoporosis in rats by promoting osteoblastogenesis. Furthermore, treatment with Kae in rat BMSCs enhanced mineralization, elevated ALP activity, increased the expression levels of Runx-2 and Osterix and increased the levels of p-S6K and decreased the levels of p-4E/BP1 and,

consistent with its ability to promote osteoblast differentiation. In contrast, treatment with rapamycin, an mTOR inhibitor, produced the opposite phenotype. Taken together, these data suggested that the protective effects of Kae in BMSCs and in the OVX rat model resulted from the induction of osteogenesis via mTOR signaling, or at least partially via the regulation of downstream effectors of the mTOR pathway.

Introduction

Osteoporosis is described as a 'silent disease', characterized by gradual bone loss that occurs in the absence of other symptoms over a period of years (1,2). Due to its increasing prevalence, osteoporosis severely affects human health and quality of life (1). Osteoporosis is a metabolic disease that is common in postmenopausal women and the elderly (1). Phenotypically, it is characterized by bone loss, bone microstructural damage, bone fragility and increased susceptibility to fractures (1). The approved medications for the prevention and treatment of osteoporosis in clinical practice are frequently ineffective and cause adverse reactions (1-5). New therapeutic strategies are therefore urgently required.

Kaempferol (Kae; Fig. 1) is a type of natural flavonoid, extracted from the rhizome of *Kaempferia galangal* L. (6). As a natural flavonol, Kae is present in a variety of Chinese medicinal herbs, plants, fruits and beans (6), and is prized for its medicinal properties, which include anti-inflammatory (7) and antitumoral (8) effects, as well as being beneficial for the treatment of diabetes (9), cardiovascular disease (10) and osteoporosis (11). Kae has been previously confirmed to be beneficial to bone microarchitecture by increasing bone density and reversing osteoporosis in ovariectomized (OVX) rats. However, the precise mechanism(s) governing these effects have not been defined.

Mammalian target of rapamycin (mTOR) is a member of the phosphatidylinositol 3-kinase-related kinase family

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Abbreviations: Kae, kaempferol; BMSCs, bone mesenchymal stem cells; mTOR, mammalian target of rapamycin; OVX, ovariectomized; Rapa, rapamycin

Key words: Kae, osteoporosis, BMSCs, mTOR, Rapa

of protein kinases (12). mTOR functions through two structurally and functionally distinct multi-protein complexes, mTORC1 and mTORC2, which are involved in cell growth, proliferation, survival, protein synthesis, autophagy and transcription (13). mTOR is also an important regulator of bone metabolism known to promote osteoblastic differentiation and increase bone matrix synthesis (14,15). In addition, mTORC1 and mTORC2 have been implicated in the regulation of bone homeostasis (16-18).

Therefore, mTOR was hypothesized to be a novel target for the development of new and effective osteoporosis therapies. The aim of the present study was to investigate whether Kae was able to enhance the osteogenic differentiation and function of bone marrow mesenchymal stem cells (BMSCs) via mTOR activation.

Materials and methods

Reagents. Kae (purity>98%) was purchased from the National Institute for the Control of Pharmaceutical and Biological Products (Beijing, China). Kae was dissolved in DMSO and diluted to 0.01% in PBS. Rapamycin (Rapa), a specific inhibitor of mTOR, was purchased from Selleck Chemicals. The Alizarin Red S (ARS) staining buffer and alkaline phosphatase (ALP) detection kits were purchased from Nanjing Jiancheng Bioengineering Institute. Anti-runt-related transcription factor 2 (Runx2; cat. no. ab23981) and anti-Osterix (cat. no. ab22552) was purchased from Abcam. Anti-eukaryotic translation initiation factor 4E-binding protein 1 (4E/BP1; cat. no. 94525), anti-phosphorylated (p)-4E/BP 1 (cat. no. 2855) and anti-ribosomal protein S6 kinase B1 (S6K1; cat. no. 9202), anti-p-S6K1 (cat. no. 9204) was obtained from Cell Signaling Technologies, Inc. Horseradish peroxidase-labeled anti-immunoglobulin G secondary antibody (goat anti-mouse IgG; cat. no. SA00001-1; and goat anti-rabbit IgG; cat. no. SA00001-2) was obtained from ProteinTech Group, Inc. Anti- β -actin antibody (cat. no. KL002) was provided by Nanjing Jiancheng Bioengineering Institute.

Animals. A total of 30 adult (age, 6-8 weeks) female Sprague-Dawley (SD) rats weighing 180-220 g were obtained from the Nanchang University Laboratory Animal Center (Nanchang, China) and maintained under a 12-h dark/light cycle at 22-25°C and 40-70% humidity. Animals were allowed access to food and water *ad libitum*. Experiments were performed according to the Guide for the Care and Use of Laboratory (National Institutes of Health), and were approved by the Ethics Committee of Nanchang University (no. 2017-0122).

Experimental groups and treatments. A total of 30 rats were randomly divided into five groups (n=6 in each group) as follows: i) Sham group, in which the abdominal cavities of the rats were opened and fat tissue around the ovaries removed; ii) OVX group, in which the rats were ovariectomized; iii) Kae treatment group (OVX + Kae), in which the OVX rats were continuously given Kae (100 mg/kg/day) via gavage for 8 weeks; iv) Kae and Rapa administration group (OVX + Kae + Rapa), in which the rats were treated as in the Kae treatment group and in addition received intraperitoneal injections of Rapa (0.2 mg/kg/day) for 8 weeks; and v) Rapa

group (OVX + Rapa), in which the OVX rats received intraperitoneal injections of Rapa (0.2 mg/kg/day) for 8 weeks. The dose and timing of Kae and Rapa administration were ascertained in preliminary experiments. Briefly, the OVX rats were given Kae at 25, 50 and 100 mg/kg/day. Bone mineral density (BMD) was detected after 8 weeks to select the working concentration. Kae at 100 mg/kg/day was the most effective concentration in improving BMD in OVX rats. Rapa was administered at 0.2 mg/kg/day according to the manufacturer's protocol (Selleck Chemicals). Subsequent experiments were performed following the 8 weeks of treatment.

Assessment of BMD and bone microarchitecture. Following treatment, the rats were sacrificed and their right femurs and tibiae were dissected. The 2D total bone mineral content was used to calculate the BMD as previously described (19). The right femurs of the rats were analyzed using dual-energy X-ray absorptiometry with a Lunar Prodigy Advance system (version 13.6; GE Healthcare).

Based on the median values of total BMD, selected trabecular microarchitecture of the femoral metaphysis was evaluated using micro-computed tomography (micro-CT; Scanco Medical AG) with Scanco image processing language software (version 5.08b; Scanco Medical AG). Scans were performed from the proximal growth plate in the distal direction (18 μ m/slice) as the distal femur has a high concentration of trabecular bone compared with the proximal and middle regions of the femur. A volume of interest (VOI) was selected, defined as the cross-sectional area spanning 100 slices from the proximal growth plate. The 2D scans were used to produce 3D reconstructions of the bone microarchitecture, which were used to measure bone morphometric parameters of the selected VOI, including the bone volume fraction [bone volume/tissue volume (BV/TV)], trabecular number (Tb.N), trabecular separation (Tb.Sp), trabecular thickness (Tb.Th) and structure model index (SMI). The operator conducting the CT analysis was blinded to the treatments associated with the specimens. All examinations were conducted according to the principles and procedures described in the most recent National Research Council publication of the Guide for the Care and Use of Laboratory Animals and the ARRIVE guidelines (20).

Isolation of rat BMSCs. BMSCs were flushed from the femurs and tibiae of normal SD and OVX rats with PBS in a biosafety cabinet, using a 5 ml syringe fitted with a needle (21G). Mononucleated cells were isolated by density gradient centrifugation at 400 x g for 20 min at room temperature in rat lymphocyte separation medium (TBD Science) at a concentration of 1.091 g/ml. Isolated cells were cultured in low-glucose DMEM (HyClone; GE Healthcare Life Sciences) containing 15% fetal bovine serum (HyClone; GE Healthcare Life Sciences). All cells were maintained in a 37°C incubator within an atmosphere containing 5% CO₂. BMSCs were identified by CD44 and CD34 labeling (21). The culture medium was regularly replaced and cells were passaged after reaching 70-80% confluence. Third-generation BMSCs were harvested for subsequent experiments.

Cytotoxicity assays. The cytotoxicity of BMSCs was determined using an MTT assay. Briefly, rat BMSCs were plated

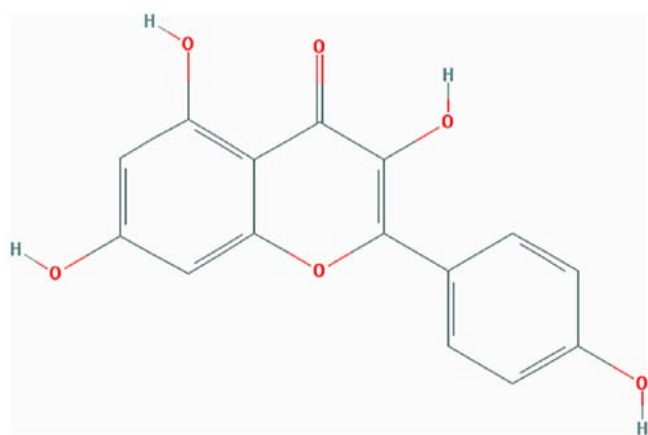


Figure 1. Chemical structure of kaempferol (Pubchem CID: 5280863).

into 96-well plates at a density of 1×10^5 cells/well. After 24 h of culture in a 5% CO_2 incubator at 37°C , adherent cells were treated with a range of Kae concentrations (0.1, 1, 10 and $100 \mu\text{M}$) for 24 h. The MTT reagent ($10 \mu\text{l}$; 10 mg/ml) was subsequently added to each well, and cells were incubated for a further 4–6 h in a 5% CO_2 incubator at 37°C . The medium was removed and $100 \mu\text{l}$ DMSO was added to each well. Plates were shaken for 10 min, and the absorbance was measured at 492 nm using a Spectra Max Paradigm microplate reader (Molecular Devices, LLC). Cytotoxicity was calculated as follows: Cytotoxicity (%) = $(1 - \text{absorbance of sample} / \text{absorbance of control}) \times 100$.

Experimental groups and treatments. Third-passage BMSCs were randomly divided into 5 groups: i) Control group, in which BMSCs derived from normal rats were treated with osteogenic induction medium (Cyagen Biosciences, Inc.) to induce osteoblast differentiation; ii) OVX group, in which BMSCs derived from OVX rats were treated in the same way as the control group; iii) OVX + Kae group, in which BMSCs derived from OVX rats were incubated with 0.1, 1, 10 or $100 \mu\text{M}$ Kae and treated in the same way as the control group; iv) OVX + Kae + Rapa group, in which BMSCs derived from OVX rats were incubated with Kae and $10 \mu\text{M}$ Rapa and then treated in the same way as the control group; v) OVX + Rapa group, where BMSCs derived from OVX rats were incubated with $10 \mu\text{M}$ Rapa and treated in the same way as the control group. The dose of Rapa was determined during preliminary experiments. Briefly, MTT was detected to evaluate the optimal concentration of Rapa. Rapa (at 0.1, 1, 10 or $100 \mu\text{M}$) was incubated with BMSCs. Rapa was found to exhibit a concentration-dependent cytotoxic effect. Therefore, $10 \mu\text{M}$ was selected. BMSCs were harvested at day 15 in all groups.

ARS staining. ARS staining was used to assess the osteogenic differentiation of BMSCs. Briefly, rat BMSCs were plated into 24-well plates at a density of 2.5×10^5 cells/well. After induction of osteogenesis, cells were fixed with 95% alcohol for 15 min at room temperature and the BMSCs were stained with ARS for 5 min at room temperature. Mature osteoblasts that differentiated from rBMSCs displayed intense brown-red staining after 5 min of ARS staining (magnification, $\times 40$). Cells were imaged at $\times 400$ magnification by light microscopy, after the

addition of 10% (w/v) cetylpyridinium chloride to precipitate calcium ions. The absorbance of each well was measured at 562 nm using a Spectra Max Paradigm microplate reader (Molecular Devices, LLC).

Alkaline phosphatase activity assay. The effects of Kae on osteoblasts were assessed using an ALP assay kit. The absorbance/optical density (OD) of each sample was measured at 492 nm using a microplate reader. ALP activity in the osteogenic induction medium (Cyagen Biosciences, Inc.) was determined as follows: ALP activity (Jinshi unit/100 ml) = $(T - B) / (S - B) \times 0.02 \text{ mg/ml} \times 100 \text{ ml} \times a$, where T = OD of the test sample; B = OD of the blank; S = OD of the standard; and a = dilution factor of the sample. According to the conditions above, a Jinshi unit is defined as the ALP activity that releases of 1 mg of phenol every 15 min after mixing with 100 ml liquid matrix at 37°C .

Western blot analysis. Cells were lysed in RIPA lysis buffer containing 50 mM Tris-HCl (pH 7.4), 150 mM NaCl, 1% sodium deoxycholate, 1% NP-40, 1 mM PMSF and 1 mM EDTA. Then, extracts were centrifuged at $24,750 \times g$ at 4°C for 15 min to remove insoluble material. Total protein concentrations were determined using a bicinchoninic acid assay (Beyotime Institute of Biotechnology), according to the manufacturer's protocol. An equivalent quantity of total protein ($40 \mu\text{g}$) per well was diluted in sample buffer containing 100 mM dithiothreitol and heated to 98°C for 5 min. Lysates were separated using 10–15% SDS-PAGE (Bio-Rad Laboratories, Inc.) and subsequently transferred to PVDF membranes. The membranes were blocked in 5% non-fat dry milk for 2 h at room temperature and incubated overnight at 4°C using primary antibodies (1:500) against β -actin, Runx2, 4E/BP 1, p-4E/BP1, S6K1 and p-S6K1. Membranes were washed in TBS with Tween-20 (TBS-T), and incubated with horseradish peroxidase-conjugated secondary antibody (1:2,000) at room temperature for 1 h. Membranes were washed three times for 20 min in TBS-T, and protein bands were visualized by enhanced chemiluminescence (Proteintech Group, Inc.). Band intensities were measured and quantitated using Quantity One software (version 4.6.6; Bio-Rad Laboratories, Inc.) and β -actin was used for normalization.

Statistical analysis. SPSS (version 20.0; IBM Corp.) was used for statistical analysis. Data are presented as the mean \pm SEM from six independent experiments. The variance homogeneity test and one-way ANOVA were performed between groups. Newman-Keuls test was used following ANOVA. $P < 0.05$ was considered to indicate a statistically significant difference.

Results

Kae ameliorates OVX-induced osteoporosis in rats. The 3D trabecular bone microarchitecture, used for the assessment of distal femoral metaphysis, was calculated from micro-CT images (Fig. 2). The mean BV/TV, Tb.N, Tb.Th and BMD values in the OVX group were significantly lower than those of the sham group, while Tb.Sp and SMI were higher ($P < 0.05$). Compared with the OVX group, BV/TV, Tb.N, Tb.Th and BMD values were significantly greater, while Tb.Sp and SMI

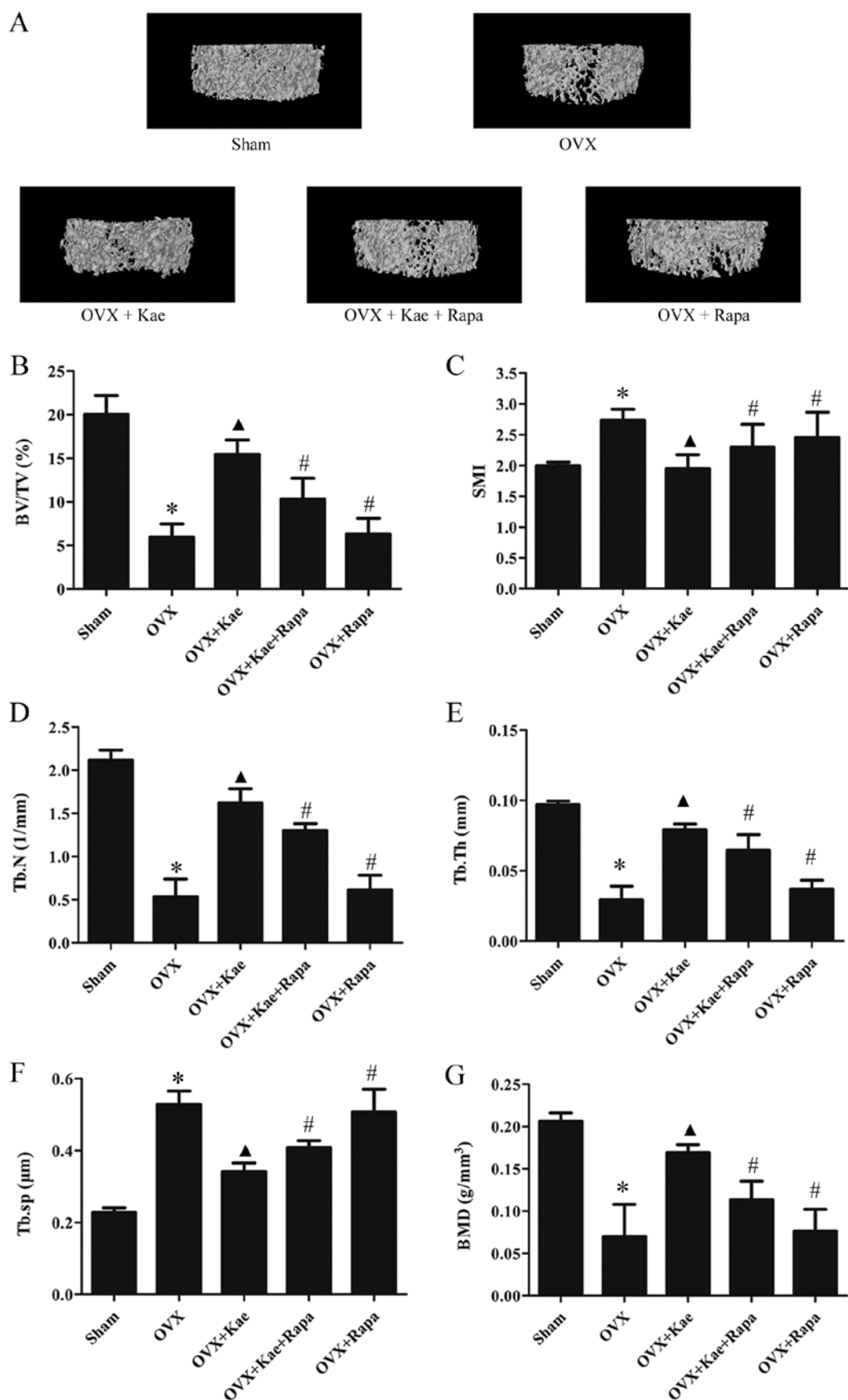


Figure 2. Effects of Kae on bone microarchitecture in rats. (A) Representative trabecular bone microarchitecture of the femoral metaphysis for each group, obtained from micro-CT images. Quantification of various parameters of bone microarchitecture for each group: (B) BV/TV, (C) SMI, (D) Tb.N, (E) Tb.Th, and (F) Tb.Sp. (G) BMD parameters measured using micro-CT in each group. The data are expressed as the mean \pm SEM (n=6). *P<0.05 vs. control group; [▲]P<0.05 vs. OVX group; [#]P<0.05 vs. OVX + Kae group. Kae, kaempferol; micro-CT, micro-computed tomography; BV/TV, bone volume/tissue volume; Tb.N, trabecular number; Tb.Sp, trabecular separation, Tb.Th, trabecular thickness; SMI, structure model index; BMD, bone mineral density; Rapa, rapamycin; OVX, ovariectomized.

were lower in the OVX + Kae group ($P < 0.05$). Thus, treating OVX rats with Kae significantly improved bone microarchitecture and mass. The opposite trend was observed in the OVX + Kae + Rapa and OVX + Rapa groups: OVX rats treated with Kae + Rapa or Rapa only exhibited lower BV/TV, Tb.N and Tb.Th values and higher Tb.Sp and SMI values compared to the OVX + Kae group ($P < 0.05$ vs. OVX + Kae).

Assessment of the optimal Kae dose that promotes osteoblast differentiation. The cytotoxicity of various concentrations of Kae was tested on BMSCs. As shown in Fig. 3, Kae displayed a concentration-dependent cytotoxic effect, demonstrating high cytotoxicity at 100 μ M. Additionally, it was observed that Kae induced osteoblast differentiation (Fig. 4). On day 15 following osteogenic induction, ARS staining revealed increased calcium deposition with increasing Kae concentrations. However, the number of osteoblasts unexpectedly decreased when cells were treated with 100 μ M Kae. From the range of concentrations tested, the optimal concentration of Kae that promoted osteoblast differentiation was 10 μ M. This concentration was therefore used in subsequent experiments.

Kae promotes osteogenic differentiation. ARS staining was used to assess the induction of osteogenesis in BMSCs. As illustrated in Fig. 5, following the induction of osteogenic differentiation, the rate of calcific nodule formation decreased in the OVX group compared to the control group ($P < 0.05$). Moreover, treatment with Kae significantly increased the number of calcified nodules in the OVX + Kae group ($P < 0.05$ vs. OVX group). However, the protective effect of Kae on osteoblasts was antagonized by the co-administration of Rapa ($P < 0.05$ vs. OVX + Kae).

Kae increases ALP expression in BMSCs. ALP activity was measured during osteogenesis. As shown in Fig. 6, the expression of ALP in the OVX group was significantly reduced compared with the control group ($P < 0.05$). In addition, ALP activity was significantly higher in the OVX + Kae group compared to the OVX group ($P < 0.05$ vs. OVX group). Pretreatment with Kae + Rapa or with Rapa alone led to reduced levels of ALP activity compared to Kae treatment alone ($P < 0.05$).

Kae upregulates Runx2 and Osterix expression. Western blot analysis was used to assess the expression of the osteogenesis-related transcription factors Runx2 and Osterix during osteogenesis. Fig. 7 demonstrates that a significant down-regulation of Runx2 and Osterix expression occurred in the OVX group compared with the control group ($P < 0.05$). Kae significantly increased the expression of Runx2 and Osterix compared with the OVX group ($P < 0.05$). However, their expression was significantly lower in the OVX + Kae + Rapa and OVX + Rapa groups compared with the OVX + Kae group ($P < 0.05$).

Kae alters the phosphorylation of 4E/BP1 and S6K1. Western blot analysis was used to determine the levels of phosphorylated and total 4E/BP1 and S6K1, important downstream regulators of the mTOR pathway. As shown in Fig. 8, the levels of p-4E/BP1 in the OVX + Kae group were significantly

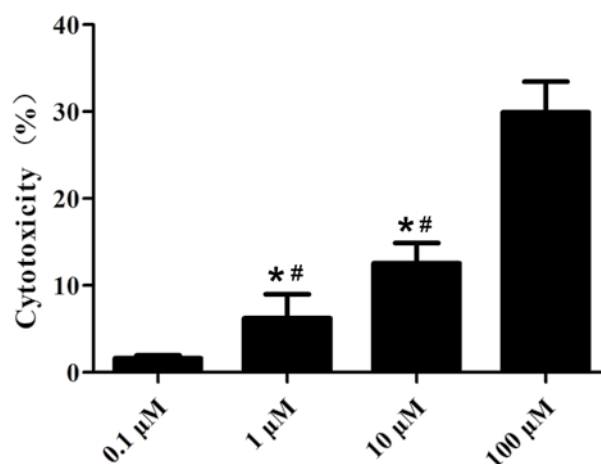


Figure 3. Effects of Kae on cell cytotoxicity. BMSCs were treated with different concentrations of Kae and cell cytotoxicity was assessed via an MTT assay. The data are expressed as the mean \pm SEM of six individual experiments. * $P < 0.05$ vs. 0.1 μ M and # $P < 0.05$ vs. 100 μ M. Kae, kaempferol; BMSCs, bone mesenchymal stem cells.

lower than in the OVX group ($P < 0.05$). However, a notably increased level of 4E/BP1 p-4EB/P1 was observed in the OVX + Kae + Rapa and OVX + Rapa groups. In addition, Kae increased S6K1 and p-S6K1 levels (Fig. 9; $P < 0.05$ vs. OVX group). Conversely, the levels of S6K1 and p-S6K1 were significantly lower in the OVX + Kae + Rapa and OVX + Rapa groups compared with the OVX + Kae group ($P < 0.05$). Taken together, these results indicated that Kae may promote osteogenesis through mTOR signaling as 4E/BP1 was activated by mTOR, whereas S6K1 exhibited the opposite trend.

Discussion

Estrogen has a significant role in bone metabolism (22), inhibiting bone resorption and stimulating bone formation (23). Flavonoids are estrogen-like chemicals used as estrogen substitutes, which may exert protective effects against postmenopausal bone loss (11,24). In addition, Kae is a phytoestrogen that possesses osteogenic properties (6). Guo *et al* (25) demonstrated that Kae stimulates osteogenic differentiation in cultured osteoblasts through estrogen receptor signaling. In addition, evidence suggests that Kae has a positive effect on bone metabolism, since it was observed to promote osteogenesis and to inhibit osteoclast activity, adipocyte formation and autophagy (26-30). From the *in vivo* assessment of bone microarchitecture, the present study verified that the bone mass parameters (BMD, BV/TV, Tb.N and Tb.Th) were significantly elevated, and the parameters of osteoporosis (SMI and Tb.Sp) reduced in OVX rats treated with Kae compared with OVX group rats. These changes in bone morphology induced by Kae in OVX rats are similar to those found in previously published research (26,27). However, the concentration of Kae used in the *in vivo* experiment of the present study was higher than that of previous studies (26,27). The similarity of the results may be associated with differences in delivery time or the vehicle solution used in the present study. Further research is required to effectively assess the effect of different doses of Kae on bone metabolism. Additionally, when BMSCs were isolated from OVX rats and osteogenesis was induced *in vitro*,

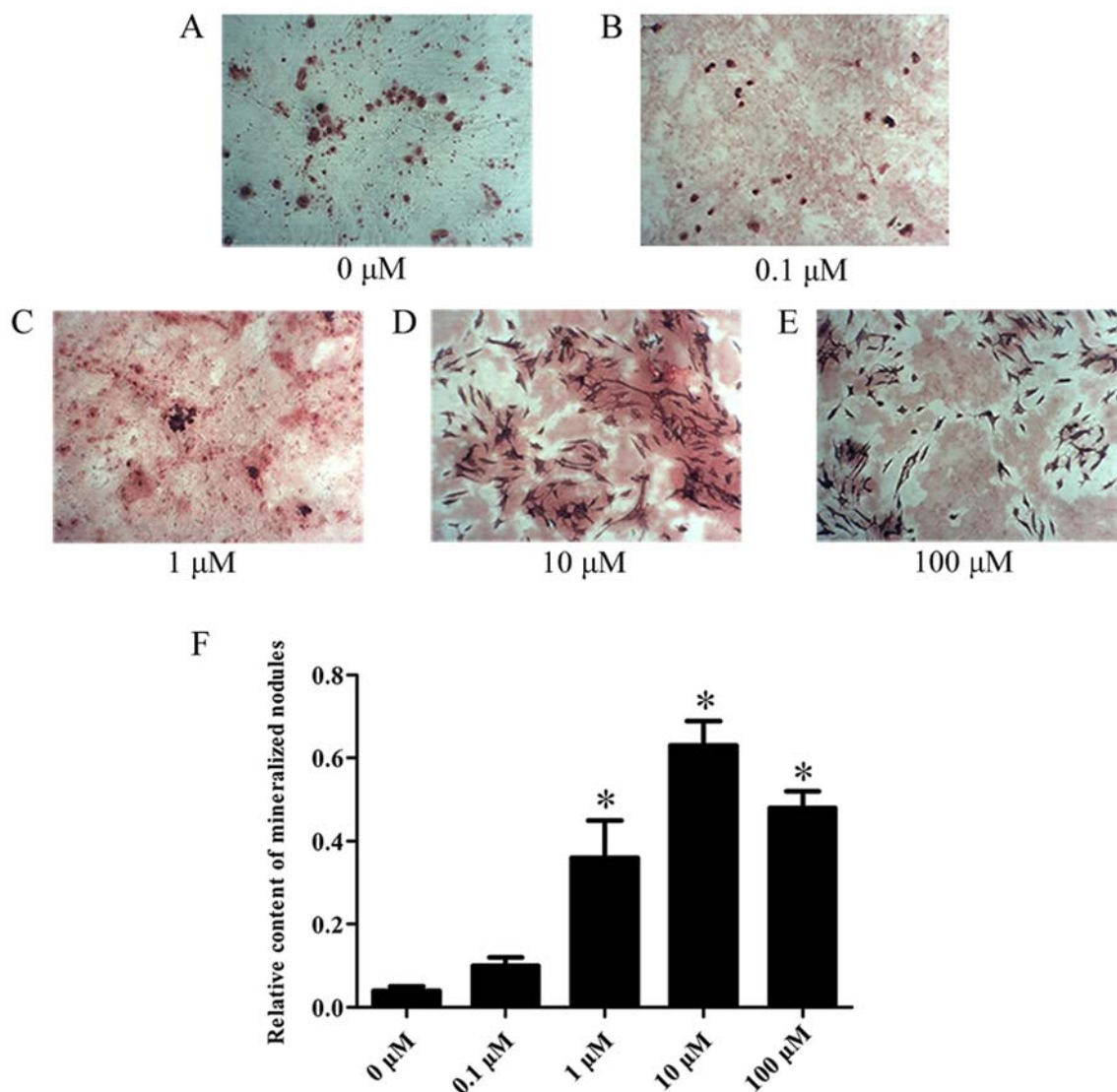


Figure 4. Effects of Kae on BMSC commitment to an osteogenic fate. ARS staining was performed to observe the osteogenic effects of Kae on rat BMSCs (0-100 μ M). ARS-stained images (magnification, x400) show the effect of the treatment with (A) 0, (B) 0.1, (C) 1, (D) 10 and (E) 100 μ M Kae. (F) Bar graph illustrating the quantitative analysis. Values are presented as the mean \pm SEM of six individual experiments. * P <0.05 vs. 0 μ M. Kae, kaempferol; rBMSCs, rat bone mesenchymal stem cells; ARS, Alizarin Red S.

Kae was found to promote osteogenic differentiation and increase ALP activity, consistent with Kim *et al* (30), although their data suggest that Kae stimulates osteogenic differentiation through increased expression of autophagy-related factors. In addition, Trivedi *et al* (26) also reported that treatment with Kae led to a reduction in adipogenic differentiation of BMCs in a Kae + OVX group, verified by a significant reduction in adipocyte number compared with an OVX group using Oil Red O staining. Both osteoblasts and adipocytes share a common precursor (MSCs), and the commitment to either fate is maintained in a dynamic balance under physiological conditions, serving a critical role in the microenvironment of the bone marrow (31-33). However, the imbalance between adipogenesis and osteogenesis has been linked to a number of pathophysiological processes, such as osteoporosis, osteopenia, obesity and aging (31-33). The observations of the present study lend support to these previously published data. Kae principally exerts its osteogenic function via promotion of bone formation, but whether it functions via inhibition of

adipogenesis requires further investigation. Together, the *in vitro* and *in vivo* experiments suggest that Kae prevented OVX-induced bone loss by promotion of osteoblast function. To determine the molecular mechanism of Kae-dependent osteogenic differentiation, an assessment of the activation of different signaling pathways was performed.

Runx-2, also termed core binding factor α -1, activates and initiates the differentiation of BMSCs into osteoblasts and regulates osteoblast maturation (34). Runx-2 is a key transcription factor during osteoblast differentiation and serves a crucial role in bone formation and reconstruction (35). Osterix, a zinc finger-containing transcription factor required for osteoblast differentiation, was first discovered by Nakashima *et al* in 2002 (36). A study revealed that the silencing of mouse Osterix induces chondrogenesis and chondrocyte differentiation; in addition, bone formation is impaired but Runx2 expression is unaffected (37). Osterix was not found to be expressed in Runx2-deficient mice, suggesting that it acts downstream of Runx2 during osteoblast differentiation (37).

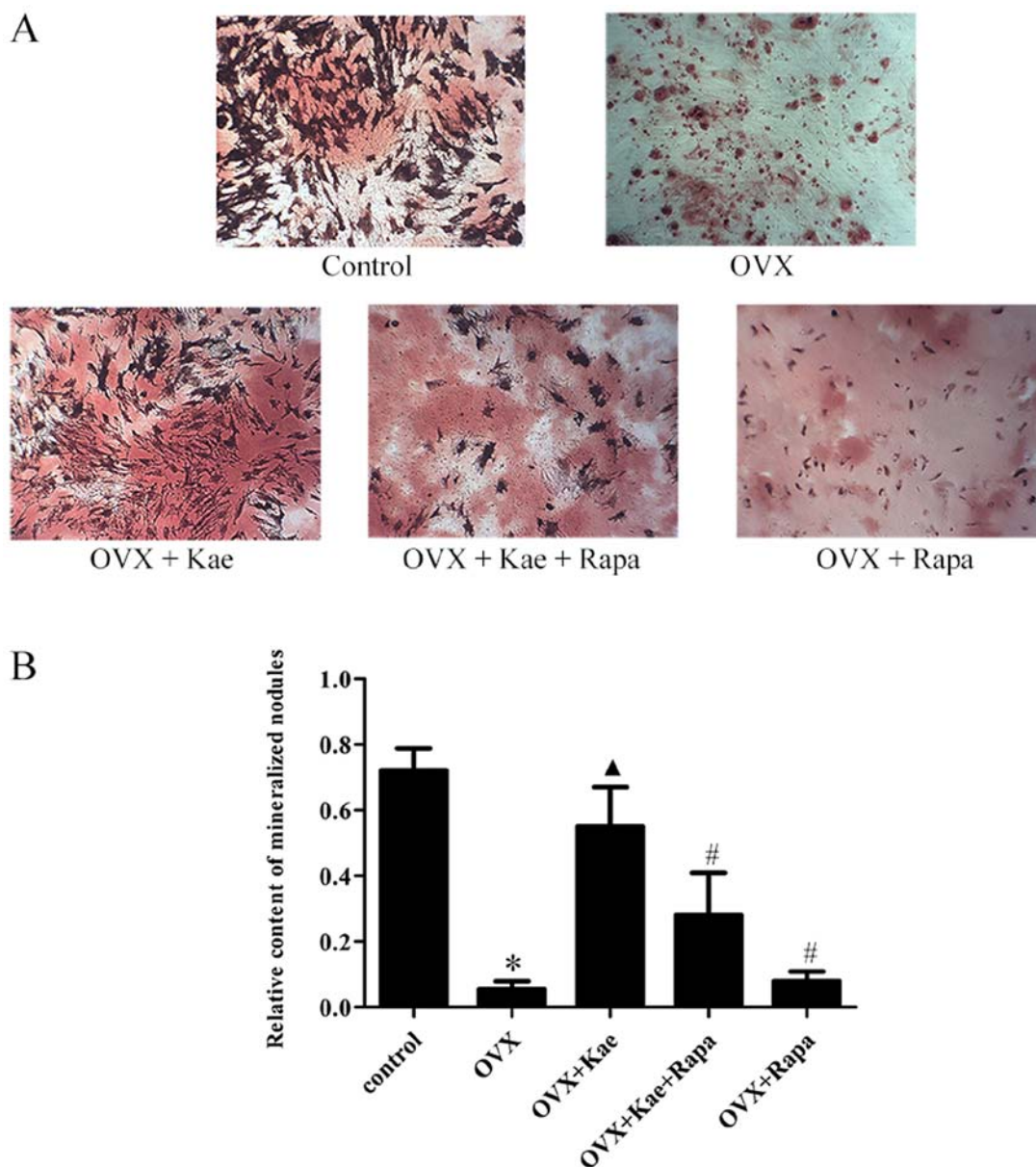


Figure 5. Kae promotes the differentiation of BMSCs into osteoblasts. ARS staining was performed to observe the osteogenic differentiation of cultured BMSCs. (A) ARS-stained images (magnification, x400) and respective (B) quantitative analysis. The data are expressed as the mean \pm SEM (n=6 for each group). *P<0.05 vs. control group; ^P<0.05 vs. OVX group; #P<0.05 vs. OVX + Kae group. Kae, kaempferol; BMSCs, bone mesenchymal stem cells; ARS, Alizarin Red S; Rapa, rapamycin; OVX, ovariectomized.

Thus, both Runx2 and Osterix are master transcription factors that promote osteoblast differentiation and bone formation. In the present study, western blot analysis was used to detect the expression levels of Runx2 and Osterix in BMSCs after the induction of osteogenesis. Kae was found to significantly upregulate Runx2 and Osterix expression. This suggested that the effect of Kae on osteogenic differentiation is mediated via Runx2 and Osterix.

mTOR is a member of the phosphatidylinositol 3-kinase-related kinase family of protein kinases (12). As a core component of intracellular signaling, mTOR regulates cell growth, proliferation and survival, protein synthesis, autophagy and transcription through two functionally and structurally distinct multi-component complexes, mTORC1 and mTORC2 (13). mTORC1 consists of mTOR, regulatory-associated protein of mTOR (Raptor) and G protein β -unit-like

protein (38). mTORC1 promotes the phosphorylation of S6K1, but inhibits the phosphorylation of 4E/BP1 (39,40). Activation of mTORC1 improves growth factor metabolism, cell growth and proliferation (39,40). mTORC2 consists of mTOR, mTOR associated protein, LST8 homolog, Raptor independent companion of mTOR complex 2 and mSIN1, phosphorylating and activating Akt (phosphorylation site, S473) and glycogen synthase kinase phosphorylation (41,42).

Studies have demonstrated that mTOR signaling has numerous key regulatory functions in various diseases, including cancer (43-46), infectious diseases (47,48), atherosclerosis, and degenerative and autoimmune diseases (49-51), and has been suggested to be a novel therapeutic target for the treatment of these diseases (43-51). Other studies have shown that mTOR signaling is also involved in the regulation of multiple aspects of skeletal development (52-54). However, controversy remains

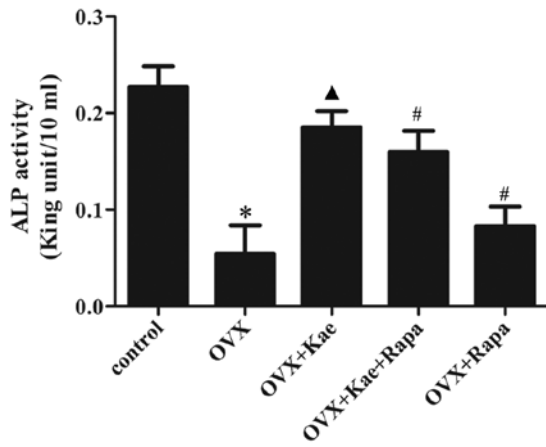


Figure 6. Kae increases the level of ALP in BMSCs. The data are expressed as the mean ± SEM (n=6 for each group). *P<0.05 vs. control group; [▲]P<0.05 vs. OVX group; [#]P<0.05 vs. OVX + Kae group. Kae, kaempferol; BMSCs, bone mesenchymal stem cells; ALP, alkaline phosphatase; Rapa, rapamycin; OVX, ovariectomized.

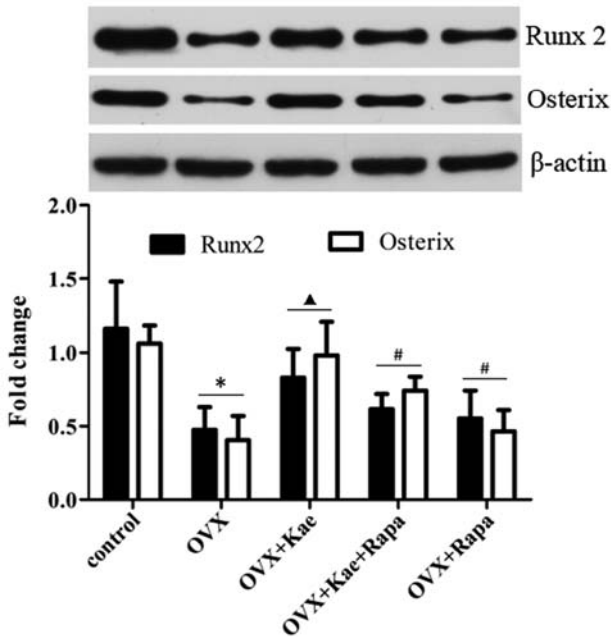


Figure 7. Kae upregulates the expression of Runx2 and Osterix. Western blot analysis was used for the assessment of Runx2 and Osterix expression, and β-actin was used as the internal control. Pretreatment with Kae induced increased expression of Runx2 and Osterix in BMSCs, which was antagonized by Rapa. The data are expressed as the mean ± SEM (n=6 for each group). *P<0.05 vs. control group; [▲]P<0.05 vs. OVX group; [#]P<0.05 vs. OVX + Kae group. Kae, kaempferol; BMSCs, bone mesenchymal stem cells; Runx 2, Runt-related transcription factor; Rapa, rapamycin; OVX, ovariectomized.

as to whether mTOR promotes (17,18,55,56) or inhibits (56-59) osteogenic differentiation. A recent study indicated that knockdown of DEP domain containing MTOR interacting protein (Deptor) promotes BMSC osteogenesis both *in vitro* and *in vivo*, with higher expression of Deptor contributing to the progression of osteoporosis (56). It is well known that over-expression of Deptor downregulates the activity of mTORC1 and mTORC2, indicating that mTOR may play a crucial role during osteogenesis. Similarly, a previous study demonstrated

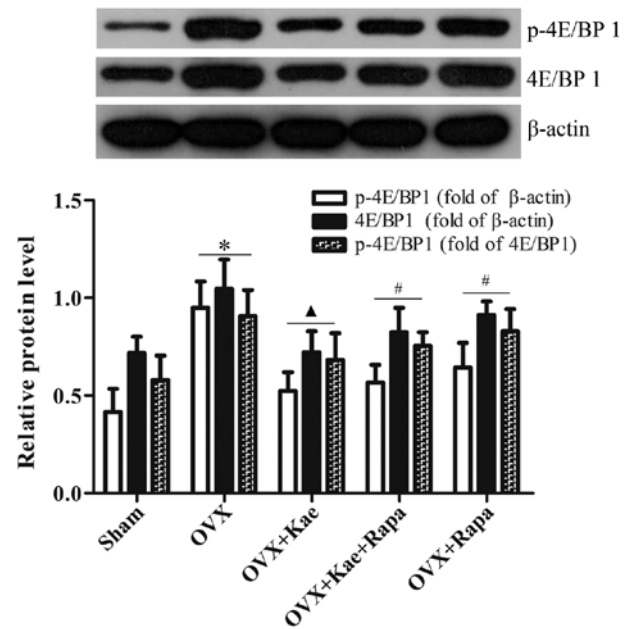


Figure 8. Kae decreases the phosphorylation of 4E/BP1. Relative levels of 4E/BP1 and p-4E/BP1 were detected by western blotting. Treatment with Kae decreased the levels of p-4E/BP1, which was antagonized by Rapa. The data are expressed as the mean ± SEM (n=6 for each group). *P<0.05 vs. respective sham group; [▲]P<0.05 vs. respective OVX group; [#]P<0.05 vs. respective OVX + Kae group. Kae, kaempferol; Rapa, rapamycin; BMSCs, bone mesenchymal stem cells; 4E/BP1, eukaryotic translation initiation factor 4E-binding protein 1; p, phosphorylated; OVX, ovariectomized.

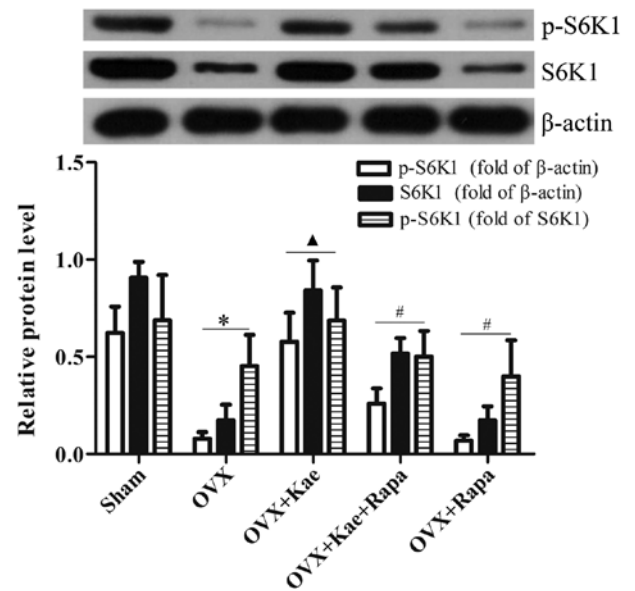


Figure 9. Kae increases the phosphorylation of S6K1. Protein levels of S6K1 and p-S6K1 were detected by western blotting. Treatment with Kae increased the levels of p-S6K1, an effect that was antagonized by Rapa. The data are expressed as the mean ± SEM (n=6 for each group). *P<0.05 vs. respective sham group; [▲]P<0.05 vs. respective OVX group; [#]P<0.05 vs. respective OVX + Kae group. Kae, kaempferol; Rapa, rapamycin; BMSCs, bone mesenchymal stem cells; S6K1, ribosomal protein S6 kinase B1; p, phosphorylated; OVX, ovariectomized.

that treatment with Rapa attenuated the expression of ALP, downregulated the expression and phosphorylation of S6K1 and inhibited osteoblastic differentiation in vascular smooth

muscle cells (VSMCs), suggesting that mTOR participates in the osteoblastic differentiation of VSMCs (18). In the present study, Kae was found to increase the expression of important downstream regulatory factors in the mTOR pathway, which also indicates that mTOR signaling may be involved in osteogenesis. However, a number of researchers have published contrary results. For instance, Dai *et al* (59,60) indicated that deletion of Raptor (an indispensable component of mTORC1) in osteoclasts leads to increased bone volume and decreased bone resorption. In other words, mTORC1 is critical for osteoclast formation. Raptor deficiency in osteoclasts impairs bone resorption without affecting bone formation, ultimately leading to increased bone mass in mice. On the other hand, Xu *et al* (58) reported that the activation of mTORC1 in B lymphocytes promotes osteoclast formation and causes osteoporosis in mice by promoting TNF superfamily member 11 expression with negative regulation of Akt and β -catenin. A separate study by Wang *et al* (61) also indicated that suppression of the mTOR signaling pathway by rapamycin was able to promote BMSC differentiation into osteoblasts in the context of degenerative scoliosis. These conflicting results may be attributed to different methodologies (different cell lines, drug concentrations or experimental design), and underline the requirement for further research. Thus, for further validation, BMSCs treated with Rapa, a specific inhibitor of mTOR, exhibited a marked decline in osteogenic activity and decreased Runx2 and Osterix expression. This treatment also reversed the positive effects of Kae exposure. Similarly, a number of studies have revealed that Kae can exert cytoprotective effects via regulation of mTOR signaling in different cell lines (62-64). Taken together, these data support the hypothesis that the mTOR-Runx2/Osterix signaling axis is associated with the osteogenic effect of Kae in OVX rats and respective BMSCs.

In summary, the present study demonstrated that Kae has a positive effect on bone formation and osteogenesis in OVX rats, which is possibly exerted via mTOR-Runx2/Osterix signaling. Since Kae naturally occurs in a variety of foods, it is likely that it may exert beneficial effects as a food supplement by improving postmenopausal osteoporosis. Further large scale randomized *in vivo* trials are now required to confirm the potential of Kae in the treating osteoporosis.

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Availability of data and materials

The datasets used and/or analyzed during the present study are available from the corresponding author on reasonable request.

Authors' contributions

JL conceived and designed the present study and revised the manuscript for important intellectual content. JZ was involved in all experiments and was a major contributor in writing the manuscript. JW modified the study design and designed the structure of the article. YL performed the experiments, analyzed the data and wrote the manuscript. JM and XL completed the data analysis. BX and ZY performed the cell culture, ARS staining, ALP activity assay, western blotting. All authors read and approved the final manuscript.

Ethics approval and consent to participate

Experiments were performed according to the Guide for the Care and Use of Laboratory (National Institutes of Health), and were approved by the Ethics Committee of Nanchang University (no. 2017-0122).

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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