

Zoledronic acid promotes osteoclasts ferroptosis by inhibiting FBXO9-mediated p53 ubiquitination and degradation

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Bisphosphonates (BPs)-related osteonecrosis of jaw (BRONJ) is a severe complication of the long-term administration of BPs. The development of BRONJ is associated with the cell death of osteoclasts, but the underlying mechanism remains unclear. In the current study, the role of Zoledronic acid (ZA), a kind of bisphosphonates, in suppressing the growth of osteoclasts was investigated and its underlying mechanism was explored. The role of ZA in regulating osteoclasts function was evaluated in the RANKL-induced cell model. Cell viability was assessed by cell counting kit-8 (CCK-8) assay and fluorescein diacetate (FDA)-staining. We demonstrated that ZA treatment suppressed cell viability of osteoclasts. Furthermore, ZA treatment led to osteoclasts death by facilitating osteoclasts ferroptosis, as evidenced by increased Fe^{2+} , ROS, and malonyldialdehyde (MDA) level, and decreased glutathione peroxidase 4 (GPX4) and glutathione (GSH) level. Next, the gene expression profiles of alendronate- and risedronate-treated osteoclasts were obtained from Gene Expression Omnibus (GEO) dataset, and 18 differentially expressed genes were identified using venn diagram analysis. Among these 18 genes, the expression of F-box protein 9 (FBXO9) was inhibited by ZA treatment. Knockdown of FBXO9 resulted in osteoclasts ferroptosis. More important, FBXO9 overexpression repressed the effect of ZA on regulating osteoclasts ferroptosis. Mechanistically, FBXO9 interacted with p53 and decreased the protein stability of p53. Collectively, our study showed that ZA induced osteoclast cells ferroptosis by triggering FBXO9-mediated p53 ubiquitination and degradation.

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2 **mediated p53 ubiquitination and degradation**

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23 **Abstract**

24 Bisphosphonates (BPs)-related osteonecrosis of jaw (BRONJ) is a severe complication of the long-
25 term administration of BPs. The development of BRONJ is associated with the cell death of
26 osteoclasts, but the underlying mechanism remains unclear. In the current study, the role of
27 Zoledronic acid (ZA), a kind of bisphosphonates, in suppressing the growth of osteoclasts was
28 investigated and its underlying mechanism was explored. The role of ZA in regulating osteoclasts
29 function was evaluated in the RANKL-induced cell model. Cell viability was assessed by cell
30 counting kit-8 (CCK-8) assay and **fluorescein diacetate** (FDA)-staining. We confirmed that ZA
31 treatment suppressed cell viability of osteoclasts. Furthermore, ZA treatment led to osteoclasts
32 death by facilitating osteoclasts ferroptosis, as evidenced by increased Fe^{2+} , ROS, and
33 malonyldialdehyde (MDA) level, and decreased glutathione peroxidase 4 (GPX4) and glutathione
34 (GSH) level. Next, the gene expression profiles of alendronate- and risedronate-treated osteoclasts
35 were obtained from Gene Expression Omnibus (GEO) dataset, and 18 differentially expressed
36 genes were identified using venn diagram analysis. Among these 18 genes, the expression of F-
37 box protein 9 (FBXO9) was inhibited by ZA treatment. Knockdown of FBXO9 resulted in
38 osteoclasts ferroptosis. More important, FBXO9 overexpression repressed the effect of ZA on
39 regulating osteoclasts ferroptosis. Mechanistically, FBXO9 interacted with p53 and decreased the
40 protein stability of p53. Collectively, our study showed that ZA induced osteoclast cells ferroptosis
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42 **Keywords:** Bisphosphonates related osteonecrosis of jaw; Zoledronic acid; ferroptosis; FBXO9;

43 p53

44 **Introduction**

45 Bisphosphonates (BPs) inhibit osteoclast activity and disrupt osteoclast mediated bone resorption
46 [1, 2]. BPs are widely used in the treatment of bone metastasis cancer[3], osteoporosis[4], and
47 multiple myeloma[5]. BRONJ is an injury of the jaw that affects patients treated with BPs. Since
48 it was reported in 2003, BRONJ has been considered as a common and important adverse side
49 effect of BPs treatment, especially nitrogen-containing bisphosphonates (Residronate,
50 Alendronate, and Zoledronic acid(ZA))[6]. There are various hypotheses for the development of
51 BRONJ, the most recognized hypothesis was bone remodeling suppression. Although significant
52 progress has been made in prevention and treatment of BRONJ base on the hypotheses, the
53 mechanisms underlying the development of BRONJ remain unclear.

54 Osteoclasts, members of the monocyte/macrophage hematopoietic, play an important role in
55 the progression of bone remodeling[8]. RAW264.7 cells and bone marrow-derived macrophages
56 (BMDMs) can be induced into osteoclasts by receptor activator of nuclear factor- κ B ligand
57 (RANKL), and has been widely used as a cell model for the study of osteoclast related diseases *in*
58 *vitro*[9]. The number and the resorptive function of osteoclasts were usually increased during the
59 process of bone remodeling [10]. Therefore, osteoclast is one of the core targets for the treatment
60 of osteoporosis and other bone-remodeling-related diseases. It is well known that ZA can lead to
61 a stronger inhibition of osteoclasts differentiation and induces the apoptosis of osteoclasts[11],
62 while the underlying mechanism of ZA in the function of osteoclasts remains unclear.

63 Ferroptosis is a recently identified type of iron-mediated cell death. Unlike other forms of

64 programmed cell death, such as apoptosis and necroptosis, ferroptosis does not involve the
65 activation of caspase protein[12, 13]. It is characterized by an increased level of lipid peroxidation
66 products and reactive oxygen species (ROS). The dysregulation of ferroptosis has been related to
67 many pathological processes, such as cancer[14], neurodegenerative diseases[15], and
68 inflammation-related diseases[16]. More and more studies showed that ferroptosis contribute to
69 the development of BRONJ. Jose *et al.* found that the levels of MDA, GSSG, and 8-oxo-dG and
70 the GSSG/GSH ratio in serum and saliva were significantly higher in patients with BRONJ
71 compared with controls[17]. Ma *et al.* demonstrated that melatonin suppresses osteoblast
72 ferroptosis and improved the osteogenic capacity of MC3T3-E1 by activating the Nrf2/HO-1
73 pathway[18]. However, whether ferroptosis was involved in the osteoclasts differentiation and
74 death induced by ZA is still unknown. In the current study, we showed that ZA inhibits the
75 osteoclasts viability in a dose-dependent manner. For the first time, we showed that ZA promotes
76 the ferroptosis of osteoclast by increasing the protein stability of p53. ZA-induced downregulation
77 of ubiquitin E3 ligase FBXO9, and FBXO9 overexpression restores cell viability inhibition of
78 osteoclast induced by ZA. Moreover, FBXO9 facilitates ubiquitination-mediated degradation of
79 p53.

80

81 **Materials and methods**

82 **Cell culture**

83 RAW264.7 cells were purchased from the ATCC (TIB-71, Manassas, VA, USA) and cultured in
84 alpha-Modified Eagle's Medium (α -MEM, Gibco, USA) with 100 U/ml penicillin, and 100 μ g/ml

85 streptomycin at 37 °C in a humidified atmosphere containing 5% CO₂ and 95% air. To BMDMs,
86 bone marrow cells were purchased from the ATCC (CRL-2420, Manassas, VA, USA) and cultured
87 in RPMI 1640 medium (Thermo Fisher Scientific, Waltham, MA, USA) for 6 days with 100 U/ml
88 penicillin, 100 µg/ml streptomycin, 10% FBS, and 10 ng/mL recombinant mouse macrophage
89 colony-stimulating factor (PeproTech). For osteoclast formation assay RAW264.7 cells were
90 seeded in 12-well plate s (1×10^4 cells/well) supplemented with 50 ng/ml RANKL (R&D Systems.)
91 for 6 days. BMDMs (1×10^4 cells/well) were cultured in the presence of M-CSF (10 ng/mL) and
92 RANKL (50 ng/mL) for 6 days.

93 **TRAP staining**

94 TRAP histochemical staining was performed to confirm the osteoclast as previously described[25],
95 by using an acid phosphatase, leukocyte (TRAP) kit (Sigma-Aldrich). Briefly, 1×10^5 BMDMs or
96 RAW264.7 induced osteoclasts were then fixed in 10% neutral-buffered formalin (NBF) solution
97 for 20 minutes, then the NBF solution was replaced with TRAP staining solution and 0.1% Fast
98 Red AS TR salt at RT. After 45 minutes cells were washed with $1 \times$ PBS for three times and
99 imaged.

100 **Measurement of cell viability**

101 Cell viability of BMDMs and RAW264.7 was assessed by using a CCK-8 and FDA assay
102 (Dojindo, Japan). For CCK-8 assay, BMDMs and RAW264.7 cells (1×10^4 cells per well) were
103 seeded in 96-well plates for 24 hours, then cells were treated with different doses of ZA (5, 10,
104 and 50µM) for 48 h. To analyze the cells death of osteoclast, BMDMs, and RAW264.7 (5×10^3
105 cells per well) were treated with 10µM of ZVAD-FMK, 2µM of Fer-1, or 10µM of necrostatin-1,

106 for 48 h with or without ZA (50 μ M). Then, a total 10 μ L of CCK-8 reagent was added to each well
107 additional 4 h at 37°C with 5% CO₂, and the absorbance at 450 nm of each well was assayed using
108 a microplate reader (BioTek Instruments). For FDA assay, after culture with different dose ZA (5,
109 10, and 50 μ M) for 48h, BMDMs and RAW264.7 cells were treated with 10 μ l of FDA solution (5
110 mg/mL; Invitrogen, CA, USA) at 37°C with 5% CO₂ for 20 minutes, then cells images were
111 obtained using a fluorescence microscope (Olympus Corporation, Japan).

112 **Fe²⁺ concentration**

113 The concentration of Ferrous iron (Fe²⁺) in BMDMs and RAW264.7 cells in the presence or
114 absence of ZA (50 μ M) was assessed using an iron assay kit (MAK025, Sigma-Aldrich, MO, USA)
115 as the manufacturer's instructions. Briefly, cell samples were incubated with 10 μ L of iron reducer
116 for 30 minutes at RT, then 100 μ L iron probe was added to trigger the reaction; thus, the absorbance
117 was measured at 593 nm.

118 **Lipid reactive oxygen species (ROS) assay**

119 Lipid ROS level in BMDMs and RAW264.7 cells in the presence or absence of ZA (50 μ M) was
120 assayed using C11-BODIPY (Invitrogen), a fluorescent-labelled oxidation sensitive probe. In
121 brief, BMDMs and RAW264.7 cells were seeded in 24-well plates (5 \times 10⁵/well) and treated with
122 ZA (50 μ M) with or without FBXO9 overexpression for 48h, then BMDMs and RAW264.7 cells
123 were cultured with C11-BODIPY probe with a final concentration of 1 μ M in at 37°C with 5% CO₂
124 for 30 minutes, then the Lipid ROS levels were assayed using flow cytometer.

125 **MDA and GSH content**

126 MDA in BMDMs and RAW264.7 cells was analyzed using a lipid peroxidation assay kit

127 (ab118970, Abcam) in the presence or absence of ZA (50 μ M). GSH content in BMDMs and
128 RAW264.7 cells was assayed using a Glutathione Assay Kit (ab65322, Abcam) according to the
129 standard protocol. Briefly, cell supernatant, 5,5' -dithio-bis 2-nitrobenzoic acid solution and the
130 reagents of kits were mixed together and incubated at RT for 10 minutes, then NADPH was added
131 into this system to trigger the reaction. the absorbance of 5-thio-2-nitrobenzoic acid was detected
132 at 412 nm.

133 **Transient transfection of FBXO9 or si-FBXO9**

134 The recombinant plasmids pcDNA-FBXO9 containing FBXO9 cDNA were sub-cloned into
135 pcDNA3.1 vector via EcoR V/Hind III sites. To overexpress FBXO9, the pcDNA-FBXO9 was
136 transfected into BMDMs cells using Lipofectamine 2000. FBXO9 Knockdown and
137 transfection were performed according to the manufacturer's instructions. Briefly, cells were
138 transfected with 10 nM of si-FBXO9 RNA (sense AUCAGAAUGACAAUCUCCUCU,
139 antisense GGAAGAUUGUCAUUCUGAUGCU
140) or si-Control RNA (sense CAGUCGCGUUUGCGACUGGUU, antisense CCAGUCG-
141 CAAACGCGACUGUU), and the cell was induced by M-CSF (10 ng/mL) and RANKL (50
142 ng/mL) for further experiment.

143 **Quantitative real-time PCR (qPCR)**

144 After treatment with 50 μ M of ZA for 48h, total RNA was isolated with Trizol reagent (Sigma-
145 Aldrich) as instructed by the manufacturer. Reverse transcriptional PCR was carried out using the
146 SMART PCR cDNA Synthesis Kits (Clontech). qPCR was carried out on ABI 7500 RealTime
147 PCR System (Applied Biosystem) with powerup SYBR green Mix (ThermoFisher). The fold

148 changes of RNA transcripts were calculated by the $2^{-\Delta\Delta C_t}$ method and the 18s was used as a
149 reference gene. The qPCR primer pairs in Table 1

150 **Western blotting analysis**

151 BMDMs-induced osteoclast (1×10^5 cells/well in 12-well plates) were treated with or without
152 ZA(50 μ M) for 48h, the total protein was isolated using RIPA Buffer (Solarbio, Beijing, China),
153 and total protein concentration was quantified by BCA protein assay kit (Beyotime). 20 μ g of
154 protein were separated by 8% SDS-PAGE and transferred to PVDF membranes (Merck Millipore,
155 Billerica, MA, USA). After blocking with 5% bovine serum albumin or 5% nonfat milk, the
156 membranes were incubated with anti-FBXO9 (1:1000, PA5-25475, Thermo Fisher Scientific),
157 p53(1.0 μ g/mL, MA5-14067, Thermo Fisher Scientific), ubiquitin (1:2000, ab134953, Abcam),
158 and GAPDH (1:5000, MA1-16757, Thermo Fisher Scientific) overnight at 4 °C. Then the
159 membranes were treated with HRP-conjugated anti-mouse or rabbit secondary antibody (1:5000)
160 for 1h at room temperature.

161 **Co-immunoprecipitation (Co-IP)**

162 BMDMs-induced osteoclast were lysed using NP40 buffer (10 mM Tris-HCl at pH 8.0, 140 mM
163 NaCl, 1.5 mM MgCl₂, 0.5% Nonidet-P40, 20 mM dithiothreitol, 500 U/mL RNAsin, and 0.5%
164 [w/v] deoxycholate), cell lysates were incubated with FBXO9 (2 μ g/ml; PA5-25475) or p53 (2
165 μ g/ml; MA5-14067) antibody for 4h, then Protein A/G beads (Thermo Fisher Scientific) were
166 added to the IP reactions and left rotating overnight at 4°C, then beads were washed by PBS
167 containing protein inhibitors for three times, then the immunoprecipitates were analyzed using
168 western blotting with FBXO9 antibody or p53 antibody.

169 **Statistical analysis**

170 All the data are shown as the mean \pm standard error of mean (SEM) from three independent
171 experiments. Statistical analysis was carried out using SPSS 19.0 software
172 (IBM Corp., NY, USA). The significance between two groups was analyzed using one-way
173 ANOVA followed by Tukey-Kramer multiple comparisons test or unpaired Student's *t*-test.
174 $p < 0.05$ was considered to indicate a statistically significant difference.

175

176 **Results**

177 **ZA treatment facilitated the ferroptosis of osteoclasts**

178 To investigate the function of ZA on osteoclasts, RAW264.7 cells and bone marrow-derived
179 macrophages (BMDMs) were pre-treated with RANKL for 6 days followed by ZA treatment in
180 different concentrations (5, 10, and 50 μ M) for 48 hours, the cell model was confirmed by TRAP
181 staining (Figure 1 A and B). Cell viability was analyzed by CCK-8 assay. As shown in Figure 1 C
182 and D, ZA treatment suppressed cell viability in a dose-dependent manner. The results from FDA
183 staining also showed that the effect of ZA on promoting cell viability of osteoclasts (Figure 1 E
184 and F). Impressively, the CCK-8 assay results showed that cell death of osteoclasts induced by ZA
185 treatment was obviously blocked by ferrostatin-1 (Fer-1, a specific inhibitor of ferroptosis) but not
186 necrostatin-1 (a specific inhibitor of necroptosis) and ZVAD-FMK (a specific inhibitor of
187 apoptosis) (Figure 1 G and H).

188 To define the role of ZA in the ferroptosis of osteoclasts, the ferroptosis signaling was evaluated
189 in osteoclasts after ZA treatment. As shown in Figure 2 A-C and F-H, ZA treatment markedly

190 increased the Fe²⁺ level, MDA content, and ROS level in a dose-dependent manner in osteoclasts,
191 differentiated from RAW264.7 cells and BMDMs, suggesting the promotion of ferroptosis
192 signaling in osteoclasts treated by ZA. Besides, ZA treatment also suppressed the levels of Gpx4
193 and GSH in a dose-dependent manner in osteoclasts (Figure 2D, E, I, and J). These results
194 demonstrate that ZA treatment facilitates the ferroptosis of osteoclasts.

195 **FBXO9 was downregulated in osteoclasts after ZA treatment**

196 To investigate the mechanism underlying ZA-induced osteoclasts ferroptosis, the differentially
197 expressed genes (DEGs) of osteoclasts induced by bisphosphonates alendronate- and risedronate-
198 treatment were obtained from GSE63009, and the common DEGs were identified by venn diagram
199 analysis. As shown in Figure 3A, eighteen common genes were identified (CFAP53, COL14A1,
200 ARSJ, ABCA9, CXorf57, GPR22, STXBP5L, MSANTD4, RRP15, UGT1A2, IRF4, TFAP2D,
201 TRHDE, ASMT, CAPS, COMMD10, VSTM4, FBXO9). The level of these 18 genes was
202 evaluated in osteoclasts treated with or without ZA using qPCR analysis. Figure 3B and C showed
203 that only FBXO9 was significantly decreased in osteoclasts after treatment with ZA. Similar to the
204 qPCR results, the results from western blotting showed that the expression of FBXO9 was
205 obviously decreased in osteoclasts after treatment with ZA. These results indicate that the FBXO9
206 was downregulated by ZA treatment (Figure 3D).

207 **FBXO9 inhibition facilitated the ferroptosis of osteoclasts**

208 To investigate the function of FBXO9 on osteoclasts, the expression of FBXO9 was down-
209 regulated by si-FBXO9 in BMDMs-differentiated osteoclasts. As shown in Figure 4 A-B, the
210 expression of FBXO9 was significantly decreased by si-FBXO9. Cell viability of osteoclasts was

211 significantly decreased in the FBXO9 knockdown group compared with the control group (Figure
212 4C). The results from FDA staining also showed the effect of FBXO9 on inhibiting cell viability
213 (Figure 4D and E).

214 To investigate the role of FBXO9 in ferroptosis of osteoclasts, the ferroptosis signaling was
215 evaluated in osteoclasts differentiated from BMDMs. As shown in Figure 4F-J, FBXO9
216 knockdown significantly increased the Fe^{2+} level, MDA content, ROS level and decreased the
217 GPX4 level, GSH content in osteoclasts. These results suggested that FBXO9 inhibition facilitates
218 the ferroptosis of osteoclasts

219 **ZA treatment facilitated the ferroptosis of osteoclasts by suppressing FBXO9**

220 To explore whether FBXO9 mediated the function of ZA in regulating the ferroptosis of
221 osteoclasts, the osteoclasts differentiated from BMDMs were treated by ZA in the presence or
222 absence of FBXO9. As shown in Figure 5A and B, the qPCR and western blotting analysis results
223 showed that the expressions of FBXO9 were decreased by ZA treatment and restored by FBXO9
224 overexpression. CCK8 results showed that cell viability of osteoclasts was decreased by ZA
225 treatment, but these effects were blocked by FBXO9 overexpression (Figure 5C). Consistently, the
226 FDA staining also showed the inhibition of ZA on osteoclasts cell viability was restored by FBXO9
227 overexpression (Figure 5D and E). Besides, the Fe^{2+} level, MDA content, and ROS level were
228 obviously increased and the GPX4 level, GSH content was significantly decreased by ZA
229 treatment, while these effects were blocked by FBXO9 overexpression (Figure 5F-J). These results
230 suggested that ZA treatment facilitates the ferroptosis of osteoclasts by suppressing FBXO9.

231 **FBXO9 inhibition facilitated the ferroptosis of osteoclasts by blocking the ubiquitin**

232 mediated-proteasome degradation of p53

233 Previous studies showed that FBXO9, an E3 ubiquitin ligase, mediated protein stability through
234 ubiquitin mediated-proteasome degradation[20]. Given that dysregulated ubiquitination has been
235 widely reported to be involved in many diseases by regulating cell ferroptosis[26, 27], the target
236 of FBXO9 was predicted by ubibrowser (<http://ubibrowser.ncpsb.org.cn/ubibrowser/>). Figure 6A
237 showed the 20 potential target genes that interacted with FBXO9. Interestingly, among these
238 genes, the p53 gene is an important regulator of ferroptosis. We next explored whether FBXO9
239 decreases p53 protein level by promoting its ubiquitination-mediated degradation. Knockdown of
240 FBXO9 in osteoclasts did not change the p53 mRNA level (Figure 6B). Fascinatingly, the protein
241 level of p53 was significantly increased after FBXO9 inhibition (Figure 6C), suggesting that
242 FBXO9 decreased p53 expression possibly by the ubiquitin-proteasome-mediated degradation.
243 Next, a reciprocal Co-IP assay was performed to confirm whether FBXO9 directly interacts with
244 p53. As shown in Figure 6D, a positive p53 signal was observed in the protein complex pulled
245 down by FBXO9 antibody. Meanwhile, FBXO9 was also detected in the co-immunoprecipitation
246 complex pulled-down by p53 antibody. Next, cycloheximide assay (CHX) was performed to detect
247 the protein stability of p53 in osteoclasts transfected with si-FBXO9. As shown in Figure 6E, the
248 protein stability of p53 was obviously increased in the FBXO9 knockdown osteoclasts. Then the
249 p53 ubiquitination was assessed through IP with FBXO9 antibody and subsequent western blotting
250 with ubiquitin antibody. Figure 6F showed that FBXO9 knockdown obviously decreased p53
251 ubiquitination in osteoclasts (Figure 6F). in conclusion, these results indicated that FBXO9 directly
252 interacts with p53 and promotes its degradation.

253

254 **Discussion**

255 BRONJ is one of the severe complications of BPs administration reported by Marx *et al.* in
256 2003[6]. It usually occurs in patients with bone metastatic cancer or osteoporosis, and undergoes
257 bisphosphonate therapy. ZA is a kind of nitrogen-containing bisphosphonates and is widely used
258 in the treatment of bone metastatic cancer and osteoporosis. Zhu *et al.* reported that ZA facilitates
259 TLR-4-mediated M1 type macrophage polarization in the development of BRONJ[28]. Huang *et*
260 *al.* demonstrated that ZA inhibited osteoclast differentiation and function by regulating the NF- κ B
261 and JNK signaling pathways[29]. However, the mechanisms underlying ZA regulates osteoclast
262 function in the occurrence of BRONJ remains unclear. In the current study, we clarified that ZA
263 promotes osteoclasts ferroptosis by inhibiting FBXO9-mediated p53 ubiquitination and
264 degradation, as evidence by (I) ZA treatment facilitated the ferroptosis of osteoclasts; (II) FBXO9
265 was downregulated in osteoclasts after ZA treatment; (III) FBXO9 inhibition facilitated the
266 ferroptosis of osteoclasts; (IV) ZA treatment facilitated the ferroptosis of osteoclasts by
267 suppressing FBXO9;(V) FBXO9 inhibition facilitated the ferroptosis of osteoclasts by blocking
268 the ubiquitin mediated-proteasome degradation of p53.

269 Although a growing body of research have explored the role of BPs in the pathogenesis of
270 BRONJ, the mechanism of BPs on the development of BRONJ is not completely understood.
271 Growing studies have demonstrated that BPs have high affinity to hydroxyapatite crystals, thereby
272 suppressing the osteoclasts resorptive ability by inducing the apoptosis of osteoclasts[30, 31].
273 Moreover, due to the lack of cytokines released by osteoclasts, the differentiation of osteoblasts

274 was blocked, thus suppressing the healing ability of bone, suggesting that the differentiation of
275 osteoclasts plays an important role in the development of BRONJ[32]. More recently, ZA has
276 been reported to inhibit osteoclast differentiation by regulating the NF- κ B and JNK signaling
277 pathways[29]. Another study has shown that ZA inhibits osteoclast differentiation by interrupting
278 RANKL/RANK pathway[33]. Consistent with previous studies, we confirmed that ZA decreased
279 cell viability of osteoclasts induced by RANKL, specifically ZA-induced cell viability decrease
280 was blocked by ferroptosis inhibitor, suggesting an important role of ferroptosis in the
281 development of BRONJ.

282 Ferroptosis is a kind of iron- and ROS-dependent form of cell death, different with necrosis,
283 apoptosis, and other forms of cell death. Right now, almost all the mechanisms of ferroptosis are
284 associated with reactive oxygen species (ROS)[12]. Given the role of ZA in regulating ROS
285 production [22-24], here we investigated whether ZA suppresses the growth of osteoclast by
286 accelerating ferroptosis. We found that the ferroptosis-related marker such as the levels of Fe²⁺,
287 MDA content, ROS level was obviously increased in the osteoclasts treated with ZA, suggesting
288 the ZA induced the ferroptosis of osteoclasts. However, the underlying mechanism of ZA-induced
289 osteoclast ferroptosis remains unknown.

290 To elucidate the mechanism of ZA-induced osteoclast ferroptosis, we compared the
291 expression profiles of osteoclasts in the presence or absence of alendronate or risedronate
292 treatment, and got 18 genes with significant differences in the osteoclasts treated by BPs. Among
293 these 18 genes, FBXO9 was identified to be significantly reduced in ZA-treated osteoclasts.
294 Further experiment showed that FBXO9 knockdown promoted the ferroptosis of osteoclasts, and

295 the ferroptosis of osteocalsts induced by ZA was blocked by FBXO9 overexpression, suggesting
296 that ZA promotes the ferroptosis of osteoclasts by downregulating the expression of FBXO9.

297 The F-box only protein 9 (FBXO9), a member of the F-box protein family, is the substrate
298 recognition subunit of skp1-cullin1-f-box E3 ligase complex and plays a key role in ubiquitination
299 and subsequent target protein degradation[19]. Liu *et al.* demonstrated that FBXO9 interacted with
300 Neurog2 and promoted its destabilization is a major contributor in directing multipotent NC
301 progenitors toward glial lineage [20]. Vanesa Fernández-Sáiz *et al.* demonstrated that, under the
302 growth factor deprivation condition, FBXO9-mediated ubiquitination of Tel2 and Tti1 inactivated
303 mTORC1, but activated the PI3K/Akt pathway to increase survival of multiple myeloma[21].
304 However, the function of FBXO9 in the development of BRONJ and the regulatory mechanism
305 remain unclear. Growing studies suggested that E3 ubiquitin ligase regulates ferroptosis by
306 degrading substrates. Yang *et al.* reported that Nedd4 ubiquitylates VDAC2/3 to suppress erastin-
307 induced ferroptosis in melanoma[26]. Another study showed that TRIM26 facilitates the
308 ferroptosis of HSCs to suppress liver fibrosis by mediating the ubiquitination of SLC7A11[35].
309 Therefore, we speculated whether FBXO9 also regulates ferroptosis by mediating the
310 ubiquitination of target genes. Interestingly, we found that p53, a key upstream regulator of
311 ferroptosis, is one of the FBXO9 targets. Our data showed that FBXO9-knockdown did not change
312 the p53 mRNA level but significantly increased the p53 protein level, suggesting that FBXO9-
313 mediated p53 expression by the ubiquitin-proteasome system. Further experiment showed that
314 FBXO9 directly interacts with p53 and the ubiquitination level of p53 was downregulated by
315 FBXO9 knockdown. In addition, the protein stability of p53 was promoted by FBXO9 knockdown.

316 These data suggesting that p53 is the direct target of FBXO9 and FBXO9-mediated p53
317 ubiquitination and degradation in osteoclast.

318 **Conclusions**

319 Taken together, the current data demonstrated that FBXO9 was downregulated in ZA-treated
320 osteoclast and promoted osteoclasts ferroptosis by inhibiting FBXO9-mediated p53 ubiquitination
321 and degradation. Our study provided a possible theoretical target for the clinical treatment of
322 BRONJ.

323 There are still some deficiencies in the current research, such as the current conclusions still need
324 to be further confirmed by clinical and animal experiments.

325 **Conflicts of Interest**

326 The authors declare no competing or financial interests.

327

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395

396 **Figure legends**

397 **Figure 1. ZA treatment facilitated the ferroptosis of osteoclasts**

398 The osteoclasts cell model induced by RANKL (50 ng/ml) treatment. (A and B) Multinucleated
399 cells were visualized by tartrate-resistant acid phosphatase (TRAP) staining.(C and D) Cell
400 viability of Raw264.7 and BMDM derived osteoclasts was assessed using CCK8 assay after
401 treatment with different concentrations of ZA (5,10, and 50 μ M) (n = 3). (E and F) Cell viability
402 of Raw264.7 and BMDM derived osteoclasts was assessed using FDA staining after treatment
403 with different concentrations of ZA (5,10, and 50 μ M) (n = 3). (G and H) Cell viability of
404 Raw264.7 and BMDM derived osteoclasts was assessed using CCK8 assay after treatment with
405 ZA for 48h (50 μ M) in the presence or absence of 10 μ M of ZVAD-FMK, 2 μ M of Fer-1, or 10 μ M
406 of necrostatin-1(n = 3). * p <0.05, ** p <0.01.

407 **Figure 2. ZA treatment facilitated the ferroptosis of osteoclasts**

408 (A-E) the level of Fe²⁺, MDA content, ROS level, the level of Gpx4, and GSH content in Raw264.7
409 derived osteoclasts was assessed by Elisa assay after treatment with different concentrations of ZA
410 (5,10, and 50 μ M) (n=3) * p <0.05, ** p <0.01, *** p <0.001. (F-J) the level of Fe²⁺, MDA content,

411 ROS level, the level of Gpx4, and GSH content in BMDM derived osteoclasts was assessed by
412 Elisa assay after treatment with different concentrations of ZA (5,10, and 50 μ M) (n=3) * p <0.05,
413 ** p <0.01, *** p <0.001.

414 **Figure 3. FBXO9 was downregulated in osteoclasts after ZA treatment**

415 (A) Venn analysis of DEGs of alendronate and risedronate-treated osteoclast. (B and C) The
416 mRNA level of 18 genes in Raw264.7 and BMDM derived osteoclasts was assessed using qPCR
417 after treatment with ZA (50 μ M) (n=3) * p <0.05, ** p <0.01. (D) The protein level of FBXO9 in
418 Raw264.7 and BMDM derived osteoclasts was assessed using western blot after treatment with
419 ZA (50 μ M)

420 **Figure 4. FBXO9 inhibition facilitated the ferroptosis of osteoclasts**

421 (A) The mRNA level of FBXO9 in BMDM derived osteoclasts was assessed using qPCR after
422 treatment with or without si-FBXO9 (n=3). *** p <0.001. (B) The protein level of FBXO9 in
423 BMDM derived osteoclasts was assessed using western blot after treatment with or without si-
424 FBXO9 (n=3). ** p <0.01. (C) Cell viability of BMDM derived osteoclasts was assessed using
425 CCK8 assay after treatment with or without si-FBXO9 (n=3). ** p <0.01. (D and E) Cell viability
426 of BMDM derived osteoclasts was assessed using FDA staining after treatment with or without si-
427 FBXO9 (n=3). * p <0.05. (F-J) the level of Fe²⁺, MDA content, ROS level, the level of Gpx4, and
428 GSH content in BMDM derived osteoclasts was assessed by Elisa assay after treatment with or
429 without si-FBXO9 (n = 3) * p <0.05, ** p <0.01.

430 **Figure 5. ZA treatment facilitated the ferroptosis of osteoclasts by suppressing FBXO9**

431 (A) The mRNA level of FBXO9 in BMDM derived osteoclasts was assessed using qPCR after

432 treatment with ZA (50 μ M) in the presence or absence of FBXO9 (n=3). ** p <0.01. (B) The protein
433 level of FBXO9 in BMDM derived osteoclasts was assessed using western blot after treatment
434 with ZA (50 μ M) in the presence or absence of FBXO9 (n=3). * p <0.05. (C) Cell viability of
435 BMDM derived osteoclasts was assessed using CCK8 assay after treatment with ZA (50 μ M) in
436 the presence or absence of FBXO9 (n=3). * p <0.05, ** p <0.01. (D and E) Cell viability of BMDM
437 derived osteoclasts was assessed using FDA staining after treatment with ZA (50 μ M) in the
438 presence or absence of FBXO9 (n=3). * p <0.05. (F-J) the level of Fe²⁺, MDA content, ROS level,
439 the level of Gpx4, and GSH content in BMDM derived osteoclasts was assessed by Elisa assay
440 after treatment with ZA (50 μ M) in the presence or absence of FBXO9 (n = 3) * p <0.05, ** p <0.01.

441 **Figure 6. FBXO9 inhibition facilitated the ferroptosis of osteoclasts by blocking the ubiquitin**
442 **mediated-proteasome degradation of p53**

443 (A) The target of FBXO9 was predicted by ubibrowser. (B) the p53 mRNA expression in the
444 FBXO9 knockdown and control cell was assessed by qPCR (n=3). (C) the protein level of p53 in
445 the FBXO9 knockdown and control cell was assessed by western blot (n=3). (D) FBXO9 directly
446 interacts with p53. The proteins from BMDM derived osteoclasts were IP with IgG or antibodies
447 against FBXO9 and p53, following by western blot analysis (n=3). (E) The stability of p53 protein
448 was regulated by FBXO9. BMDM derived osteoclasts treated with or without si-FBXO9 in the
449 presence of cycloheximide (CHX, 25 ug/ml) for various times as indicated and cell lysates were
450 then assessed by western blot (n=3). ** p <0.01. (F) The cell lysates isolated from scramble and si-
451 FBXO9 infected BMDM derived osteoclasts were immunoprecipitated with anti-p53 antibody,
452 then analyzed by western blot using ubiquitin antibody (n=3).

Table 1 (on next page)

sequence

sequence

- 1 CFAP53:
- 2 forward primer:5'-GACAAAATGAGAGAGAGAACCAAGT-3'
- 3 reverse primer: 5'-TCCCTGAACTGCTGGTCTAAC-3'
- 4 COL14A1
- 5 forward primer:5'-ACTGGTTTTTCACGGGTGTTC-3',
- 6 reverse primer: 5'-TAAGTCGAGGAGAGGCAAGC-3'
- 7 ARSJ
- 8 forward primer:5'- CTGAGATAAAGACGCCCACC-3',
- 9 reverse primer: 5'- ATAGAATGCTGAAGTCCCGTG-3'
- 10 ABCA9
- 11 forward primer:5'- CAGAGGGAGTGAAGAGAAAGC -3',
- 12 reverse primer: 5'- GCTCTGTGTTTGTGAAAGTGG -3'
- 13 CXorf57:
- 14 forward primer:5'- GCAGTATAGGGAACAAAAGCG-3',
- 15 reverse primer: 5'- TGCTTGAGATGTTGAGGGAC-3'
- 16 GPR22:
- 17 forward primer:5'- CCACTGTCATACCCACTAAGC-3',
- 18 reverse primer: 5'- ATGCAGTAAAGTACCAGGACG-3'
- 19 STXBP5L:
- 20 forward primer:5'- GATCAAGTGACCTGTACCAGC-3',
- 21 reverse primer: 5'- ATTTACATGGTCTGAGGTGGG-3'
- 22 MSANTD4
- 23 forward primer:5'- CAGAGGTCAAAGTGGAAAGAGG-3',
- 24 reverse primer: 5'- ATCAATGTGAGGGAAGTCAGG-3'
- 25 RRP15
- 26 forward primer:5'- GAAATGCTGTGCAGAGTGAAG-3',
- 27 reverse primer: 5'- TCCTGCTTCCTTAACCTTTTCG-3'
- 28 UGT1A2
- 29 forward primer:5'- TCTGCGTTCTCTTTCCTGTG -3',
- 30 reverse primer: 5'- AGCATGTTCTGGACCCTTG -3'
- 31 IRF4
- 32 forward primer:5'- AACAAAGCTAGAAAGAGACCAGAC-3',
- 33 reverse primer: 5'- TCACCAAAGCACAGAGTCAC-3'
- 34 TFAP2D
- 35 forward primer:5'- AAAGATGATCCTAGCCACCAAG-3',
- 36 reverse primer: 5'- TGTGTTAAGTGCCTCTGGATG-3'
- 37 TRHDE
- 38 forward primer:5'- AGGAAGGCTTTGCTCACTAC -3',
- 39 reverse primer: 5'- CTGTGATACTGGATGGGAACTG -3'
- 40 ASMT
- 41 forward primer:5'- GAAGTGGGACAGGAAGTGAG -3',

42 reverse primer: 5'- CGGGAACAGGAAGTGGC -3'
43 CAPS
44 forward primer:5'- AGCTCGAAGACACAATCCG -3',
45 reverse primer: 5'- TCCATGTCCACTGCAAAGAG -3'
46 COMMD10
47 forward primer:5'- AGTGGGATGGCAGCTTAAC-3',
48 reverse primer: 5'- TCGAACAGCTCCTTGTGATTG-3'
49 VSTM4
50 forward primer:5'- CCTGGCAGTCTGTGTTTCA-3',
51 reverse primer: 5'- CTCTTACCCTTCTGTGGCTG-3'
52 FBXO9
53 forward primer:5'- ATGAGAGTCCGGCTGAGAGA-3',
54 reverse primer: 5'- AGAGCTTCTTCCTGCTCTGC-3'
55 18s
56 forward primer:5'- CTCAACACGGGAAACCTCAC-3',
57 reverse primer: 5'- CGCTCCACCAACTAAGAACG-3'

Figure 1

ZA treatment facilitated the ferroptosis of osteoclasts

ZA treatment facilitated the ferroptosis of osteoclasts

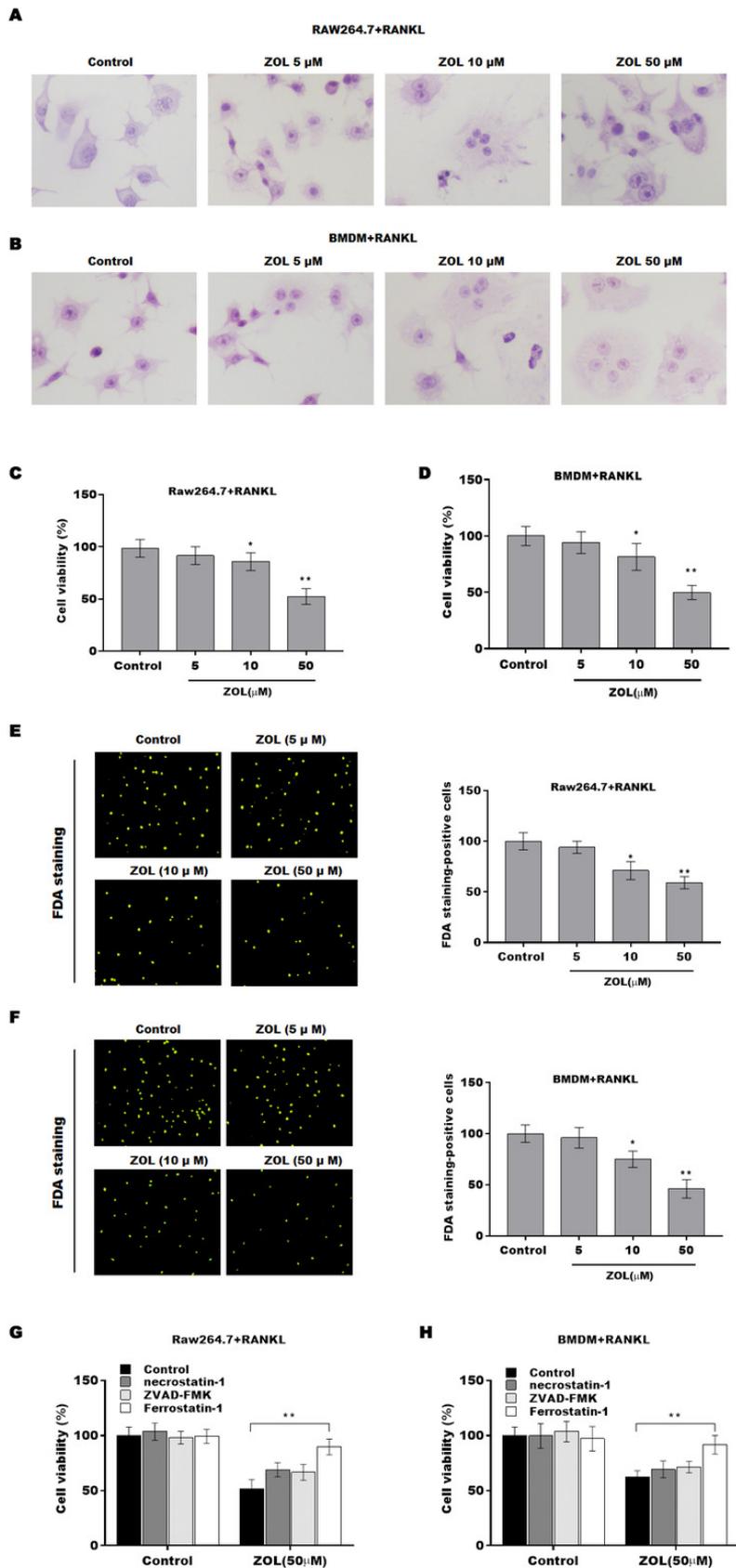


Figure 2

ZA treatment facilitated the ferroptosis of osteoclasts

ZA treatment facilitated the ferroptosis of osteoclasts

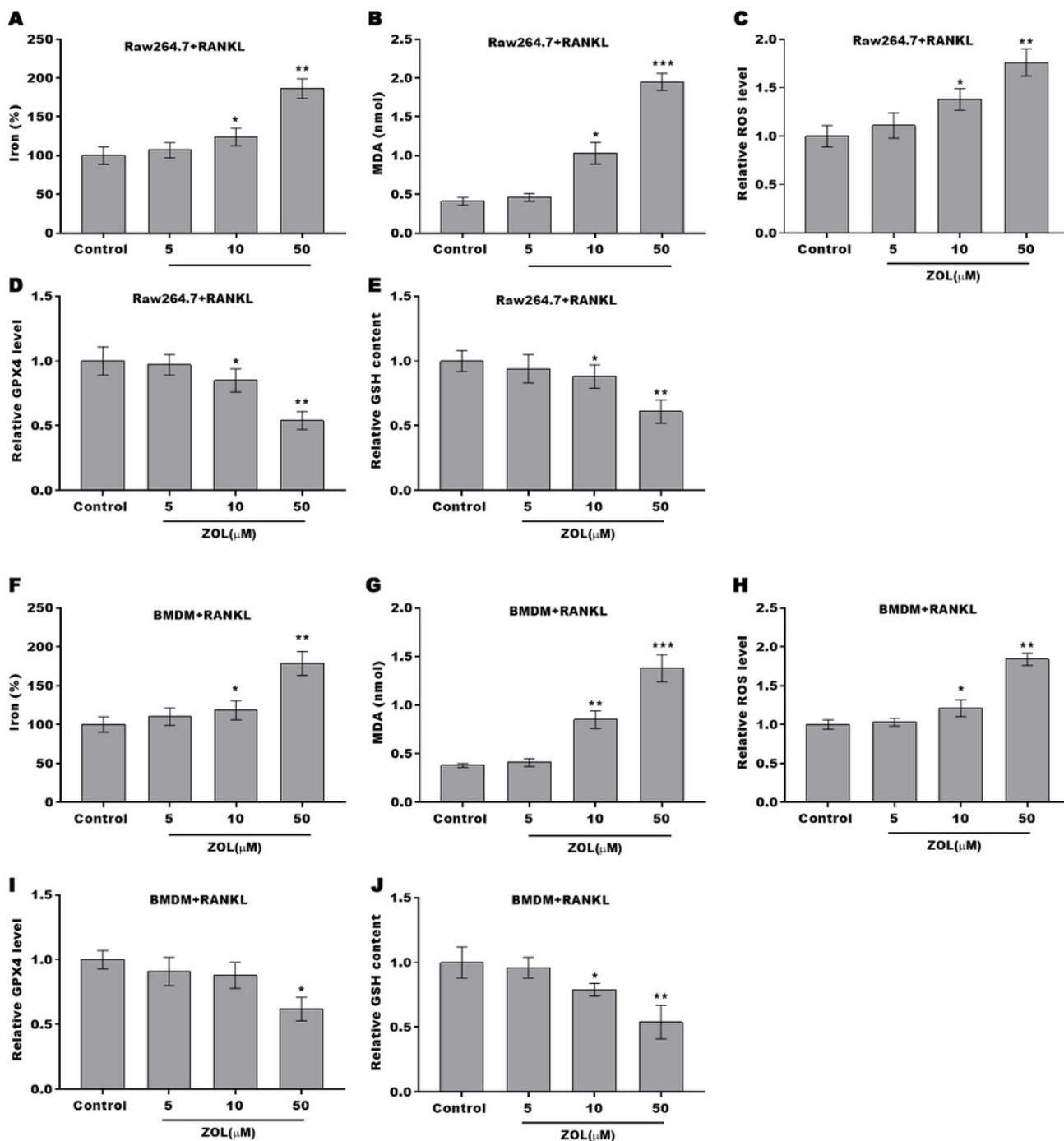


Figure 3

FBX09 was downregulated in osteoclasts after ZA treatment

FBX09 was downregulated in osteoclasts after ZA treatment

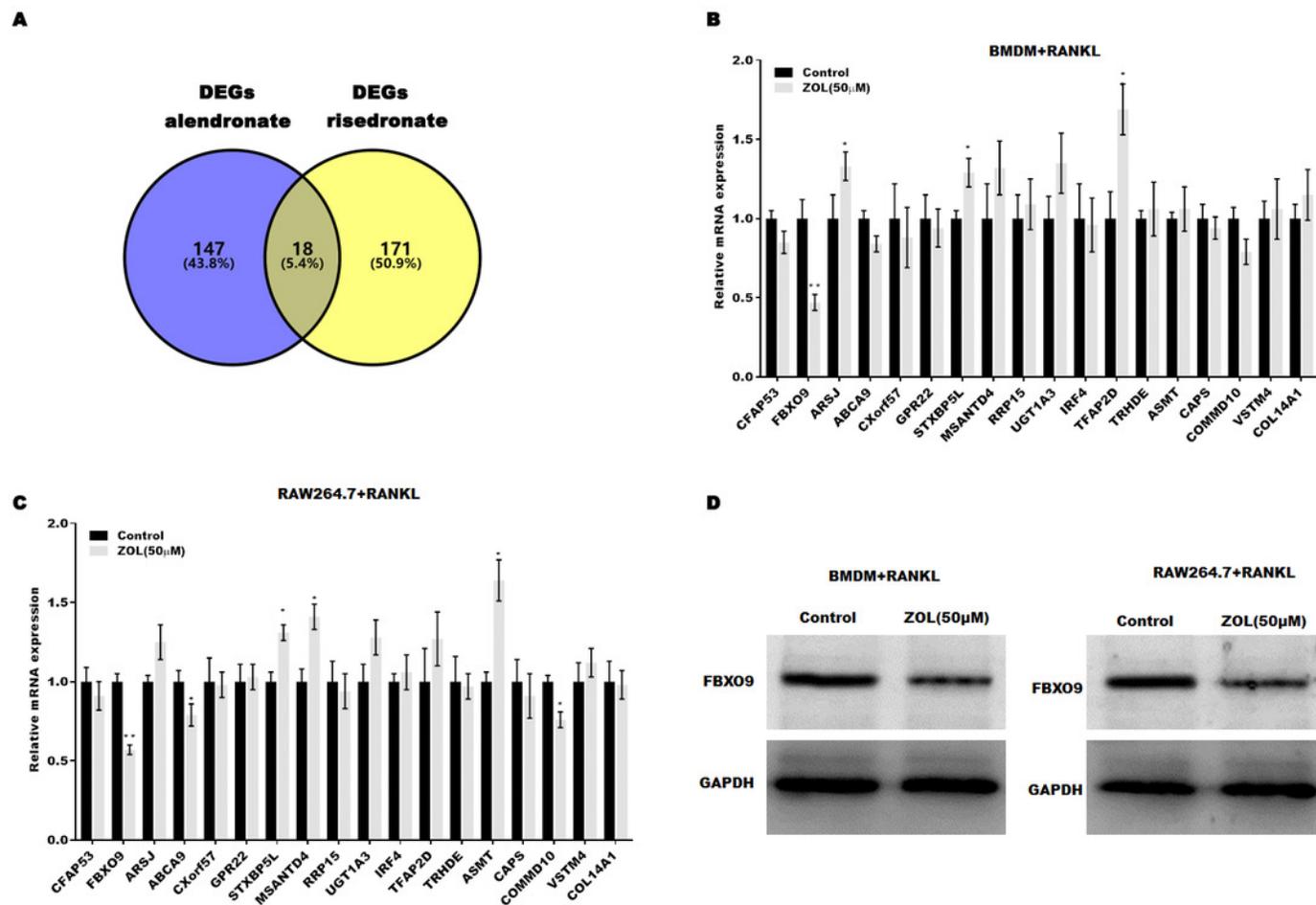


Figure 4

FBXO9 inhibition facilitated the ferroptosis of osteoclasts

FBXO9 inhibition facilitated the ferroptosis of osteoclasts

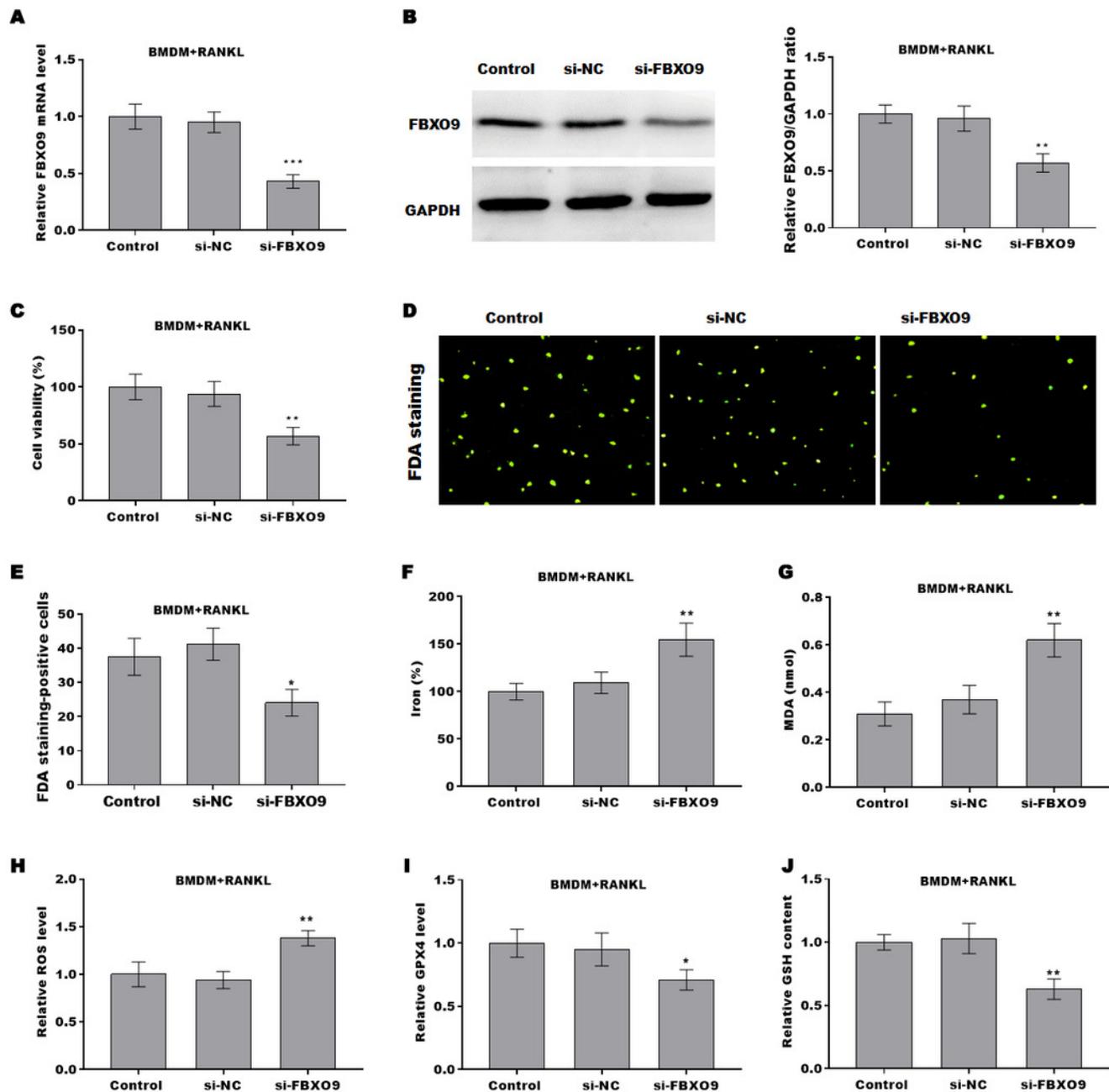


Figure 5

ZA treatment facilitated the ferroptosis of osteoclasts by suppressing FBXO9

ZA treatment facilitated the ferroptosis of osteoclasts by suppressing FBXO9

