

1 **A small molecule, Odanacatib, inhibits inflammation and bone loss caused**
2 **by endodontic disease**

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25 **Running Head:** Inhibition of Ctsk prevents periapical tissue damage

26

27 **Abstract**

28 Periapical disease, an inflammation disease mainly caused by dental caries, is
29 one of the most prevalent infectious diseases of mankind, affecting both children
30 and adults. The infection travels through the root leading to inflammation, bone
31 destruction, and severe pain for the patients. Therefore, the development of a
32 new class of anti-periapical disease therapies is necessary and critical for
33 treatment and prevention. A small molecule, Odanacatib (ODN), which is a
34 Cathepsin K (Ctsk) inhibitor, was investigated to determine its ability to treat this
35 disease in a mouse model of periapical disease. While Ctsk was originally found
36 in osteoclasts as an osteoclast-specific lysosomal protease, we were surprised to
37 find that ODN can suppress bacteria-induced immune response as well as bone
38 destruction in the lesion area. The X-ray and MicroCT (μ CT) showed that ODN
39 treatment had significant bone protection effects at different time points.
40 Immunohistochemical and immunofluorescent staining show that ODN treatment
41 dramatically decreased F4/80⁺ macrophages and CD3⁺ T-cells in the lesion
42 areas 42 days after infection. Consistent with these findings, qRT-PCR and
43 ELISA analysis showed low levels of pro-inflammatory mRNA (*Tumor necrosis α* ,
44 *Interleukin 6* and *Interleukin 23 α*) and correspondent cytokines expression in
45 ODN-treated disease group. The mRNA level of *Toll-like receptor 4, 5, and 9* also

46 largely decreased in the ODN-treated disease group. Our results demonstrated
47 that ODN can inhibit endodontic disease development, bone erosion, and
48 immune response. These results indicate that application of this small molecule
49 offers a new opportunity to design effective therapies that could prevent
50 periapical inflammation and revolutionize current treatment options.

51

52 **Keywords**

53 Endodontic disease, Cathepsin K, Inflammation, Bone resorption, Dental caries,
54 Osteoclast

55

56 **Introduction**

57 Periapical lesions result from microbial infection of the dental pulp tissue by
58 autogenous oral microflora, which induce inflammation in the pulp tissue (1). This
59 inflammatory process, much like periodontitis, increases in magnitude in the
60 apical region of the root canal system and subsequently in the periapical area,
61 leading to periapical bone resorption as the infection spreads throughout the
62 canal system towards the apical foramen and into the adjacent bone (2).
63 Furthermore, periapical lesions that present with radiographic bone lesions are
64 preceded by necrotic pulp tissues. Unlike periodontal disease inflammation (3, 4),
65 endodontic lesions exhibit differences in the character of the immune response
66 (5).

67

68 Despite many studies aimed to alleviate the effects of oral disease, there is still
69 an urgent need to improve the health of millions with periapical disease who
70 suffer from oral bacterial infection-induced periapical inflammation, oral bone
71 erosion and the potential loss of teeth. The discovery of the critical roles of
72 Cathepsin K (Ctsk) in osteoclastic bone resorption is the fruit of decades of
73 investigation on bone biology and disease (6, 7). Ctsk is the only gene for which
74 an essential role in bone resorption has been clearly demonstrated in both mice
75 and humans, and current research data shows that Ctsk is also indispensable in
76 the immune system (8). Recent studies demonstrated that cathepsins are
77 required for toll-like receptor 9 expressed in dendritic cells, which plays an
78 essential role in innate recognition of microbial products and activation of
79 defense responses (9). Our research led to the surprising and pivotal realization
80 that Ctsk also has significant functions in the immune system (10, 11), so we
81 termed Ctsk as an “osteimmune gene”. The study of bone and the immune
82 system, which are termed osteoimmunology, represents a new perspective
83 reconsidering the multiple biological events in bone and the immune system in
84 recent years (12, 13).

85

86 The success of the novel treatment of inflammatory diseases like periodontitis
87 and rheumatoid arthritis (RA) has impressively demonstrated the clinical benefit
88 can be gained from therapeutic intervention of specific protein by using small
89 molecules (14). Odanacatib (ODN) is a potent, orally active selective inhibitor of
90 Ctsk being developed for the treatment of postmenopausal osteoporosis.

91 Treatment with ODN decreases bone resorption by selectively inhibiting
92 proteolysis of matrix protein by Ctsk without affecting other osteoclast activities or
93 osteoclast viability (15). However, there is still no research activity using Ctsk
94 inhibitors applied for oral inflammatory diseases considering its dual functions in
95 bone and immune system. In addition, according to the knowledge of the
96 investigators of current studies, there are still questions of what functions Ctsk
97 has in activating immune cells, such as macrophages or dendritic cells (Antigen-
98 presenting cells, APCs), and what leads to T cell activation and autoimmunity
99 causing the subsequent production of inflammatory cytokines. Therefore, based
100 on the cited research and our previous results, the current investigation was
101 conducted with respect to the likelihood that Ctsk has a significant role in the
102 activation of macrophages and TLRs-mediated innate and adaptive immune
103 responses in the pathogenesis of periapical lesions by using a specific Ctsk small
104 molecule inhibitor ODN.

105

106 **Materials and Methods**

107 **Animals.** 105 seven- to eight-week-old male wild-type (WT) BALB/cJ mice,
108 purchased from the Jackson Laboratory were used for introducing a periapical
109 mouse model. Mice were divided into 4 groups at each time point (7, 21 and 42
110 days): (1) normal group with Ctsk inhibitor (no bacterial infection) (3.606
111 mg/kg/wk, n=7 mice); (2) normal group without Ctsk inhibitor (no bacterial
112 infection) (n=7 mice); (3) bacterial infection group treated with Ctsk inhibitor
113 (3.606 mg/kg/wk, n=7 mice) or (0.7212 mg/kg/wk, n=7 mice); (4) bacterial

114 infection group without Ctsk inhibitor (n=7 mice). The animals were maintained in
115 the University of Alabama at Birmingham animal facility and were given distilled
116 water and allowed to feed freely. All experimental protocols were approved by
117 the NIH and the Institutional Animal Care and Use Committee of the University of
118 Alabama at Birmingham and completed within 16 weeks after birth. Approval for
119 the animal protocol related to this study (Animal Protocol Number 131209236)
120 was renewed by the University of Alabama at Birmingham (UAB) Institutional
121 Animal Care and Use Committee (IACUC) on August 27, 2013.

122

123 **Administration of Cathepsin K (Ctsk) inhibitor.** Male mice in the normal and
124 periapical disease group were treated with ODN and compared with the
125 untreated normal and disease groups. Treated mice were administrated orally at
126 3.606 mg/kg/wk or 0.7212 mg/kg/wk (5-fold lower dose) of pharmacological
127 grade ODN (selleckChem, USA) in dimethyl sulfoxide (DMSO) (Sigma-Aldrich)
128 from one week before the establishment of the disease model to the end of
129 sample harvest. The ODN dose was chosen to match the treatment of human by
130 convention to a proper dose for mouse (16). The control mice were administrated
131 orally with DMSO.

132

133 **Pulp exposure and bacterial infection.** The left and right first molar pulp
134 exposure was performed as described (10, 17). In brief, mice were anesthetized
135 with peritoneal injection of 62.5mg/kg ketamine and 12.5mg/kg xylazine according
136 to body weight. The dental pulps of the mandibular first molars were exposed

137 with ¼ round carbide bur powered by a variable speed electric rotary hand piece
138 (Osada Electric, Los Angeles, CA) under a surgical microscope (model MC-M92;
139 Seiler, St Louis, MO). After removing the roof of the pulp chamber, the size of the
140 exposure was approximately 0.5-1.0mm in diameter. We used stainless steel
141 hand files #8 (Dentsply/Maillefer, Johnson City, TN) and stainless steel rotary
142 files #15 (Dentsply/Maillefer, Johnson City, TN) to establish canal patency.
143 Eradication of infected microorganisms was not completed in this therapy.

144

145 Bacterial culture and infection procedure protocols were conducted as described
146 (10, 17). In brief, exposed pulps were infected with a mixture of four common
147 human endodontic pathogens, including *Prevotella intermedia* [American Type
148 Culture Collection (ATCC) 25611; Manassas, VA], *Fusobacterium nucleatum*
149 (ATCC 25586), *Peptostreptococcus micros* (ATCC 33270), and *Streptococcus*
150 *intermedius* (ATCC 27335). All four species of bacteria were cultured under strict
151 anaerobic conditions (80% N₂, 10% H₂, and 10% CO₂), inoculated to broth, and
152 cultivated over seven days. Microbes were harvested, resuspended, and
153 evaluated for cell concentration of each species via optical density reading. The
154 wave length used for cell number counting is 600nm (One OD unite equals
155 6.67*10⁸ particles), the cell density per ml can be obtained by calculation. Then
156 the cell density of each species was adjusted to 10¹⁰ per ml for bacterial
157 inoculation. The four organisms were then mixed for a total of 10¹⁰ cells of each
158 bacterial species/ml PBS in 3% methylcellulose. Ten microliters of the
159 polymicrobial solution was placed inside the access opening of each molar and

160 carried to the periapical tissues using a #8 endodontic file. After exposed pulps
161 were infected with the mixture of four common human endodontic pathogens,
162 access openings were left open to the oral environment for 24 hours. All
163 exposure sites were then filled with self-cure composite temporary filling.

164

165 **Harvest and preparation of samples.** Animals were sacrificed by CO₂
166 inhalation on day 7, 21 and 42 after the initial infection. The mandibles were
167 removed and hemisected. After removal of soft tissue, the jaw samples from the
168 left side were fixed in 4% formaldehyde for 24 hours then stored in 70% ethanol
169 prior to X-ray and μ CT exposure and measurement analysis of bone loss. The
170 jaw samples from the right side were fixed in 4% paraformaldehyde and prepared
171 for histological analysis. The right sections were prepared for histology analysis
172 according to standard protocol with modification for samples that were prepared
173 for paraffin sectioning. In brief, samples specified for paraffin section were fixed
174 in 4% formaldehyde for 24 hours, washed with PBS, and decalcified in 10%
175 EDTA for 25 days (EDTA replenished each day). For the samples applied for
176 RNA and protein extraction, the periapical bone tissues surrounding the mesial
177 and distal root were extracted from the mandible together with surrounding bone
178 in a block specimen using a surgical microscope for total RNA and cytokine
179 enzyme linked immunosorbent assays (ELISA). For the RNA extraction,
180 periapical tissues were rinsed in chilled PBS, weighed (3-5mg/tissue), and
181 immediately put into RNA $\text{later}^{\text{®}}$ ICE (Invitrogen, USA) overnight at 4 °C and then
182 stored at -80 °C until applied for RNA extraction. For ELISA assay, the periapical

183 tissues were rinsed in chilled PBS, weighed (3-5mg/tissue) and immediately
184 frozen at -80 °C until applied for protein extraction.

185

186 **Micro-computed tomography (μCT) analysis.** Micro-computed tomography
187 (μCT) scans were evaluated for bone loss as described with modification (10,
188 17). Briefly, the two-dimensional picture that showed the largest target root canal
189 was selected as the baseline image. For each sample, an approximate total of
190 101 micro-tomographic slices with an increment of 12 μm were acquired in front
191 of and behind the baseline image. From the three-dimensional stack of μCT
192 images, a “pivot” section was the periapical area including root area and furcation
193 area, where the inflammation may cause the bone resorption around the apex of
194 the tooth. The segmentation parameter is 0.7/1/210.

195

196 **Histological analysis.** Samples were fixed by 4% paraformaldehyde for 24
197 hours, decalcified by 10% EDTA for 25 days with a new solution each day,
198 dehydrated by 50%, 70%, 95%, 100% alcohol for 2 hours each, washed in
199 paraffin acetone for 30 min in a 60 °C incubator, washed in chloroform for 30 min
200 in 60 °C incubator, infiltrated with paraffin in a 60 °C incubator 3x for 1hr each
201 and embedded in paraffin. To detect bone-resorbing osteoclasts, Tartrate-
202 resistant acid phosphatase (TRAP) staining was performed. Tissue sections
203 were de-paraffinized and hydrated through xylenes and graded alcohol series,
204 preincubated with 50 mM sodium acetate and 40 mM potassium sodium tartrate
205 buffer for 20 min, and incubated with TRAP substrate solution.

206

207 **Immunofluorescence analysis.** We performed immunofluorescence analysis as
208 we have previously described (18), with the exception that we used rat polyclonal
209 anti-CD3 (Abcam, Cambridge, MA) as the primary antibodies and observations
210 were performed by epifluorescence in a Zeiss Axioplan microscope (Carl Zeiss
211 Microscopy, LLC, USA). Nuclei were visualized with 1 µg/ml DAPI (4',6-
212 diamidino-2-phenylindole) (Sigma Aldrich, USA). The experiments were set in
213 triplicate on three independent occasions.

214

215 **Immunohistochemistry analysis.** We performed immunohistochemistry using
216 corresponding sections of mandibular tissue from the normal group (with or
217 without inhibitor) and the disease group (with or without inhibitor), then the slides
218 were analyzed by immunohistochemistry for expression and localization of
219 proteins of the macrophage marker F4/80 (Rat monoclonal, 1:200) (eBioscience,
220 San Diego, CA), TLR4 (Goat polyclonal, 1:200), TLR5 (Rabbit polyclonal, 1:200)
221 (Santa Cruz, USA) and TLR9 (Rabbit monoclonal, 1:500) (Sigma-Aldrich, USA).
222 Vector stain ABC kit Anti-Rat, Goat and Rabbit IgG Peroxidase Polymer
223 Detection Systems along with a DAB kit (Vector Laboratories, Burlingame, CA)
224 as a substrate were used for the peroxidase-mediated reaction. The experiments
225 were set in triplicate on three independent occasions.

226

227 **RNA extraction and Real-time quantitative PCR (qRT-PCR).** For RNA
228 extraction, the prepared samples were transferred to the tube pre-filled with

229 beads (Nextadvance Company, USA) and homogenized using a Blender (Bullet
230 Blender[®], Nextadvance Company, USA). The RNA extraction was performed
231 with the standard procedure using Trizol reagent (Invitrogen, USA). The
232 extracted RNA was applied for reverse transcription using Vilo[®] master kit
233 (Invitrogen). Real-Time quantitative PCR (qPCR) was performed as described
234 (10, 17) using primers purchased from Invitrogen as listed (**see Table S1 in**
235 **supplemental material**). Briefly, cDNA fragments were amplified by Sybrgreen[®]
236 Fast Advanced Master Mix (Applied Biosystems, Foster City) and detected by a
237 Step-One real-time PCR system (Applied Biosystems, Foster City). The mRNA
238 expression level of the housekeeping gene *β-actin* was used as an endogenous
239 control and enabled calculation of specific mRNA expression levels as a ratio of
240 *β-actin*. Experiments were repeated at least three times.

241

242 **Protein extraction and cytokine for Enzyme-linked immunosorbent assay**
243 **(ELISA)**. For protein extraction, the frozen periapical tissue samples were
244 transferred to a tube prefilled with beads (Nextadvance Company, USA) and
245 homogenized using a Blender (Bullet Blender[®], Nextadvance Company, USA).
246 The tissue fragments were dispersed in 300ul of lysis buffer. The mixture was
247 centrifuged at 1,2000 RPM for 10 min, and the supernatant was collected and
248 stored at -80 °C until assay. ELISA was used as described (19, 20) to evaluate
249 the effect of inhibition of Ctsk on the levels of TNF- α , IL-6 and IL-23 α in
250 inflammatory periapical tissues extracted from the normal group (with or without
251 inhibitor) and the disease group (with or without inhibitor). Briefly, assays for

252 cytokines in extracts employed commercially available ELISA kits that are
253 obtained from the following sources: TNF- α (eBioscience, USA), IL-6
254 (eBioscience, USA) and IL-23 α (Biolegend, USA). All assays were conducted in
255 accordance with the manufacturer's instructions. Results were expressed as pg
256 cytokine/ml tissue.

257

258 **Statistical analysis and data quantification analysis.** Experimental data are
259 reported as mean \pm SD of triplicate independent samples. All experiments were
260 performed in triplicate on three independent occasions. The figures are
261 representative of the data. Data were analyzed with the two-tailed Student's t-test.
262 Mann-Whitney U test was used for the non-parametric test. P values <0.05 or U
263 values > 1.96 were considered significant. Data quantification analysis were
264 performed using the NIH Image J Program as described (10, 17).

265

266 **Results**

267 **Inhibition of Ctsk exhibits bone protective effects of the endodontic lesion**
268 **area at different time points.** Our previous study on AAV-mediated Ctsk
269 inhibition in our periapical lesion mouse model revealed that Ctsk is important for
270 bone resorption and inflammation in periapical lesions, but not for exploring the
271 possible mechanism (10). Thus, it is important to determine the trend change of
272 osteoclasts and inflammation in the periapical lesion area by inhibition of Ctsk
273 and its possible mechanism at different time points. To test the possible role
274 during the progress of periapical lesions mediated by Ctsk, we applied the

275 periapical mouse model established by our previous study (10, 17). Samples
276 from the normal group with or without inhibitor (only images of day 42 showed in
277 **Fig. 1**) and from the bacterial infected group with or without inhibitor (3.606
278 mg/kg/wk) were analyzed by X-ray and μ CT at days 7, 21 and 42 after infection
279 (**Fig. 1**). X-ray imaging of the first molar tooth of the normal group with and
280 without inhibitor at different time points showed that there was no reduced bone
281 density area (**see Fig. S1 in the supplemental material**). The X-ray and μ CT
282 test showed low bone density area at furcation and periapical area in the
283 bacterial infection group without inhibitor at 7, 21 and 42 days (**Fig. 1B-D**). As the
284 time increased, the low density of bone regions in the disease group without
285 inhibitor was widening (red arrows). On the contrary, there was no obvious
286 reduced bone density area in the infected disease group with inhibitor (3.606
287 mg/kg/wk) at different time points (**Fig. 1B-D**). To further quantify the bone
288 resorption in the periapical lesion area, we applied quantification of μ CT to
289 compare the difference between different groups (**Fig. 1E**). The quantification
290 analysis showed a significant difference in the disease group with or without
291 inhibitor at 7, 21 and 42 days (**Fig. 1E**). The different dose of Ctsk inhibitor (3.606
292 mg/kg/wk) or 5-fold lower (0.7212 mg/kg/wk) seemed to have a dose dependent
293 bone protective effect by μ CT analysis (**see Fig. S2 in the supplemental**
294 **material**). We also conducted histological analysis by Hematoxylin and Eosin
295 (H&E) for the periapical lesion area for the different groups (**Fig. 2**). The H&E
296 stain showed that the remaining bone area in mesio, distal and furcation areas in
297 the disease group with inhibitor (3.606 mg/kg/wk) is much higher than that in the

298 disease group without inhibitor at 7, 21 and 42 days (**Fig. 2A and B**). We also
299 found that the distribution of mononuclear inflammatory cells was evident in the
300 disease group without inhibitor at different time points (red arrows) (**Fig. 2A**).
301 Although the bone area decreased in the disease group without inhibitor, there
302 was no significant change at the first 21 days. However, the remaining bone area
303 in the disease group without inhibitor was still significantly less than the inhibitor
304 treated group at 42 days (**Fig. 2C**). There was no significant change in the
305 normal group with or without inhibitor (**see Fig. S3A, C in the supplemental**
306 **material**). Above all, the results showed that Ctsk is critical for bone resorption
307 during the progress of periapical lesions.

308

309 **Inhibition of Ctsk exhibits decreased osteoclast numbers at endodontic**
310 **and furcation areas at different time points.** By applying TRAP staining, we
311 found that the number of osteoclasts decreased in the disease group with
312 inhibitor (3.606 mg/kg/wk) compared to the disease group without inhibitor at
313 different time points (**Fig. 3A, B**). We also noticed that the number of osteoclasts
314 increased significantly in the disease group without inhibitor compared to the
315 inhibitor group at 7 days (**Fig. 3C**), and the osteoclast numbers remained at a
316 high level during the progress of periapical lesion in our current study. There was
317 no significant difference of osteoclast number in the normal group with or without
318 inhibitors at 42 days (**see Fig. S3B, D in the supplemental material**).

319

320 **Ctsk may have effects on immune response and have functions in immune**
321 **cells at the periapical lesion area.** The macrophage is important for the host
322 immune response during the periapical bacterial infection (2). In order to
323 determine whether inhibition of Ctsk has effects on the expression of
324 macrophages, we conducted IHC for F4/80, a specific marker of macrophages
325 (Fig. 4). The results showed that the F4/80 positive macrophages increased
326 significantly in the disease group without inhibitor (Fig. 4C). We also conducted
327 the immunofluorescence analysis for the periapical lesion area for the CD3
328 positive T cells in different groups (Fig. 5). We then merged each set of images
329 for comparison and location (Fig. 5A). These images showed significantly less
330 visible CD3 positive T cells (white arrow) in the disease group with inhibitor
331 (3.606 mg/kg/wk) than that of the disease group without inhibitor, which indicated
332 that inhibition of Ctsk decreased the number of T cells in the periapical lesion
333 area (Fig. 5C).

334

335 **Inhibition of Ctsk reduced the expression of pro-inflammatory and**
336 **osteoclast marker genes as well as TLRs and cytokines in the periapical**
337 **lesion at different time points.** To evaluate the effect of Ctsk inhibition on the
338 mRNA levels of *Tumor Necrosis Factor- α* (*TNF- α*), *Interleukin 6* (*IL-6*), *Interleukin*
339 *6* (*IL-23 α*) and *Cathepsin K* (*Ctsk*), which are related to inflammatory and
340 osteoclast status in periapical tissues, qRT-PCR was used as described (10, 17,
341 19). The relative mRNA expression levels showed that the expression of Ctsk was
342 inhibited at different time points in the inhibitor (3.606 mg/kg/wk) treated disease

343 group, but not in the untreated disease group (**Fig. 6A**). Proinflammatory gene
344 expression also increased in the disease group at 7, 21 and 42 days (**Fig. 6A**).
345 The result of TLRs expression showed that *TLR9* increased significantly starting
346 from day 21 in the disease group (**Fig. 6A**). The expression of *TLR4* and *TLR5*
347 was also detected at different time points since *TLR4* and *TLR5* are responsible
348 for recognition of bacteria pathogen lipopolysaccharide (LPS) and flagellin (21).
349 Although *TLR4* increased at disease group without inhibitor at 7 and 21 days,
350 there was no significant difference between disease group and inhibitor treated
351 group at 42 days (**Fig. 6A**). The IHC stain confirmed the qRT-PCR result (**see**
352 **Fig. S4 in the supplemental material**). The expression of *TLR5* decreased
353 significantly in inhibitor treated disease group starting from 21 days (**Fig. 6A**).
354 IHC stain of *TLR5* showed *TLR5* expression decreased significantly in the
355 inhibitor treated disease group at 42 days (**see Fig. S5 in the supplemental**
356 **material**), which was consistent with the qRT-PCR result. Although the gene
357 expression level of *IL-6* and *IL-23 α* decreased in the disease group treated with
358 or without inhibitor (3.606 mg/kg/wk) at 42 days, cytokine expression was still
359 higher when compared to the normal group. The ELISA result showed that TNF-
360 α , *IL-6* and *IL-23 α* reached the highest levels in the disease group without
361 inhibitor at 42 days (**Fig. 6B**). Since *Ctsk* has been shown to possibly play a
362 critical role in the *TLR9* related pathway (9), we also conducted IHC of *TLR9* to
363 explore whether *Ctsk* affects the *TLR9* expression in periapical disease at 42
364 days after infection. The results showed that *TLR9* was expressed abundantly in

365 the periapical lesion area in the disease group without inhibitor but not in the
366 inhibitor treated group (**Fig. 7**).

367

368 **Discussion**

369 Periapical lesions result from pulp necrosis induced by bacterial infection. The
370 infection produces an inflammatory reaction first within the pulp and,
371 subsequently, in the periapical region after egress of microorganisms, their by-
372 products and/or altered pulp tissues from the infected root canal. The periapical
373 lesion is characterized by the presence of numerous inflammatory cells, which
374 then lead to immune inflammation of periapical tissue. Generally, the body's
375 immune defense system gradually removes the pathogens after the removal of
376 bacterial infections, but during the process of obliterating foreign microorganisms,
377 inflammatory factors produced by the body could also damage the surrounding
378 normal tissue and cause periapical bone tissue absorption. If this process
379 continues, teeth will eventually become loose or fall out. Bone resorption is
380 mediated by bone resorptive cytokines produced by the inflammatory response.
381 Consequently, the question is particularly important of how to control an
382 excessive inflammatory response in the inflammatory process while inhibiting
383 corresponding bone resorption.

384

385 Cathepsin K, a lysosomal cysteine protease, is abundantly expressed in
386 osteoclasts. This fact, as well as its enzymatic properties, implies that it is a key
387 factor in normal bone remodeling and in pathological processes, such as

388 osteoporosis, osteoarthritis and inflammatory mediated bone resorption disease
389 such as periapical disease (10, 11). Recent studies showed *Ctsk* also has
390 important functions in immune response (9, 22, 23). Our previous study has
391 already demonstrated that knocking down *Ctsk* by AAV gene therapy can
392 significantly reduce the bone resorption and inflammation (10). However, the
393 main drawback of using AAV as vectors is that the small size of their genome
394 significantly limits the amount of genetic material it can carry (24). Small
395 molecule can overcome this by having several advantages: a greater universe of
396 treatable diseases, lower cost with greater ease of manufacturing, and patient
397 preference for an oral availability (14, 25). ODN is a highly selective, potent, and
398 reversible inhibitor of *Ctsk* and inhibits OC-mediated bone resorption in vitro, but
399 does not reduce the number of osteoclasts, and appears to reduce bone
400 resorption while preserving bone formation (26, 27). ODN is currently being
401 developed as an orally bioavailable therapeutic for the treatment of
402 postmenopausal osteoporosis. In a clinical trial, 36 months of once weekly ODN
403 treatment increased lumbar spine and total hip bone mineral density (BMD) and
404 reduced bone turnover markers in postmenopausal women with low BMD (15).

405

406 Since *Ctsk* has been shown to have functions in both osteoclast and immune
407 cells such as macrophages and dendritic cells (9, 11, 28), whether inhibition of
408 *Ctsk* can affect not only osteoclasts but also immune response is still unclear.
409 This study proposed its novel role in endodontic inflammation lesions. Patients
410 with endodontic disease suffer from both inflammation-induced tissue damage

411 and bone loss (2), a single target that can dramatically improve both conditions is
412 ideal. In current study, we tested the inhibition of Ctsk using this specific small
413 molecule inhibitor ODN by oral administration in periapical lesion mouse model to
414 determine whether Ctsk plays dual roles in periapical disease to simultaneously
415 reduce inflammation and bone resorption at different time points.

416

417 Due to the nature and progression of endodontic infection, changing micro-
418 environmental conditions are selected for more anaerobic and virulent species of
419 bacteria (29). We induced periapical disease by inoculating the root canal system
420 of the mandibular first molar with a mixture of four common endodontic pathogens:
421 *Prevotella intermedia* [American Type Culture Collection (ATCC) 25611;
422 Manassas, VA], *Fusobacterium nucleatum* (ATCC 25586), *Peptostreptococcus*
423 *micros* (ATCC 33270), and *Streptococcus intermedius* (ATCC 27335) as
424 described (10, 17). Our results showed that bone resorption at different time
425 points was much more advanced in the periapical periodontitis group without
426 inhibitor than that in the periapical periodontitis group with inhibitor, and
427 meanwhile we found that bone resorption increased as time increased in different
428 groups, but this trend was more obvious in the periapical disease group without
429 inhibitor (**Fig. 1**). Further quantitative analysis by μ CT confirmed the bone
430 protective effects at 7, 21 and 42 days by inhibition of Ctsk (red arrow) (**Fig. 1**).
431 To determine the trends of bone resorption and the infiltration of inflammatory
432 cells in the periapical lesion area, we also conducted H&E staining to detect the
433 changes of bone resorption in different groups at 7, 21 and 42 days after

434 infection (**Fig. 2**), which was consistent with the results of X-ray and μ CT. We
435 also want to explore the proper dose used for treatment of periapical disease in
436 mice, so we chose two different dose 3.606 mg/kg/wk and 0.7212 mg/kg/wk,
437 which was recalculated from the dose used for clinical trial on human by
438 considering the difference of body surface area between human and mice (16).
439 The result showed that the 3.606 mg/kg/wk had a better bone protection effect at
440 42 days after infection (**see Fig. S2 in the supplemental material**). These
441 results showed that the bone resorption was more notable in the periapical
442 disease group without inhibitor than that in the inhibitor treated disease group at
443 different time points. Inflammatory cell infiltration in the periapical disease group
444 without inhibitor was increasingly more serious at 7, 21 and 42 days, from which
445 we can see a large number of blue-stained monocytes distributed in the apical,
446 mesial/distal and furcation area (red arrow) (**Fig. 2A**). In the inhibitor treated
447 periapical disease group we can see that the trabecular bone area was basically
448 unchanged and infiltration of inflammatory cells was significantly reduced. Above
449 all, inhibition of Ctsk has significant bone protection effects and prevents
450 inflammatory cells infiltration.

451

452 To investigate the mechanism of the effect of Ctsk inhibition *in vivo*, TRAP
453 staining and immunohistochemistry (IHC) staining was applied. Studies showed
454 that periapical inflammatory cells play an important role in the development of
455 periapical periodontitis, which secrete a variety of cytokines (IL-1 α , TNF- α , IL-6
456 and IL-11), and are involved in the adjustment of the activation and apoptosis of

457 corresponding effector cells such as osteoclast, thus controlling the bone
458 resorption (30-32). In our previous study, we showed that Ctsk knockdown can
459 abolish the extra-cellular acidification and then inhibit bone resorption, while
460 osteoclast differentiation is only slightly affected *in vitro* (10, 11). ODN has also
461 been shown not to affect the osteoclast numbers *in vivo* (33). Therefore, the
462 number of osteoclasts of various groups *in vivo* at different time points was
463 determined in this experiment to clarify the possible mechanism. We found that
464 the number of osteoclasts did not change in the normal control group (**see Fig.**
465 **S3 in the supplemental material**), but significantly reduced at different time
466 points in the lesion area of the inhibitor treated periapical periodontitis group
467 compared to that of the untreated periapical disease group (**Fig. 3**). This
468 indicated that the immune response may also be affected by inhibition of Ctsk
469 since pre-osteoclast activation is due to RANKL secreted by osteoblasts, T cells
470 and B cells, and these cells can be activated by proinflammatory cytokines and
471 chemokines such as IL-1, IL-6, TNF α and IL-17 (34). Studies indicated that
472 activated macrophages are the source of bone-resorbing cytokines in periapical
473 disease (35), which can be activated by T-lymphocytes or by bacterial endotoxin,
474 as part of innate immunity. In order to fully demonstrate the function of Ctsk in
475 periapical disease immune response, the present study also examined whether
476 inhibition of Ctsk affects the expression of macrophages by detecting the
477 macrophage marker F4/80 (**Fig. 4**). The results showed that the number of
478 macrophages increased significantly in the periapical disease group without
479 inhibitor at 42 days, but not in the inhibitor treated group (**Fig. 4**), which indicated

480 that Ctsk may play an important role in both osteoclasts and immune responses
481 (**Fig. 3, 4**).

482

483 Since macrophage expression decreased in the inhibitor treated disease group
484 and to further investigate the effect of Ctsk function in adaptive immune
485 response, immune fluorescence analysis was applied. We determined the
486 number of CD3 positive T cells in the periapical disease area from different
487 groups at 42 days (**Fig. 5**). The results showed that the CD3 positive T cells
488 decreased significantly in the inhibitor treated infection group, which indicated
489 that the adaptive immune response has been impaired by inhibition of Ctsk. As
490 the effector cell of inflammation, T cells can produce a variety of effector
491 molecules (36, 37). Therefore, the number of T cells is an important indicator of
492 the severity of local inflammatory response. As a T-cell surface marker, CD3 is
493 commonly used to detect T cell population including $\alpha\beta$ T cells and $\gamma\delta$ T cells.
494 The results showed that the number of CD3⁺ T cells was significantly reduced in
495 the inhibitor treated periapical disease group than that in the disease group (**Fig.**
496 **5A**). T cells showed aggregated distribution in the periapical periodontitis group,
497 while the inhibitor treated periapical periodontitis group only showed minor T cell
498 distribution, which indicated that the immune system-mediated inflammatory
499 reaction is suppressed in Ctsk inhibited disease groups. The result can be an
500 explanation of the reduction of the number of osteoclasts from another aspect
501 (**Fig. 3**). T cells and macrophages decreased, which might result in a lower

502 proinflammatory cytokine production, which might finally cause lower osteoclasts'
503 activation and differentiation.

504

505 In order to confirm this hypothesis, the relative mRNA level was quantified by
506 Sybrgreen based qRT-PCR by using the primer set as listed (**see Table S1 in**
507 **the supplemental material**). Our results showed that the *Ctsk* gene reached the
508 highest level at 42 days in the periapical disease group *in vivo*, but not in the
509 inhibitor treated disease group, which showed no obvious change of *Ctsk*
510 expression (**Fig. 6A**). To further investigate the mechanism of the protective
511 effect of inhibition of *Ctsk* in the mouse model of periapical disease, we tested
512 the expression level of pro-inflammatory maker genes including *TNF- α* , *IL-6* and
513 *IL-23 α* . The *TNF- α* and *IL-6* genes' expression levels increased significantly in
514 the periapical disease group at day 7 compared to that in the inhibitor treated
515 disease group (**Fig. 6A**). The expression of *IL-23 α* reached the highest level at
516 day 21. The protein level expression of cytokines, including *TNF- α* , *IL-6* and *IL-*
517 *23*, all increased in the periapical disease group at different time points. Although
518 the proinflammatory cytokine also increased in the inhibitor treated disease
519 group, the level is much lower when compared to the disease group (**Fig. 6B**).
520 Cytokines such as *TNF- α* , *IL-6* and *IL-23 α* are soluble mediators and released
521 from immunocompetent cells in the periapical inflammatory processes. These
522 cytokines can stimulate bone resorption in periapical lesion areas (38). The
523 proinflammatory cytokines such as *TNF- α* has been reported to have a key role
524 in regulating the inflammatory response (39). *IL-6* is produced at the site of

525 inflammation and also plays a key role in the acute phase response, which is
526 also shown by our qRT-PCR results (**Fig. 6A**) (40). As receptors recognize
527 several major bacterial pathogenics, Toll-like receptors (TLRs) are important in
528 immune response. Currently 12 TLR family members have been found in
529 humans and other mammals, but not all TLRs are expressed in mammals. For
530 example, TLR11 and 12 are expressed in mice, but not in humans (41). TLR4, 5
531 and 9 can specifically identify different types of bacterial antigens, which play an
532 important role in bacterial infection. The study from Beklen et al. showed that
533 except TLR7 and TLR8, expression levels of TLR1 to TLR10 in human gingival
534 surface is higher in periodontitis than healthy people (42), suggesting that TLRs
535 play a more significant role in the activation and development of periodontitis.
536 Therefore, TLR4, 5 and 9 were selected as the research object in this experiment
537 to detect the changes in the inhibition of Ctsk in the periapical disease model.
538 The relationship between Ctsk and TLRs has not been fully clarified yet. The
539 recent studies demonstrate that Ctsk has critical role in TLRs functions. In
540 rheumatoid arthritis disease, Asagiri et al. demonstrated that Ctsk might have
541 effect on DCs cytokine expression through TLR9 *in vitro* (9). Hirabara et al.
542 showed Ctsk inhibitor might inhibit TLR-4 dependent Ctsk expression in human
543 fibroblast *in vitro* (43). Ctsk has also been shown to be involved in development
544 of psoriasis-like skin lesions through TLR7 activation (44). In our current study,
545 we showed that TLR5 and 9 all decreased in the inhibitor treated disease group
546 compared to the disease group at different time points (**Fig. 6A**). The IHC stain of
547 TLR5 and 9 further confirmed the qRT-PCR result (**see Fig. S5 in supplemental**

548 **material and Fig. 7).** The TLR4 expression increased in the disease group at 7
549 and 21 days, however, decreased at 42 days. There was no significant difference
550 at 42 days between disease group and inhibitor treated disease group (**see Fig.**
551 **S4 in supplemental material and Fig. 6).**

552

553 In summary, we investigated the possible mechanism of the effect of Ctsk in
554 simultaneously targeting osteoclasts and immune cells. The inhibition of Ctsk by
555 the specific small ODN molecule reduced bone resorption by inhibition of the
556 osteoclast function and inflammation simultaneously and in turn decreased
557 osteoclast activation and differentiation, which exhibits a promising therapeutic
558 method on periapical inflammation disease.

559

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734 **Figure Legends**

735 **Figure 1. Inhibition of Ctsk by specific inhibitor reduced infection-stimulated**
736 **endodontic bone resorption by Micro-computed tomography (μ CT) analysis.**
737 **(A)** X-ray and μ CT analysis of endodontic lesion area of the normal group in
738 natural course with or without inhibitor (3.606 mg/kg/wk) at 42 days. **(B-D)** X-ray
739 and μ CT analysis of endodontic lesion area of bacterial infected group with or
740 without inhibitor (3.606 mg/kg/wk) at 7, 21 and 42 days. There was an obvious
741 reduced bone density area compared to inhibitor treated disease group at
742 different time points (Black arrows). The X-ray and μ CT analysis showed
743 increased low bone density area at furcation and periapical area at 42 days
744 compared with 7 days and 21 days. **(E)** Quantification of bone volume/tissue
745 volume (BV/TV) measured for periapical lesion area of different groups. *:
746 $P < 0.05$, **: $P < 0.01$, ***: $P < 0.001$. N = 3, repeated three times. N.S: No
747 Significance. Left column of panel **A-D** are X-ray images. Middle column of panel
748 **A-D** are typical slice 2D μ CT images, while the right column of panel **A-D** are 3D
749 Reconstitutions images.

750

751

752 **Figure 2. Inhibition of Ctsk by specific inhibitor showed bone protection**
753 **effects at different time points by histology analysis. (A)** H&E staining of the
754 periapical mesio, distal root sections and furcation area sections from the
755 bacterial infected group with or without inhibitor (3.606 mg/kg/wk) at different time
756 points (7, 21 and 42 days). Red arrows indicated the inflammatory monocytes
757 infiltration. The red dot area indicated the remaining bone in the lesion area. **(B)**
758 Quantification of remaining bone area at 7, 21 and 42 days of these three areas.
759 **(C)** Time course analysis of the bone resorption at bacterial infected group with
760 or without inhibitor (3.606 mg/kg/wk). *: $P < 0.05$, **: $P < 0.01$, ***: $P < 0.001$. N.S:
761 No Significance. N = 3, repeated three times.

762

763

764 **Figure 3. Inhibition of Ctsk by specific inhibitor showed reduced TRAP⁺ cells in**
765 **periapical lesion area at different time points by histology analysis. (A)** TRAP
766 staining of the periapical mesio, distal root sections and furcation area sections
767 from the bacterial infected group with or without inhibitor (3.606 mg/kg/wk) at
768 different time points (7, 21 and 42 days). **(B)** Quantification of TRAP⁺ cell
769 numbers at 7, 21 and 42 days of these three areas. **(C)** Time course analysis of
770 TRAP⁺ cell numbers at bacterial infected group with or without inhibitor (3.606
771 mg/kg/wk). The TRAP⁺ cell number increased in bacterial infected group without
772 inhibitor at different time points, and the TRAP⁺ cell number in infected group
773 with inhibitor (3.606 mg/kg/wk) was significant lower compared to infected group
774 without inhibitor. *: $P<0.05$, ***: $P<0.001$. N.S: No Significance. N = 3, repeated
775 three times.

776

777

778 **Figure 4. Inhibition of Ctsk by specific inhibitor showed reduced F4/80⁺ cells of**
779 **periapical lesion area at 42 days by immunohistological analysis. (A)**
780 Immunohistochemistry stains of F4/80⁺ (Brown) macrophages in periapical lesion
781 area in normal group and bacterial infected group with or without inhibitor (3.606
782 mg/kg/wk) at 42 days. Normal serum served as negative control. **(B)** Higher
783 magnification view of infection disease group at 42 days. Red arrows indicated
784 F4/80⁺ macrophages. **(C)** Quantification of F4/80⁺ macrophages in alveolar area
785 of normal group and bacterial infected group with or without inhibitor (3.606
786 mg/kg/wk) at 42 days. **: $P<0.01$, ***: $P<0.001$. N = 3, repeated three times.

787

788

789 **Figure 5. Inhibition of Ctsk by specific inhibitor decreased the number of T**
790 **cells in periapical lesion area at 42 days. (A)** Immunofluorescence staining of
791 CD3⁺ (red) T cells in periapical lesion area in the normal group and the bacterial
792 infected group with or without inhibitor (3.606 mg/kg/wk) at 42 days. **(B)** Normal
793 serum served as negative control at the same area (Without primary antibody).

794 (C) Quantification of CD3⁺ cells analysis demonstrated that inhibition of Ctsk has
795 significantly reduced expression of CD3⁺ T cells in bacterial infected group with
796 inhibitor (3.606 mg/kg/wk) at 42 days. ***: $P < 0.001$. N.S: No Significance. N = 3,
797 repeated three times.

798

799

800 **Figure 6. Inhibition of Ctsk by specific inhibitor reduced the expression of pro-**
801 **inflammatory genes as well as TLRs genes and cytokines in the periapical**
802 **lesion at different time points. (A)** qRT-PCR of osteoclast genes (i.e. *Ctsk*) and
803 pro-inflammatory genes (i.e. *TNF- α* , *IL-6* and *IL-23 α*) as well as TLRs genes (i.e.
804 *TLR4*, 5 and 9) in the periapical lesion from normal group and bacterial infected
805 group with or without inhibitor (3.606 mg/kg/wk) at 7, 21 and 42 days. *β -actin* was
806 used as an endogenous control. (B) Expression of *TNF- α* , *IL-6* and *IL-23 α* in the
807 periapical lesion at 7, 21 and 42 days detected by ELISA. Significance was
808 compared between disease group with and without inhibitor at different time
809 points. *: $P < 0.05$, **: $P < 0.01$, ***: $P < 0.001$. N.S: No Significance. N = 4, repeated
810 three times.

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812

813 **Figure 7. Inhibition of Ctsk by specific inhibitor shows reduced expression of**
814 **TLR9 in periapical lesion area at 42 days by immunohistological analysis. (A)**
815 Immunohistochemistry stains of TLR9⁺ (Brown) cells areas in normal group and
816 bacterial infected group with or without inhibitor (3.606 mg/kg/wk) at 42 days.
817 Normal serum served as negative control. (B) Higher magnification view of
818 infection disease group at 42 days. Red arrows indicated TLR9⁺ cells. (C)
819 Quantification of TLR9⁺ cells in frucation, mesio and distal periapical area of
820 normal group and bacterial infected group with or without inhibitor (3.606
821 mg/kg/wk) at 42 days. *: $P < 0.05$, **: $P < 0.01$, ***: $P < 0.001$. N.S: No Significance.
822 N = 3, repeated three times.

823













