

Remodeling Articular Immune Homeostasis with an Efferocytosis-inspired Nanoimitator Mitigates Rheumatoid Arthritis

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1 **Remodeling Articular Immune Homeostasis with an Efferocytosis-inspired**
2 **Nanoimitator Mitigates Rheumatoid Arthritis**

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30 **Abstract**

31 Massive intra-articular infiltration of the pro-inflammatory macrophages is a prominent
32 feature of rheumatoid arthritis (RA) lesions, which are thought to underlie articular
33 immune dysfunction, severe synovitis and ultimate joint erosion. Here we report an
34 efferocytosis-inspired nanoimitator (EINI) for in situ targeted reprogramming of the
35 synovial inflammatory macrophages (SIMs) and thus thwarting their autoimmune
36 attack and reinstating articular immune homeostasis, which mitigates RA. The EINI
37 consisted of a drug-based core with an oxidative stress-responsive phosphatidylserine
38 (PtdSer) corona and a shell of P-selectin-blocking motif, low molecular weight heparin
39 (LMWH). When systemically administrated, the LMWH on the EINI first bound to P-
40 selectin overexpressed on endothelium in subsynovial capillaries, which functioned as
41 an antagonist disrupting neutrophils synovial trafficking. Due to the high dysregulation
42 of the synovial microvasculature, the EINI subsequently enriched in joint synovium
43 where the shell was exfoliated upon the reactive oxygen species stimulation, and PtdSer
44 corona was then exposed. In an efferocytosis-like manner, the PtdSer-coroneted core
45 was in turn phagocytosed by SIMs, which synergistically terminated the SIMs-initiated
46 pathological cascades and serially reconstructed the intra-articular immune homeostasis,
47 conferring a chondroprotection effect. These findings demonstrate that SIMs can be
48 precisely remodeled via the efferocytosis-mimetic strategy, which holds great potential
49 for RA treatment.

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56 **Key words:** Efferocytosis-inspired nanoimitator; Rheumatoid arthritis; Synovial
57 inflammatory macrophage; Articular immune homeostasis; Chondroprotection

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59 **Introduction**

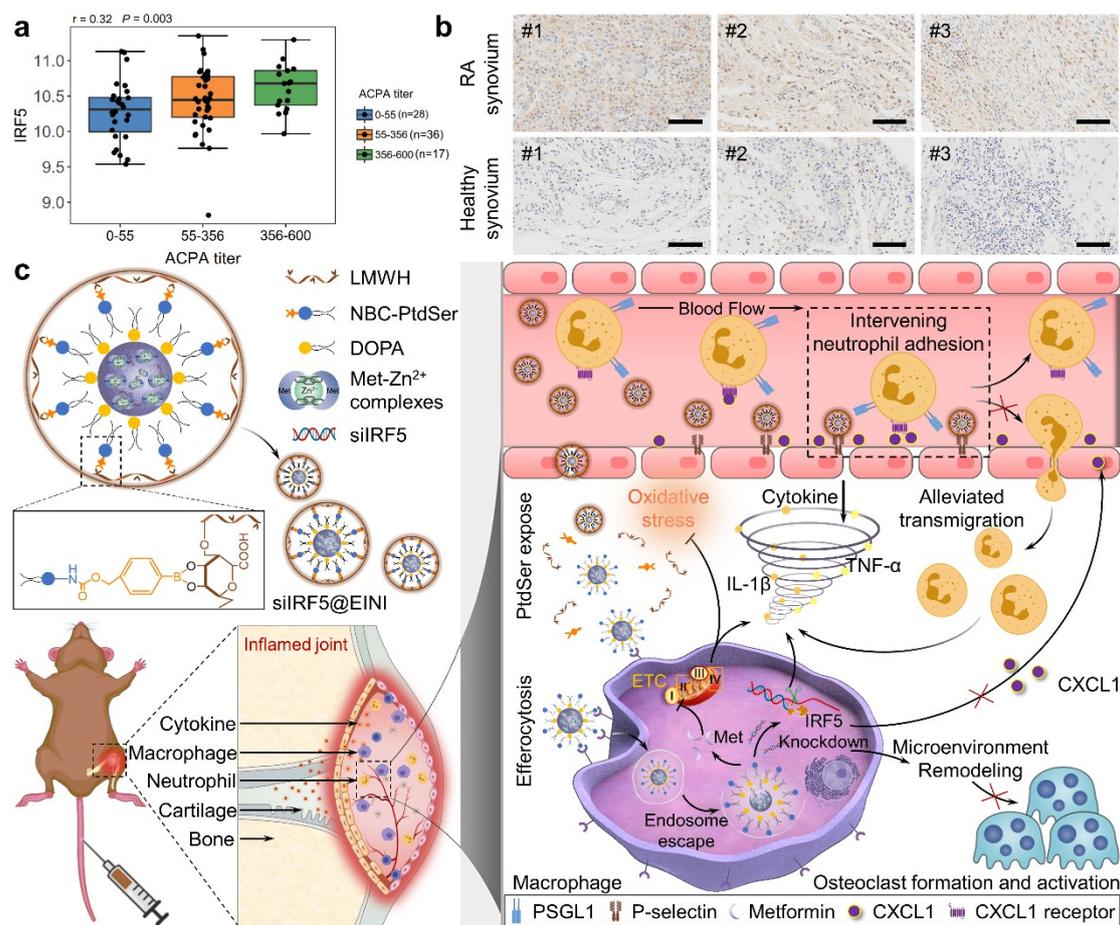
60 Rheumatoid arthritis (RA), a widespread and devastating systemic autoimmune
61 disease, characterized by articular immune dysfunction with serious synovitis and joint
62 erosion that causes progressive disability^{1, 2}. Over the past two decades, multiple
63 immunosuppressants have been approved by the U.S. FDA for RA treatment³. Despite
64 combining use of these immunosuppressants, about one-third of patients with RA fail
65 to reach sustained clinical remission⁴. Moreover, the continuous use of
66 immunosuppressive therapeutics often weakens the immune system and do increase the
67 risk of infections⁵. The pathogenesis of RA involves the complicated inflammatory
68 networks, including the multiplicity of cytokine targets and a complex crosstalk of
69 inflammatory cells^{6, 7}. Among these, synovial inflammatory macrophages (SIMs) play
70 a pivotal role in orchestrating the cytokine environment, which are thought to underlie
71 articular immune dysfunction, synovitis and ultimate joint erosion^{8, 9}. As such,
72 terminating the SIMs-initiated cascades and serially reinstating articular immune
73 homeostasis may be a reversible approach for RA therapy, which remains largely
74 unexplored.

75 Patients with RA can be divided into two major subsets based on the presence versus
76 absence of anti-citrullinated protein antibodies (ACPAs)¹⁰. The ACPA⁺ subset of
77 disease constitutes approximately two-thirds of all cases of RA and generally has a more
78 severe disease course¹¹. Interferon regulatory factor 5 (IRF5) is a master regulator in
79 defining the classical inflammatory phenotype of macrophages¹² and translates various
80 signals related to the SIMs in RA synovium^{13, 14}. In humans, polymorphisms in the IRF5
81 genes have been associated with an increased risk of developing autoimmune disease
82 including RA^{15, 16}. Bioinformatically, we found that the expression of IRF5 revealed a
83 positive correlation with the ACPAs titer in RA synovium (Fig. 1a). Consistently, in
84 patients with ACPA⁺ RA, the expression level of IRF5 protein was obviously elevated
85 (Fig. 1b). Targeted silencing of IRF5 in SIMs may therefore be an efficient strategy that
86 could facilitate the anti-inflammation polarization of macrophages and thus abort the
87 SIMs-initiated cascades in ACPA⁺ RA.

88 Precisely targeting intra-articular macrophages is of utmost importance for
89 efficiently regulating SIMs-initiated cascades. The phagocytosis of the apoptotic cells,
90 called efferocytosis, is one of the critical innate functions of macrophages, which
91 maintains tissue homeostasis^{17, 18}. During efferocytosis, phosphatidylserine (PtdSer)
92 exposure on the outer leaflet of the plasma membrane in the apoptotic cells is a key
93 "eat-me" signal for macrophages¹⁹. PtdSer coronation may therefore enhance the
94 targeted-internalization of nano-formulation by macrophages. However, after systemic
95 administration, PtdSer-coronated nano-formulation may be up-taken by the pan-
96 macrophages, which results in severe adverse effects. Of note, intra-articular oxidative
97 stress caused by high level of reactive oxygen species (ROS) is one of the typical
98 characteristics of RA lesions^{20, 21}. As such, ROS-responsive exfoliation may be an
99 efficient way to manipulate the locoregional exposure of PtdSer corona of the designed
100 nano-formulation intra-articular, which enabled the specific-phagocytosis of the nano-
101 formulation by SIMs.

102 Here, we sought to develop an efferocytosis-mimetic self-deliverable nanoimitator
103 for in situ targeted reprogramming of the SIMs and thus thwarting their autoimmune
104 attack and reinstating articular immune homeostasis for RA-reversible treatment (Fig.
105 1c). The siIRF5-carrying efferocytosis-inspired nanoimitator (siIRF5@EINI) was for
106 the first time sequentially assembled by a drug-based core with an oxidative stress-
107 responsive phosphatidylserine (PtdSer) corona and an outer shell of low molecular
108 weight heparin (LMWH). With the shielding of LMWH, the siIRF5@EINI was
109 endowed with the stealth property in circulation, an enhanced inflamed regions
110 retention, and a blocking function of P-selectin that retards the articular trafficking of
111 neutrophils (NEs). When systemically administrated, the LMWH on the nanoimitator
112 first bound to P-selectin overexpressed on endothelium in subsynovial capillaries,
113 which functioned as an antagonist disrupting NEs synovial trafficking. Due to the
114 leakage of the intra-articular blood vessel, the siIRF5@EINI subsequently enriched in
115 joint synovium where the shell was exfoliated upon the ROS stimulation, and PtdSer
116 corona was then exposed. In an efferocytosis-like manner, the PtdSer-coronated core

117 was in turn phagocytosed by SIMs. We demonstrated that the siIRF5@EINI
 118 synergistically terminated the SIMs-initiated pathological cascades and reconstructed
 119 the intra-articular immune homeostasis, and ultimately conferred a chondroprotection
 120 effect and restored the joint function, all of which were comprehensively investigated
 121 with in a RA murine model.



122

123 **Fig. 1 Efferocytosis-inspired nanoimitator in situ targeted silencing of IRF5 in**
 124 **SIMs for precision RA therapy.** **a** The positive correlation between the IRF5 gene
 125 expression levels and the anti-citrullinated protein antibodies (ACPAs) titer in the RA
 126 synovium ($r = 0.32$, $P = 0.003$). **b** Immunohistochemistry staining of IRF5 in synovium
 127 from RA patients and healthy donors (3 cases each group). Scale bar = 50 μ m. **c**
 128 Schematic illustration shows that the efferocytosis-inspired nanoimitator terminates the
 129 SIMs-initiated pathological cascades, synergistically restores the articular immune
 130 homeostasis, and ultimately reverses bone erosion.

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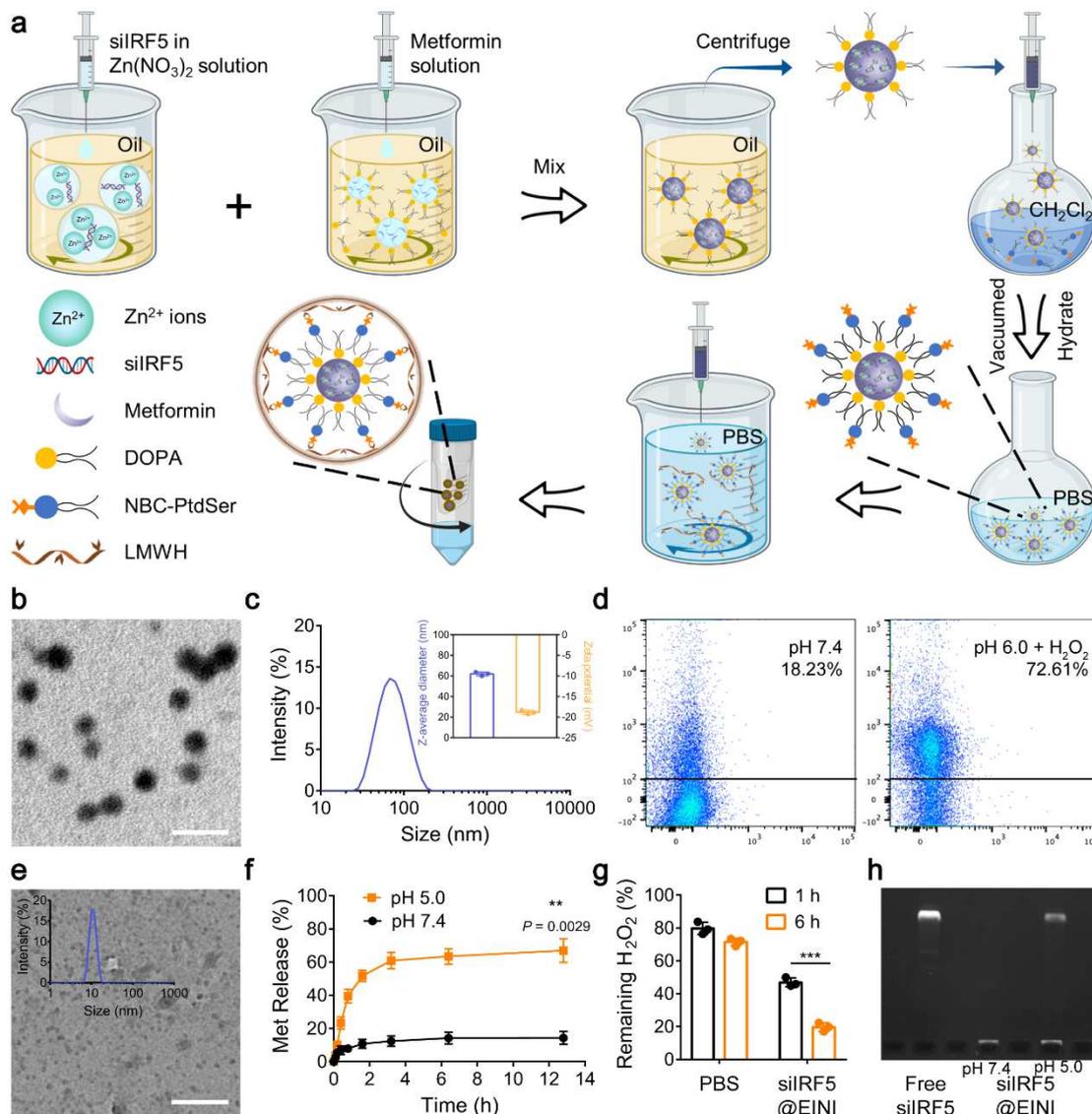
132 **Results**

133 **Preparation and characterization of EINI**

134 Via conjugating 4-nitrophenyl 4-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)
135 benzyl carbonate (NBC) with PtdSer, we first synthesized a ROS-responsive PtdSer-
136 NBC conjugate (Supplementary Fig. 1, 2). The efferocytosis-inspired nanoimitator was
137 prepared (Fig. 2a). Briefly, dioleoylphosphatidic acid (DOPA) stabilized siIRF5-laden
138 metformin-Zn²⁺-based nanocore was constructed using the reverse microemulsion
139 method, and then coronated with PtdSer-NBC. Subsequently, the corona of nanocore
140 was cloaked with LWMH via a reversible boronate ester linker by the reaction of cis-
141 diols and phenylboronic acid groups, and the siIRF5-laden efferocytosis-inspired
142 nanoimitator (siIRF5@EINI) was thus achieved. Under transmission electron
143 microscopy (TEM) (Fig. 2b), the obtained nanoimitator showed a spherical morphology
144 with a hydrodynamic diameter of ~72 nm (Fig. 2c). After dual coating with PtdSer-
145 NBC and LWMH, the zeta potential of the nano-formulation decreased from 44.5 ± 0.4
146 mV to -18.7 ± 0.03 mV (Fig. 2c and Supplementary Fig. 3). We next investigated the
147 stimulation-triggered presenting of PtdSer of the siIRF5@EINI in an RA-mimicking
148 microenvironment. As shown in Fig. 2d and Supplementary Fig. 4, 72.61% of the
149 PtdSer-presenting nanoparticles were determined to band to fluorescein isothiocyanate
150 (FITC)-annexin V in the solution of 0.1 mM H₂O₂, which was merely 18.23% in PBS
151 control group, indicating oxidative stress in RA tissues could convert PtdSer-NBC of
152 the nanoimitator into PtdSer.

153 Next, we evaluated the acid-triggered drug release of the nanoimitator in vitro. The
154 morphology of nanoimitator exhibited significant degradation into ultrasmall species in
155 acidic environments as shown by the TEM images (Fig. 2e). Structural disassembly
156 was further confirmed with dynamic light scattering (DLS) assay as shown in the
157 embedded image of Fig. 2e. Consistently, we found that under acidic conditions (pH
158 5.0), the release of metformin from the nanoimitator increased compared to that under
159 physiological conditions (pH 7.4) (Fig. 2f). With electrophoresis assay (Fig. 2h), siIRF5
160 release profile was also monitored. At lysosomal pH of 5.0, siIRF5 burst released from

161 the nano-formulation, while at physical pH of 7.4, we did not detect any release of
 162 siIRF5 from the siIRF5@EINI. The ROS-responsive exfoliation of the EINI was further
 163 determined with the H₂O₂ scavenging assay. Fig. 2g revealed a time-dependent ROS
 164 elimination of the nanoimitator upon the ROS stimulation, implying the ROS-
 165 stimulated deshielding of the nanoimitator in the RA pathological condition.



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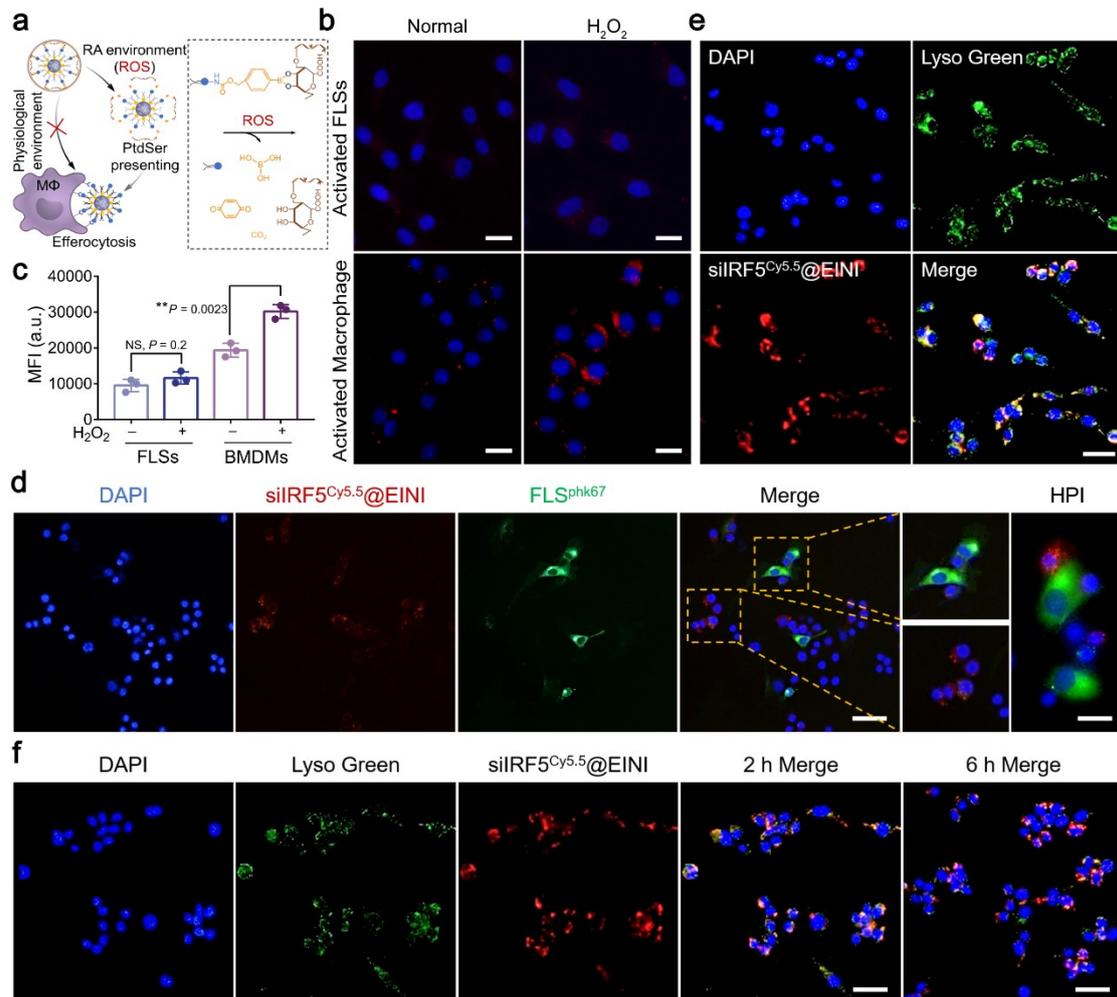
167 **Fig. 2 Preparation and characterization of siIRF5@EINI.** **a** Schematic of the
 168 preparation of nanoimitator by a modified reverse microemulsion method. **b**
 169 Transmission electron microscope image of siIRF5@EINI. Scale bar = 100 nm. **c**
 170 Hydrodynamic size and zeta potential determined with DLS. Data are presented as the
 171 mean \pm s.d. ($n = 3$ independent experiments). **d** Flow cytometry analysis of PtdSer-
 172 presenting. ($n = 3$ independent experiments). **e** TEM images of the nanoimitator after

173 triggered degradation under acidic conditions (pH 5.0). Scale bar = 100 nm. **f** Drug
174 release of the nanoimitator in PBS at different pH values. Data are presented as the
175 mean \pm s.d. ($n = 3$ independent experiments). ****** $P < 0.01$. **g** ROS scavenging ability of
176 nanoimitator. Results are reported as the mean \pm s.d. ($n = 3$ independent experiments).
177 P value was calculated by comparing with group PBS. ******* $P < 0.001$. **h** Electrophoretic
178 gel assay showing the pH-responsive release of siIRF5 from the nanoimitator. The
179 experiments were repeated three times, independently.

180 **Cellular engulfment specificity of the nanoimitator in vitro**

181 PtdSer exposure triggers the efferocytosis of macrophages (Fig. 3a). We next
182 determined the engulfment and intracellular trafficking of the nano-formulation in vitro
183 using bone marrow derived macrophages (BMDMs) that were isolated from mouse
184 (Supplementary Fig. 5). BMDMs were pre-activated with TNF- α and then treated with
185 the nanoimitator that was pre-conditioned with the 0.1mM H₂O₂. As shown in Fig. 3b,
186 significant fluorescence was detected in BMDMs, indicating that abundant
187 nanoimitators were engulfed into BMDMs. With fibroblast-like synoviocytes (FLSs) as
188 a control, we also determined the cellular engulfment specificity of the nanoimitator.
189 Minimal Cy5.5 signal was detected in FLSs (Fig. 3b). The results indicate that the EINI
190 was specifically engulfed by macrophages when ROS existed. Moreover, under an RA-
191 mimetic condition, an increased fluorescent intensity was determined in macrophages
192 compared to that under physiological microenvironment. A similar trend was found in
193 flow cytometry quantification (Fig. 3c). With a co-culture system, we further compared
194 the uptake of the nanoimitator by the pre-activated macrophages or FLSs under an RA-
195 mimetic condition. Consistently, fluorescence visualization with confocal laser
196 scanning microscope (CLSM) also indicates the specific engulfment of the PtdSer-
197 coroneted core by macrophages rather than FLSs (Fig. 3d). Next, we wished to ascertain
198 whether the siIRF5-laden nanoimitator would escape lyso/endosome and release the
199 payloads in cytoplasm after efficient up-taken by the macrophages. Co-localization
200 with CLSM revealed the nanoimitators were internalized by the cell via an endocytosis
201 pathway and then escaped from lyso/endosome, during which the payloads were

202 released into cytoplasm (Fig. 3e, f and Supplementary Fig. 6). Collectively, these results
 203 indicated that ROS-responsive exposure of PtdSer in the siIRF5@EINI induced an
 204 efficient phagocytosis by macrophages, and the lysosomal pH-triggered dissociation of
 205 the nanocarrier resulted in a successful lyso/endosome escape.



206
 207 **Fig. 3 Cellular uptake and intracellular distribution of the nano-formulation.** **a**
 208 Mechanistic explanation for the specific engulfment of the nanoimitator by
 209 macrophages. **b** Fluorescent images of macrophages and FLSs after incubation with
 210 siIRF5@EINI that with or without H_2O_2 pre-treating. Cells were activated with $TNF-\alpha$
 211 before nano-formulation incubation. Scale bar, 20 μm . **c** Flow cytometric analysis of
 212 nanoparticle binding to macrophages or FLSs. Data are presented as the mean \pm s.d. (n
 213 = 3 independent experiments). $**P < 0.01$. NS, not significant. **d** Representative images
 214 of H_2O_2 -pretreated siIRF5@EINI (red) engulfed by macrophages (not labeled)/FLSs
 215 (green) in a co-culture pattern. Scale bar, 50 nm. The right panel shows the

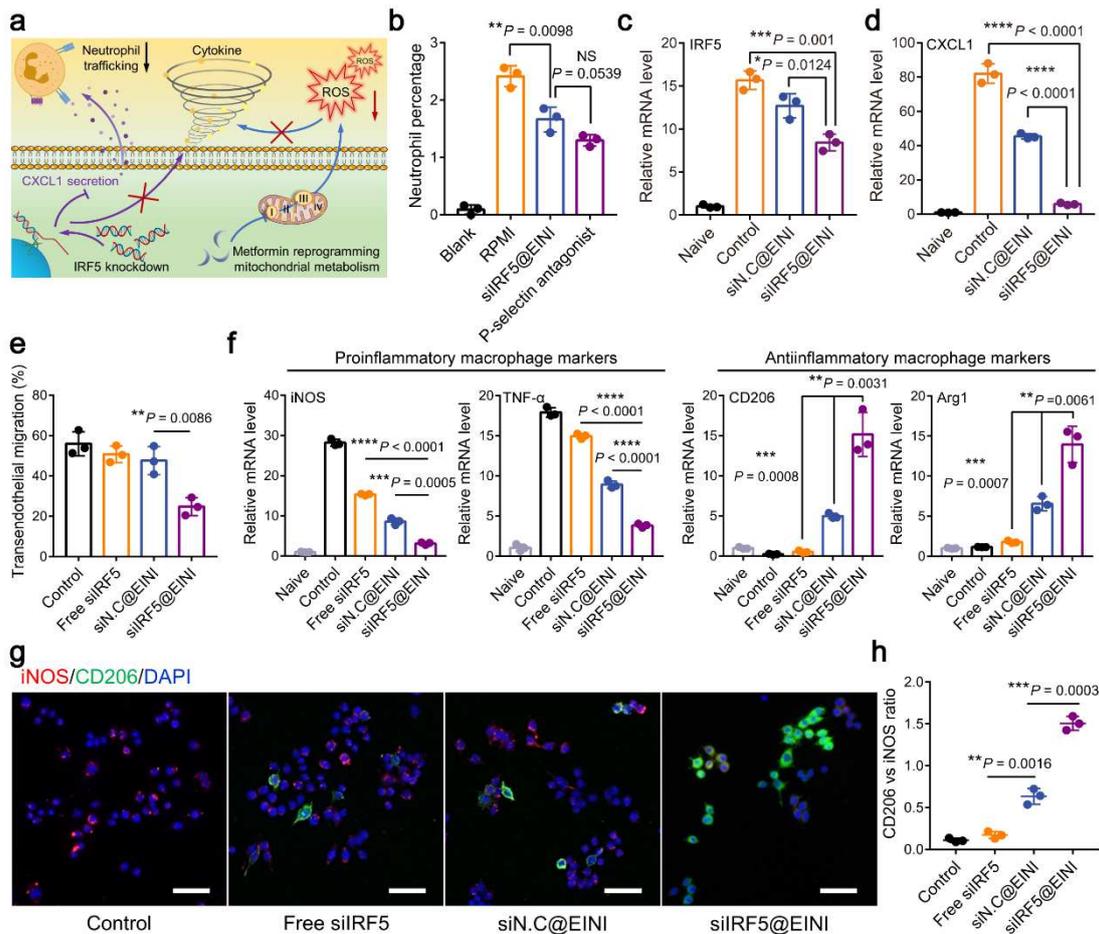
216 representative high power images (HPI). Scale bar, 20 μm . **e** Observation of engulfment
217 pathway of the nanoimitator. Scale bars, 20 μm . **f** Fluorescent visualization of siRNA
218 localization in macrophages 1 or 6 hours after incubation with siIRF5@EINI. Scale bar,
219 50 μm . In all datasets, the experiments were repeated three times, independently.

220 **Inflammatory modulation of the nano-formulation**

221 Comprehensively regulating the production of pro-inflammatory cytokines,
222 oxidative stress, and recruitment of NEs is essential for reestablishing articular immune
223 homeostasis (Fig. 4a). LMWH can block P-selectin initiated cell adhesion cascade, and
224 thus inhibit NEs migration into the inflamed regions²². We next evaluated the intervene
225 effect of the nanoimitator with a monolayer model of TNF- α activated human umbilical
226 vein endothelial cells (HUVECs). The results showed that the adhesion of NEs to
227 HUVECs significantly decreased after pre-incubation with the nanoimitator,
228 comparable to that treated with P-selectin antagonist (Fig. 4b and Supplementary Fig.
229 8). We further examined the gene silencing efficiency of the nano-formation after up-
230 taken by macrophages. As shown in Fig. 4c, the relative level of IRF5 mRNA from
231 TNF- α activated BMDMs revealed a dramatically decrease after treatment with the
232 siIRF5@EINI, which is well consistent with the protein data determined by western
233 blot (Supplementary Fig. 9). IRF5 plays a critical role in the secretion of multiple
234 chemokines including macrophage-derived chemokine CXCL1, a neutrophil
235 attractant¹⁴. We also detected the secreting level of CXCL1 from the treated
236 macrophages. As shown in Fig. 4d, treatment with siIRF5@EINI dramatically
237 downregulated macrophage-derived CXCL1 expression (Fig. 4d). Furthermore,
238 neutrophil migration was also examined. We found that IRF5 ablation significantly
239 inhibited the transendothelial migration of NEs (Fig. 4e). Therefore, we inferred that
240 the nanoimitator may function as an antagonist disrupting the synovial trafficking of
241 neutrophils.

242 Macrophages are phenotypically plastic²³. The expression of IRF5 in macrophages
243 defines the classical inflammatory phenotype¹². We next evaluated the modulative
244 effect of the nano-formulations on the phenotype conversion of the macrophages. As

245 shown in [Fig. 4f](#), treatment of the siIRF5@EINI reduced the mRNA expression level
246 of pro-inflammatory markers including tumor necrosis factor- α (TNF- α) and inducible
247 nitric oxide synthase (iNOS), meanwhile significantly increased the levels of anti-
248 inflammatory markers like mannose receptor (CD206) and arginase-1 (Arg-1). Similar
249 results were obtained in the immunofluorescence-stained images of BMDMs ([Fig. 4g](#),
250 [h](#)). Notably, treatment with metformin-single laden nano-formulation demonstrated a
251 strong elevation of M2-like subpopulation with highly expressed CD206. Concurrent
252 RNAi-mediated gene silencing and metformin-mediated metabolism reprogramming
253 further increased the immunoregulatory phenotype conversion of macrophages.
254 Metformin can target the transport chain of mitochondrial electron and thus reprogram
255 mitochondrial metabolism^{24, 25}. As shown in [Supplementary Fig. 10a, b](#), metformin-
256 related treatment significantly prevented the over-zealous ROS, which synergistically
257 leads to immunoregulatory phenotype conversion. ROS generating efficiency was also
258 evaluated by the determination of the ratio of NAD⁺/NADH. As shown in
259 [Supplementary Fig. 10c](#), a significant reduction of the ratio of NAD⁺/NADH was
260 detected in macrophages treated with metformin-related formulation. These results
261 suggested the siIRF5@EINI could boost the re-programming of macrophages to an
262 immunoregulatory phenotypes and thus significantly reduce macrophage-derived pro-
263 inflammatory cytokines, which is conducive to reestablishing the articular immune
264 homeostasis.



265

266 **Fig. 4 Inflammatory modulation of the nanoimitator.** **a** Schematic illustration of

267 siIRF5@EINI-induced inflammatory regulation. **b** Quantitative analysis of the adhered

268 neutrophils on HUVECs monolayers. The culture medium RPMI was used as a negative

269 control. Data are presented as the mean \pm s.d. ($n = 3$ independent experiments). ** $P <$

270 0.01. NS, not significant. **c**, **d** IRF5 (**c**) and CXCL1 (**d**) mRNA expression level in

271 activated macrophages treated with each formulation. Data are shown as the mean \pm

272 s.d. ($n = 3$ independent experiments). * $P < 0.05$, *** $P < 0.001$, **** $P < 0.0001$.

273 **e** The ratio of transendothelial neutrophils compared with all neutrophils after intervention

274 with different treatments on the HUVECs monolayer model. The monolayers of

275 HUVECs were stimulated with 10 ng/ml TNF- α . Data are shown as the mean \pm SD (n

276 = 3 independent experiments). ** $P < 0.01$. NS, not significant. **f** qRT-PCR analysis of

277 the phenotypical changes of the macrophages that were pre-treated with the

278 nanoimitator in vitro. Data are presented as the mean \pm s.d. ($n = 3$ independent

279 experiments). ** $P < 0.01$, *** $P < 0.001$, **** $P < 0.0001$. **g**, **h** Immunostaining of the

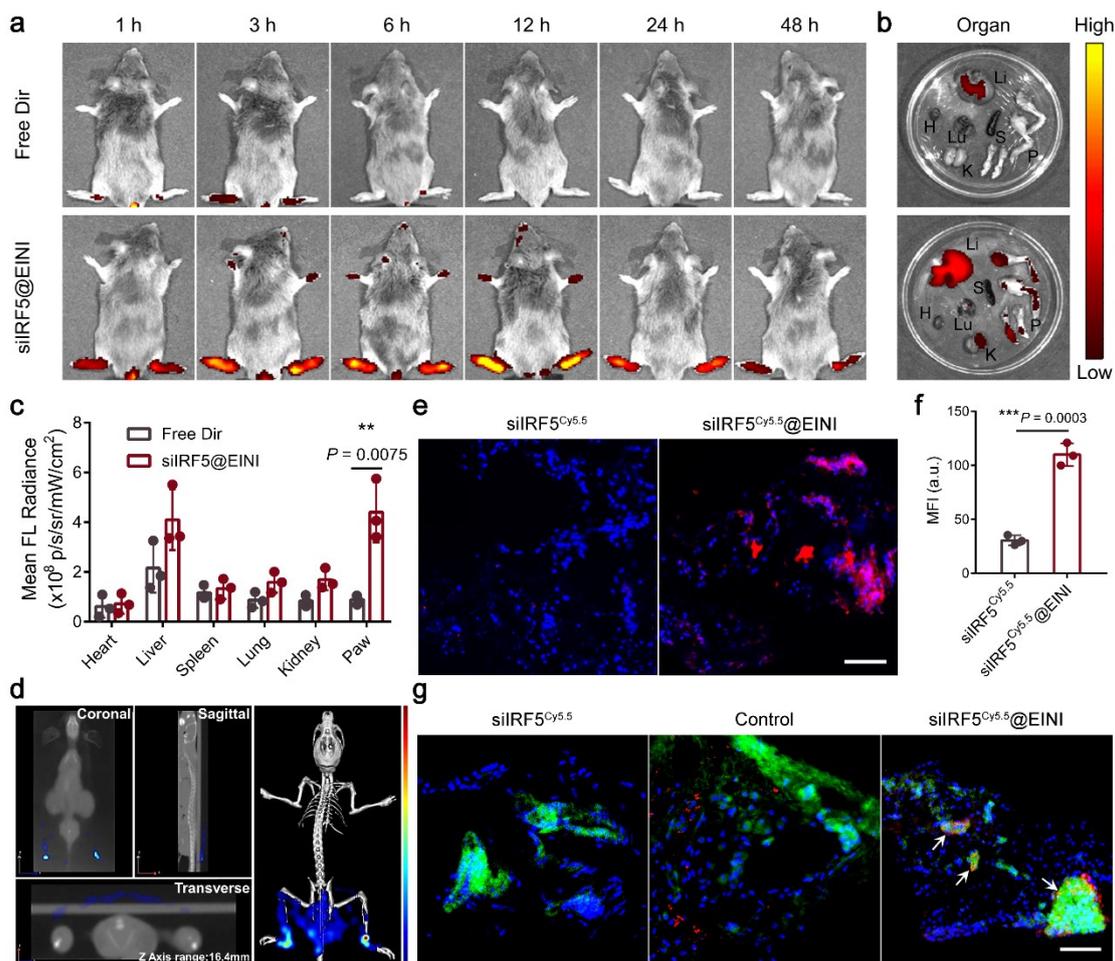
280 macrophages with different phenotypes (**g**) and the corresponding quantification of the
281 relative fluorescence intensity (**h**). Cells were pre-treated with TNF- α and then
282 incubated with each formulation for 24 h. Data are presented as the mean \pm s.d. ($n = 3$
283 independent experiments). ** $P < 0.01$, *** $P < 0.001$. Scale bars, 50 nm.

284 **Articular localization of the nano-formulation in vivo**

285 A collagen-induced arthritis (CIA) murine model was first established as previously
286 reported^{26, 27}. In vivo biodistribution and pharmacokinetic properties of the self-
287 deliverable nanoimitator were next evaluated. As shown in Fig. 5a, the EINI efficiently
288 deposited in the inflamed joints of the diseased mice 48 h post intravenous
289 administration. The major organs and paws of the mice from each group were collected
290 48 hours post-injection and subjected to fluorescence analysis ex vivo. The EINI
291 showed prominently enriched in the arthritic area of the CIA mice (Fig. 5b).
292 Quantification of the fluorescent intensity further confirmed that a significant increase
293 in mean fluorescence intensity (MFI) in EINI-treated mice than that in the free drug-
294 treated group (Fig. 5c). The articular-targeted accumulation of the nano-formulation
295 was also imaged using a hybrid microcomputed tomography-fluorescence tomography
296 (μ CT-FLT) device. As depicted in Fig. 5d, the EINI achieved an increased accumulation
297 in the inflamed joints. The retention profile of the nanoimitator in the synovium of
298 arthritic joints was further evaluated. 72 h after each treatment, the CIA mice were
299 sacrificed and the synovium cryosections were conducted. Fig. 5e, f revealed the EINI
300 preferentially enriched in the inflammatory synovial sites compared to that from free
301 dye-treated group. Further immunofluorescent staining demonstrated Cy5.5-labeled
302 siIRF5 co-localized with the synovial macrophages (Fig. 5g), indicating that the EINI
303 could efficiently accumulate intra-articular and specifically target synovial
304 macrophages in situ.

305 The potential toxicity of the systemic administrated nano-formulation was also
306 tested. No adverse effect was detected in hematological assay of the treated mice in
307 terms of white blood cell (WBC) and red blood cell (RBC) count and the hemoglobin
308 (HGB) (Supplementary Fig. 11a). Splenocytes were also isolated and evaluated. As

309 shown in [Supplementary Fig. 11b](#), no obvious apoptosis was detected in the treated
 310 mice. Histological analysis of organs coincidentally revealed the biosafety of the
 311 nanoimitator in vivo ([Supplementary Fig. 11c](#)).



312
 313 **Fig. 5 EINI selectively enriched in the inflamed joints of the CIA mice.** **a** In vivo
 314 fluorescence images of CIA models taken at different time points post injected with Dir
 315 labeled EINI and free Dir, respectively. **b** The fluorescence images of the excised major
 316 organs and paws harvested from the mice at 48 h post injection. **c** Quantification of
 317 fluorescence intensity in the inflamed joints and major organs of the CIA mice. Data
 318 are the mean \pm s.d. ($n = 3$ independent experiments). $**P < 0.01$. **d** μ CT-FLT imaging
 319 at 48 h after intravenous administration with the nanoimitators ($n = 3$ independent
 320 experiments). **e** Trafficking profile of nanoimitator in the joint synovium. Red
 321 represents nanoimitator and blue represents nuclei. Scale bars, 50 μ m. **f** Quantitative
 322 analysis of the accumulation of the nano-formulation in the synovium of the treated

323 mice. Data are presented as the mean \pm s.d. ($n = 3$ independent experiments). *** $P <$
324 0.001. **g** Representative images of the nanoimitator taken up by synovial macrophages
325 in vivo. Blue, DAPI-stained cell nuclei; Red, siIRF5^{Cy5.5}; Green, FITC conjugated anti-
326 F4/80 antibodies labeled macrophages. Arrow indicates the co-localization of the
327 nanoimitator and the SIMs labelled with anti-F4/80 antibodies. ($n = 3$ independent
328 experiments). Scale bars, 50 μ m.

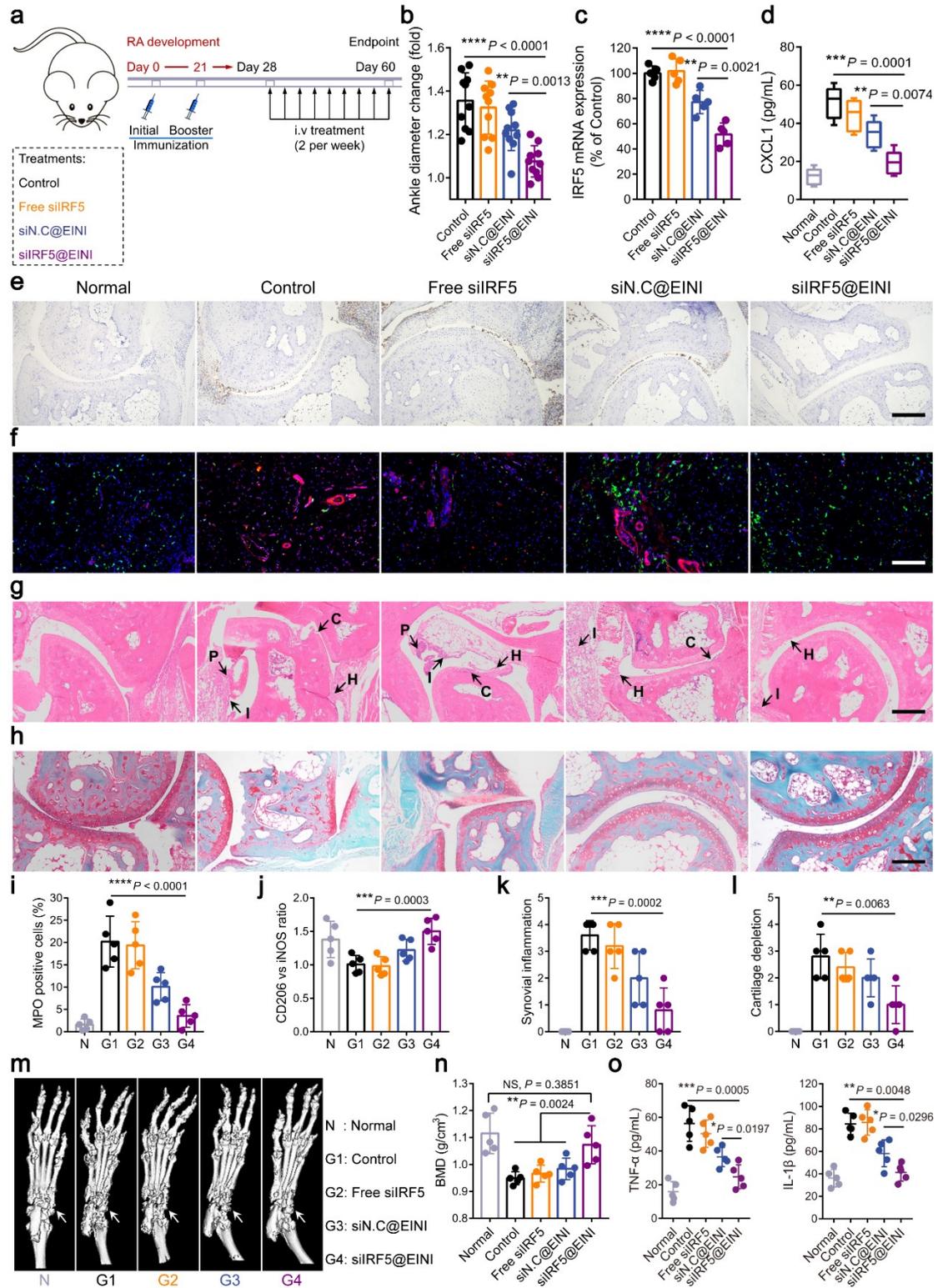
329 **EINI-enabled reestablishment of immune homeostasis and chondroprotection in** 330 **vivo**

331 We next investigated the therapeutic potent of the nano-formulations using the
332 abovementioned CIA mouse model (Fig. 6a). At the study endpoint, the transverse
333 ankle diameter of mice injected with siIRF5@EINI significantly smaller than the PBS
334 group, indicating a reduction inflammation and oedema formation (Fig. 6b and
335 Supplementary Fig. 12). We subsequently performed the gait analysis of the treated
336 mice using a catwalk system. As shown in Supplementary Fig. 13, siIRF5@EINI
337 treatment markedly ameliorates the abnormality of the footprint patterns of the CIA
338 mice compared with that of control groups. IRF5 gene silencing efficiency was also
339 confirmed in vivo with the CIA mice. Treatment with siIRF5-laden nano-formulation
340 significantly downregulated its expression in macrophages that were isolated from the
341 arthritic joint (Fig. 6C and Supplementary Fig. 14). Consistently with the in vitro results,
342 IRF5 silencing reduced the secretion of macrophage-derived CXCL1 in the inflamed
343 joint (Fig. 6d), which in turn decreased the recruitment of NEs. We further detected the
344 infiltration of NEs in inflamed joint by immunohistochemical staining with anti-
345 myeloperoxidase (MPO). As shown in Fig. 6e and Fig. 6i, the siIRF5@EINI-treated
346 group showed minimal MPO expression with few positively stained cells in the
347 inflamed sites, while overwhelming MPO-positive cells were detected in the control
348 joints. Competitively binding P-selectin on vascular endothelial cells can decrease the
349 infiltration of NEs into arthritic joints, and thus promotes the restoration of the articular
350 immune microenvironment^{22, 28}. More importantly, treatment with the siIRF5@EINI
351 synergistically induced phenotypic alteration of synovial macrophages (Fig. 6f and Fig.

352 6j) and subsequently decreased the unchecked infiltration of NEs in the RA joints which
353 was conducive to reconstruct the articular immune homeostasis.

354 We next wished to ascertain whether systemic injection of the efferocytosis-inspired
355 nanoimitator would control synovitis and reverse bone erosion in CIA mouse model. In
356 hematoxylin and eosin (H&E)-stained sections, an almost completely recovered
357 articular cavity with a clear interface without obvious synovitis and articular cartilage
358 degeneration was detected in the siIRF5@EINI-treated mice (Fig. 6g and Fig. 6k),
359 while severe pathologic changes including extensive inflammatory cell infiltration,
360 synovial hyperplasia, pannus formation, and cartilage erosion were observed in the
361 control group. Pannus formation in the synovial intimal lining features the pathological
362 changes of RA²⁹. Synovitis creates an oxidative stress microenvironment that actively
363 induces a tumor-like invasive pannus and further exacerbates joint damage²⁰. As shown
364 in Supplementary Fig. 15a, the angiogenesis and leukocyte extravasation in the
365 synovial sublining was significantly reduced in the siIRF5@EINI-treated groups
366 compared with the others. These pathogenetic changes in the synovium implied the
367 reversion of pannus progression. Additionally, safranin O staining showed optimal
368 preservation of cartilage structure of the mice in the siIRF5@EINI-treated group (Fig.
369 6h and Fig. 6l). Consistent with the therapeutic efficacy results, tartrate-resistant acid
370 phosphatase (TRAP) staining of osteoclasts further demonstrated that siIRF5@EINI
371 treatment significantly decreased the number of osteoclasts in RA joints and achieved
372 the slightest bone erosion (Supplementary Fig. 15b). CT was also performed to assess
373 bone erosion. As shown in Fig. 6m, the reconstructed micro-CT images revealed that
374 both of the ankle and the toe joints of the mice in the control groups suffered the most
375 serious bone corrosion. Quantitative analyses of bone mineral density (BMD)
376 demonstrated that the mean BMD in siIRF5@EINI-treated mice was almost completely
377 recovered comparable with the healthy mice (Fig. 6n). To evaluate the potential
378 systemic immune response, we further examined the serum levels of TNF- α and IL-1 β
379 in CIA mice. As summarized in Fig. 6o, these inflammation-related cytokines decreased
380 after treatment, indicating effective reduction of inflammation at the systemic level.

381 These results collectively suggested that the efferocytosis-inspired nanoimitator could
 382 effectively repolarize SIMs to an immunoregulatory phenotype in situ, serially reinstate
 383 the articular immune homeostasis and ultimately terminated the bone erosion.



384

385 **Fig. 6 EINI-enabled reestablishment of immune homeostasis and**

386 **chondroprotection in vivo. a** The therapeutic regimen with a CIA mouse model. **b**
387 Changes of the hind ankle in diameter on day 60 after CIA induction compared to that
388 on day 0. Data are presented as the mean \pm s.d. ($n = 5$ biologically independent animals).
389 $**P < 0.01$, $****P < 0.0001$. **c** Quantification of IRF5 expression in synovial
390 macrophages after each treatment. Data are presented as the mean \pm s.d. ($n = 5$
391 biologically independent animals). $**P < 0.01$, $****P < 0.0001$. **d** Secreting level of
392 CXCL1 in synovial tissue isolated from inflamed joint of the treated mice. Data are
393 presented as the mean \pm s.d. ($n = 5$ biologically independent animals). $**P < 0.01$, $***P$
394 < 0.001 . **e** The infiltration of the neutrophils in arthritic joints in each group. Joint
395 sections in each group were stained with anti-myeloperoxidase (MPO) antibodies.
396 Scale bar 200 μm . ($n = 5$ biologically independent animals). **f** Immunostaining of iNOS
397 and CD206 (red: iNOS; green: CD206; blue: DAPI; scale bar: 50 μm) to determine the
398 phenotype of SIMs in the CIA models treated with different formulations. ($n = 5$
399 biologically independent animals). **g, h** Histological analysis with H&E (scale bar 200
400 μm) (**g**) and safranin O staining (scale bar 100 μm) (**h**) of the joint tissues that excised
401 from the mice after different treatments. ($n = 5$ biologically independent animals). H,
402 synovium hyperplasia; I, immune cell infiltration; P, pannus formation; C, cartilage
403 damage. **i-l**, The percentage of MPO-positive cells in panel e (**i**) and the ratio of the
404 fluorescence intensity of CD206 versus iNOS in panel f (**j**). The histopathological
405 evaluation of synovial inflammation (**k**) and cartilage depletion (**l**) were obtained from
406 the images in (g) and (h), respectively. Data are presented as the mean \pm s.d. ($n = 5$
407 biologically independent animals). $**P < 0.01$, $***P < 0.001$, $****P < 0.0001$. **m**
408 Representative micro-CT images used for quantification of bone erosion of mice. **n**
409 Quantification of bone mineral density determined by micro-CT. Data are presented as
410 the mean \pm s.d. ($n = 5$ biologically independent animals). $**P < 0.01$. NS, not significant.
411 **o** Concentration profiles of TNF- α and IL-1 β in serum of CIA mice treated with
412 different groups. Data are presented as the mean \pm s.d. ($n = 5$ biologically independent
413 animals). $*P < 0.05$, $**P < 0.01$, $***P < 0.001$. NS, not significant.

414

415 **Discussion**

416 The current standard-of-care therapies in clinic for patients with rheumatoid
417 arthritis (RA) are palliative, which are still largely relying on multiple
418 immunosuppressants³. Despite trying multiple different immunosuppressants, about
419 one-third of RA patients fail to achieve disease remission⁴. Most available
420 immunosuppressants therapies suppress the immune system systemically, which
421 increase the risk of infections⁵. Although the precise mechanism of RA remains elusive,
422 the latest clinical pathogenic insights indicate that extensive infiltration of
423 inflammatory cells occurred during RA formation and aggravation³⁰. Of these cells,
424 synovial inflammatory macrophages (SIMs) are key players that involved in various
425 proinflammatory cytokine secretion, leading to the unchecked neutrophil
426 transmigration and initialing osteoclast-dependent cartilage destruction and bone
427 erosion^{8, 9, 31}. To precisely reprogram SIMs for RA treatment, we developed the
428 efferocytosis-inspired nanoimitator (EINI) for terminating the SIMs-initiated
429 pathological cascades, which could serially reconstruct the intra-articular immune
430 homeostasis, conferring a chondroprotection effect, and ultimately restore the joint
431 function.

432 RA is a systemic autoimmune disease¹. Symmetrical inflammatory polyarthritis is
433 the primary clinical manifestation, usually beginning in the small joints of the hands
434 and the feet, spreading later to the larger joints³². Tracking and targeting inflamed joint
435 synovium is of utmost importance for the treatment of this systemic autoimmune
436 disease. Efferocytosis is one of the critical innate functions of macrophages, which
437 maintains tissue homeostasis^{17, 18}. PtdSer exposure on the outer leaflet of the plasma
438 membrane in the apoptotic cells is a key "eat-me" signal for macrophages¹⁹. The
439 recognition of PtdSer by macrophages mainly depends on its head group³³. We
440 demonstrated that PtdSer-coroneted core displayed a boosted internalization by
441 macrophages both in vitro (Fig. 3b) and in vivo (Fig. 5g). Surficial painting of LMWH
442 on the PtdSer-coroneted core subdued the innate vulnerability of PtdSer to phagocytosis
443 of the mononuclear phagocyte system in blood stream, thus avoiding rapid clearance

444 (Fig. 5a). Moreover, the LMWH motif endowed the EINI with an enhanced retention
445 in the inflamed regions (Fig. 5b and Fig. 5c) and a competitively inhibiting function
446 towards P-selectin²² that disrupts the unchecked articular trafficking of NEs (Fig. 6e).
447 Due to the leakage of the intra-articular blood vessel, the EINI subsequently enriched
448 in joint synovium where the shell was exfoliated upon the ROS stimulation, and PtdSer
449 corona was then exposed. In an efferocytosis-like manner, the PtdSer-coroneted core
450 was in turn phagocytosed by SIMs (Fig. 5g).

451 Polymorphisms in IRF5 genes are associated with an increased risk of developing
452 RA (Fig. 1a, b)¹⁵. IRF5 translates various signals related to the pro-inflammatory
453 macrophages in joint synovium, including the secretion of neutrophil attractants-
454 chemokine CXCL1¹⁴. IRF5 deficiency supported a reduction in macrophage-derived
455 CXCL1 and limits neutrophil infiltration into the inflamed joint. Our results
456 demonstrated that IRF5 targeted-silencing with our EINI could repolarize SIMs to anti-
457 inflammatory phenotype (Fig. 4g and Fig. 6f), and terminated the SIMs-initiated
458 pathological cascades, thus inhibiting the unchecked recruitment of NEs (Fig. 4e and
459 Fig. 6e) and the initiation of the osteoclast-dependent cartilage destruction and bone
460 erosion (Fig. 6h, n and Supplementary Fig. 15b). In addition, the metformin in the
461 nanoimitator can prevent complex I-derived ROS and thus promote the SIMs-targeted
462 reprogramming^{24, 34}. Our data indicates the dissociated metformin from the EINI
463 significantly decreased the production of ROS in SIMs (Supplementary Fig. 10), which
464 further blocked the ROS-mediated inflammatory signal and was conducive to
465 reconstructing the intra-articular immune homeostasis synergistically.

466 In sum, our findings establish that synovial inflammatory macrophages (SIMs) can
467 be targeted reprogrammed for RA reversible treatment with the proposed efferocytosis-
468 inspired nanoimitator (EINI). By manipulation of the shielding and the exposure of
469 PtdSer on the nano-formulation, we demonstrated the SIMs in RA joints could be
470 precisely targeted in situ. Functionally, the EINI effectively repolarized SIMs to an
471 immunoregulatory phenotype in situ, serially terminating the SIMs-initiated
472 pathological cascades and ultimately reinstated the articular immune homeostasis. Our

473 work therefore provides a regulating strategy of macrophage heterogeneity for RA
474 reversible treatment, which may be extended to various macrophage-involving
475 autoimmune diseases, such as atherosclerosis, idiopathic pulmonary fibrosis and
476 inflammatory bowel disease.

477 **Materials and Methods**

478 **Materials**

479 Dioleoylphosphatidic acid (DOPA) and dioleoyl phosphatidylserine (PtdSer) were
480 purchased from A.V.T. (Shanghai) Pharmaceutical Co.,Ltd. (Shanghai, China).
481 Metformin, cyclohexane, Igepal CO-520 and zinc nitrate hexahydrate
482 ($Zn(NO_3)_2 \cdot 6(H_2O)$) were obtained from Sigma-Aldrich (St. Louis, MO, USA). Low
483 molecular weight heparin (LMWH, MW 3800~5000, Mw/Mn (PDI) = 1.34) was
484 purchased from Melonepharma (Dalian, China). Recombinant murine macrophage
485 colony stimulating factor (M-CSF) (315-02) and recombinant human TNF- α (300-01A)
486 were purchased from PeproTech (Rocky Hill, USA). RPMI 1640 medium, Dulbecco's
487 modified Eagle medium (DMEM), modified Eagle medium, trypsin EDTA, type I
488 collagenase, fetal bovine serum (FBS), and PBS were obtained from Gibco (USA). DiR
489 (D12731) and PKH67 were purchased from Invitrogen (USA). The anti-CD206
490 antibody, anti-iNOS antibody, anti-Ly-6G antibody and anti-myeloperoxidase (MPO)
491 antibody were purchased from BioLegend. All siRNAs used in CIA model were
492 synthesized by Genepharma (Shanghai, China), and the sequences used were as follows:
493 (i) IRF5: 5'-dTdT-CUG CAG AGA AUA ACC CUG A-dTdT-3' (sense) and 5'-dTdT
494 UCA GGG UUA UUC UCU GCA G dTdT-3' (antisense). A dye was introduced to the
495 5'-end of the antisense strand of siIRF5. (ii) Negative control scrambled siRNA (siN.C).

496 **Human synovium specimens**

497 Human synovial tissue sections were obtained from ACPA⁺ RA patients ($n = 3$) and
498 healthy donors ($n = 3$) of Department of orthopaedic surgery at Shandong Provincial
499 Hospital Afliliated to Shandong First Medical University. All samples were obtained with
500 written informed consent and collected using a standard protocol approved under the
501 Review Board protocol of Shandong First Medical University (Approval no. 2019-272).

502 **Cell culture**

503 Primary macrophages were derived from bone marrow-derived macrophages
504 (BMDMs) of 7-8-week-old mice which were maintained in DMEM supplemented with
505 10% FBS and 1% penicillin/streptomycin. The femurs were dissected from the mouse.
506 Then, the bone marrow was rinsed with 1× PBS to isolate bone marrow cells, which
507 was followed by erythrocyte lysis in ACK lysis buffer. After centrifugation, the cells
508 were resuspended in media with M-CSF (20 ng/ml) to induce the maturation of
509 macrophages. After 72 hours of incubation, macrophage differentiation medium was
510 discarded wash plates with sterile PBS to remove non-adherent cells. The adherent
511 BMDMs were harvested and could be used in subsequent experiments for at least 7
512 days after induction. The BMDMs were collected after detachment with 0.05% trypsin
513 EDTA and centrifugation at 400g for 5 min, and then the BMDMs were stained with
514 PerCP-Cy5.5 conjugated anti-CD11b and FITC conjugated anti-F4/80 antibodies. The
515 induction of mature macrophages was evaluated by flow cytometry (Beckman Coulter,
516 USA).

517 Mouse neutrophils were collected from whole blood of C57BL/6 mice using the
518 Percoll gradient method³⁵. Briefly, before isolation of neutrophils, 1 mg/kg LPS was
519 injected intraperitoneally into the mice to activate neutrophils in vivo. After 12 h,
520 conduct the second injection. Three hours after the second injection, blood was
521 collected by submandibular bleeding. Pooled blood was centrifuged (3,220g, 5 min,
522 4 °C) and the buffy coat on top was added into a Percoll mixture solution consisting of
523 78%, 69% and 52% (v:v) Percoll in PBS, followed by centrifugation at 1500g for 30
524 min. The cell from the interface of the 69% and 78% gradient layers and the 78%
525 fractions were recovered to a new tube. ACK lysis buffer was then added to the sample
526 to lyse the erythrocytes. Neutrophils were purified by washing with ice-cold 1×PBS
527 three times, and resuspend cells by culture medium (fig. S7).

528 **Synthesis of NBC-PtdSer**

529 The NBC-PtdSer was synthesized according to the procedure in [Supplementary](#)
530 [Scheme 1, 2](#). First, synthesis of NBC: 4-(Hydroxymethyl)phenylboronic acid pinacol

531 ester (0.25 g, 1.1 mmol) was dissolved in 10 ml of dry THF. Triethylamine (0.3 ml, 2.15
532 mmol) was added followed by 4-nitrophenyl chloroformate (0.27 g, 1.3 mmol) and the
533 reaction was allowed to stir at room temperature for 12 h. The reaction was diluted with
534 ethyl acetate and washed with 1.0 M HCl followed by saturated NaHCO₃. The organic
535 layer was dried over MgSO₄, filtered and concentrated. NBC was purified on a silica
536 gel column eluting with 5% EtOAc in hexanes to give 0.26 g (0.65 mmol, 61 % yield)
537 as a white solid. The ¹H NMR spectrum of the product was obtained on a Bruker
538 Advance DRX-400 spectrometer (Germany) with tetramethylsilane (TMS) as internal
539 standard.

540 Then, synthesis of NBC-PtdSer: PtdSer (100 mg, 0.1344 mmol) and NBC (64.4 mg,
541 0.1613 mmol) were dissolved in dry DMF (3 ml) under an N atmosphere. After adding
542 triethylamine (56 µl, 0.4032 mmol), the reaction was stirred 12 h before being quenched
543 by pouring into 100 ml of 0.5 M HCl. The aqueous layer was extracted with 3 × 25 ml
544 chloroform. The combined organic layer was washed with 5 × 100 ml water and once
545 with brine. The collected organic phase was dried over Na₂SO₄, filtered, and
546 concentrated under reduced pressure. The resulting crude was subjected to silica gel
547 column chromatography. Gradient elution from 100% chloroform to 20% MeOH in
548 chloroform was needed to purify the product as a white solid.

549 **Preparation and characterization of nano-formulation**

550 siIRF5@EINI were prepared using a modified water-in-oil reverse micro-emulsion
551 protocol. Briefly, 500 µl of 30 mM Zn(NO₃)₂·6(H₂O) and 100 µl siIRF5 (siRNA at an
552 input concentration of 200 nM) were dispersed in 20 ml cyclohexane/Igepal CO-520
553 (71/29V/V) solution to form a very well dispersed water-in-oil reverse micro-emulsion.
554 To prepare the metformin loaded microemulsion, metformin solution (100 µl, 30 mM)
555 and DOPA (100 µl, 35 mM) were added into a separate 20 ml oil phase. After mixing
556 the above two microemulsions for 30 min to form the condensed cores, 40 ml of ethanol
557 was added and the mixture was centrifuged at 12,500g for 20 min to remove
558 cyclohexane and the surfactant. To prepared PtdSer-coroneted core, the pure pellets
559 were suspended in 750 µl chloroform and mixed with 4 mg NBC-PtdSer. And then

560 evaporating the chloroform, 4 ml of 1× PBS was added to form PtdSer-coronated core.
561 Additionally, the corona of nanocore was cloaked with LWMH via a reversible boronate
562 ester linker by the reaction of cis-diols and phenylboronic acid groups, and the
563 efferocytosis-inspired nanoimitator was achieved.

564 The morphology and size of the nanoimitator was observed under a Hitachi H-7650
565 TEM (Hitachi, Japan) after negative staining with uranyl acetate solution (2%, wt/wt).
566 The particle size distribution and zeta potential were measured by dynamic light
567 scattering (DLS) using a Malvern Zetasizer Nano ZS. The PtdSer-presenting was
568 validated by flow cytometry. Briefly, the nanoimitator was resuspended in PBS at pH
569 7.4 were incubated with or without H₂O₂ (0.1 mM). Then, these nano-formulations
570 were incubated with FITC-Annexin V at the present of Ca²⁺ for 15 min respectively
571 and the binding rate of FITC-Annexin V was detected by flow cytometry. To quantify
572 metformin release, the nanoimitator was resuspended in PBS at pH 5.0 or pH 7.4. At
573 predetermined time points, aliquots from each group were centrifuged to pellet the
574 nanoparticles, and the concentration of released metformin in the supernatant was
575 measured using an HPLC system (Agilent 1100, USA) under the following
576 chromatographic conditions: column, C18 column (5×250 mm, particle size 5 μm);
577 mobile phase, methyl alcohol-15 mM KH₂PO₄ (60:40, v/v); flow rate, 1.0 ml/min;
578 detection wavelength, 233 nm.

579 **Gel retardation assay**

580 Free siIRF5 and siIRF5@EINI (in 1×PBS) was separately mixed with loading
581 buffer and loaded into 3% wt agarose gel with NA-Green (Beyotime Biotechnology
582 Co., Ltd, China). Electrophoresis was conducted in 1×tris-acetate EDTA (TAE) buffer
583 at 80 V for 30 min. To investigate the pH responsiveness, siIRF5@EINI was
584 resuspended in PBS at pH 5.0 or pH 7.4 for 1 h. After that, the samples were analyzed
585 by gel electrophoresis. The resulting gels were analysed using a UV illuminator (IS-
586 2200; Alpha Innotech, San Leandro, CA, USA) to show the location of siIRF5.

587 **Macrophages targeting abilities of EINI in vitro**

588 BMDMs or FLSs were seeded in 6-well tissue culture plates at 50% confluency and
589 cultured overnight. Cell culture medium was changed and 10 ng/ml recombinant human
590 TNF- α was added to stimulate 4 h. The nanoimitator was then added into the cells with
591 or without H₂O₂ (0.1 mM) pre-treating. After incubation, cells were washed three times
592 with ice-cold PBS and fixed with 4% paraformaldehyde, then the cells were incubated
593 at 37 °C for 10 min with 10 nM DAPI, and imaged with an EVOS inverted fluorescence
594 microscope (Thermo Fisher Scientific). For flow cytometric analysis, cells were
595 scraped and collected after PBS wash, then analysed with a Beckman Coulter Gallios
596 flow cytometer. The results were analyzed using FlowJo software. To verify the specific
597 macrophages homing of the nano-formulation, PKH67-labeled FLSs were cocultured
598 with the unlabeled macrophages at a ratio of 1:1. Cells were stimulated with 10 ng/ml
599 TNF- α for 4 h at 37 °C. H₂O₂ pretreated nanoimitator was added to the medium for 2
600 hours. The uptake of the nanoimitator was observed by CLSM.

601 **siRNA tracking and release in the cell**

602 To investigate endocytic pathways and intracellular release behavior of siIRF5,
603 macrophages were treated with the siIRF5@EINI that were pre-conditioned with the
604 0.1mM of H₂O₂. At predetermined time, the treated cells were rinsed with PBS and
605 incubated at 37 °C for 30 min with 100 nM LysoTracker Green in cell culture medium.
606 After incubation, cells were washed with ice-cold PBS, fixed with 4% PFA 30 min at
607 4 °C, and followed by DAPI staining. The fluorescence imaging was performed by
608 CLSM.

609 **Evaluation of the effects of nanoimitator on the adhesion and transmigration** 610 **neutrophils**

611 For assaying the intervene effect on cell adhesion, 2 \times 10⁵ HUVECs were seeded in
612 6-well plate and pre-stimulated by incubation for 6 h with TNF- α (10 ng/ml). Then
613 treated with RPMI medium, P-selectin antagonist (Human P-selectin/CD62P antibody,
614 R&D Systems) and the nanoimitator for 30 min, respectively. 1 \times 10⁶ activated
615 neutrophils were seeded on top of the treated endothelial monolayer. After 30 min at
616 37 °C under gentle shaking, each well was washed twice with PBS to remove the non-

617 adhered cells and labeled with FITC-labeled anti-Ly-6G antibody. Finally, evaluation
618 of neutrophils adhered to HUVEC monolayers using CLSM and flow cytometry.

619 The inhibitory effects of the nanoimitator on the migration of neutrophils were
620 evaluated by transwell assays. Prior to the experiment, the isolated neutrophils were
621 pre-stimulated with 10 μ M fMLF for 15 min at 37 °C. The activated macrophages were
622 treated with different formulations (the free siIRF5, 0.1 mM H₂O₂-pretreated
623 siN.C@EINI and siIRF5@EINI) for 24 h. Cell media was collected and the bottom of
624 transwell chambers was filled with cell media from each formulation. Then, 1×10^6 of
625 preactivated neutrophils were seeded on 3 μ m polycarbonate membrane. After
626 incubation for 1 h, the neutrophils that migrated to the lower chamber were stained with
627 crystal violet and counted. The results were expressed as a percentage of the total
628 number of neutrophils added.

629 **Gene silencing assay**

630 IRF5 gene silencing efficiency of the nanoimitator was investigated by quantitative
631 real-time PCR (qRT-PCR). Macrophages were seeded in a six-well plate (1×10^6 cells
632 per well) in growth medium for 24 hours. Cells were treated with TNF- α (10 ng/ml) for
633 6 h. The cells were incubated with the desired nano-formulations in 2 ml of fresh media.
634 After 48 h, the cells were washed with PBS and the total RNA was extracted using the
635 RNeasy Mini Kit (Qiagen) according to the manufacturer's instructions. Real-time
636 quantitative PCR was performed to measure levels of mRNAs in macrophages
637 encoding IRF5. β -actin was chosen as an internal housekeeping gene to normalize the
638 IRF5 mRNA. The mRNA expression level was calculated based on comparative Ct
639 method ($2^{-\Delta\Delta C_t}$). IRF5 blockade would have a major impact on reducing secretion level
640 of macrophage-derived CXCL1, hence the CXCL1 mRNA expression was then
641 evaluated by qRT-PCR. Real-time PCR was performed using specific primers
642 (supplementary Table 1).

643 **Phenotype switching of macrophages**

644 For fluorescence immunostaining, macrophages were stimulated with TNF- α (10

645 ng/ml) to induce inflammation. Then, H₂O₂-pretreated nano-formulations were added
646 to the media, and PBS and free siRNA were prepared in parallel for comparison. After
647 4 hours, the cells were incubated for another 24 hours. Cells were fixed with 4% PFA
648 for 15 min and permeation with 0.1% Triton X 100 (room temperature, 10 min), the
649 cells were blocked with normal rat serum (37 °C, 1 h) and incubated with anti-iNOS or
650 anti-CD206 antibody at 4 °C overnight, followed by counterstaining with DAPI for 10
651 min. Fluorescence imaging was performed with confocal microscopy. qRT-PCR
652 analysis was performed with the same procedure as mentioned above. The primer
653 sequences were shown in supplementary Table 1.

654 **Collagen-induced arthritis (CIA) model and treatment protocols**

655 Autoimmune collagen-induced arthritic (CIA) mouse model was established by the
656 following protocol. Immunization type II collagen (CII) lyophilized powder was
657 dissolved in acetic acid (0.1 mol/l) and blended with an emulsion of complete Freund's
658 adjuvant (CFA) in equal volume. Employed homogenizer to emulsify the mixture
659 adequately in an ice bath. Next, healthy DBA/1 mice were intradermally injected the
660 adjuvant (100 µl) at the tail root. The 2 times the dose immunization boost was given
661 on day 21. The control group administrated solvent (acetic acid) injection.

662 To study therapeutic efficacy with CIA mice, the treatments began at day 7 after the
663 second immunization. The CIA animals were randomly assigned into four groups (1
664 nmol of siRNA dose per mouse, 10 mg/kg metformin-equivalent dose, $n = 5$). 200 µL
665 PBS (G1); free siIRF5 solution (G2); siIN.C@EINI (G3) and siIRF5@EINI (G4) were
666 administrated by tail vein twice a week from day 28 to 60.

667 **Study of the biodistribution of EINI in CIA model**

668 After boosting immunization for 7 days, DiR labeled nanoimitator was
669 intravenously injected to CIA mice via tail vein. At predetermined time, the real-time
670 imaging of nanoparticles in arthritis area was observed by the IVIS system (CRI, Inc.,
671 excitation: 748 nm, optical filter: 780 nm). At 48 h post injection, these mice were
672 sacrificed to harvest the major organs (heart, liver, spleen, lung and kidneys) and
673 inflamed paws, quantitatively assaying the clearance ratio of nanoformulations in vivo.

674 Specially, fluorescence molecular tomography computed tomography (FMT-CT)
675 on day 2 postinjection to identify nanoimitator localization in the inflamed regions. For
676 observation of the specific homing of the nanoimitator to inflamed synovium, 100 μ l
677 of Cy5.5 labeled nano-formulation were injected into CIA mice via the tail vein, with
678 free Cy5.5 labeled siIRF5 as a control. After 72 h, the mice were euthanized, and the
679 synovium was obtained and fixed with 4% PFA 48 h at 4 °C. After dehydration and
680 embedded in Tissue-Tek OCT Compound (Sakura Finetek) for frozen sectioning. And
681 then the slides were stained with anti-F4/80 antibodies, followed by DAPI staining. The
682 fluorescence imaging was performed by CLSM.

683 **qRT-PCR analysis of IRF5 expressions from collected synovial macrophages**

684 Fresh synovial tissues were digested with 1 mg/ml type I collagenase in HBSS, and
685 incubated at 37 °C, 5% CO₂ in a humidified atmosphere for 30-45 min. Disaggregated
686 tissue elements were passed through a 70 μ m cell strainer. Synovial macrophages
687 (CD45+, CD11b+, F4/80^{high}, Ly6G-, Ly6C^{Low})³⁶ were sorted on a BECKMAN Moflo
688 Astrios EQ (BECKMAN Coulter). Next, total RNA was isolated by following
689 manufacturer instructions. Synthesis of cDNA and qRT-PCR was performed as
690 previously described in vitro analysis.

691 **Histological analysis of inflamed joints**

692 Histopathology was evaluated to determine the features of RA including neutrophils
693 migration, macrophage phenotype, angiogenesis, synovitis, and cartilage destruction.
694 At study endpoints, mice were euthanized and hind ankle joints were collected for H&E
695 staining, safranin-O or TRAP staining. To detect infiltration of neutrophils, joint
696 sections were dewaxed, and stained with primary antibodies anti-MPO (Bioss
697 Antibodies). Biotinylated anti-rabbit IgG was used as the secondary antibody for
698 chromagen development. For evaluation of phenotype switching of macrophages,
699 tissue sections were stained with antibodies anti-iNOS or anti-CD206. Tissue sections
700 were then stained with DAPI mounting solution, and observed using CLSM.

701 Dissected the hind paw joints were evaluated with micro-computed tomography

702 (micro-CT) imaging. Micro-CT scanning parameters were voxel size, 9 μm ; voltage,
703 90 kV; current, 88 μA ; exposure time, 4 min. To compare the therapeutic effect in each
704 group, three-dimensional images were reconstructed using Mimics software
705 (Materialise) and the mean bone mineral density (BMD) of the joints, were calculated
706 by software CTAn (Skyscan).

707 **Histological scoring of mouse arthritic joints**

708 Histopathological scoring was performed as described grading scheme³⁷. Briefly,
709 joints of arthritic mice were assigned scores of 0 to 4 for inflammation based on H&E
710 staining, according to the following criteria: 0 = normal; 1 = minimal infiltration of
711 inflammatory cells in periarticular area; 2 = mild infiltration; 3 = moderate infiltration;
712 and 4 = marked infiltration. Cartilage depletion was identified by diminished Safranin
713 O staining of the matrix and was scored on a scale of 0 to 4, where 0 = no cartilage
714 destruction (full staining with Safranin O), 1 = localized cartilage erosions, 2 = more
715 extended cartilage erosions, 3 = severe cartilage erosions, and 4 = depletion of entire
716 cartilage. Histologic evaluations were performed in a blinded manner.

717 **Statistical analysis**

718 Results are presented as mean \pm s.d. Error bars represent the s.d. of the mean from
719 independent samples. Two-tailed Student's t-test was applied to test the statistical
720 significance of difference between two groups. For multiple comparisons, the data were
721 analyzed using a one-way analysis of variance (ANOVA). All statistical analyses were
722 carried out with Prism software package (PRISM 8.3.0; GraphPad Software). Statistical
723 significance was set at $*P < 0.05$, $**P < 0.01$, $***P < 0.001$, $****P < 0.0001$.

724 **Data and materials availability:** The data that support the plots within the paper and
725 other findings of this study are available from the corresponding author upon reasonable
726 request. IRF5 mRNA expression level in patients with RA retrieved from the
727 ArrayExpress (<https://www.ebi.ac.uk/arrayexpress/>) under Accession code E-MTAB-
728 6141³⁸.

729

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843 **Author contributions:** X.J. and S.Z. conceived the study and designed experiments.
844 S.Z. and Q.C. characterized properties of the materials. S.Z., Q.C., C.T. and Z.L.
845 performed the animal experiments. Z.M., C.C., J.Z., Y.L. P.S., R.Z., Z.Y., Y.W. and M.H.
846 contributed to data analysis and interpretation. S.Z. crafted all the figures and wrote the
847 manuscript. X.J., M.M., W.L., X.W., G.Y., B.S., and J.L. edited and revised the
848 manuscript and supervised the research.

849 **Competing interests:** The authors declare that they have no competing interests.

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Supplementary Information for
Remodeling Articular Immune Homeostasis with an Efferocytosis-inspired
Nanoimitator Mitigates Rheumatoid Arthritis

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Yan Wang¹, Xia Wei⁵, Jun Li⁵, Wei Li^{2,3}, Mohnad Abdalla¹, Gongchang Yu⁶, Bin Shi⁶,
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899 derived ROS.

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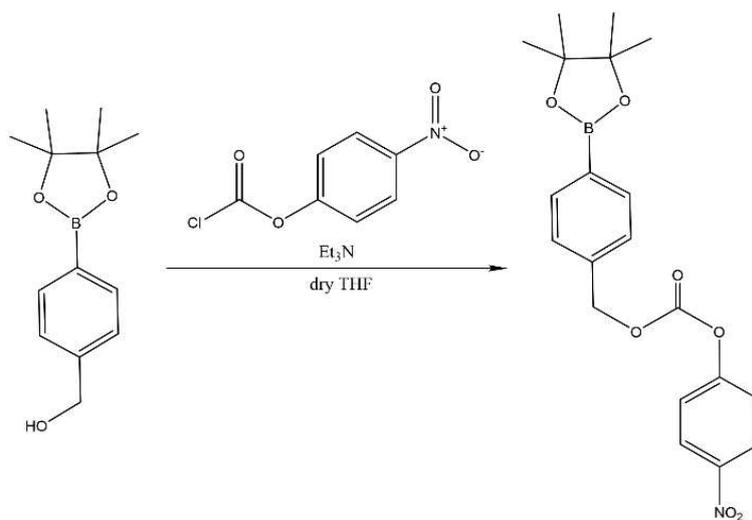
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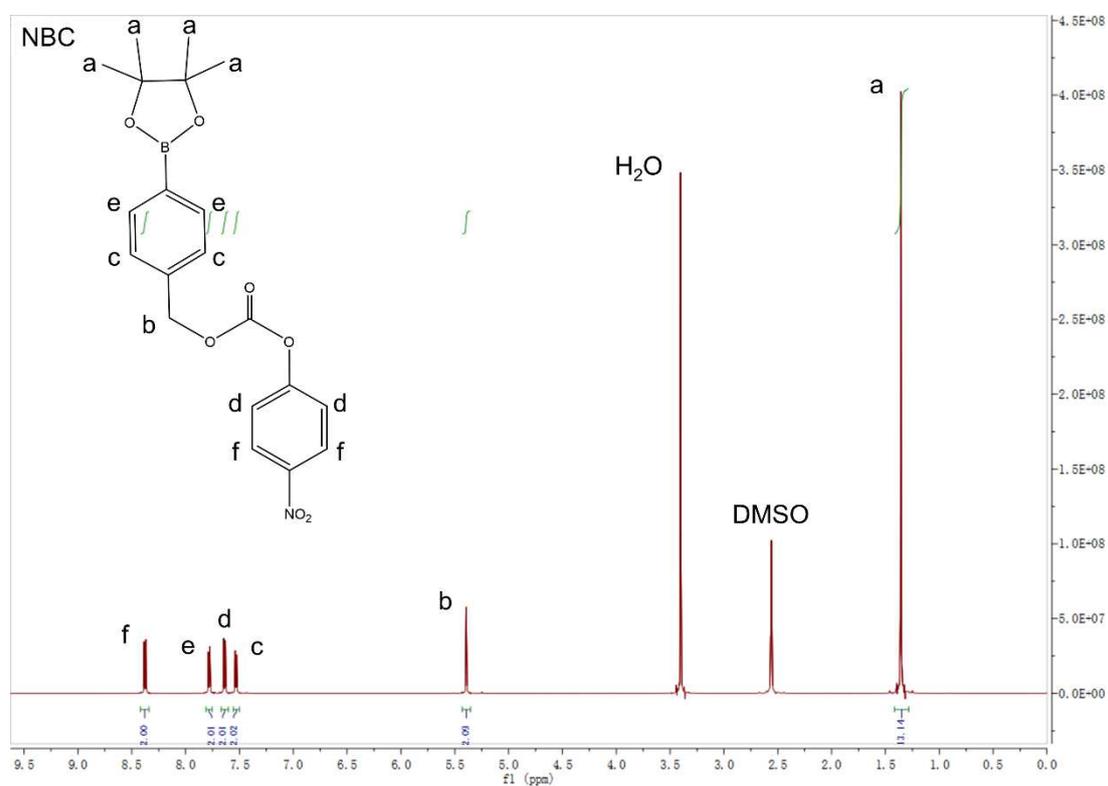
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907 analyses of the TRAP-stained osteoclasts.

908 Supplementary Table 1. List of primers used for real-time PCR.

909 **Supplementary Scheme 1. Synthetic of NBC.**



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912 **Supplementary Figure 1. ¹H NMR spectrum of NBC in DMSO-d₆.**

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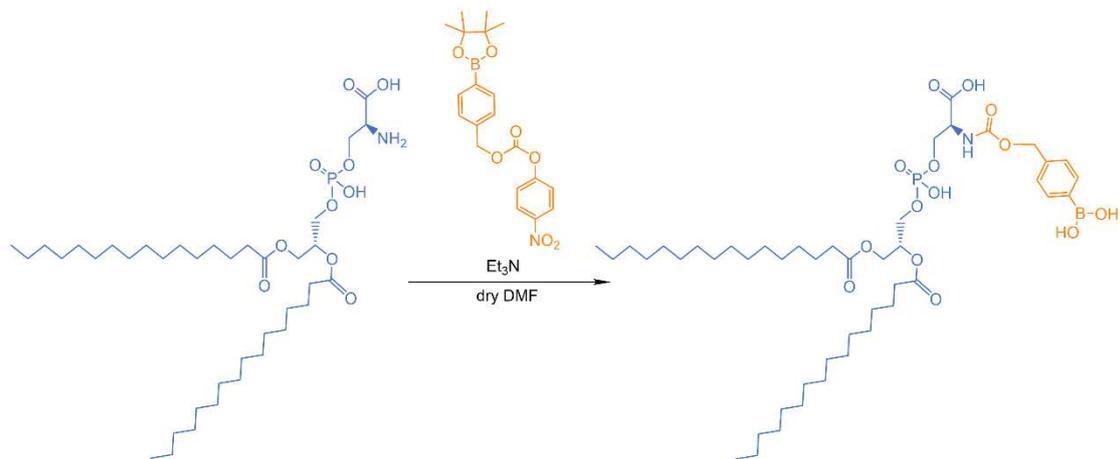
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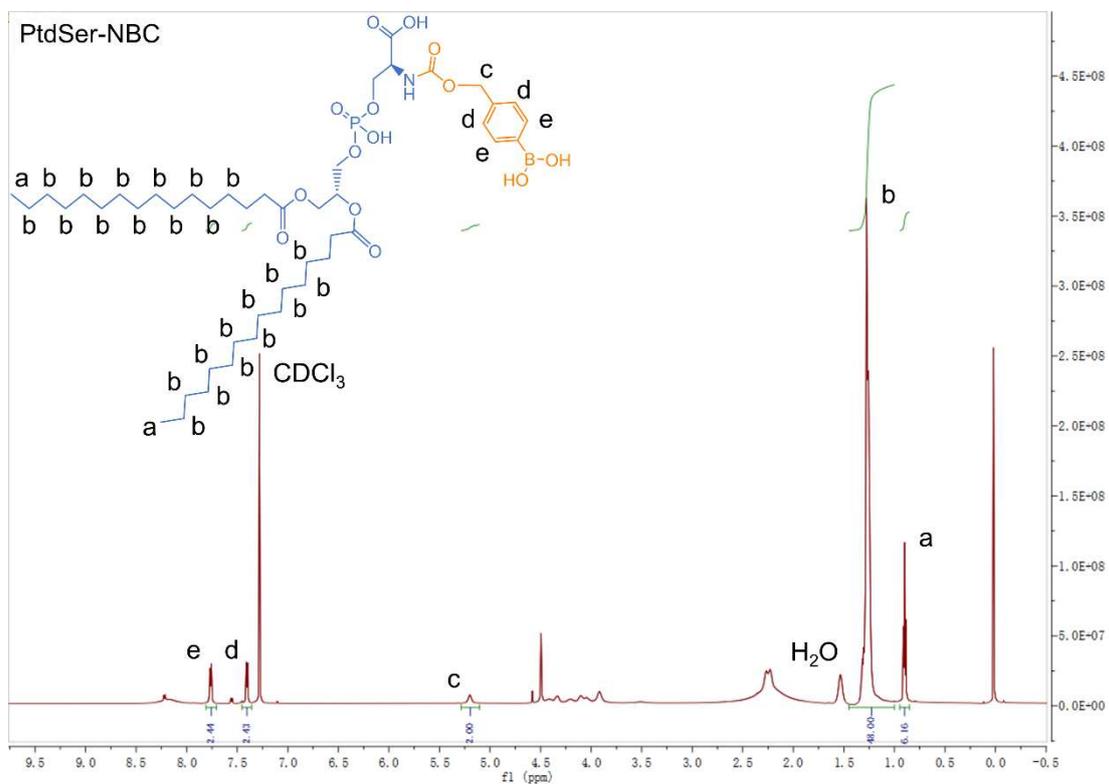
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920 **Supplementary Scheme 2. Synthetic route developed to produce PtdSer-NBC.**



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Supplementary Figure 2. ¹H NMR spectrum of PtdSer-NBC in CDCl₃.

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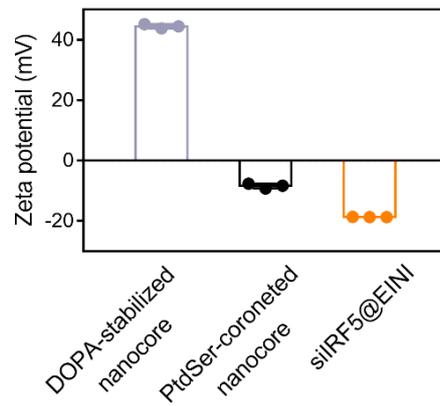
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935 **Supplementary Figure 3. Zeta potential of DOPA-stabilized nanocore, PtdSer-**
936 **coroneted nanocore and siIRF5@EINI.**

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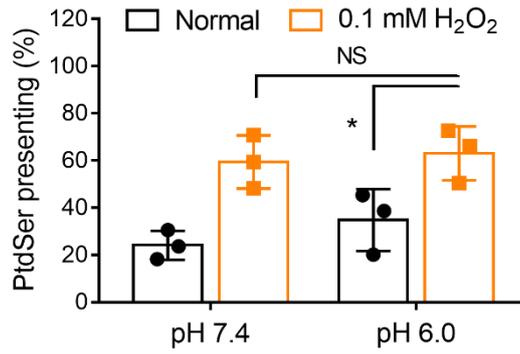
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966 **Supplementary Figure 4. Flow cytometry analysis of PtdSer-presenting.** The
 967 nanoimitator was resuspended in PBS at pH 7.4 or pH 6.0 respectively, and was
 968 incubated with or without H₂O₂ (0.1 mM). Data are presented as the mean ± s.d. (*n* = 3
 969 independent experiments). ***P* < 0.01. NS, not significant.

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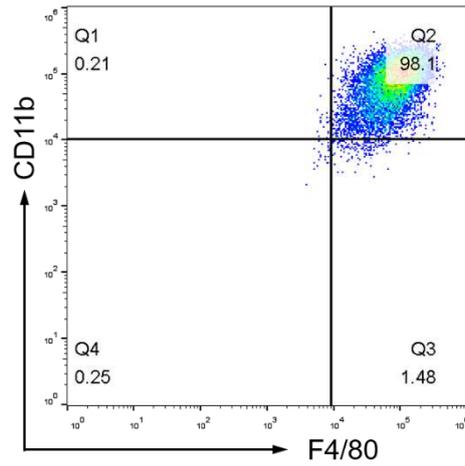
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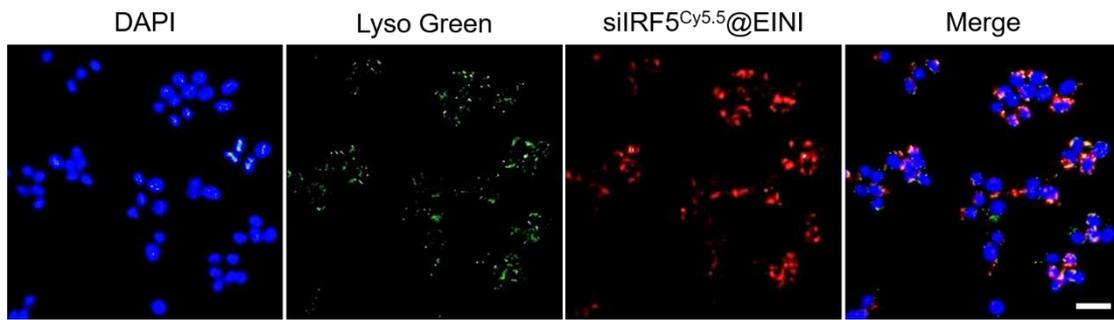
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999 **Supplementary Figure 5. The expression levels of CD11b and F4/80 in BMDMs**
1000 **analyzed by flow cytometry.**

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1030 **Supplementary Figure 6. Fluorescent visualization of siRNA localization in**
1031 **macrophages 6 hours after incubation with nanoimitator. Scale bars, 50 nm.**

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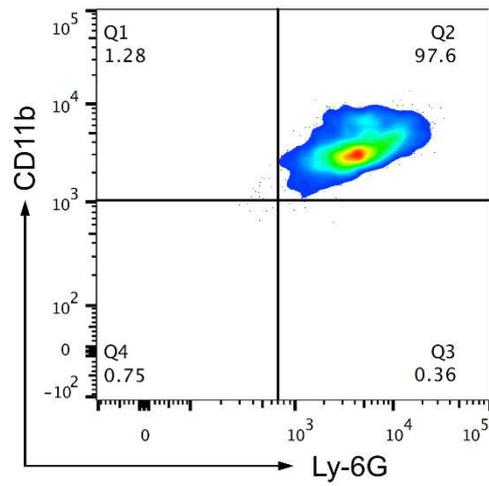
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1066 **Supplementary Figure 7. Flow cytometric analysis of the purity of neutrophils**
1067 **doubly stained with FITC-conjugated Ly-6G and PerCP-Cy5.5-conjugated**
1068 **CD11b antibodies.**

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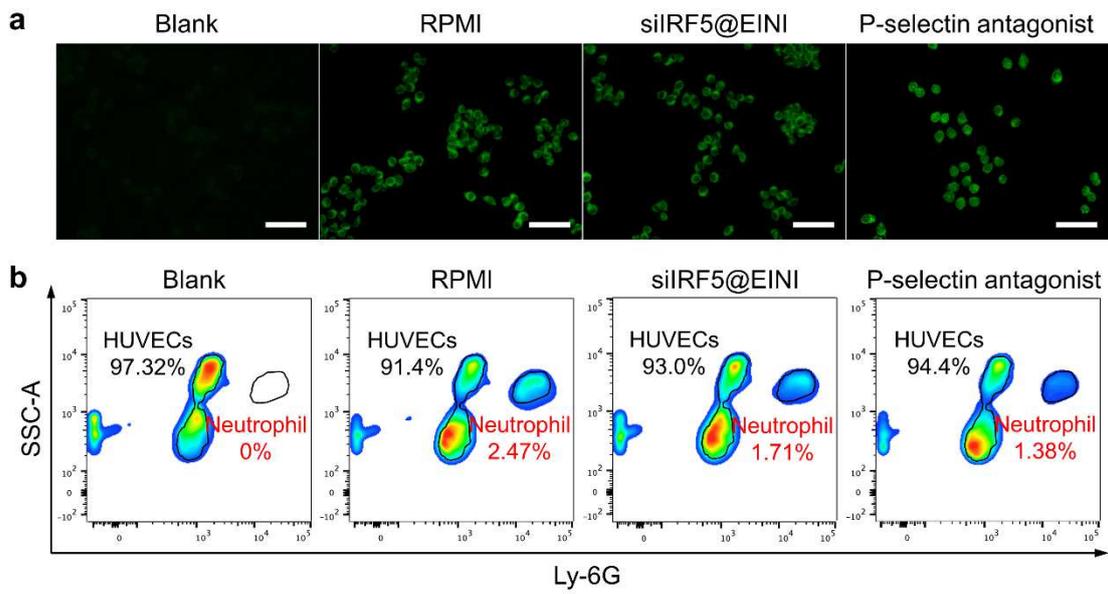
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1098 **Supplementary Figure 8. Effective intervention by nanoimitator on inflammatory**
1099 **neutrophils. a,** Evaluation of neutrophil adhesion to a HUVECs monolayer after
1100 incubation with nanoimitator. Scale bars, 50 nm. **b,** FACS analysis of neutrophil
1101 adhesion to HUVECs. All values are expressed as the mean \pm s.d. ($n = 3$ independent
1102 experiments).

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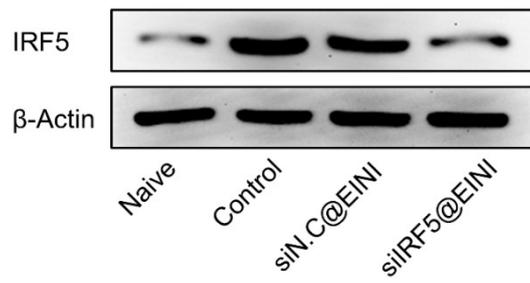
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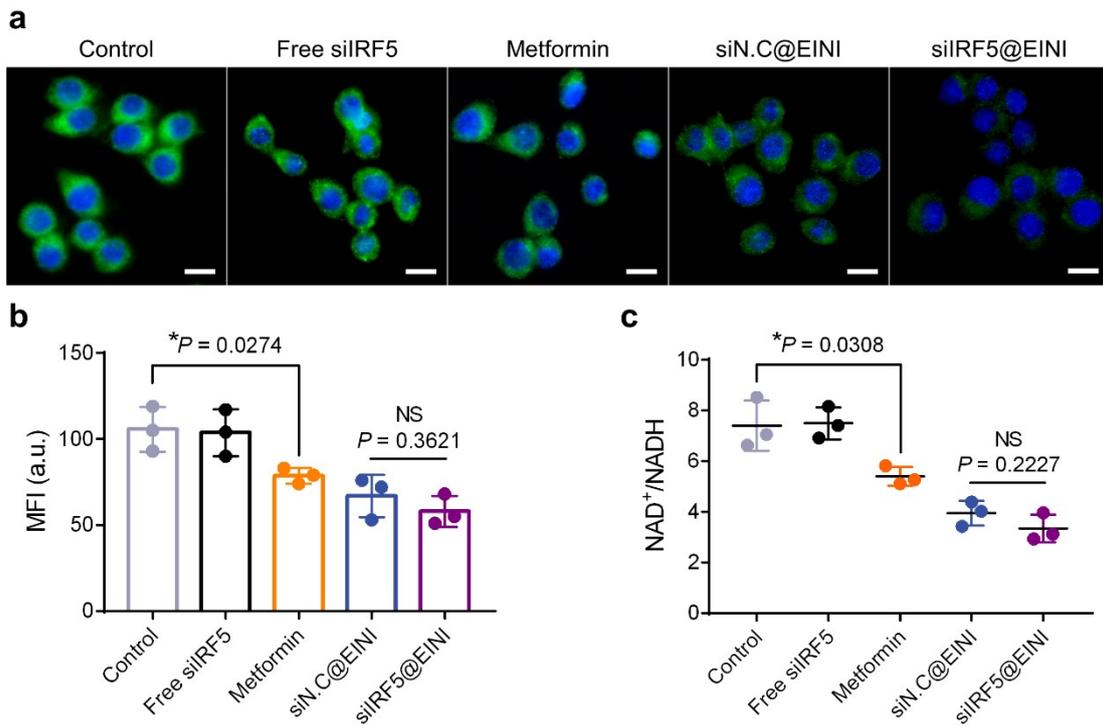
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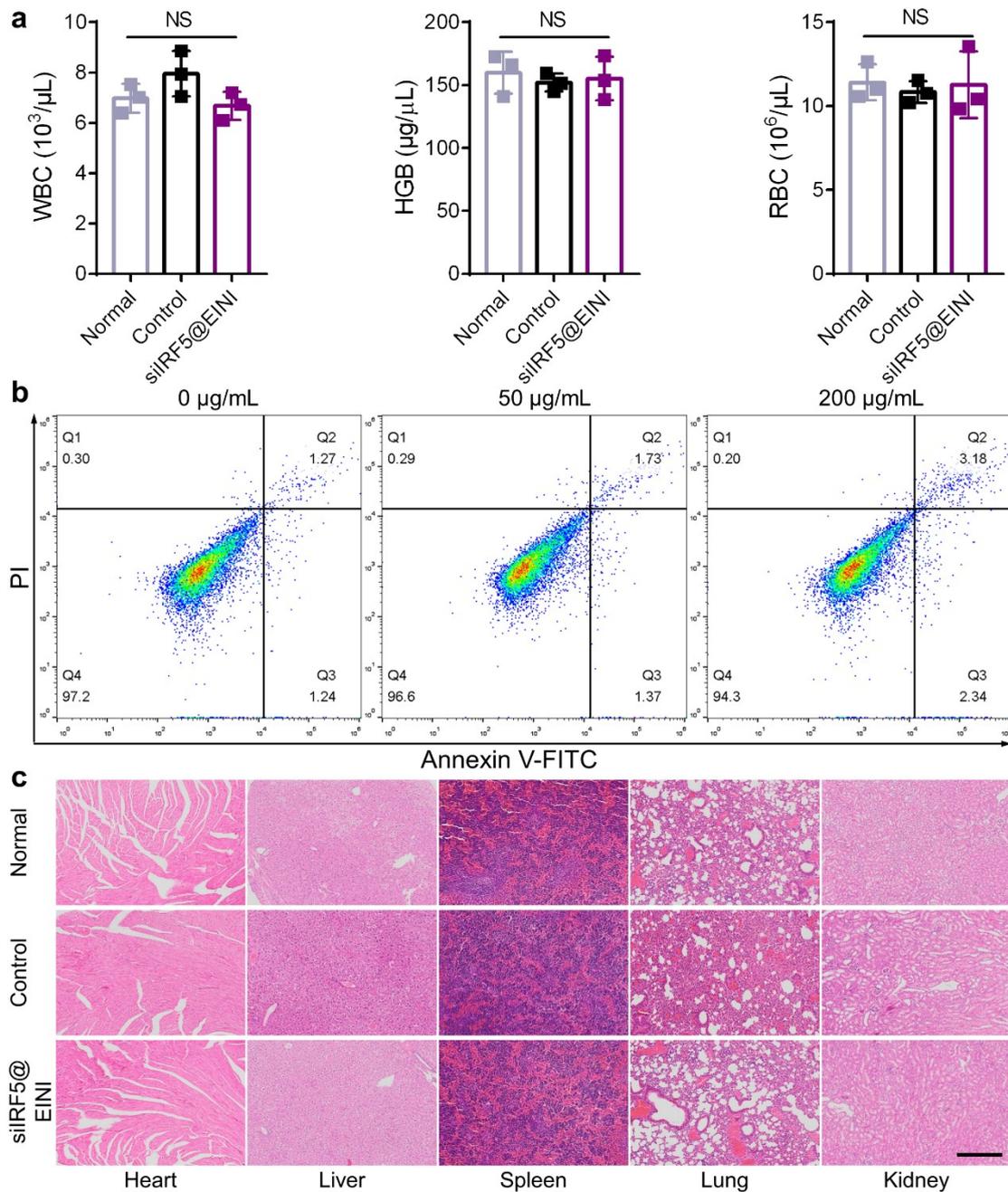
1125 **Supplementary Figure 9. IRF5 expression in macrophages treated with different**
1126 **formulations, detected by western blot.**

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1162 **Supplementary Figure 10. Metformin reprogramming mitochondrial metabolism**
 1163 **in macrophages, which act as mitochondrial complex I (CI) inhibitor directly**
 1164 **inhibit CI-derived ROS. a**, Fluorescence images and the quantification result (**b**) of
 1165 intracellular ROS generation in macrophages stained with DCFH-DA. The scale bars
 1166 were 20 μ m. **c**, The levels of oxidized and reduced nicotinamide adenine dinucleotide
 1167 ($NAD^+/NADH$) were measured in macrophages using a colorimetric assay. All values
 1168 are expressed as the mean \pm s.d. ($n = 3$ biologically independent animals). * $P < 0.05$.
 1169 NS, not significant.

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Supplementary Figure 11. Systemic toxicity evaluation of siIRF5@EINI. **a**, During the experimental cycle, mice were followed up to sacrifice on day 60, the change of white blood cell (WBC), hemoglobin (HGB) and red blood cell (RBC) were measured. Data are presented as the mean \pm s.d. ($n = 3$ independent experiments). NS, not significant. **b**, Apoptosis of T cells extracted from the spleens of mice that received i.v. injections of siIRF5@EINI during the experimental cycle. Mice injected with PBS were used as a negative control. **c**, Histological sections of major organs at 60th day. Scale bar, 50 μm . ($n = 3$ biologically independent animals).

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1193 **Supplementary Figure 12. Representative lesions image from macroscopic**
1194 **observation of the hind paws in different treatment groups. Scale bar, 3 mm. (*n* =**
1195 **5 biologically independent animals).**

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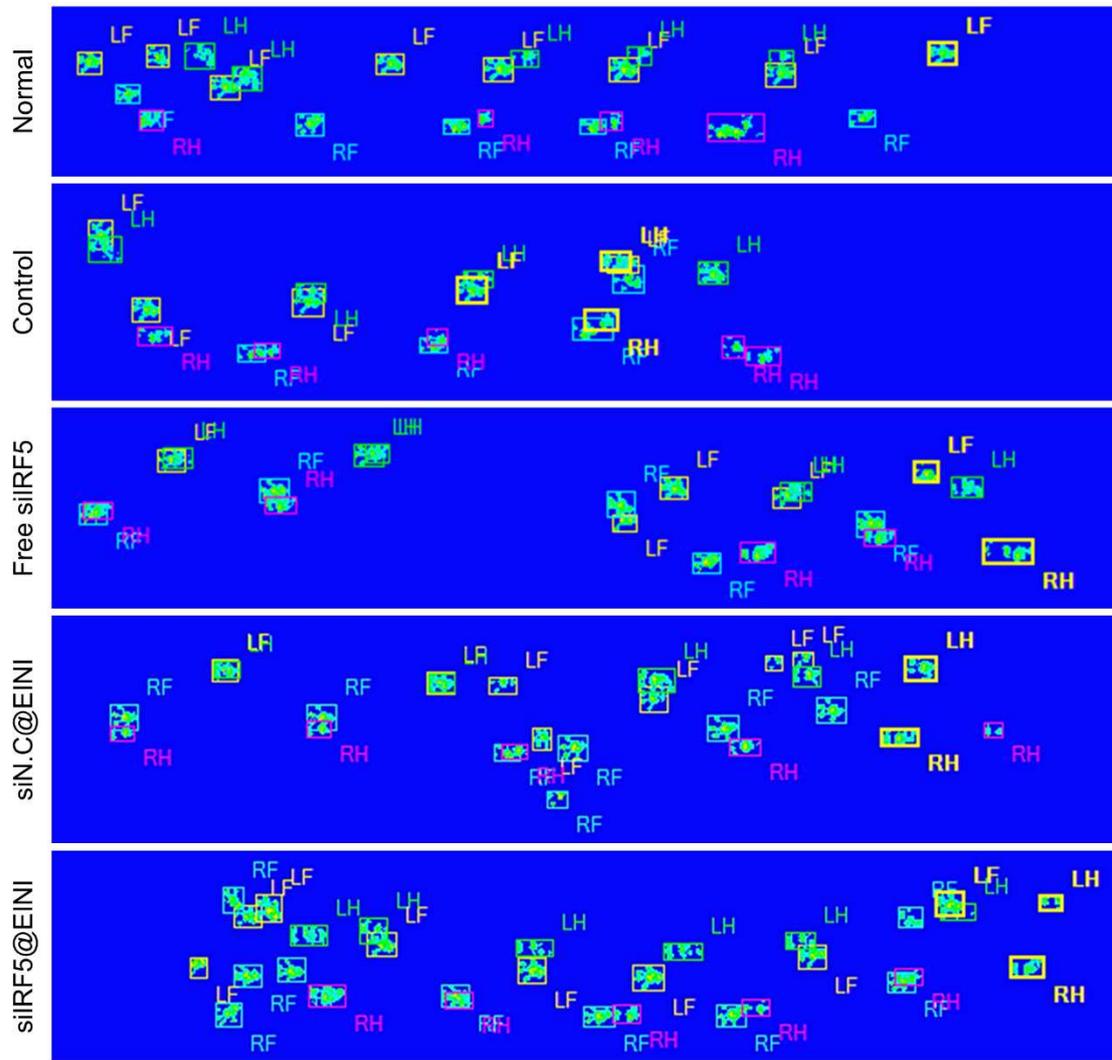
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Supplementary Figure 13. Representative signal images of the footprint assay.

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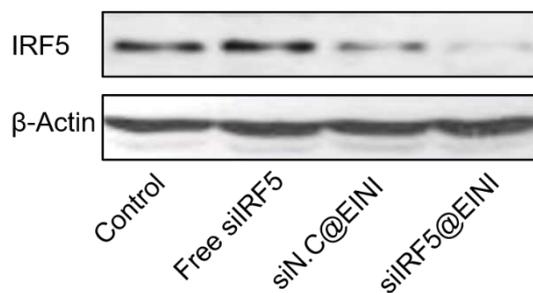
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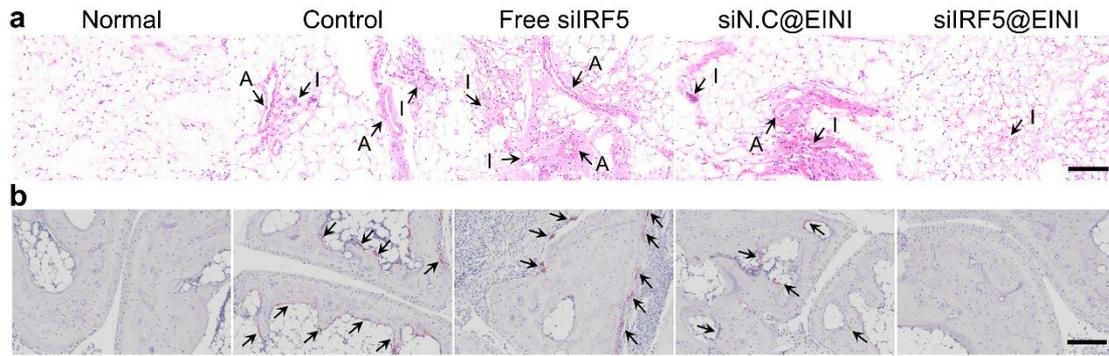
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1249 **Supplementary Figure 14. Western blot images of IRF5 protein levels in the**
1250 **synovial macrophages in different formulations treated group.**

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1284 **Supplementary Figure 15. H&E staining of synovium and immunohistochemical**
1285 **analyses of the TRAP-stained osteoclasts. a**, H&E staining of synovium extracted
1286 from mice of different treatment groups. I, immune cell infiltration; A, angiogenesis.
1287 Scale bars = 50 μ m. ($n = 5$ biologically independent animals). **b**, Immunohistochemical
1288 analyses of the TRAP-stained osteoclasts in the joint tissues from rats receiving the
1289 indicated treatment. Scale bar = 100 μ m. ($n = 5$ biologically independent animals).

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Supplementary Table 1. List of primers used for real-time PCR

IRF5	Forward	5'-CATTCAGCGGGAAGTCAAGA-3'
	Reverse	5'-TGTCTGCCGACCAAGAAAGC-3'
CXCL1	Forward	5'-CACCCAAACCGAAGTCATAGC-3'
	Reverse	5'-GGGGACACCTTTTAGCATCTTT-3'
iNOS	Forward	5'-AGCCAAGCCCTCACCTACTT-3'
	Reverse	5'-CTCTGCCTATCCGTCTCGTC-3'
TNF- α	Forward	5'-TGGAAGTGGCAGAAGAGGCAC-3'
	Reverse	5'-AGGGTCTGGGCCATAGAAGTGA-3'
Arg1	Forward	5'-TGCTTAGCTCTGTCTGCTTTGC-3'
	Reverse	5'-GAACACGGCAGTGGCTTTAAC-3'
CD206	Forward	5'-GGAGGCTGATTACGAGCAGT-3'
	Reverse	5'-CATAGGAAACGGGAGAACCA-3'
β -actin	Forward	5'-CTACAATGAGCTGCGTGTGG-3'
	Reverse	5'-CAGGTCCAGACGCAGGATGGC-3'

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