

1 **In silico, in vitro, and in vivo models reveal EPHA2 as a target for decreasing**  
2 **inflammation and pathological endochondral ossification in osteoarthritis.**

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28

29 **Abstract**

30 Low-grade inflammation and pathological endochondral ossification are processes  
31 underlying the progression of osteoarthritis, the most prevalent joint disease  
32 worldwide. In this study, data mining on publicly available transcriptomic datasets  
33 revealed EPHA2, a receptor tyrosine kinase associated with cancer, to be associated  
34 with both inflammation and endochondral ossification in osteoarthritis. A  
35 computational model of cellular signaling networks in chondrocytes predicted that in  
36 silico activation of EPHA2 in healthy chondrocytes increases inflammatory mediators  
37 and triggers hypertrophic differentiation, the phenotypic switch characteristic of  
38 endochondral ossification. We then evaluated the effect of inhibition of EPHA2 in  
39 cultured human chondrocytes isolated from individuals with osteoarthritis and  
40 demonstrated that inhibition of EPHA2 indeed reduced inflammation and hypertrophy.  
41 Additionally, systemic subcutaneous administration of the EPHA2 inhibitor ALW-II-41-  
42 27 attenuated joint degeneration in a mouse osteoarthritic model, reducing local  
43 inflammation and pathological endochondral ossification. Collectively, we  
44 demonstrate that pharmacological inhibition of EPHA2 with ALW-II-41-27 is a  
45 promising disease-modifying treatment that paves the way for a novel drug discovery  
46 pipeline for osteoarthritis.

47

48      Keywords: Chondrocyte hypertrophy, inflammation, kinase inhibitor, virtual cell, drug

49      target

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71 **Introduction**

72 Osteoarthritis (OA) stands as the most widespread and debilitating  
73 musculoskeletal condition worldwide, characterized by progressive joint  
74 alterations leading to pain and functional limitations [1, 2]. In OA, joint  
75 inflammation ensues, resulting in the loss of cartilage lining the joint surface and  
76 the formation of bony outgrowths known as osteophytes at the joint edges.  
77 Current pharmacological treatments mainly focus on symptom management,  
78 primarily using pain relievers that do not impede disease advancement.  
79 Therefore, there is an imperative need to discover drugs capable of modifying  
80 OA progression.

81 Chondrocyte hypertrophy, a phenotype preceding cartilage calcification and  
82 eventual replacement by bone, plays a crucial role in OA pathogenesis. Under  
83 normal circumstances, articular chondrocytes resist hypertrophic changes.  
84 However, in OA, the onset of hypertrophic differentiation accelerates cartilage  
85 breakdown [3]. Similarly, joint margins are established through the growth of an  
86 initial cartilage template that undergoes endochondral ossification and is replaced  
87 by bone [4]. The inflammatory environment in OA, characterized by synovitis,  
88 involves heightened macrophage activation and the release of inflammatory  
89 cytokines like tumor necrosis factor-alpha (TNF- $\alpha$ ) [5, 6]. Activation of  
90 inflammatory signaling pathways not only leads to cartilage matrix degradation  
91 but also triggers chondrocyte hypertrophy [7, 8].

92

93 Hence, we propose that targeting a key regulator governing both chondrocyte  
94 hypertrophy and inflammation could serve as an effective therapeutic approach  
95 for OA.

96

97 **Methods**

98 ***In silico* simulations**

99 A computational model of the intracellular signaling pathways regulating  
100 articular chondrocyte phenotypes was leveraged and completed with information  
101 about EPHA2 [9, 10]. In silico experiments involved setting targeted variables to  
102 specific values (0 for inhibition, 1 for activation) for 1,000 computing steps,  
103 followed by allowing variables to evolve freely until a new stable state was  
104 reached, simulating a bolus treatment effect. This process was repeated 100 times,  
105 and outcomes were averaged to compute final profiles and standard deviations.  
106 Four perturbations were applied: (1) EPHA2 activation, (2) pro-inflammatory  
107 cytokine activation, (3) combined activation of EPHA2 and pro-inflammatory  
108 cytokines, and (4) EPHA2 blockade with pro-inflammatory cytokine activation.

109 The model and associated code are available via the following GitHub repository:  
110 [https://github.com/Rapha-L/Virtual\\_Chondrocyte\\_for\\_EPHA2\\_study](https://github.com/Rapha-L/Virtual_Chondrocyte_for_EPHA2_study)

111

112 **Evaluation of ALW-II-41-27 in OA chondrocytes**

113 *In vitro* validation was conducted using chondrocytes isolated from human  
114 articular cartilage obtained from OA donors (2 females, 1 male, aged 61, 64, and

115 69 years). Cartilage was obtained with implicit consent as waste material from  
116 patients undergoing total knee replacement surgery, approved by the medical  
117 ethical committee of the Erasmus MC, University Medical Center, Rotterdam  
118 (protocol number MEC-2004-322). Cartilage chips were subjected to protease  
119 and collagenase B digestion to isolate chondrocytes, which were then expanded  
120 in monolayer culture. For redifferentiation, chondrocytes were cultured in a 3D  
121 alginate bead model [11, 12]. After two weeks, cells were treated with 10  $\mu$ M of  
122 ALW-II-41-27, vehicle (DMSO), and/or TNF- $\alpha$  for 24 hours. Medium and  
123 alginate beads were harvested for further analyses.

124

## 125 **Animal model**

126 All animal experimentation procedures were conducted in compliance with the  
127 Animal Ethical Committee of Erasmus University Medical Center (License  
128 number AVD101002015114, protocol number 16-691-06). Twelve-week-old  
129 male C57BL/6 mice (C57BL/6J0laHsd, 27.01 g  $\pm$  2.05 g; Envigo,  
130 Cambridgeshire, UK) were group-housed in individually ventilated cages and  
131 maintained on a 12-hour light/dark cycle with unrestricted access to standard diet  
132 and water at the Experimental Animal Facility of the Erasmus MC. Mice were  
133 randomly assigned to two experimental groups (N=8 per group): Control and  
134 ALW-II-41-27-treated mice.

135 For all procedures, mice were anesthetized using 3% isoflurane/0.8 L O<sub>2</sub>/min  
136 (Pharmachemie BV, Haarlem, the Netherlands). Osteoarthritis (OA) was induced

137 unilaterally by intra-articular injections of 60 µg Monoiodoacetate (MIA)  
138 (Sigma-Aldrich, St. Louis, USA) in 6 µl of saline (0.9% NaCl; Sigma-Aldrich)  
139 on day 0. Injections were administered following a 3-4 mm dermal incision made  
140 to the right knee at the height of the patellar tendon, using a 50 µl syringe  
141 (Hamilton, Bonaduz, Switzerland) and 30G needle (BD Medical, New Jersey,  
142 USA).

143 ALW-II-41-27 was administered via Alzet micro-osmotic pumps (Durect  
144 Corporation, CA, USA) model 1004, with delivery rates of 0.11 µl/hour,  
145 implanted subcutaneously on the back of the mice, slightly posterior to the  
146 scapulae, immediately after the intra-articular injections. Osmotic pumps were  
147 filled with dimethyl sulfoxide: polyethylene glycol alone (55:45 ratio, vehicle-  
148 treated group, N=8 mice) or containing 6 mg of ALW-II-41-27 dissolved in  
149 vehicle (treated group, N=8 mice), resulting in a dose of 6.6 µg/hour. A third  
150 group of N=8 mice received osmotic pumps delivering a dose of 1.7 µg/hour of  
151 ALW-II-41-27. A third group of N=8 mice received the implantation of osmotic  
152 pumps delivering a dose of 1.7 µg/ hour of ALW-II-41-27. In the figures 5 and 6  
153 we report the 6.6 µg/ hour dose of ALW-II-41-27. Synovial thickness, Krenn  
154 score, and osteophyte size for all groups can be found in supplementary figure 6.  
155 Mice were euthanized in accordance with Directive 2010/63/EU by cervical  
156 dislocation under isoflurane anesthesia 14 days following MIA injection. Knee  
157 specimens were fixed in 4% formalin (v/v) for 1 week, decalcified in 10% EDTA

158 for 2 weeks, and embedded in paraffin. Coronal sections of 6  $\mu\text{m}$  were cut for  
159 analysis.

160

161 **Data and statistical analyses**

162 Statistical analysis was conducted using GraphPad Prism 9.0 and IBM SPSS 24  
163 (IBM). Each *in vitro* experiment comprised a minimum of 3 biological replicates  
164 and was replicated using cells obtained from 3 donors. The *in vivo* study was  
165 designed to ensure equal-sized groups, employing randomization and blinded  
166 analyses. The stated group size represents the number of independent values used  
167 for statistical analysis. Sample size for the *in vivo* study was determined based on  
168 the distribution of weight over the hind limbs as read-out parameter. Based on  
169 previous studies, we consider an increase of 13% (standard deviation of 10%) in  
170 weight distribution on the affected limb in time in the therapy groups as relevant  
171 in our study [13]. Sample size calculation was performed with a statistical power  
172 of 80% and a significance level of 0.05, resulting in N=8.

173 For statistical analysis, a linear mixed model with Bonferroni's multiple  
174 comparisons test was utilized.

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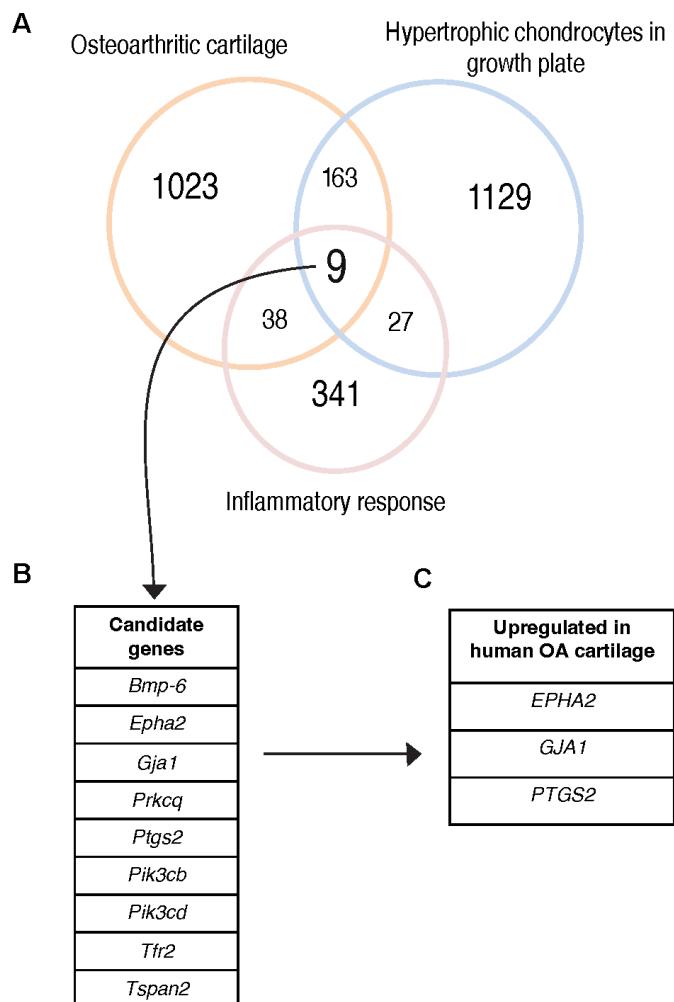
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180 **Results**

181 ***EPHA2* is as an inflammation-related gene upregulated in hypertrophic  
182 chondrocytes and osteoarthritic cartilage**

183 To identify a new therapeutic target for OA linked with chondrocyte hypertrophy  
184 and inflammation, we conducted data analysis on two publicly available murine  
185 microarray datasets [14, 15]. Differentially expressed genes (DEGs) in the  
186 articular cartilage of mice with OA induced by destabilization of the medial  
187 meniscus (DMM) were compared to those from mice undergoing sham surgery,  
188 identifying OA-related genes. By intersecting this set with DEGs in the  
189 hypertrophic zone versus the proliferative zone of the mouse growth plate, we  
190 identified 172 genes differentially expressed in OA associated with chondrocyte  
191 hypertrophy. Among these, nine genes were associated with the gene ontology  
192 inflammatory response [16] (Figure 1 A & B). Subsequently, we examined the  
193 expression of these nine genes in a human microarray dataset obtained from OA  
194 and healthy articular cartilage [17], finding that three genes were also upregulated  
195 in human OA cartilage (Figure 1 C). While *GJA1* and *PTGS2* have been  
196 previously studied in the context of OA [18], *EPHA2*, a tyrosine kinase receptor,  
197 has an undisclosed role in OA. *Epha2* was 30-fold upregulated in OA versus sham  
198 mouse articular cartilage (adjusted p-value= 2E-02) [15], and 3-fold upregulated  
199 in human OA versus healthy cartilage (adjusted p-value= 1E-08) [17].  
200 Additionally, *Epha2* was 19-fold higher in the hypertrophic compared to the  
201 proliferative zone of the murine growth plate (adjusted p-value= 1E-07) [14].

202



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204

205 **Figure 1. Identification of EPHA2 as a novel target for OA associated with**  
206 **inflammation and chondrocyte hypertrophy.** (A) The Venn diagram illustrates the  
207 genes related to the inflammatory response, genes that were differentially regulated  
208 in murine osteoarthritic cartilage (DMM vs sham) and genes differentially expressed  
209 in the murine growth plate (proliferative vs hypertrophic zone). The number of genes  
210 in each dataset is represented, together with those that overlapped. (B) List of 9  
211 targets that overlapped in the three databases. (C) Target genes upregulated in  
212 human OA compared with healthy articular cartilage. DEGs with a fold change greater  
213 than 3, and an adjusted p value lower than of 0.05, were considered for the analysis.

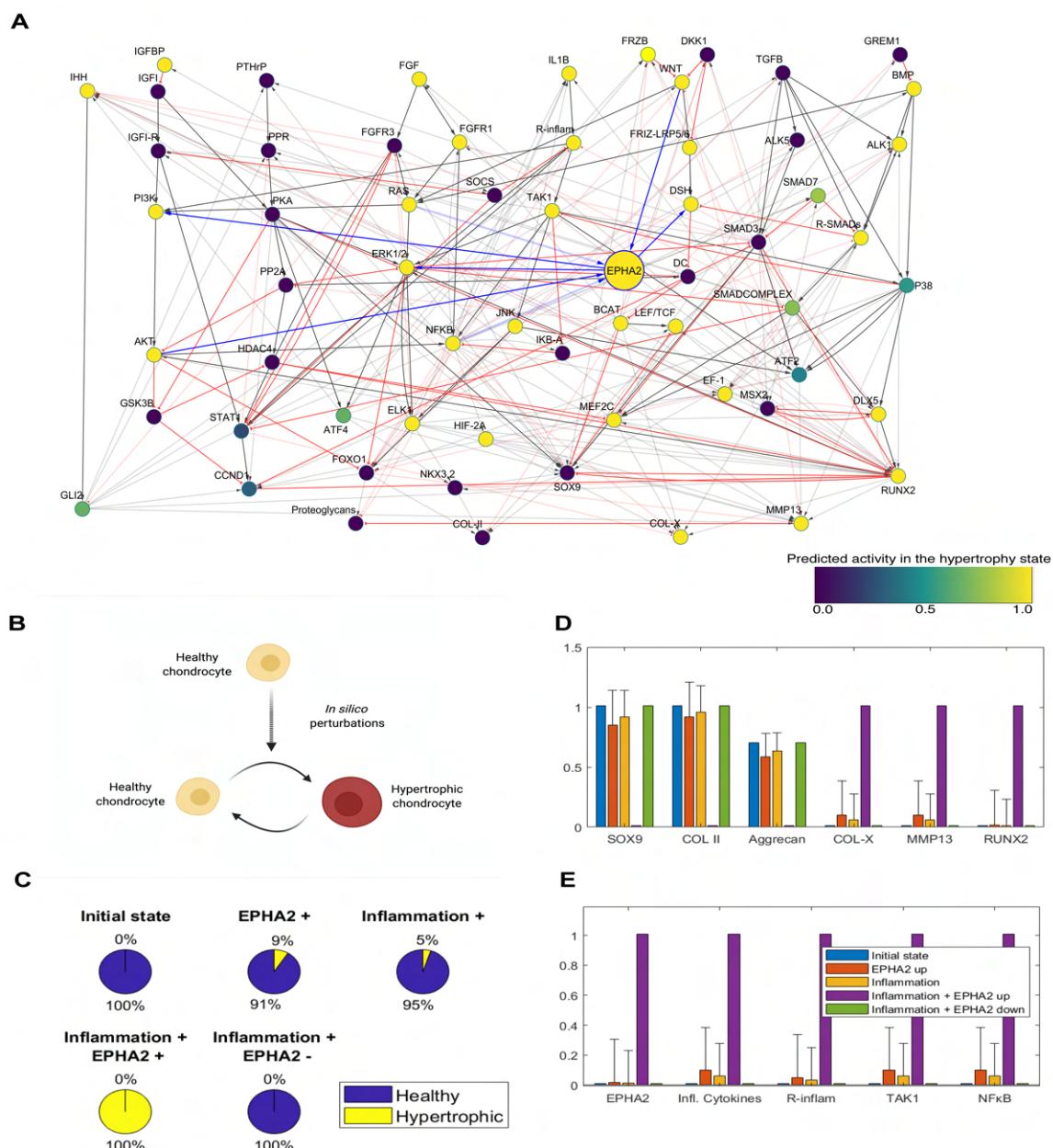
214

215 **EPHA2 triggers inflammatory signaling activation and hypertrophy in a**  
216 **virtual chondrocyte**

217 To assess the role of EPHA2 on chondrocyte phenotype we utilized a  
218 computational model representing two cellular states: healthy and hypertrophic  
219 chondrocytes (Figure 2 A) [9, 19]. EPHA2 exhibited activity in the hypertrophic  
220 state but remained inactive in the healthy state (supplementary figure 1).  
221 Activation of EPHA2 led to a more pronounced transition of healthy  
222 chondrocytes towards the inflammatory hypertrophic state compared to  
223 activation of inflammatory cytokines (9% and 5% for 'EPHA2+' and  
224 'Inflammation+', respectively; Figure 2 B & C). Consequently, both conditions  
225 prompted a decline in anabolic markers (collagen type II, Aggrecan: blue vs  
226 orange and yellow bars; Figure 2 D) and an increase in hypertrophic and  
227 inflammatory markers (Figure 2D & E). Full activation of both EPHA2 and

228 inflammatory cytokines synergistically induced a complete transition of healthy  
229 chondrocytes towards the inflammatory hypertrophic state ('Inflammation+  
230 EPHA2+'; Figure 2 C). Intriguingly, inhibiting EPHA2 (Inflammation+ EPHA2-  
231 ) abolished this cellular state switching, accompanied by a significant reduction  
232 in the activity of hypertrophic and inflammatory entities (green vs violet and  
233 yellow bars; Figure 2 D & E). These computational findings substantiate the  
234 hypertrophic and inflammatory role of EPHA2 in chondrocytes.

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237

238 Figure 2. EPHA2 activation induces chondrocyte hypertrophy and inflammation  
 239 in silico (A) Integration of EPHA2 in the regulatory network of articular  
 240 chondrocytes. Solid lines denote protein interaction while modulation at the gene  
 241 level is in dotted lines. Activating interactions are in grey with delta arrows and  
 242 inhibitory interactions are in red with a half circle. Connections with EPHA2 are

243 denoted in blue. All components of the model are represented with a variable that  
244 is named with capital letters. The variable represents neither the protein activation  
245 level nor the gene expression but a product of both (global activity). The color of  
246 the nodes in the network denotes the global activity of the variables in the  
247 hypertrophic state, with dark blue being 0 and yellow being 1. (B) Effect of  
248 gradually increasing the value of the EPHA2 input (from 0 to 1) in the scenario  
249 with EPHA2 activation and Inflammation, on the chondrocyte state transition.  
250 The initial state being the healthy (SOX9 +). (C) Activation of EPHA2 in the  
251 healthy state promotes the transition to a hypertrophic phenotype. Forced  
252 activation (+) or inhibition (-) of the entities from the initial healthy state is  
253 shown. Inflammation represents the activation of the variables related to  
254 inflammatory cytokines and their receptors, as input in the model. Pie charts  
255 represent the predicted percentage of perturbations leading to a transition to the  
256 hypertrophic phenotype or remaining in the healthy phenotype. (D) Average  
257 activity of the chondrogenic markers, being type II collagen, Aggrecan and SRY-  
258 box transcription factor (SOX9), and of hypertrophic markers, being Runt-related  
259 transcription factor 2 (RUNX2), Matrix Metallopeptidase (MMP13) and type 10  
260 Collagen. (E) Average activity of EPHA2 and variables associated with  
261 inflammation. The bars represent the mean of the results of a hundred in silico  
262 experiments (+ standard deviations, +SD). There is no SD for the initial state  
263 (blue condition) as it denotes the starting point before the in silico perturbation is  
264 applied. Figure created with Cytoscape, MATLAB and Biorender.com

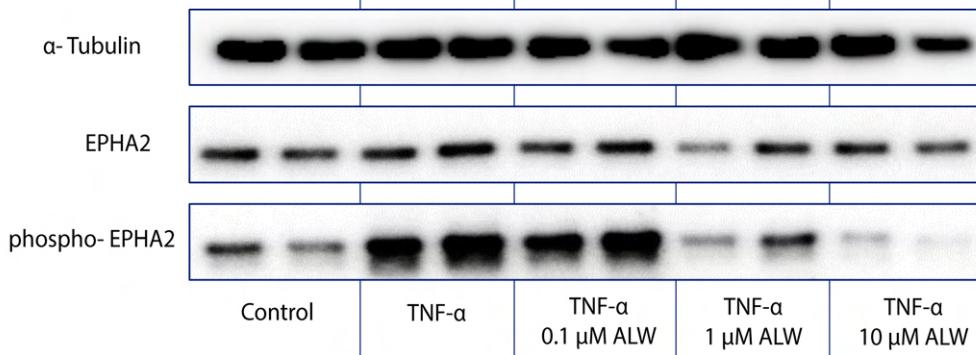
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266 **ALW-II-41-27 reduces human OA-derived chondrocyte inflammation**

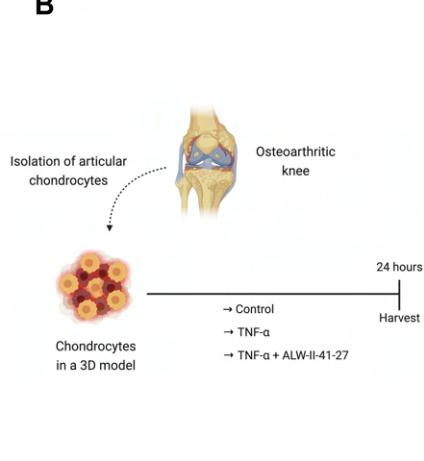
267 To investigate the potential of pharmacologically inhibiting EPHA2 to alleviate  
268 inflammation in OA, we utilized the tyrosine kinase inhibitor ALW-II-41-27, a  
269 type II kinase inhibitor known for its high selectivity for EPHA2 [20, 21].  
270 Initially, we validated the ability of ALW-II-41-27 to reduce TNF- $\alpha$ -induced  
271 phosphorylation of EPHA2 in a dose-dependent manner (Figure 3 A). Notably,  
272 the concentration of 10 micromolars exhibited the highest efficacy in decreasing  
273 both EPHA2 phosphorylation and the expression of catabolic enzymes MMP1  
274 and MMP13 in OA cartilage explants (supplementary figure 3). Subsequently,  
275 we examined the effects of ALW-II-41-27 at a concentration of 10 micromolars  
276 on TNF- $\alpha$ -treated human chondrocytes from OA donors (Figure 3 B). Treatment  
277 with ALW-II-41-27 significantly inhibited TNF- $\alpha$ -induced inflammatory  
278 responses, as evidenced by a reduction in the secretion of nitric oxide metabolites  
279 (Figure 3 C). Furthermore, ALW-II-41-27 administration mitigated the TNF- $\alpha$ -  
280 induced expression of the inflammatory cytokine interleukin (IL)-6, known to be  
281 a downstream target of TNF- $\alpha$  (Figure 2 D & E) [22]. Moreover, ALW-II-41-27  
282 treatment countered the TNF- $\alpha$ -induced upregulation of the cartilage matrix-  
283 degrading enzymes MMP1 and MMP13 (Figure 3 F & G). These findings  
284 demonstrate the anti-inflammatory potential of ALW-II-41-27 in TNF- $\alpha$ -  
285 stimulated OA chondrocytes.

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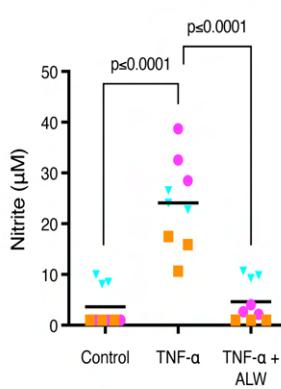
**A**



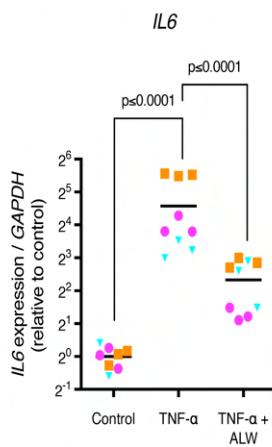
**B**



**C** Nitrite in the medium

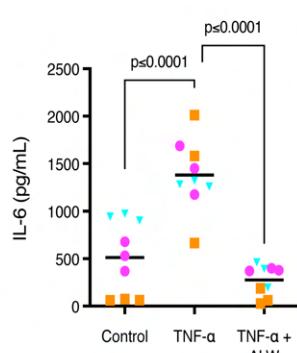


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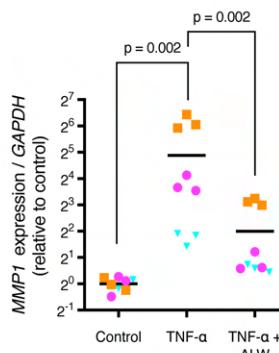
**E**

IL-6 in the medium



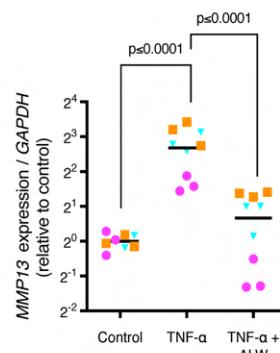
**F**

*MMP1*



**G**

*MMP13*



289 Figure 3. ALW-II-41-27 decreases TNF- $\alpha$  induced catabolism and inflammation  
290 in human chondrocytes. (A) ALW-II-41-27 decreases TNF $\alpha$ -induced  
291 phosphorylation of EPHA2 in a dose-dependent manner. (B) Experimental set-  
292 up to evaluate the anti-inflammatory capacity of ALW-II-41-27 (10  $\mu$ M) in  
293 human OA chondrocytes cultured with 10 ng/mL TNF- $\alpha$ . (C) Evaluation of  
294 nitrite in the medium as a marker for inflammatory activity as determined by  
295 Griess reagent. (D, F-G) Gene expression of IL6, MMP1 and MMP13 determined  
296 by qPCR. The average of control, per donor, is set to 1. (E) IL-6 in the medium  
297 determined by ELISA. Experiments were performed in triplicate, with cells from  
298 three donors. Donors are represented with different colors and symbols: violet  
299 circles (donor 1), blue triangles (donor 2) and orange squares (donor 3). The  
300 horizontal line in the graphs represents the mean. Data were analyzed with the  
301 linear mixed model with Bonferroni's multiple comparisons test. Figure created  
302 with Biorender.com

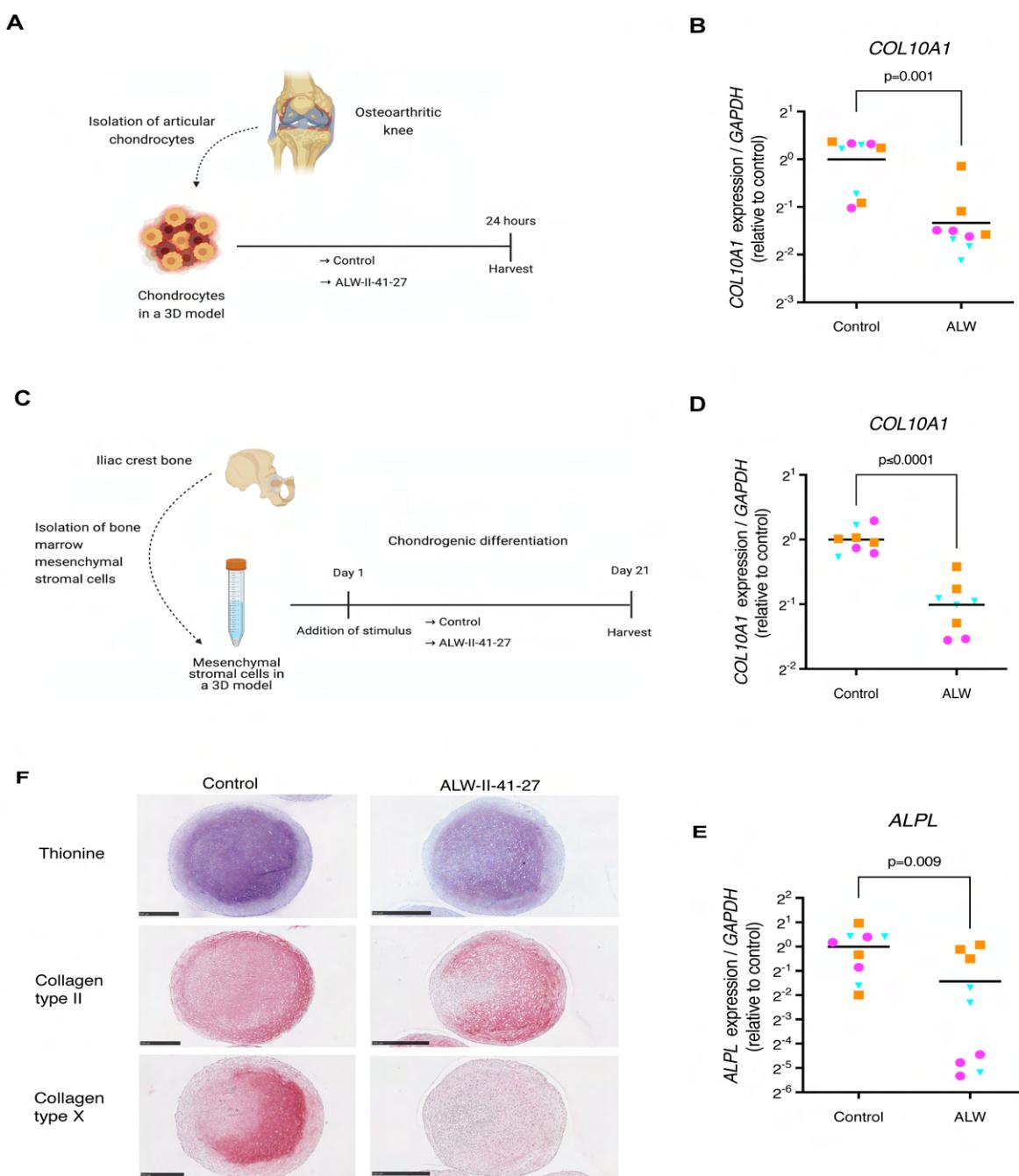
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#### 304 **ALW-II-41-27 decreases chondrocyte hypertrophy**

305 We then proceeded to evaluate whether pharmacological inhibition of EPHA2  
306 with ALW-II-41-27 could impede hypertrophy in human OA-derived articular  
307 chondrocytes (Fig. 4 A). ALW-II-41-27 decreased *COL10A1* expression (Figure  
308 4 B), suggesting a reduction in hypertrophy. Given that chondrocyte hypertrophy  
309 poses a significant obstacle to stable hyaline cartilage tissue engineering [23], we  
310 further investigated the effect of ALW-II-41-27 on cartilage tissue engineered

311 constructs from mesenchymal stromal cells (MSCs), well-known to become  
312 hypertrophic and prone to ossify when implanted *in vivo* [24-27] (Figure 4 C).  
313 Gene expression analysis indicated that the addition of ALW-II-41-27 mitigated  
314 the hypertrophic markers *COL10A1* and *ALPL* (Figure 4 D & E). Likewise, when  
315 treated with ALW-II-41-27, the cells exhibited reduced deposition of type X  
316 Collagen (Figure 4 F and supplementary figure 4). Intriguingly, neither  
317 glycosaminoglycan nor type II Collagen deposition showed reduction, suggesting  
318 that ALW-II-41-27 primarily targeted hypertrophy.

319



320

321

322 Figure 4. ALW-II-41-27 decreases chondrocyte hypertrophy. (A) Experimental  
323 set-up to evaluate the capacity of ALW-II-41-27 (10  $\mu$ M) to modulate  
324 hypertrophy in human OA chondrocytes. (B) Gene expression of the hypertrophic  
325 marker *COL10A1* in cultured OA chondrocytes determined by qPCR. (C)

326 Experimental set-up to evaluate how ALW-II-41-27 (100 nM) affects  
327 hypertrophy in tissue engineered cartilage from MSCs. (D, E) Gene expression  
328 of COL10A1 and ALPL in MSC-generated cartilage determined by qPCR. (F)  
329 Histological analysis of tissue engineered cartilage derived from MSCs; Thionine  
330 staining showing glycosaminoglycans (violet), and immunohistochemistry of  
331 type II and type X collagen (red/pink). Experiments were performed with 3  
332 replicates, for each of the three donors. Donors are represented with different  
333 colors and symbols: violet circles (donor 1), blue triangles (donor 2) and orange  
334 squares (donor 3). For gene expression analysis, the average of control replicates  
335 is set to 1 per donor. The horizontal line in the graphs represents the mean. For  
336 statistical analysis, the linear mixed model with Bonferroni's multiple  
337 comparisons test was performed. Figure created with Biorender.com

338

339 **ALW-II-41-27 treatment reduces joint inflammation and pathological  
340 endochondral ossification *in vivo***

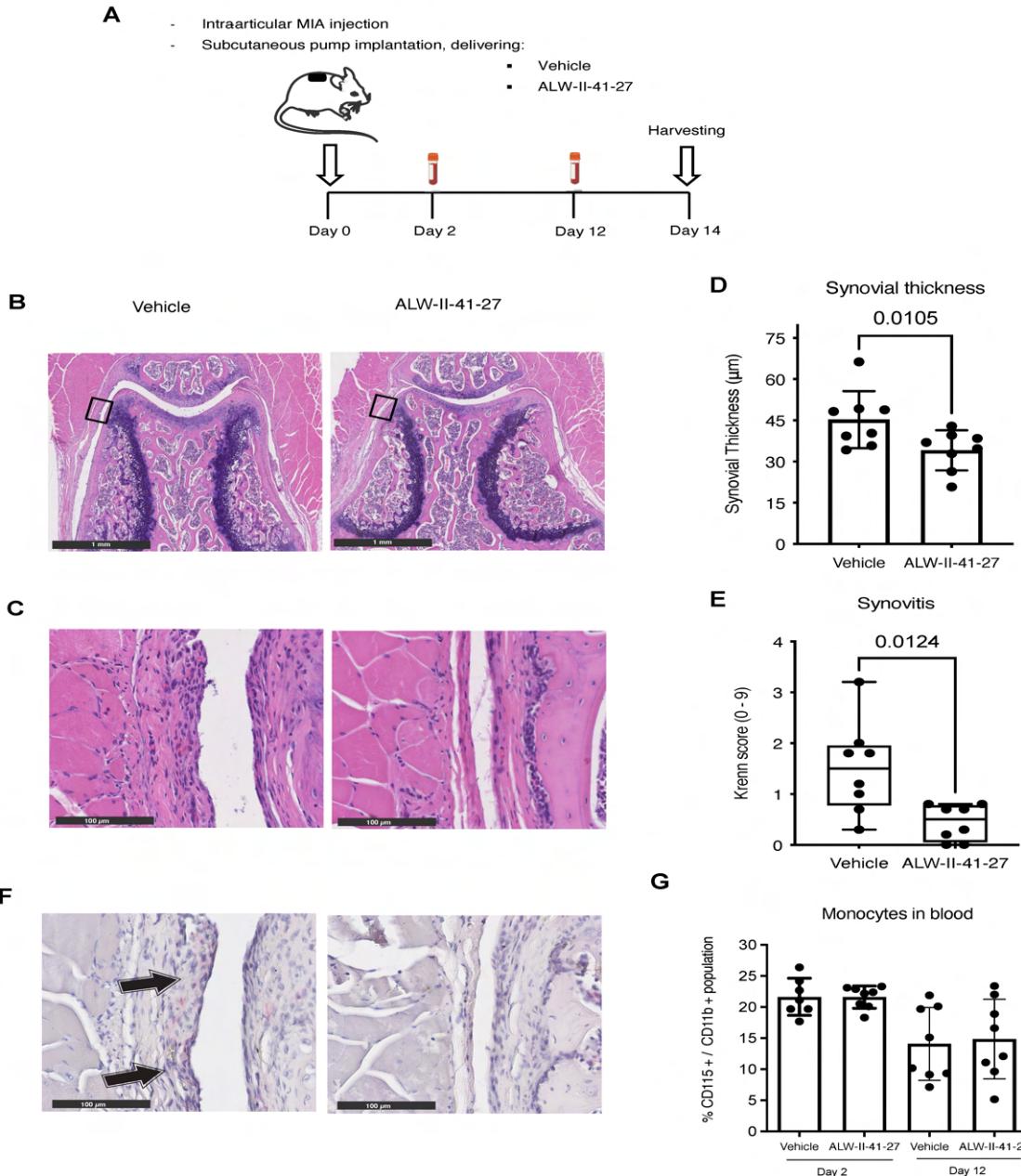
341 EPHA2 inhibition reduced hypertrophy and inflammation *in silico* and its  
342 pharmacological inhibition with ALW-II-41-27 in cell culture *in vitro* confirmed  
343 these results. Subsequently, we sought to assess the potential therapeutic efficacy  
344 of ALW-II-41-27 *in vivo* using a mouse model of joint pain and degeneration  
345 induced by intra-articular injection of monoiodoacetate (MIA) [28]. ALW-II-41-  
346 27 was administered subcutaneously with a controlled delivery system (Figure 5  
347 A). Notably, the total body weight remained unaffected by ALW-II-41-27

348 administration throughout the 14-day study period, indicating no significant  
349 adverse effects on the general health of the animals (supplementary figure 5).

350 Treatment with ALW-II-41-27 led to a reduction in synovial membrane thickness  
351 and synovitis compared to vehicle-treated mice (Figure 5 B, C, D & E). Synovitis  
352 is governed by macrophages, the crucial regulators of OA progression and  
353 primary mediators of the inflammatory response [29, 30]. Macrophages were  
354 solely observed in the synovial lining of vehicle-treated mice, suggesting  
355 effective attenuation of joint inflammation (Figure 5 F). Moreover, there were no  
356 discernible alterations in peripheral blood monocyte levels at day 2 and 12  
357 (Figure 5 G), indicating selective reduction of local joint inflammation without  
358 affecting systemic immune cells.

359 Despite joint pain being a significant OA symptom often associated with  
360 inflammation [31], no significant difference in weight distribution between limbs  
361 was detected between ALW-II-41-27-treated and vehicle-treated mice,  
362 suggesting no apparent effect on pain (supplementary Figure 6).

363



364

365

366 Figure 5. ALW-II-41-27 treatment attenuates joint inflammation. (A)  
367 Experimental set-up of the in vivo experiment. Intra-articular injection of  
368 monoiodoacetate (MIA, 60  $\mu$ g in 6  $\mu$ l of saline) was applied to the right knee of  
369 mice for each experimental group (N=8) to induce OA. An osmotic pump was  
370 implanted on the back of the mice, slightly posterior to the scapulae, which

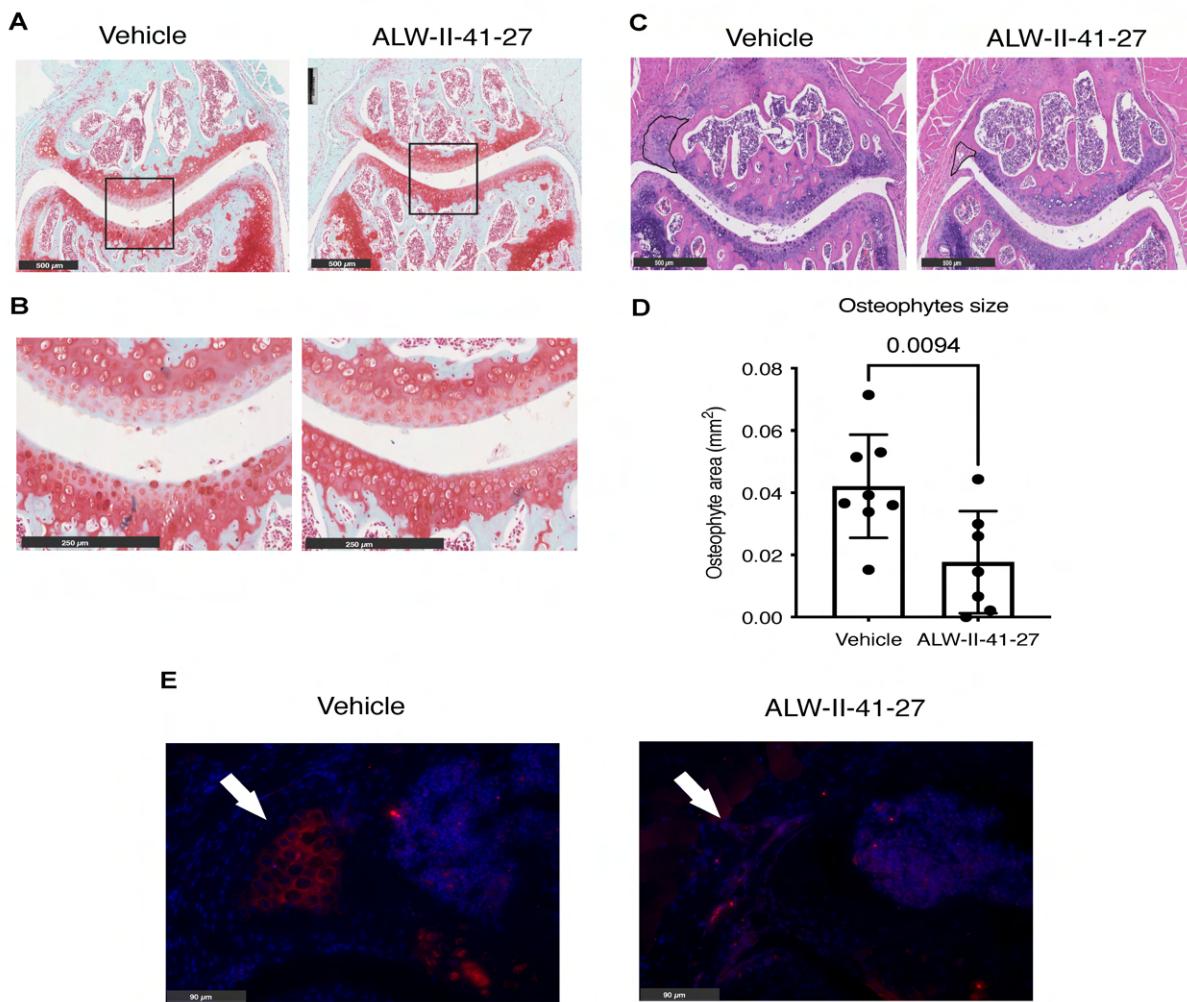
371 continuously delivered vehicle or ALW-II-41-27 in a dose of 6.6  $\mu$ g/ hour. At  
372 day 2 and 12 peripheral blood was harvested. At day 14 mice were euthanized to  
373 assess the effects of ALW-II-41-27 on the degenerated joint. (B) Hematoxylin  
374 and Eosin staining of knees (patellofemoral region) from vehicle-treated and  
375 ALW-II-41-27-treated mice. Black square indicates the region of magnification  
376 for the image below (C) showing the synovial lining where synovial thickness  
377 and Krenn score were determined. (D) Synovial thickness is represented by the  
378 mean  $\pm$  SD. (E) Krenn score illustrated with box-and-whiskers plots, with line  
379 indicating the median and error bars spanning maximum to minimum values. (F)  
380 Immunohistochemistry of F4/80 (pink) showing macrophages in the synovial  
381 lining. Arrows indicate positive staining. (G) Percentage of monocytes present in  
382 the peripheral blood of mice at day 2 and 12, respect to the myeloid cell  
383 population. In all graphs, each dot represents data of an individual mouse (N=8).  
384 For statistical analysis, the linear mixed model with Bonferroni's multiple  
385 comparisons test was performed. Figure created with Biorender.com

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387 Cartilage degeneration, evidenced by proteoglycan loss, was observed in all mice  
388 irrespective of ALW-II-41-27 treatment (Figure 6 A & B), suggesting that ALW-  
389 II-41-27 was not able to rescue matrix degeneration *in vivo*. However,  
390 histological assessment revealed significantly smaller osteophytes, particularly at  
391 the lateral side of the patella, in ALW-II-41-27-treated mice (Figure 6 C & D).  
392 Additionally, type X collagen deposition was reduced in the knees of ALW-II-

393 41-27-treated mice (Figure 6 E), suggesting effective targeting of endochondral  
394 ossification in the injured joint.

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398 Figure 6. ALW-II-41-27 treatment attenuates pathological endochondral  
399 ossification (A) Safranin-O / Fast Green staining of vehicle-treated and ALW-II-  
400 41-27-treated mice knees with magnification of the patellofemoral region. Black  
401 square indicates the region of magnification for the image below (B) showing the  
402 central part of the patellofemoral articular cartilage. (C) Hematoxylin and Eosin  
403 stain of vehicle-treated and ALW-II-41-27-treated mice knees with magnification  
404 of the patellofemoral region. Osteophyte's diameter is indicated with a black line  
405 in the lateral side of the patella. (D) Osteophyte area adjacent to the lateral side  
406 of the patella is represented by the mean  $\pm$  SD. Each dot represents data of an  
407 individual mouse (n=8). For statistical analysis, the linear mixed model with  
408 Bonferroni's multiple comparisons test was performed (E) Immunofluorescence  
409 of type X collagen (red) and DAPI (blue) in the lateral side of the patellofemoral  
410 region. Arrow indicates an osteophyte in the lateral side of the patella.

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420 **Discussion**

421 There is an urgent unmet need for effective therapies for OA patients. Here, we  
422 show that EPHA2 is a promising drug target for OA and we report the small  
423 molecule ALW-II-41-27 as a disease-modifying OA drug (DMOAD),  
424 specifically targeting inflammation and pathological endochondral ossification  
425 (Fig. 7A). To find targets associated with inflammation and chondrocyte  
426 hypertrophy we have used a particular sequence of studies that involved *in silico*,  
427 *in vitro* and *in vivo* analyses (Figure 7 B). For the *in silico* analyses we leveraged  
428 previously published large gene expression dataset depositories and narrowed  
429 them down to one target of interest. The role of the identified target on  
430 inflammation and chondrocyte hypertrophy was further investigated through *in*  
431 *silico* experiments using a computational model of cellular signaling networks  
432 controlling chondrocyte phenotypes. These *in silico* studies served as the starting  
433 point for a series of *in vitro* experiments using different cell models, which were  
434 followed by an *in vivo* study. Our study illustrates the efficacy of this  
435 experimental approach in uncovering a novel target for specific biological  
436 processes in osteoarthritis

437 While previous research has implicated EPHA2 as a key player in diseases like  
438 cancer [32-35] and irritable bowel disease [36], our study marks the first to  
439 underscore its significance in OA. Other tyrosine kinases, including fibroblast  
440 growth factor receptor (FGFR) 1, Fyn and vascular endothelial growth factor  
441 receptor (VEGFR), have been implicated to promote chondrocyte hypertrophy

442 [37-40]. Our findings reveal that the tyrosine kinase EPHA2 not only contributes  
443 to hypertrophy but also to inflammation, making it a compelling target for  
444 mitigating pathological mechanisms in OA.

445 We have identified a promising compound, ALW-II-41-27, demonstrating  
446 potential as a disease-modifying osteoarthritic drug (DMOAD). To date, a wide  
447 variety of DMOADs targeting chondrocyte hypertrophy have been tested in  
448 experimental pre-clinical studies [41-43]. It is unclear, however, whether those  
449 agents have been subjected to further research. Regulatory agencies, such as the  
450 Food and Drug Administration (FDA) or the European Medicines Agency  
451 (EMA), have not yet approved any existing disease-modifying pharmacological  
452 intervention for OA [44]. Considering the pre-clinical data of ALW-II-41-27 to  
453 modulate OA pathogenesis and its extensive pharmacological analysis in other  
454 conditions [32-36], it is expected that clinical trials involving this compound may  
455 pose lower risks with a higher likelihood of success.

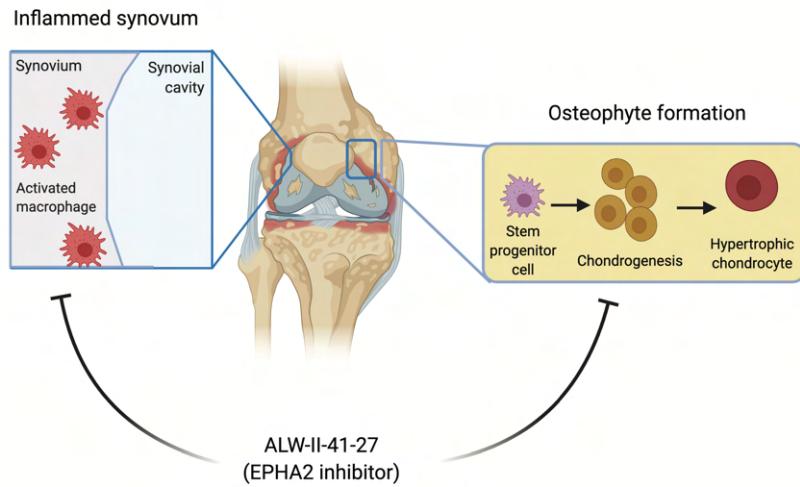
456 In addition to cartilage hypertrophy, inflammation of the synovium plays a crucial  
457 role in OA pathology [45]. Our results indicate that ALW-II-41-27 exhibits anti-  
458 inflammatory properties, aligning with its previously observed effects in a model  
459 of irritable bowel syndrome [36]. EPHA2 is not limited to chondrocytes but is  
460 also expressed in various other cell types found in the synovium, including  
461 fibroblasts, monocytes, and macrophages [46, 47]. Thus, the observed anti-  
462 inflammatory mechanism in our *in vivo* setting may also be linked to the action  
463 of ALW-II-41-27 on these cell types.

464 Our study has certain limitations. Treatment initiation coincided with OA  
465 induction in our research. This decision was influenced by the progressive nature  
466 of the disease, posing a challenge for a DMOAD to reverse extensive joint  
467 structural changes in end-stage OA. Hence, maximizing the pharmacological  
468 benefits of ALW-II-41-27 to reduce inflammation and osteophytosis might be  
469 more effective if administered during the earlier stages of the disease. Our study  
470 did not demonstrate that *in vivo* administration of ALW-II-41-27 prevented  
471 cartilage loss or pain in the MIA mouse model. Further investigation using  
472 alternative experimental animal models is warranted to determine whether ALW-  
473 II-41-27 specifically targets hypertrophy and inflammation, or if its effects extend  
474 to preventing cartilage degeneration and alleviating pain. Additionally, exploring  
475 the potential effects of ALW-II-41-27 on post-traumatic OA or other non-  
476 chemically induced forms of OA is essential.

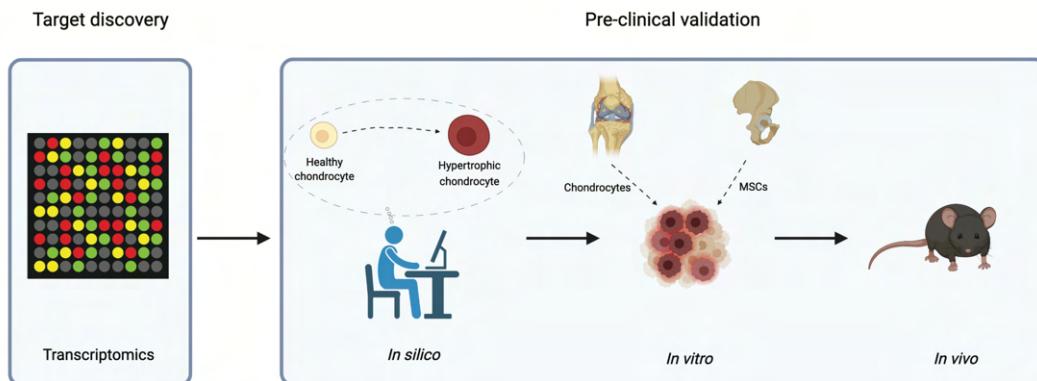
477 In conclusion, our findings suggest that EPHA2 contributes to the pathogenesis  
478 of OA. The use of ALW-II-41-27 to inhibit EPHA2 showed promising results in  
479 mitigating inflammation and pathological endochondral ossification across  
480 various models, including *in silico* simulations, *in vitro* experiments with patient-  
481 derived cells, and a mouse model. These results underscore the potential of ALW-  
482 II-41-27 as a candidate drug for modifying the course of OA, warranting further  
483 investigation.

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A



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487 Figure 7. Graphical representation of main findings and experimental approach.

488 (A) ALW-II-41-27 attenuates synovitis and osteophyte formation. (B) Drug

489 discovery pipeline combining transcriptomic datasets, in silico, in vitro and in

490 vivo models. Figure created with Biorender.com

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492 All authors approved the final version of the manuscript

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494 **Ethical statement**

495 Animal experiments were approved by the medical ethical committee of the  
496 Erasmus MC, protocol EMC 16-691-06. Human articular cartilage was obtained  
497 with the approval of Erasmus MC, protocol MEC-2004-322 and mesenchymal  
498 stromal cells MEC-2014-16.

499

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505

506 **Supplementary methods**

507 Description of further procedures can be found in supplementary methods.

508

509 **Competing interest statement**

510 The author(s) declared the following potential conflicts of interest with respect to  
511 the research, authorship, and/or publication of this article: M.G. Chambers is an  
512 employee of Eli Lilly but they did not fund this research.

513

514 **References**

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