

# Hydroquinone, a cigarette smoke compound, affects cartilage homeostasis through activation of the aryl hydrocarbon receptor pathway

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15 **stress<sup>5</sup>, IL-1 $\beta$ <sup>6</sup>.**

16

17 **Abstract**

18 Exposure to cigarette smoke has a proven detrimental impact on different aspects of  
19 human health. Increasing evidences link smoking to degeneration of joint tissues. However,  
20 the toxic mechanisms elicited by the different components of cigarette smoke have not been  
21 fully elucidated yet. We have previously shown that exposure to hydroquinone (HQ), a pro-  
22 oxidant chemical present in cigarette smoke, can promote joint tissue degradation in murine  
23 models of rheumatoid arthritis through the activation of the aryl hydrocarbon receptor (AhR)  
24 pathway. Osteoarthritis (OA) is a chronic debilitating articular disease characterized by  
25 progressive degradation of the articular cartilage, whose onset and progression have also  
26 been associated with smoking. In this work we aimed to investigate the effect of HQ  
27 exposure on articular chondrocytes and how it affects cartilage homeostasis. Cell viability,  
28 gene expression, oxidative stress and inflammatory parameters were quantified in  
29 primary articular chondrocytes exposed to HQ in presence or absence of IL-1 $\beta$  pre-  
30 stimulation. HQ stimulation downregulated phenotypic markers genes such as SOX-9 and  
31 Col2a1, whereas upregulated the expression of the catabolic enzymes MMP-3 and  
32 ADAMTS5. HQ also promoted oxidative stress and reduced proteoglycan content. HQ  
33 exacerbated the pro-inflammatory effects mediated by the IL-1 $\beta$  co-stimulation. Finally, we  
34 showed that HQ-degenerative effects were mediated by the activation of AhR. Together, our  
35 findings address the harmful effects of HQ in the articular cartilage health, providing novel

36 evidence surrounding the toxic mechanisms of environmental pollutants underlying the onset  
37 of articular diseases.

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39

## 40 **Introduction**

41 The exposure to environmental pollutants can severely affect and compromise human  
42 health. Cigarette smoke is considered the major environmental risk factor for the  
43 developmental and progression of chronic inflammatory diseases such as chronic obstructive  
44 pulmonary disease (COPD), cardiovascular diseases and psoriasis (Stampfli and Anderson,  
45 2009).

46 Hydroquinone (HQ) is a chemical compound representing 3 % of the particle matter  
47 phase of cigarette smoke (Bodnar et al., 2012). This xenobiotic is a benzene metabolite which  
48 can mediate immune- and mielotoxicity (Mcgregor, 2007; Li et al., 2018). Exposure to HQ  
49 has been also associated with increased apoptosis and development of oxidative stress in  
50 immune cells (Cho et al., 2008; Lee et al., 2012). In vitro studies showed that HQ alone is  
51 responsible for 10 % of oxidative stress-mediated cellular toxicity when cells are exposed  
52 to cigarette smoke (Stabbert et al., 2017).

53 We have recently shown that exposure to HQ aggravated joint damage in experimental  
54 animal models of rheumatoid arthritis (RA) through the activation of the aryl hydrocarbon  
55 receptor (AhR) pathway by promoting inflammation, an increased influx of immune cells,  
56 synovial proliferation and oxidative stress (Heluany et al., 2018a,b; Heluany et al., 2021).

57 Exposure to cigarette smoke has also been associated with increased pain and cartilage  
58 loss in both osteoarthritis (OA) and in degenerative disc diseases (Goldeberg, Scott and  
59 Mayo, 2000; Fogelholm and Alho, 2001; Amin et al., 2007). So far, the main research focus  
60 has been the investigation of the role of nicotine in the disease progression, often with  
61 inconclusive or contrasting results (Felson and Zhang, 2015).

62 OA is the most widespread and disabling type of chronic and degenerative articular  
63 disease, characterized by low-grade inflammation, breakdown of the articular cartilage,  
64 abnormal remodeling of the subchondral bone and growth of bone spurs, also known as  
65 osteophytes (Berenbaum, 2013; Mobasheri et al., 2017). The chondrocyte is the only cell  
66 type populating the articular cartilage and plays an essential role in maintaining cartilage  
67 homeostasis. OA progression is tightly linked to the loss of chondrocyte function, due to the  
68 deregulation of several signaling cascades and the acquisition of a hypertrophic  
69 phenotype (Akkiraju and Nohe, 2015; Thomas et al., 2007). In addition, the upregulation of  
70 pro-inflammatory cytokines such as interleukin-1 $\beta$  (IL-1 $\beta$ ) has been strongly associated with  
71 loss of chondrocyte homeostasis and exacerbation of articular cartilage damage in  
72 OA (Sellan and Berenbaum, 2010).

73 In this work we investigated the effect of HQ on articular chondrocytes to elucidate how it  
74 can affect the articular cartilage health. We stimulated bovine articular chondrocytes and  
75 cartilage explants with HQ to investigate its effects on cell viability, modulation of phenotypic  
76 genes and oxidative stress. Our results showed that HQ exposure altered cell proliferation,  
77 down-regulated the expression of the phenotypic markers SRY-Box transcription factor 9

78 (SOX-9), type II collagen (Col2a1) and type X collagen (Col-X), while up-regulating the  
79 expression of catabolic markers matrix metalloproteinase 3 (MMP-3) and A Disintegrin and  
80 Metalloproteinase with Trombospondin motifs 5 (ADAMTS5). HQ exposure also promoted  
81 oxidative stress and potentiated the pro-catabolic effects of IL-1 $\beta$ . We also showed that HQ  
82 activity is mediated through the activation of the AhR signaling pathway. Together, our  
83 findings suggest that the AhR activation in articular chondrocytes could be an important  
84 mechanism through which environmental pollutants compromise cartilage homeostasis.

85

## 86 **Material and Methods**

### 87 *Isolation and culture of bovine articular chondrocytes and cartilage explants*

88 Bovine articular cartilage explants were isolated from the metatarsal and metacarpal  
89 joints of adult cows purchased within less than 24 hours of death from a local abattoir. The  
90 articular cartilage was dissected in sterile conditions and processed for chondrocyte  
91 isolation or explant cultures as previously described (Nalesso et al., 2011). Bovine articular  
92 chondrocytes (BACs) were cultured with complete medium (CM; DMEM F12 supplemented  
93 with 10 % fetal bovine serum (FBS) and 1 % of antibiotics penicillin and streptomycin) at  
94 37 °C in a 5 % CO<sub>2</sub> incubator. Cells at passage 1 to 3 were used for the experiments  
95 described in this manuscript.

96

### 97 *MTT assay*

98 Cell integrity and viability were measured through the 3-(4,5-dimethylthylthiazol-2-yl)-  
99 2,5 diphenyltetrazolium bromide (MTT) method. Briefly,  $1 \times 10^4$ /well BACs were seeded in 96-  
100 well plates and cultured in CM for 24 hours. Thereafter, cells were washed with PBS and  
101 cultured with DMEM F12 supplemented with 0.1 % of BSA for 24 hours. BACs were  
102 treated as described in the individual experiments. Then, 0.5 mg/mL of MTT solution was  
103 added to each well and incubated for 3 hours in the dark at 37 °C. Thus, the medium  
104 was removed, and the blue formazan crystals were dissolved in 200  $\mu$ L of DMSO. The  
105 optical density reading was recorded at 570 nm in a plate reader (Clario Star, BMG  
106 LABTECH).

107

### 108 *Gene expression*

109 For gene expression analyses,  $7.5 \times 10^4$  BACs were plated per well in 24-well plates  
110 and cultured with CM for 24 hours. Then the cells were washed with PBS and treated as  
111 described in the Results section. Total RNA was extracted from BACs using TRIzol reagent  
112 (Invitrogen), according to the manufacturer's instructions. Three hundred and fifty nanograms  
113 of total RNA from each sample were reverse transcribed to cDNA using a High Capacity  
114 cDNA Reverse Transcription Kit (Applied Biosystems). Quantitative PCR was performed with  
115 hot-start DNA polymerase (Qiagen), as previously described (Nalesso et al.,  
116 2011). Primer sequences are listed in Table 1. All data were normalized to internal control of  
117  $\beta$ -actin values. All experiments were performed in a BioRad PCR system (CFX96TM Optics  
118 Module).

119

120

121 *Micromass cultures*

122 BACs were plated at a density of  $25 \times 10^4$  cells per well in complete medium in 24-well  
123 plates. Micromass cultures were prepared as described before (De Bari et al., 2001).  
124 The micromasses were stimulated as described in the individual experiments with HQ  $\pm$  IL-  
125  $1\beta$  (20 ng/mL; RP0106B-025 Kingfisher Biotech). After 48 hours, the culture media were  
126 collected and the micromasses were washed with PBS and fixed with cold methanol.

127

128 *Alcian blue staining*

129 The micromasses were stained overnight with Alcian blue, as previously described  
130 (De Bari et al., 2001). After proteoglycan extraction with 6 M guanidine hydrochloride (Sigma-  
131 Aldrich), the absorbance was read at 630 nm with a spectrophotometer (Clario Star, BMG  
132 LABTECH). The absorbance values were normalized to protein content. Images of the  
133 micromasses were acquired at room temperature with a stereomicroscope (SZTL 350  
134 Stereo Binocular Microscope, VWR®).

135

136 *Safranin-O staining*

137 Bovine articular cartilage explants were stimulated ex vivo with HQ as described in the  
138 Results section. At the end of the exposure period after 72 hours, the culture media was  
139 collected while the cartilage specimens were fixed in 4 % paraformaldehyde (PFA) and  
140 paraffin embedded. Three-micrometre thick sagittal sections were stained with safranin-O as  
141 described before (Nalesso et al., 2011), and images were taken using the same settings on  
142 an optical microscope (Nikon). Images were acquired through 10x magnification objective  
143 lenses and the SO-staining was quantified by using the ImageJ software (NIH). For such, 4  
144 fields per slide (totalling 16 images per experimental group) were considered for the analysis  
145 and the mean of intensity from all groups was generated and used for the statistical analysis.

146

147 *DMMB assay*

148 Release of glycosaminoglycans (GAGs) by micromasses and cartilage explants upon  
149 exposure to HQ was measured in the culture media by using the Dimethyl methylene Blue  
150 (DMMB) assay (adapted from Farndale et al., 1982). Briefly, 200  $\mu$ L of the DMMB solution  
151 was added to 20  $\mu$ L of the culture media in a 96-well plate. Subsequently, the absorbance  
152 was measured at 525 nm with a spectrophotometer (Clario Star, BMG LABTECH).

153

154 *Quantification of nitric oxide by Griess assay*

155 Fifty microliters of Sulphanilamide Solution (1 % in 5 % phosphoric acid; Promega)  
156 was added to 50  $\mu$ L of the culture media from micromass cultures of BACs or bovine  
157 cartilage explants treated with HQ  $\pm$  IL- $1\beta$  (20 ng/mL). After a 10 minutes incubation at RT,  
158 50  $\mu$ L of N-1-naphthylenediamine dihydrochloride solution (NED, 0.1 % in water; Promega)  
159 was added to the mix. After a further 10 minutes incubation, the absorbance was measured  
160 at 550 nm with a spectrophotometer (Clario Star, BMG LABTECH).

161

162 *Quantification of Reactive oxygen species by DCFH*

163 The intracellular accumulation of reactive oxygen species (ROS) was quantified in  
164 BACs cultures using the fluorescent probe CM-H<sub>2</sub>DCFDA. Ten thousand chondrocytes/well  
165 were seeded in 24-well plates and stimulated with HQ for 24 hours. The cells were then

166 incubated with 10  $\mu$ M CM-H2DCFDA for 30 minutes at 37 °C in the dark. The cells were  
167 resuspended in PBS and 10,000 events were acquired in a MACSQuant flow cytometer.  
168 Results are presented as arbitrary units of fluorescence.

169

#### 170 *Reporter assay*

171 BACs were seeded at a density of 7 x 10<sup>4</sup> cells/well in CM on a 24-well plate. The  
172 cells were co-transfected with 450 ng/well of pGL4.43 [luc2P/XRE/Hygro] luciferase reporter  
173 vector (Promega) and with 50 ng/well of the control vector expressing *Renilla reniformis*  
174 luciferase by using Lipofectamine (Thermo Fisher) following the manufacturer instructions.  
175 After the transfection, the cells were stimulated with CM (vehicle), or with the AhR ligand 6-  
176 formylindolo(3,2b)carbazole (FICZ, 10  $\mu$ M) or with HQ (10 or 25  $\mu$ M) for 24 hours. Luciferase  
177 activity was determined using the Dual luciferase reporter assay system (Promega). Firefly  
178 luciferase activity was normalized by the Renilla luciferase activity. Data are expressed as  
179 fold increase of relative luminescence units in comparison to vehicle.

180

#### 181 *Statistical analysis*

182 One-way ANOVA with Tukey-Post test was used to compare the statistical differences  
183 between multiple groups, and a two-tailed *t*-test was used for comparisons between two  
184 groups. Values of *p* < 0.05 were considered statistically significant. Data are expressed as  
185 the mean  $\pm$  standard error of the mean (SEM). Statistical analyses were performed using  
186 GraphPad Prism version 8.0 (GraphPad Software, CA, USA).

187

## 188 **Results**

### 189 ***In vitro exposure to HQ halts articular chondrocyte growth but does not induce*** 190 ***apoptosis***

191 To investigate whether exposure to HQ could affect chondrocyte viability, we treated  
192 bovine articular chondrocytes with different concentrations of HQ (1  $\mu$ M up to 100  $\mu$ M) and  
193 monitored cell growth across time by MTT assay.

194 HQ treatment slowed cell growth at concentrations between 10  $\mu$ M to 100  $\mu$ M over a  
195 96h time course in comparison to vehicle control (Figure 1A). While affecting cell growth, HQ  
196 did not alter cell viability under the concentrations of 1 to 50  $\mu$ M observed at the 72h time  
197 point (Figure 1B) and did not induce activation of apoptosis, as shown by the unaltered ratio  
198 of Bax/Bcl2 expression at mRNA levels (Figure 1C) (Karaliotas et al., 2015).

199

### 200 ***HQ modulates the expression of phenotypic markers in articular*** 201 ***chondrocytes***

202 We further investigated whether HQ treatment could influence the expression of  
203 phenotypic markers in articular chondrocytes, as these markers are altered during the  
204 progression of OA (Dell'Accio et al., 2001; Nalesso et al., 2011; Nalesso et al., 2017).

205 The mRNA expression of the differentiation markers SOX-9, Col2a1 and Col-X was  
206 significantly downregulated upon incubation with HQ for 24 hours (Figure 2A-C). Furthermore,  
207 the expression of the matrix remodelling enzyme MMP-3 and of the aggrecanase ADAMTS5  
208 was instead upregulated (Figure 2D, E). The catabolic activity of HQ was further confirmed



209 by a reduced amount of highly sulphated GAGs in chondrocyte micromasses upon treatment  
210 with HQ (Figure 2F, G).

211 Nitric oxide (NO) can suppress the synthesis of proteoglycans (PGs) and stimulates  
212 MMPs activity (Amin and Abramson, 1998). We therefore tested whether HQ could elicit NO  
213 production in articular chondrocytes. Indeed, HQ exposure for 48 hours significantly  
214 increased the nitrite production, as evaluated by Griess reaction (Figure 2H).

215 Finally, as HQ can induce the generation of ROS in different biological  
216 systems (Heluany et al., 2018b; Heluany et al., 2021; Mao et al., 2019), and the induction of  
217 oxidative stress in chondrocytes has been strongly associated with cartilage damage in OA  
218 (Henrotin, Bruckner and Pujol, 2003), we tested whether the HQ exposure could influence  
219 the production of ROS in BACs cultures. Our data showed that an incubation of chondrocytes  
220 with HQ for 24 hours triggered a significant increase in the levels of ROS in comparison to  
221 non-stimulated cells (Figure 2I).

222 Altogether, these results suggest that HQ alone can promote alteration in the  
223 chondrocyte metabolism and oxidative stress, which are associated with activation of tissue  
224 degenerative mechanisms.

225

### 226 ***HQ enhanced the pro-catabolic activity of IL-1 $\beta$***

227 IL-1 $\beta$  is a pro-inflammatory cytokine overexpressed in the articular cartilage during the  
228 progression of OA, promoting cartilage degradation processes (Sellan and Berenbaum,  
229 2010). Thus, to investigate whether inflammation could influence the catabolic activity of  
230 HQ, BACs were pre-stimulated with IL-1 $\beta$  at 20 ng/mL for 4 hours before being exposed to  
231 different concentrations of HQ for 24 hours or 48 hours. While HQ does not exacerbate the  
232 pro-catabolic effects of IL-1 $\beta$  at gene expression levels (Figure 3A, B), pre-stimulation with  
233 IL-1 $\beta$  increased GAG content reduction and NO generation induced by HQ, in a dose  
234 dependent fashion (Figure 3C-F).

235 These results were further confirmed on ex-vivo cultures of bovine cartilage explants.  
236 While the IL-1 $\beta$  pre-treatment did not influence the GAG content and the release induced by  
237 HQ (Figure 4A-B, D), HQ and IL-1 $\beta$  showed to have a cumulative effect on nitrite production  
238 (Figure 4C).

239 Altogether, our findings suggest that the pro-inflammatory stimuli can exacerbate  
240 the catabolic effects of HQ in the articular cartilage.

241

### 242 ***HQ activates the AhR pathway in articular chondrocytes.***

243 We previously demonstrated that HQ mediates its toxic effect in the joints through  
244 activation of the AhR pathway in murine experimental models of rheumatoid  
245 arthritis (Heluany et al., 2018a,b; Heluany et al., 2021). Thus, to investigate whether HQ  
246 signals through this pathway in the articular chondrocytes, we measured the expression  
247 levels of AhR, of the aryl hydrocarbon receptor nuclear translocator (ARNT) and of the  
248 endpoint target gene in the AhR pathway Cyp1a1 (Figure 5A) in BACS upon exposure to  
249 different concentrations of HQ. All the genes were upregulated in chondrocytes upon  
250 incubation with HQ for 24 hours and their modulations were rescued by the pre-incubation  
251 with  $\alpha$ -naphthoflavone ( $\alpha$ NF, 10  $\mu$ M), an AhR antagonist (Figure 5B-D). (Nguyen et al.,  
252 2015).

253 In the activated stage, the AhR translocates from the cytoplasm to the nucleus and  
254 forms a complex with the ARNT, which identifies and integrates with the xenobiotic responsive  
255 elements (XRE) of target genes, regulating their expression (Ma and Baldwin,  
256 2000). Activation of the pathway upon HQ stimulation was therefore further confirmed with an  
257 XRE-reporter assay in articular chondrocytes stimulated for 24 hours with HQ. As shown in  
258 the Figure 5E, treatment with HQ increased the intensity of the binding activity to the AhR  
259 gene reporter in a concentration-dependent manner. Finally, we confirmed that both the  
260 phenotypic changes as well as the pro-catabolic activity induced by HQ were mediated via  
261 the activation of the AhR pathway, as the HQ-mediated downregulation of SOX-9 and  
262 upregulation of MMP-3 were rescued by the pre-incubation with the AhR  
263 antagonist  $\alpha$ NF (Figure 5F, G).

264

## 265 Discussion

266 The effect of cigarette smoke on the health of joint tissues has not been fully elucidated  
267 yet. In a few observational studies, exposure to cigarette smoke has been shown to trigger  
268 contrasting effects, as smoking can both increase or reduce the incidence of OA, depending  
269 on the cohort examined and the methodology used for the analysis (Davies-Tuck et al.,  
270 2009; Racunica et al., 2007).

271 In an attempt to further elucidate how cigarette smoke can impact the health of joint  
272 tissues at molecular levels, here we focused our attention on investigating the role of HQ, a  
273 major component of cigarette smoke on articular cartilage health, as we have previously  
274 shown its contribution to the degeneration of other tissues in the joints in murine models of  
275 inflammatory arthritis (Heluany et al., 2018a,b; Heluany et al., 2021).

276 Our data showed that HQ decreases chondrocyte viability in a dose and time  
277 dependent manner. We and others previously associated the toxic effect of HQ with activation  
278 of apoptosis in different biological systems (Lee et al., 2012; Li et al., 2018), however, we did  
279 not detect any variation in apoptotic markers in articular chondrocytes upon HQ exposure  
280 within the timeframe of our analysis.

281 In OA cartilage, the articular chondrocytes undergo phenotypic changes, characterized  
282 by the downregulation of phenotypic lineage markers and the upregulation of pro-hypertrophic  
283 genes, as well as of matrix remodelling enzymes, which contribute to the progressive loss of  
284 the biomechanical properties of the tissue, and ultimately lead to its degradation (Arner,  
285 2002; Poole et al., 2001). Our data suggest that HQ could heavily contribute to many of these  
286 phenomena. We showed that HQ exposure promoted the downregulation of phenotypic  
287 markers, while also inducing the upregulation of MMP-3, a metalloprotease, whose activity is  
288 upregulated in OA, which can degrade several types of collagens and matrix proteins in the  
289 articular cartilage (Bortoluzzi et al., 2018). The enhancement of matrix remodelling activity  
290 induced by HQ was also supported by the reduction of GAG staining and increased GAG  
291 release upon exposure of chondrocytes 2D cultures to the xenobiotic.

292 Oxidative stress can be induced through several mechanisms and can contribute to  
293 cellular metabolic decline as well as to promote degenerative mechanisms (Suantawee et al.,  
294 2013). The induction of oxidative stress in chondrocytes has been strongly associated with  
295 increased cartilage degradation in severe cases of OA (Henrotin, Bruckner and Pujol, 2003).  
296 Moreover, nitrite production has also been shown to contribute to oxidative stress and

297 degeneration of tissue integrity in the joints (Abramson, 2008). Previous data showed that  
298 exposure to environmental pollutants could promote ROS and NO generation in chondrocytes  
299 (Lee and Yang, 2012). HQ exposure has been associated with oxidative damage and could  
300 affect the oxidative balance in other biological systems (Heluany et al., 2018b; Heluany et al.,  
301 2021; Peng et al. 2013; Pons et al. 2010). Indeed, here we confirmed the pro-oxidative effect  
302 of HQ on the articular chondrocytes, which could contribute to the overall phenotypic  
303 changes induced by the xenobiotic on these cells.

304 Interleukin 1 $\beta$  (IL-1 $\beta$ ) is a major pro-inflammatory trigger in OA that stimulates  
305 catabolic changes, suppresses anabolic pathways and decreases matrix synthesis (Jenei-  
306 Lanzl, Meurer and Zaucke, 2019). In our study, IL-1 $\beta$  and HQ had a synergistic effect in  
307 reducing proteoglycan content and in promoting oxidative stress in the articular chondrocytes,  
308 suggesting that exposure to cigarette smoke and environmental pollutants could potentiate  
309 inflammatory processes involved in the degradation of the articular cartilage in OA.  
310 Interestingly, similar synergistic effects were observed in synoviocytes derived from RA-  
311 patients co-stimulated with HQ and TNF- $\alpha$  (Heluany et al., 2021).

312 AhR is a ligand-dependent transcription factor that translocates to the nucleus upon  
313 activation by xenobiotics and pollutants (Julliard, Fechner and Mezrich, 2014). Here we  
314 confirmed that after HQ exposure, AhR translocates from cytoplasm to the nucleus, forms a  
315 heterodimer with the AhR nuclear translocator (ARNT) and promotes the transcription of  
316 target genes such as Cyp1a1. The activation of the AhR pathway has been shown to mediate  
317 several detrimental effects, such as aggravation of articular diseases, cancer promotion and  
318 endocrine disruption (Denison and Nagy, 2003; Nakahama et al., 2011; Nguyen et al.,  
319 2015). Several studies showed that this receptor plays a major role in exacerbating  
320 rheumatoid arthritis in smokers (Kobayashi et al., 2008; Nguyen et al., 2015; Talbot et al.,  
321 2018). We have also previously showed the HQ-mediated cytotoxicity in joint diseases  
322 through the activation of the AhR pathway (Heluany et al., 2018a,b; Heluany et al.,  
323 2021). Here we show that AhR and its downstream effectors are overexpressed by HQ in  
324 articular chondrocytes, and mediate the HQ catabolic effects, suggesting that xenobiotics and  
325 pollutants can directly influence the health of the articular cartilage.

326 In summary, our data demonstrated a clear detrimental effect of HQ on articular  
327 cartilage homeostasis and shed novel insight on how environmental pollutants can  
328 exacerbate the degenerative effect of pro-inflammatory mechanisms underlying the onset of  
329 articular diseases such as OA.

330

### 331 **Conflict of interests**

332 *The authors declare that the research was conducted in the absence of any commercial or financial*  
333 *relationships that could be construed as a potential conflict of interest.*

334

### 335 **Author Contributions**

336 CSH: Conceptualization, data collection, analyses, interpretation and manuscript writing, revision and  
337 proof-read. ADP: Data collection, manuscript proof-read. ND: Data collection, manuscript proof-read.  
338 SHPF: Conceptualization, supervision, interpretation and manuscript writing revision and proof-read.  
339 GN: Conceptualization, supervision, interpretation and manuscript writing, revision and proof-read.

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342



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539 Table 1: PCR primer sequences used for quantitative gene expression analysis.

Gene	Sense primer	Anti-sense primer	Annealing temperature (°C)
β-actin	AGCAGTCGGTTGGATCGAGCA	GGGAAGGCAAAGGACTTCCTGTAAC	55
SOX-9	GCTCTGGGCAAGCTCTGGAGACT	GGCGCGGCTGGTACTTGTAGTCC	60
Col2a1	ACGTCCAGATGACCTTCCTG	GGATGAGCAGAGCCTTCTTG	55
COL-X	AAAGGTCTAAGTGGCCCTTTTGTC	GAGGTTTCATGACAAAAGCACCTTGC	55
ADAMTS5	ACGGGACCGTCATGAACTAC	CTTTTGGAGCCGACTTCTTG	55
MMP-3	TGTGTGTCTTGCCCACTAGC	TGCCTGTTGCAGAATGCTAA	55
BAX	GCCCTTTTGCTTCAGGGTTTC	CATCCTCTGCAGCTCCATGT	60
Bcl-2	GGGGTCATGTGTGTGGAGAG	TCCACAAAGGCGTCCCAG	60
AhR	AGCAGCGCTAACATCACCTA	GGGACTGGCTTGACAGTTTTTC	60
ARNT	CAGGTGCACCCAGATGATGT	AGATCCAGGATACGCCCTGT	60
Cyp1a1	ACCTTGATCACTAACGGCCA	GCCTCCTTGTTACATGCTCT	60

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### Figure legends:

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#### **Figure 1: In vitro HQ exposure reduces viability but does not promote cell death.**

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Stimulation of BACs with increasing concentrations of HQ (1-100 μM) halted cell growth as evaluated by MTT assay (A) and presented in representative images after 72 hours of treatment (B). HQ treatment did not promote apoptosis (C) as the BAX/Bcl-2 mRNA ratio was not different across treatments and over time. Data represent mean ± SEM from three independent experiments and were analysed by one-way ANOVA. 16H: p= 0.0037 CM vs. 100 μM. 24H: p= 0.0215 CM vs. 1 μM; p= 0.0220 CM vs. 5 μM; p= 0.0010 CM vs. 10 μM; p= 0.014 CM vs. 25 μM; p = 0.0001 CM vs. 50 and 100 μM. 72H: p= 0.0049 CM vs. 25 μM; p = 0.0001 CM vs. 50 and 100 μM. 96H: p= 0.0262 CM vs. 10 μM; p = 0.0001 CM vs. 25, 50 and 100 μM.

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**Figure 2: HQ exposure modulated phenotypic markers in articular chondrocytes.** HQ stimulation of monolayer cultures of BACs downregulated the phenotypic markers SOX-9 (A), Col2a1 (B), Col-X (C) but upregulated the catabolic enzymes MMP-3 (D) and ADAMTS5 (E). HQ exposure reduced the production of highly sulphated GAGs (F-G) over a 48h incubation on BACs in micromass cultures, as measured by Alcian Blue quantification. HQ also increased nitrite production (H, quantified by Griess reaction) and ROS generation (I, quantified by DCFH-DA assay) in micromass cultures. qPCR results were normalized to the housekeeping gene β-actin and expressed as fold change to culture media. Data represent mean ± SEM from three independent experiments and were analysed with one-way ANOVA and unpaired t test.

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**Figure 3: HQ treatment enhances the pro-catabolic activity of IL-1β in articular chondrocytes.** Pre-incubation of monolayer cultures of BACS with IL-1β (20 ng/mL) for 4 hours before stimulation with HQ (10, 25 or 50 μM) for 48 hours, did not alter the expression



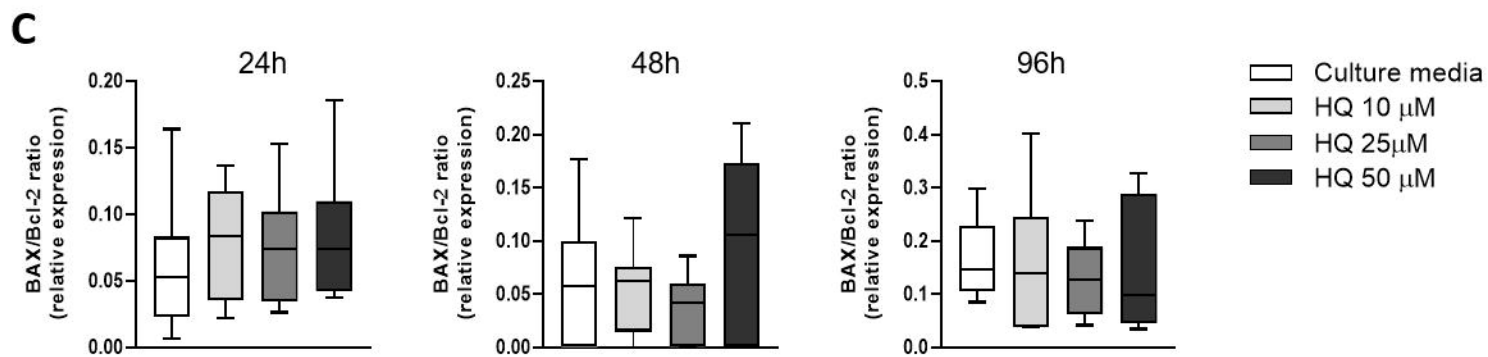
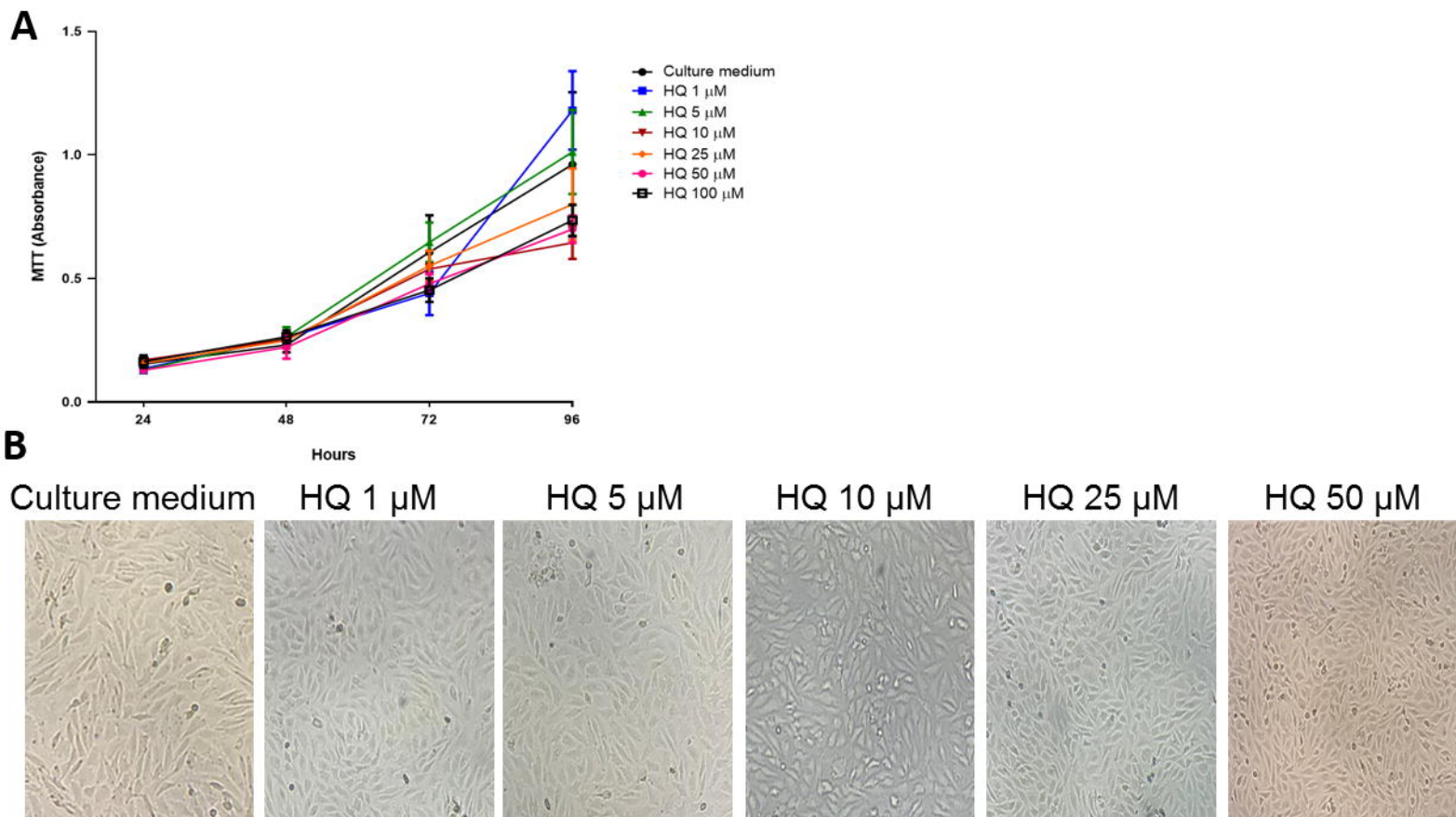
569 of the phenotypic marker Col2a1 **(A)** and of the catabolic enzyme MMP-3 **(B)** at mRNA level.  
570 Co-stimulation of IL-1 $\beta$  and HQ reduced the accumulation of highly sulphated GAGs **(C-**  
571 **D)**, as assessed by quantification of Alcian blue staining. The co-treatment also enhanced the  
572 release of GAGs **(F)** and increased the nitrite production **(G)** in micromass cultures of BACs,  
573 as measured respectively by DMMB assay and Griess reaction, respectively. qPCR results  
574 were normalized to the housekeeping gene  $\beta$ -actin and expressed as fold change to culture  
575 media. Data represent mean  $\pm$  SEM from three independent experiments and  
576 were analysed with one-way ANOVA and unpaired t test.

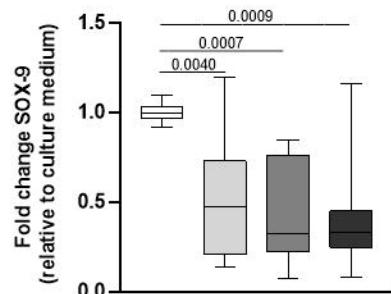
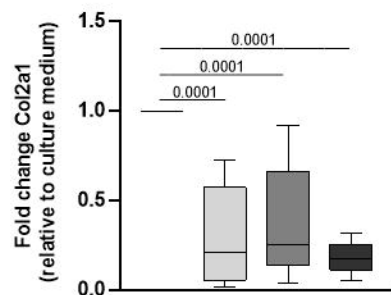
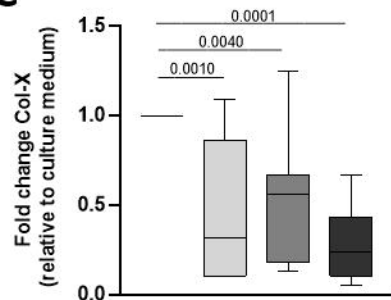
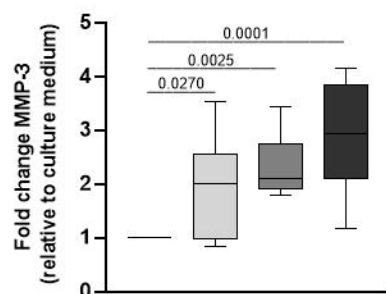
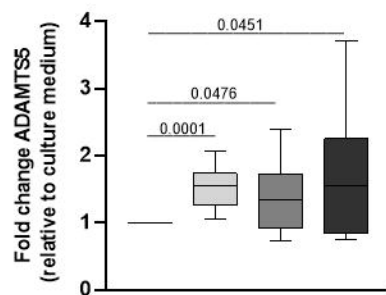
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578 **Figure 4: HQ treatment increases the pro-inflammatory activity of IL-1 $\beta$  in articular**  
579 **cartilage explants.** Bovine articular cartilage explants were pre-incubated with CM or with  
580 IL-1 $\beta$  (20 ng/mL) for 4 hours and subsequently incubated with HQ (10, 25 or 50  $\mu$ M) for 72  
581 hours. The co-stimulation did not alter GAGs synthesis and release **(A-B, D)**, but increased  
582 nitrite production **(C)** in comparison to stimulation with IL-1 $\beta$  alone. 10X magnification. Data  
583 represent  $\pm$  SEM from four independent experiments and were analysed with one-way  
584 ANOVA.

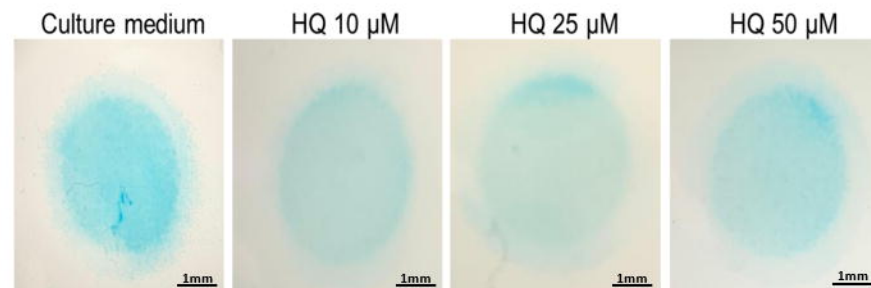
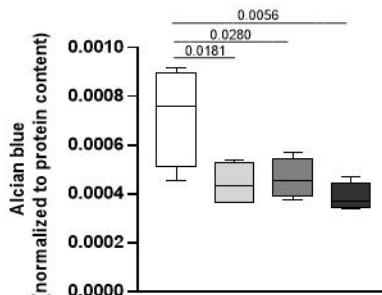
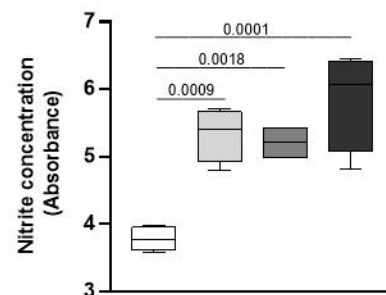
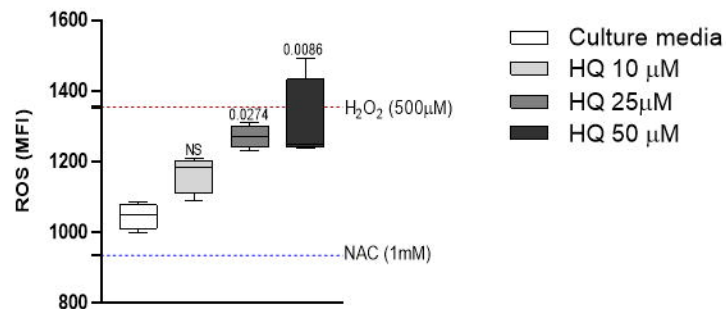
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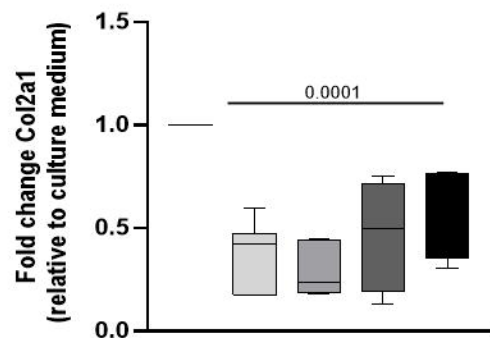
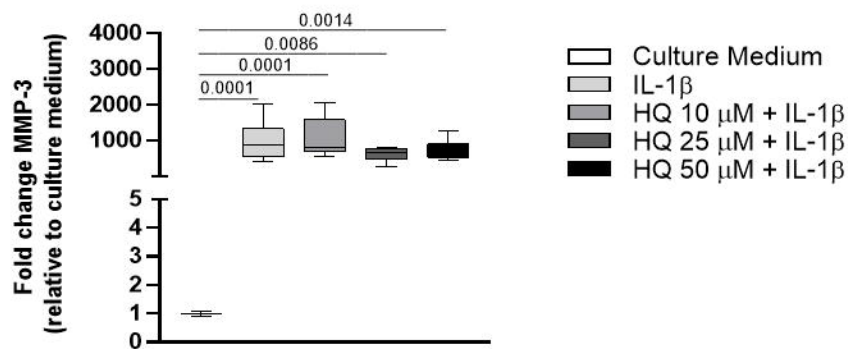
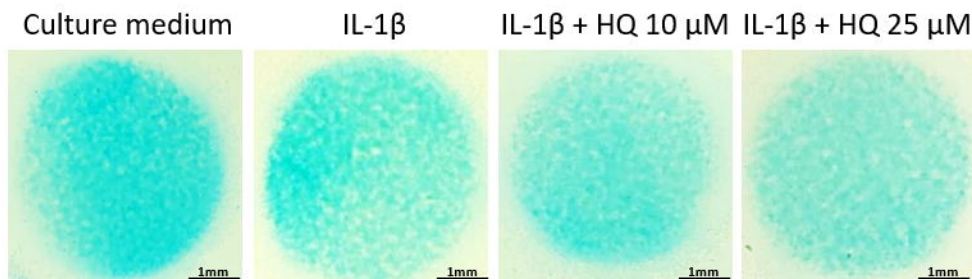
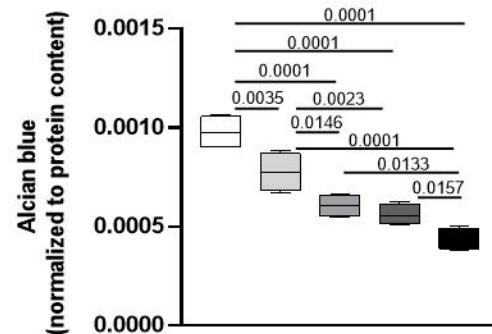
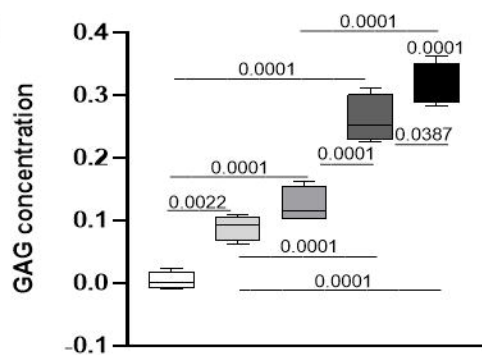
586 **Figure 5: HQ activates the AhR pathway in articular chondrocytes: (A)** Schematic  
587 representation of the AhR pathway. Modulation of components of the AhR pathway was  
588 determined by measuring mRNA levels of AhR, ARNT and Cyp1a1 in BACs incubated with  
589 CM or with HQ (10, 25 or 50  $\mu$ M) in absence or presence of  $\alpha$ NF (10  $\mu$ M) for 24 hours **(B-**  
590 **D)**. Results were normalized to the housekeeping gene  $\beta$ -actin and are expressed as fold  
591 change to CM. **(E)** Activation of the AhR receptor by HQ was confirmed through a luciferase-  
592 based reporter assay. Inhibition of AhR with  $\alpha$ NF rescued the downregulation of SOX-  
593 9 **(F)** and the upregulation of MMP-3 **(G)** mediated by HQ. Data represent mean  $\pm$  SEM from  
594 three independent experiments and were analysed by one-way ANOVA. \* $p$ <0.05, \*\* $p$ <0.01,  
595 \*\*\* $p$ <0.001 and \*\*\*\* $p$ <0.0001 vs. CM; # $p$ <0.05, ## $p$ <0.01, ### $p$ <0.001 and  
596 #### $p$ <0.0001 vs. IL-1 $\beta$ .



**A****B****C****D****E**

Culture media  
 HQ 10  $\mu$ M  
 HQ 25  $\mu$ M  
 HQ 50  $\mu$ M

**F****G****H****I**

**A****B****C****D****E****F**