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Hydrogen phosphate-mediated acellular biomineralisation within a dual crosslinked 1 2 hyaluronic acid hydrogel Ziyu Gao,^{a,b} Layla Hassouneh,^a Xuebin Yang,^a Juan Pang,^c Paul D. Thornton,^{*,b} Giuseppe 3 Tronci*^{,a,d} 4 ^a Biomaterials and Tissue Engineering Research Group, School of Dentistry, St. James's 5 University Hospital, University of Leeds, UK. E-mail: <u>g.tronci@leeds.ac.uk</u>. 6 ^b School of Chemistry, University of Leeds, Leeds, UK. E-mail: p.d.thornton@leeds.ac.uk. 7 ^c School of Material Engineering, Jinling Institute of Technology, Nanjing, China. 8 ^d Clothworkers' Centre for Textile Materials Innovation for Healthcare, School of Design, 9 10 University of Leeds, UK. 11 12 13 Highlights 1. Non-toxic acellular design of a dual crosslinked hyaluronic acid (HA) hydrogel 14 2. Hydrogen phosphate ions form physical crosslinks with cystamine crosslinked HA 15 3. Salt treatment is non-toxic and generates cell-migrating aggregated structures 16 17 4. Physical crosslinks act as a nucleation site for hydroxyapatite mineralisation 18 5. Hydroxyapatite crystals accomplished across the hydrogel in simulated body fluid 19 20 **Graphical abstract** 21



23 ABSTRACT

The creation of hyaluronic acid (HA)-based materials as biomineralisation scaffolds for cost-24 effective hard tissue regenerative therapies remains a key biomedical challenge. A non-toxic 25 and simple acellular method to generate specific hydrogen phosphate (HPO $_4^{2-}$) interactions 26 within the polymer network of cystamine-crosslinked HA hydrogels is reported. Reinforced dual 27 28 crosslinked hydrogel networks were accomplished after 4-week incubation in disodium phosphate-supplemented solutions that notably enabled the mineralisation of hydroxyapatite 29 (HAp) crystals across the entire hydrogel structure. HPO_4^{2-} -cystamine-crosslinked HA hydrogen 30 bond interactions were confirmed by attenuated total reflectance Fourier transform infrared 31 spectroscopy (ATR-FTIR) and density functional theory (DFT) calculations. HPO42-mediated 32 physical crosslinks proved to serve as a first nucleation step for acellular hydrogel 33 mineralisation in simulated body fluid allowing HAp crystals to be detected by X-ray powder 34 diffraction ($2\theta = 27^{\circ}$, 33° and 35°) and visualised with density gradient across the entire 35 hydrogel network. On a cellular level, the presence of aggregated structures proved key to 36 inducing ATDC 5 cell migration whilst no toxic response was observed after 3-week culture. This 37 mild and facile ion-mediated stabilisation of HA-based hydrogels has significant potential for 38 39 accelerated hard tissue repair in vivo and provides a new perspective in the design of dual 40 crosslinked mechanically competent hydrogels.

Keywords: Hyaluronic acid; Hydrogel; Hydrogen phosphate interaction; Cystamine crosslinking;
Biomineralisation.

43

44 **1. Introduction**

As one of the main components of extracellular matrix (ECM), hyaluronic acid (HA) has been applied widely in medicine, for example as a lubricant for osteoarthritis treatment [1] [2], wound dressing material to support healing [3] and as post-operation adhesive [4]. Recently, HA hydrogels have been developed as implants to support cell growth and aid regeneration of soft tissues including derm [3] [5], mucosa [4] [6] and tendon [7] [8], due to the biocompatibility, biodegradation profile and mechanical properties of HA. The advantageous features of HA in biology, as well as its chemical structure, which can be selectively targeted to fabricate mechanically competent bioinspired scaffolds, have also been leveraged to support the regeneration of bone. However, this has frequently required either severe or sophisticated synthetic approaches to address the mechanical and compositional requirements of bone. Although many methods have been investigated [9] [10], mild non-toxic routes enabling the fabrication of drug-free bone-like HA-based architectures have not yet been fully realized.

Ionic interactions, particularly salt effects, enable biomacromolecule crosslinking in a mild 57 58 manner that avoids chemical synthesis and/or extensive energy radiation [11][12]. The Hofmeister effect details the extent that protein solubility is altered by the presence of 59 different salts in an aqueous environment, and may be used to design protein-based hydrogels 60 61 with enhanced compressive and tensile properties [13] [14]. The mechanism of salt effect on nonelectrolytes in aqueous solutions has been explored in (i) hydration theories, (ii) 62 electrostatic theories, (iii) Van der Waals forces, and (iv) internal pressure concepts [15]. 63 64 However, non-specific ion-mediated interactions may be applied universally in macromolecules [16][17][18]. Barrett hypothesized that a particular salt could act as either a stabilizer (i.e. 65 kosmotrope) or a destabilizer (i.e. chaotrope) for a specific macromolecule [19]; for instance, 66 67 alginate may be particularly well stabilized by calcium [20][21][22]. Leveraging aforementioned 68 salt effects, we hypothesised that phosphate groups may enable the generation of additional 69 physical crosslinks in a chemically crosslinked HA hydrogel network bearing amide net-points, on the one hand, and act as nucleation sites to accomplish hydrogel biomineralisation in near-70 physiologic conditions, on the other hand. Phosphate groups were selected as the most 71 72 common component of buffer salts and since they are known to mediate protein denaturation [23], the stabilisation of HA-based electrospun fibres [24] and biomineralisation [25]. 73

Despite the crucial role of HA in the ECM of biological tissues and the unique functions of phosphate groups in hard tissue repair, the interaction of phosphate ions with HA-based hydrogels has only partially been studied, suggesting limited control of molecular interactions and macroscopic effects [26]. Attempts to characterise the interaction between HA and the phosphate head group in phospholipid model membranes have been made through differential scanning calorimetry (DSC), fluorescence spectroscopy, small-angle X-ray scattering (SAXS), infrared spectroscopy (IR) and atomic force microscopy (AFM) [27]. However, the resulting

phosphate ion-HA interaction was too insignificant to be observed by the above-mentioned methods. This underlines the experimental challenge in designing phosphate ion-mediated dual crosslinked HA-based hydrogel systems as a biomineralisation template for the direct build-up of bioinspired, mechanically competent HA matrices for hard tissue repair.

85 Other than phosphate-HA interactions, the integration of hybrid micromorphologies has attracted great interest in bone regeneration [28], and has been pursued in HA-based hydrogels 86 aiming to realize bioinspired bone-like nanocomposites [10]. The in situ precipitation of calcium 87 phosphate was reported on the surface of HA hydrogels, yielding a calcium phosphate 88 nanocomposite on the outer layer of the hydrogel scaffold [9]. Ion diffusion methods have also 89 90 been studied for mineralisation, including an electrophoresis approach [29] and a doublediffusion system [30]. However, only amorphous hydroxyapatite (HAp) was observed in the 91 electrophoresis approach, whereas only calcium phosphate minerals were obtained via the 92 sophisticated double-diffusion system. Consequently, accomplishing time-efficient and 93 controllable formation of HAp crystals with native patterns and growing density is still a great 94 challenge in the design of hierarchical 3-dimensional (3D) structures that mimic human bones 95 [31]. Constructing a secondary crosslinked structure by including HPO₄²⁻ in the hydrogel matrix 96 97 may provide microchannels within the network that enable HAp formation, and consequently 98 bone repair.

In this work, two HA-based hydrogels that contained either cystamine- or ethylenediamine-99 induced crosslinks were designed and assessed in phosphate-supplemented aqueous solutions 100 101 and a range of salts that partially comprise the Hofmeister series, with the aim of developing a simple method to induce both dual crosslinking and HAp mineralisation across the hydrogel 102 structure. We hypothesised that non-toxic phosphate-binding amide crosslinks could be 103 104 introduced during the crosslinking reaction to control the swelling and mechanical properties of 105 the HA-based hydrogels and lay down the foundation of a new bioinspired HA-based structure. The increased segment length of, and the presence of disulfide bridges in, cystamine-106 crosslinked (with respect to ethylenediamine-crosslinked) HA chains were hypothesised to 107 minimise steric hindrance and enhance the yield of physical crosslinking and acellular 108 109 biomineralisation during hydrogel incubation in phosphate-supplemented aqueous solutions.

Incubation of the hydrogels in aqueous solutions supplemented with hydrogen phosphate 110 111 $(HPO_4^{2^-})$ generated hydrogen bonds acting as physical crosslinks, thereby yielding a very stable macrostructure with customisable mechanical properties. The mineralisation process of $HPO_4^{2^-}$ -112 conditioned HA hydrogels was monitored in conventional simulated body fluid (c-SBF), whereby 113 unique hierarchical structure and gradients of HAp mineral were recorded across the entire 114 hydrogel and confirmed by X-ray computed microtomography (µCT). The simplicity and 115 mildness of this dual crosslinking and mineralisation approach enable method transferability to 116 other biopolymers and offers great promise for the creation of drug-free bioinspired materials 117 for cost-effective bone regenerative therapies. 118

119

120 2. Materials and methods

121 **2.1. Materials**

Hyaluronic acid sodium salt (molecular weight: 1,200 kDa, cosmetic grade) was purchased 122 from Hollyberry Cosmetic, 4-(4,6-dimethoxy-1,3,5-triazin-2-yl)-4-methyl-morpholinium chloride 123 124 (DMTMM) and 2-(N-morpholino) ethanesulfonic acid (MES) were purchased from Fluorochem. tris(hydroxymethyl)aminomethane 125 $(NH_4)_2SO_4$, $Na_2HPO_4 \cdot 7H_2O_1$ (TRIS), cystamine dihydrochloride and ninhydrin reagent were purchased from Alfa Aesar. Na₂SO₄, CH₃COONa 126 (NaAc), NaHCO₃, KCl, K₂HPO₄·3H₂O, MgCl₂·6H₂O, CaCl₂, and ethylenediamine were ordered 127 from VWR. Phosphate buffered saline (PBS) was purchased from Lonza.2,4,6-128 trinitrobenzenesulfonic acid (TNBS), alamarBlue[™] Cell Viability Reagent, CellTracker[™] Green 5-129 chloromethylfluorescein diacetate (CMFDA) dye and the LIVE/DEAD[™] cell stain kit were 130 131 purchased from ThermoFisher Scientific. All other reagents were purchased from Sigma-Aldrich. Unless specified, all the general reagents were analytical grade. 132

133

134 **2.2. Hydrogel preparation**

HA hydrogels were fabricated according to our previous method [32]. HA powder was dissolved in MES buffer solution (0.1 M, pH 5.5) at room temperature in 2 wt.% concentration. DMTMM (2 equivalents per HA repeat unit) was then added at 37 °C to activate the carboxyl groups of HA. Following 1-hour activation at 37 °C, either cystamine or ethylenediamine was added with a molar ratio of 0.4 moles relative to the moles of each HA repeat unit. The stirring speed was increased to 1000 rpm for 5 minutes, and either 0.6 g or 0.8 g of the reacting solution was cast into 24-well plates. HA hydrogels were obtained after 2-hour incubation at 37 °C. Cystamine and ethylenediamine crosslinked HA hydrogels were named as C2-40 and E2-40, whereby C and E signify HA crosslinking with cystamine and ethylenediamine, respectively; 2 is the wt.% of HA in the hydrogel-forming solution, whilst 40 is the mol.% of each crosslinker added with respect to HA's carboxylic groups.

146

147 2.3. TNBS assay and determination of polymer crosslinking

148 Polymer crosslinking density was indirectly assessed via determining the concentration of 149 unreacted amine groups presented by the crosslinkers (either cystamine or ethylenediamine) in each hydrogel using the 2,4,6-trinitrobenzene sulfonic acid (TNBS) assay [32]. 0.8 g of freshly 150 151 synthesised hydrogel was freeze-dried without deionised water washing. Each dry network was immersed in 2 mL NaHCO₃ solution (4 wt.%) at 40 °C for 30 minutes to remove any unreacted 152 cystamine or ethylenediamine. 1 mL of the supernatant was collected and incubated in dark (40 153 °C, 3 hours, 120 rpm) with 1 mL TNBS solution (0.5 wt.% in deionised water). 3 mL HCl (6 N) was 154 155 added to the incubated solution, and the temperature raised to 60 °C for 1 hour to terminate 156 the reaction. After cooling to room temperature, the sample solutions were diluted with 5 mL of deionised water. The unreacted TNBS was washed out by extraction with 20 mL diethyl ether 157 (×3). 5 mL of the retrieved sample solution was incubated in hot water to evaporate any diethyl 158 ether and diluted with 15 mL of deionised water. Finally, 2 mL of each solution was analysed by 159 UV-Vis spectroscopy at 346 nm. Quantification of any cystamine or ethylenediamine residue 160 was carried out by comparison with a cystamine or ethylenediamine calibration curve. 161

162

163 2.4. Hydrogel swelling tests

Various ion-hydrogel interactions were compared through changes in swelling ratio. Each replicate of prepared C2-40 and E2-40 hydrogels of known wet weight (ω_0) was individually immersed in either (NH₄)₂SO₄, Na₂SO₄, Na₂HPO₄·7H₂O, CH₃COONa (NaAc), NaCl or deionised water (50 mL solution). Swelling tests in PBS buffer solution (LONZA) and conventional

168 simulated body fluid (c-SBF) were also carried out. The wet weight (ω_t) was recorded at 169 different time points for up to 4 weeks. All the single-salt solutions used were prepared with 50 170 mM concentration and replaced by fresh solution every week with the same volume. The c-SBF 171 solution was prepared as reported previously [33]. Briefly, all the salts were added to 960 mL deionised water in the following order: 8.036 g NaCl, 0.352 g NaHCO₃, 0.225 g KCl, 0.230 g 172 K₂HPO₄·3H₂O, 0.311 g MgCl₂·6H₂O, 40 mL HCl (1.0 M), 0.293 g CaCl₂, 0.072 g Na₂SO₄, 6.063 g 173 174 TRIS. The pH of the solution was buffered at pH 7.4 by adding HCl (1.0 M). The swelling ratio 175 was calculated via eq. 1, as reported below:

176 Swelling ratio =
$$\frac{\omega_t}{\omega_0} \times 100$$
 (1)

177

178 **2.5. Hydrogel stability tests**

Hydrogels were incubated for 4 weeks in either the Na₂HPO₄-supplemented solution or deionised water. Following incubation, retrieved samples were washed by immersing in deionised water (×3) to remove any free salts and then freeze-dried. The relative mass of the hydrogel was calculated according to eq. 2 by measuring the dry weight of the freeze-dried freshly synthesized (ω_0) and retrieved (ω_d) samples, as reported below:

184 Relative mass
$$=\frac{\omega_d}{\omega_0} \times 100$$
 (2)

185

186 **2.6. Hydrogel compression tests**

Hydrogel compression properties were measured using a Bose ELF 3200 apparatus with a 0.02 mm/s compressive rate. All replicates were cut into 3 mm diameter cylinders. Compression stress and strain of either initial or salt-treated C2-40 and E2-40 hydrogels were evaluated and compared.

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2.7. Morphology study of the hydrogel network following salt treatment

Hydrogel morphology was observed using a HITACHI 3400 scanning electron microscope (SEM) under 20 kV voltage with gold coating. All hydrogels were treated with different salts for 4 weeks and flushed with deionised water. SEM analysis was carried out on freeze-dried hydrogel networks. Samples were carefully transferred into 6-well cell culture plates and frozen at -20 °C prior to lyophilisation, to minimise lyophilisation-induced sample shrinking. During the
 course of incubation, the hydrogel structures were also observed by optical microscopy (Zeiss)
 at different time points after various treatments.

200

201 2.8. Mechanistic study

 HPO_4^{2-} interaction with the HA hydrogels was investigated via attenuated total reflectance 202 Fourier transform infrared spectroscopy (ATR-FTIR, Bruker spectrometer) at room temperature 203 and density functional theory (DFT) calculations. The optimised structures were obtained by 204 DFT calculations at b3lyp/6-31G(d) level carried out using Gaussian 16 program [34]. The 205 binding energy between the HA repeat unit and HPO₄²⁻ was simply calculated as $[\Delta E = E_{total} - (E_{HA})]$ 206 + E_{HPO4})], in which the single point energy was calculated at b3lyp/6-311+G(d,p) level. For 207 display, blue dashed lines indicated the hydrogen bonds, oxygen (O) atoms were depicted in 208 red, nitrogen (N) in blue, sulfur (S) in yellow, carbon (C) in grey, hydrogen (H) in white and 209 phosphorus (P) in pink. All the atoms which were involved in hydrogen bond formation are 210 depicted as spheres. 211

212

213 2.9. Cell adhesion study

ATDC 5 chondrocytes (chondrogenic cell line) were used as non-mineralising joint resident 214 cells of the bone-cartilage interface. The initial C2-40 network (which was proven to mediate 215 secondary interactions with phosphate ions) was washed by sterile deionised water (×3) and 216 basal cell culture medium (BM) (×3). BM was composed of 50 vol.% Dulbecco's modified eagle's 217 medium (DMEM, D6546) and 50 vol.% Ham's nutrient mixture F12 (12-615), and supplemented 218 by 5 % fetal calf serum (FCS) and 1% penicillin and streptomycin (PS). The final concentration of 219 phosphorus in BM was 0.884 mM. Cells were labelled by CellTrackerTM Green (CMFDA) and re-220 suspended in medium with a cell density of 2×10^5 cells/mL. 100 µL cell suspension (2×10^4 cells) 221 was injected on the surface of each hydrogel (n=3). 2 mL of BM was added into each well after 222 3 hours seeding. Cell attachment and growth was observed and recorded after 48 hours by 223 224 fluorescence/optical microscopy (Zeiss).

225 To study the influence of Na_2HPO_4 on cell migration, BM was replaced by Na_2HPO_4 treated 226 medium (TM) after 1-week of culture. The cell culture in TM was named as "conditional cell 227 culture" and this culture time started when the medium was replaced. TM was prepared from the basal medium via supplementation of sterile Na₂HPO₄·7H₂O powder to achieve a final 228 229 concentration of 1.884 mM (1 mM increase in phosphate compared with BM). In the control group, the medium was replaced by fresh BM. ATDC 5 cell attachment and growth were 230 231 investigated by fluorescence/optical microscopy (Zeiss), cell migration was studied via Laser scanning confocal microscopy (LEICA TCS SP8, excitation wavelength 488 nm). All the samples 232 were washed by sterile PBS (×3) to remove any dead cells and impurities before calcein-AM 233 234 staining.

235

236 **2.10. Acellular mineralisation**

237 Hydrogel C2-40 was selected for the biomineralisation study given its capability to mediate secondary interactions with phosphate ions (confirmed by swelling and compression 238 measurements). After 4-week immersion in Na₂HPO₄ solution (50 mM, 1.0 L, 37 °C), C2-40 239 hydrogels (n=3) were transferred into an excess of deionised water for 24 hours to remove any 240 241 free phosphate salt, whereby the deionised water was replaced for three times during this time 242 period. Washed hydrogels were then soaked in 200 mL calcium chloride (10 mM) for another 24 hours [35]. Calcium-treated C2-40 samples were flushed by deionised water to remove any 243 surface salt and subsequently soaked in 1.0 L c-SBF for mineralisation at 37 °C for 2 weeks. Non-244 Na₂HPO₄-treated C2-40 samples were immersed in CaCl₂ (10 mM, 200 mL) for 24 hours and 245 underwent the same mineralisation procedure as a control group. The mineral structure was 246 confirmed by X-Ray powder diffraction (XRD) at room temperature in the range of 2q of 20°-247 248 60°. Freeze-dried initial and mineralised C2-40 networks, as well as mineralised C2-40 networks 249 after being burnt at 1000 °C for 30 minutes, were measured. The 3D structure of mineralisation was investigated by X-ray computed microtomography (µCT) (Skyscan 1072, Bruker, Kontich, 250 Belgium). Samples were scanned at 100 kVp, 100 mA, and 11.19 µm pixels, with a 1-mm 251 aluminium plus copper filter and a scanning time of around 60 minutes. A reconstruction 252 253 software program (NRecon; SkyScan) was used to convert the raw data into bitmap (bmp) files.

3D alignment and registration of samples were done using Data Viewer software (v1.4.3; Bruker microCT). Both CTan and CTvol (v1.10.11.0; Bruker microCT) software were used for the 3D structural analysis.

257

258 **2.11. Statistical analysis**

All the results were analysed with at least three replicates ($n \ge 3$). The results are presented as mean±SD. The significant difference was calculated through One-way ANOVA analysis with a pvalue at 0.05, which was considered as significant. These were labelled as *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001.

263

264 3. Results and discussion

265 **3.1. Hydrogel crosslinker density**

The crosslinker density was evaluated by the TNBS assay (Figure S1, Supp. Inf.) to determine the quantity of ethylenediamine or cystamine included within the covalent network formed [32]. When adding 40 mol.% of either cystamine or ethylenediamine, approximately 25 mol.% of crosslinker reacted with HA during gel formation (**Table 1**), ensuring that a comparable crosslink density was accomplished in both hydrogel networks regardless of the crosslinker used.

272

Table 1. Composition of HA hydrogels crosslinked with either cystamine (C2-40) or ethylenediamine (E2-40). TNBS
 assay was employed to quantify the crosslinker quantity in the HA network. Results are presented as Mean±SD.

HA concentration (wt.%)	Crosslinker quantity (mol.% of -COOH)		
	Added in	Reacted	
2.0	40.0	25.30±0.85	
2.0	40.0	25.27±0.01	
	HA concentration (wt.%) 2.0 2.0	HA concentration (wt.%)Crosslinker quant Added in2.040.02.040.0	

275

276 **3.2. Swelling behaviour of HA hydrogels**

Hydrogel swelling equilibrium was reached after 1 day for both C2-40 and E2-40 hydrogels following incubation in single salt-supplemented solutions (**Figure 1, a&b**). The swelling ratio

(SR) of C2-40 samples was found to be in the region of 150 wt.% in all salt solutions, whilst a
swelling ratio of 425 wt.% was measured in deionised water.



281

Figure 1. Effect of salt-supplemented solution on hydrogel swelling and compressive properties. (a-b): Swelling ratio of C2-40 (a) and E2-40 (b) hydrogels in $(NH_4)_2SO_4$, Na_2SO_4 , Na_2HPO_4 , NaAc, NaCl and deionised water (H_2O) . Insert graphs: Swelling Ratio (%, Y axis) profile over time (days, X axis). (c): Swelling ratio of C2-40 and E2-40 hydrogels in PBS and c-SBF. (d-e): Compression strain at break measured with hydrogel C2-40 following synthesis ('Original', surface flushed by deionised water before testing) and either 1-day (d) or 4-week (e) incubation in single salt-supplemented solutions. Statistical analysis is presented with respect to the H₂O group and labelled as **p < 0.01, ***p < 0.001, ****p < 0.0001. Data are presented as Mean±SD.

289

The interaction of selected salts with C2-40 resulted in decreased hydrogel swelling, following 290 the order $Na_2HPO_4 > (NH_4)_2SO_4 = Na_2SO_4 > NaCl > NaAc$. For sample E2-40, lower swelling ratio 291 values (~325 wt.%) were observed in deionised water, compared with hydrogel C2-40. 292 293 However, anion-HA hydrogel interactions were obvious in the 4-week swelling study and followed the order $Na_2SO_4 \ge (NH_4)_2SO_4 > NaCl > NaAc > Na_2HPO_4$. When comparing the two 294 295 hydrogels, the most striking difference is observed in the swelling behaviour in Na₂HPO₄-296 supplemented solutions. Following incubation of sample C2-40 in Na₂HPO₄, a significant decrease of SR was observed from 123±2 wt.% (1 day) to 62±1 wt.% (28 days). In contrast, the 297

SR of E2-40 in Na₂HPO₄ increased from 140±5 wt.% (1 day) to 161±4 wt.% (28 days). This observation was hypothesised to reflect the specific hydrogen bond interaction between HPO_4^{2-} and the cystamine-crosslinked HA chains.

As observed in the Na₂HPO₄-supplemented solution, the swelling ratio of samples C2-40 in 301 PBS solution (Figure 1c) presented a similar decreasing trend over time (SR: 101±3 wt.% (1 day) 302 \rightarrow 76±2 wt.% (28 days)), supporting the hypothesis that phosphate ions lead to a reduction in 303 hydrogel swelling. However, the swelling ratio of ethylenediamine crosslinked hydrogel (E2-40) 304 was stable (\sim 120 %) for the first 7 days before marginally decreasing over the next 21 days (SR: 305 119±1 wt.% (7 days) \rightarrow 114±2 wt.% (28 days)) (Figure 1c). Based on the significant decrease in 306 307 SR measured in hydrogel C2-40 following incubation in both Na₂HPO₄ and PBS solution, the swelling ratio was also recorded in c-SBF to further elucidate any HPO₄²⁻-mediated interaction 308 with cystamine-crosslinked HA. As expected, a similar but slower decrease in hydrogel swelling 309 310 was recorded in c-SBF over time, which is likely due to the different phosphate concentrations across the selected solutions (Table 2). 311

Table 2. Swelling ratio of C2-40 hydrogels following 4-week incubation in phosphate-supplemented solutions. The
 results are presented as Mean±SD.

Solution Name	Phosphate concentration (mM)*	Swelling ratio (%)
Na ₂ HPO ₄	50	62±1
PBS	6.658	76±2
c-SBF	1.001	87±1

314 **Concentration of hydrogen phosphate and dihydrogen phosphate.*

The significant difference in swelling ratio of hydrogel C2-40 was therefore attributed to the interactions between HPO_4^{2-} ions and cystamine-crosslinked hyaluronic acid, offering a new dimension for adjusting the swelling of the hydrogel by altering the chemical composition of the crosslinker.

Other than the swelling behaviour, the stability of C2-40 hydrogel was determined by quantifying its relative mass following 4-week incubation in the Na_2HPO_4 -supplemented aqueous solution (Figure S2, Supp. Inf.). Although a decrease in mass was observed in Na_2HPO_4 treated C2-40 networks with respect to deionised water-treated controls, a relative mass of

³¹⁵

324 67.7 ± 1.8 wt.% was measured compared to 71.7 ± 0.4 wt.% for the hydrogel controls, verifying 325 good material stability and limited Na₂HPO₄ impact.

326

327 **3.3. Compressive properties of salt-treated hydrogels**

Both C2-40 and E2-40 hydrogels were reinforced by ions to some degree, whereby stiffer 328 networks and varying values of compression strain (Figure 1, d&e) and stress at break (Table 329 330 S1) were measured, which further proved the effect of salts on hydrogel mechanical properties. Among the salt-treated samples, the most interesting phenomenon was observed in C2-40 331 hydrogels incubated in the Na₂HPO₄ environment, whereby the lowest value of compression 332 333 strain at break (77±0.3 %) was recorded after 1 day before decreasing to 54±1.5 % after 4-week treatment. All the other groups formed a relatively stable network (Figure 1, d&e). This 334 observation further supported the development of selective, strong HPO₄²⁻-mediated physical 335 crosslinks in the C2-40 hydrogel following salt treatment, so that the mechanical behaviour of 336 the resulting dual crosslinked hydrogel network could be adjusted from elastic to stiff. This 337 variation in mechanical behaviour was also supported by the trends of compression stress at 338 break measured in Na_2HPO_4 -treated and water-incubated groups after 1-day and 4-week 339 treatment (Figure S3&S4). The additional interactions between HPO₄²⁻ groups and the 340 341 cystamine-crosslinked HA network were therefore investigated as a means to induce acellular biomineralisation of cystamine-crosslinked HA hydrogel. 342

343

344 **3.4. Morphology of salt-treated hydrogels**

345 To study whether hydrogel surface morphology is affected by the salt treatment, freezedried hydrogels were inspected by SEM after 4-week incubation in Na₂HPO₄-supplemented 346 solution. Crystal-like salts were not observed by SEM in both samples C2-40 (Figure 2) and E2-347 40 (Figure S5), suggesting that salts diffused into the hydrogel and attached to the network, in 348 agreement with the salt-enhanced compression properties and decreased swelling ratio (Figure 349 1). All retrieved samples exhibited comparable porous-like surfaces, indicating minimal impact 350 of the incubation process with either salt-supplemented incubating media (Figure 2 a-f) or salt-351 352 free deionised water (Figure 2 g). This, together with the non-detection of crystal-like salts on

- 353 the hydrogel surface, suggests that any interaction of salt species, i.e. phosphate groups, with
- 354 HA's covalent network occurred at the molecular rather than microscopic scale.



Figure 2. SEM images of freeze-dried C2-40 networks following 4-week incubation in aqueous solutions. (a):
(NH₄)₂SO₄ (including zoomed-in image below); (b): Na₂SO₄; (c): Na₂HPO₄; (d): NaAc; (e): NaCl; (f): PBS; (g): H₂O.
Scale bar of (a-g): 500 μm. Scale bar of zoomed-in image of (a): 50 μm.

To further elucidate the extent of the above-mentioned ion interactions, hydrogels were incubated for three weeks in the presence of Na₂HPO₄ (50 mM), PBS solution and c-SBF. Aggregation of the hydrogel surface was observed in retrieved samples (**Figure 3**) after 3-week treatment in either Na₂HPO₄ (**Figure 3a**) or c-SBF (**Figure 3c**), whilst no visible effect was seen in hydrogels incubated for three weeks in PBS. On the other hand, when the incubation time in PBS solution was extended from 3 weeks to 3 months, aggregated structures with regular gaps were clearly visible in C2-40 hydrogels (**Figure 3 d&e**), as highlighted by the red arrows.

367 Given the absence of crystal-like aggregates via previous SEM analysis, aforementioned microscale effects are likely attributed to the development of strong interactions between 368 hydrogel C2-40 and HPO_4^{2-} ions, whereby the decreased yield of aggregation in PBS with 369 respect to Na₂HPO₄ is attributed to the slow formation of hydrogen bonds and the decreased 370 371 concentration of phosphate ions (Table 2) in the former compared to the latter medium. This aggregation mechanism provided the opportunity to create reinforced dual crosslinked 372 hydrogel networks in near-physiological conditions (as indicated by previous compression tests 373 in Figure 1 d&e) and an easy and stable method to build up HPO₄²⁻ nucleation sites in the 374 hydrogel for subsequent acellular biomineralisation. 375



Figure 3. Optical images of hydrogel C2-40 following either 3-week incubation in Na₂HPO₄ (a), PBS (b), and c-SBF
 (c), or 3-month incubation in PBS buffer (d and e). Deep aggregation of the network was indicated with red arrows.

376

381 **3.5. Mechanistic study of HPO**²⁻-induced physical crosslinking

The development of physical crosslinks between HPO₄²⁻ and cystamine-crosslinked HA was further supported by density functional theory (DFT) calculations. Three models of cystaminecrosslinked HA (C-HA) were optimised according to their energy minimum configuration. As presented in **Figure 4a**, the most stable structure was achieved in model C-HA3 (ΔE_{C-HA3} = -170.751 kcal/mol), whilst increased total interaction energies were measured with the other two models (ΔE_{C-HA1} = -162.075 kcal/mol, ΔE_{C-HA2} = -169.501 kcal/mol).



Figure 4. (a) DFT calculations of the hydrogen bond interaction between HPO₄²⁻ ions and cystamine-crosslinked hyaluronic acid (C-HA1, C-HA2 and C-HA3). In all models, oxygen (O) atoms were presented in red, nitrogen (N) in blue, sulfur (S) in yellow, carbon (C) in grey, hydrogen (H) in white and phosphorus (P) in pink. (b) Molecular structure of computed crosslinkers. (c) IR spectrum of Na₂HPO₄ (top), Na₂HPO₄-treated C2-40 network (middle) and E2-40 network (bottom) following 4-week treatment.

In the most stable model C-HA3, three atoms of oxygen (O) in the HPO_4^{2-} species engages in hydrogen bonds with the NH (1, 5) and OH (2, 3) groups of crosslinked HA, whilst the OH group in HPO_4^{2-} forms hydrogen bonds with the O atom of HA (4).

398 To investigate the influence of both the disulfide bridge and the number of carbon atoms in the crosslinking chain, the same binding sites as in C-HA were calculated in HA structure models 399 crosslinked with either 1,6-hexanediamine (6 carbon atoms), butane-1,4-diamine (4 carbon 400 atoms) or ethylenediamine (2 carbon atoms), and abbreviated as H-HA, B-HA, E-HA, 401 respectively. As presented in Figure 4b, the strongest interaction in the H-HA structure was 402 403 obtained in model H-HA3 with a ΔE_{H-HA3} = -162.149 kcal/mol (Figure S6 and Table S2), which was 8.602 kcal/mol lower than the one recorded in model C-HA3 (ΔE_{C-HA3} = -170.751 kcal/mol). 404 Although no direct binding contribution of the S-S bridge was observed, the optimised structure 405

and the reduced binding energy proved an indirect effect. In B-HA models, a ΔE_{B-HA3} of -167.491 406 407 kcal/mol was calculated in the most stable configuration, hinting at a lower interaction 408 compared to the model of 1,6-hexanediamine-crosslinked HA. Since butane-1,4-diamine is two carbon atoms shorter than 1,6-hexanediamine, the lower interaction measured in model B-HA3 409 with respect to H-HA3 suggests that the crosslinker length affects the development of HPO_4^{2-} -410 mediated physical crosslinks in the HA crosslinked chain. This observation is supported by the 411 energy calculations in model E-HA, describing HA chains crosslinked with ethylenediamine as 412 the shortest crosslinker of the three. Only one stabilised structure was obtained in this work, 413 with a final ΔE_{E-HA} of -155.330 kcal/mol. Nevertheless, the lack of stable configurations of E-HA 414 is against the development of HPO_4^{2-} -mediated hydrogen bonds in ethylenediamine-crosslinked 415 HA, thereby supporting the role of the crosslinker length in the development of phosphate ion-416 HA secondary structures. 417

Experimentally, a band corresponding to a P=O vibration was observed in the IR spectrum of 418 the Na₂HPO₄-treated networks. New peaks at 1230 cm⁻¹ and 1202 cm⁻¹ were displayed by both 419 C2-40 and E2-40 samples, which reflect the 1261 cm⁻¹ and 1185 cm⁻¹ peaks of Na₂HPO₄ (Figure 420 4c). C2-40 and E2-40 hydrogels were washed with deionised water for 24 hours to remove any 421 free Na₂HPO₄ residue and freeze-dried prior to IR measurement. The existence of a shifted peak 422 related to the P=O vibration provided strong evidence for hydrogen bond formation between 423 P=O and cystamine-crosslinked HA units. The most interesting phenomenon was the almost 424 disappearance of the original 1700 cm⁻¹ peak in the IR spectrum of the Na₂HPO₄-treated sample 425 C2-40 (Figure S7), which is attributed to the amide linkage of HA (position 5, Figure 4a) [32] and 426 which is still clearly visible in the IR spectrum of sample E2-40 following the same salt 427 treatment. The hydrogen bond between the HPO_4^{2-} ion and the nitrogen atom (N) of the amide 428 bond (position 5, Figure 4a) may shift this peak to 1640 cm⁻¹. This result strongly supports the 429 mechanism of multiple hydrogen bonds formed between HPO_4^{2-} ions and the cystamine-430 crosslinked HA chains. 431

As the most stable interaction was obtained when the phosphate-amide site binding occurred, in agreement with Barrett's work on hyaluronic acid solutions [19], we propose that minimising steric hindrance by adjusting the length of the crosslinker is critical to providing

proper access to HPO₄²⁻ ions and enabling coordination and physical crosslinking with amide bonds. Furthermore, the introduction of disulfide bridges in the HA network provided HAcrosslinked chains with increased flexibility and increased opportunities for developing secondary interactions with phosphate groups [36]. This potential intermolecular interaction may induce the rearrangement of the disulfide bonds and hydrophilic-hydrophobic sites so that detectable effects can be observed at the macroscale and influence the material properties as shown in our results.

442

443 **3.6. Cell adhesion study during HPO**²⁻ treatment

444 Following the results obtained in acellular conditions, an *in vitro* study was carried out with ATDC 5 chondrocytes. Chondrocytes were selected as non-mineralizing joint-resident cells, 445 aiming to investigate both the material-induced cell response and any cell culture-induced 446 effect on the material morphology. After 2-day cell culture in basal medium, some aggregated 447 HA network was already observed on the surface of freshly synthesised hydrogel C2-40 (Figure 448 S8), in line with the presence of phosphate groups in the cell culture medium (0.844 mM in 449 BM). The aggregation kinetics were accelerated with respect to previously discussed acellular 450 451 conditions, an observation which can be explained by considering the multiple ingredients in 452 cell culture medium and cell metabolism. At the cellular level, the fluorescently labelled live cells aligning along the aggregated structure are visible (Figure S8), whereby the weak 453 fluorescence is likely due to the quenching of the cell-labelling dye following cell growth. 454

After 1-week cell attachment and migration in basal medium, the conditional cell culture was carried out by replacing the medium with either fresh TM (1.884 mM phosphate) as testing group or BM (0.884 mM phosphate) as the control group. After 3 days of conditional cell culture and consequent calcein-AM staining, few fluorescent cells were observed via 3D confocal microscopy in either the BM or the TM group (**Figure 5**; higher resolution images are available in Figure S12 and Figure S13, Supporting Information).





Figure 5. Conditional culture of ATDC 5 cells. Cells adhesion study on the surface of C2-40 hydrogels in either basal
 medium (BM) or Na₂HPO₄ treated medium (TM) after 3 days (first and second row). Cells after 3-week conditional
 culture on C2-40 hydrogel surface (third row) and TCPs (bottom row) in either BM or TM group. Live labelling was
 presented in green and dead labelling was indicated in red. Scale bar: 100 μm.

To confirm this, C2-40 hydrogels without cells were set as a blank control, whereby only one 467 468 fluorescent dot with a maximum length of 10 µm was observed in the confocal image (Figure 469 S9). This observation is unlikely to be related to living cells and is mostly attributed to impurity or fluorescence from HPO_4^{2-} aggregation, as the cells observed in the hydrogels were 470 approximately 30 μ m in length and 10 μ m in width (Figure 5, first and second row; higher 471 resolution images are available in Figure S12, Supporting Information). When ATDC 5 cells were 472 independently seeded on the surface of each initial C2-40 hydrogel in both BM and TM, most of 473 the cells were found to adhere to the tissue culture plates (TCPs) rather than attach to the 474 hydrogel networks. This observation suggests that a tighter network may help to minimise cell 475 attachment and reduce the rate of degradation [37]. 476

477 Live&dead staining results proved that Na_2HPO_4 treatment was non-toxic during a 3-week 478 conditional cell culture period after comparing with BM groups, regardless of the hydrogel or

TCP surface (Figure 5, third and bottom row; higher resolution images are available in Figure 479 S13, Supporting Information). These results demonstrate that the HPO₄²⁻-induced hydrogel 480 aggregation process provides a regular channel for cell attachment and growth on the HA 481 482 hydrogel surface (Figure 5, TM group), unlike the freshly synthesised C2-40 hydrogel. The acellular fabrication of previously described salt-mediated microstructures (Figure 3) can be 483 proposed as the first stage of the hydrogel biomineralisation process. Here, the absence of cells 484 is key to minimizing the risk of cell aggregation on the hydrogel surface, which could otherwise 485 induce steric effects and delay HAp crystallisation. 486

487

488 **3.7. Characterisation of HAp growth within the HA-based hydrogel**

Both Na₂HPO₄ and non-Na₂HPO₄-treated C2-40 replicates were transferred into deionised 489 water for 24 hours to remove any free salt, and then further treated with 200 mL CaCl₂ (10 mM) 490 for 24 hours. The calcium concentration was chosen from a study on milk as one of the main 491 sources for calcium supplementation [35]. All the samples were flushed with deionised water 492 493 before the mineralization process, which was subsequently carried out in 1.0 L c-SBF at 37 °C. Remarkably, a homogeneous HAp phase was formed in the HA network C2-40 (Figure 6a, left), 494 495 with full mineralisation observed across the whole hydrogel structure (Figure S10). No visible mineral was observed in the control group obtained without HPO_4^{2-} treatment, ensuring that 496 the hydrogel surface remained transparent (Figure 6a, right). To further characterise the 497 mineralised structure of Na₂HPO₄-conditioned hydrogel, XRD diffraction was carried out (Figure 498 **6b**). A clear stacking structure corresponding to HAp was observed after burning the sample at 499 1000 °C for 30 minutes. 500



501

Figure 6. (a) Optical graphs captured at the end of the c-SBF incubation with both the HPO_4^{2-} -treated hydrogel C2-40 (left) and the corresponding HPO_4^{2-} -free hydrogel control C2-40 (right). (b) XRD spectra of the mineralised C2-40 after burning at 1000 °C (top, blue), the freeze-dried mineralized network (middle, black) and the initial freeze-dried C2-40 network (bottom, purple). (c-d) 3D reconstruction of the mineralised hydrogel C2-40: length-height direction (top) and height-width direction (bottom).

Some diffraction was recorded for the freeze-dried network at $2\theta = 27^{\circ}$, 33° and 35°, again 508 corresponding to the HAp phase [38], whilst no peak was observed in the initial C2-40 network. 509 510 In addition to XRD spectra and digital macrographs, μ CT was carried out as a non-damaging technique to visualise the 3D macrostructure of the mineralised C2-40 composite obtained 511 following 2-week incubation in c-SBF (Figure S11). The cross-sectional image clearly reveals the 512 decreasing HAp density from the top to the bottom side of the sample, in agreement with the 513 514 results obtained from the µCT 3D reconstructed models (Figure 6c&d), and in contrast to the 515 ion distribution surrounding the gel surface or limited formation of minerals [9][29]. This result demonstrates the high potential of the HAp-mineralized C2-40 hydrogel as a scaffold for hard 516 tissue repair, particularly as gradient hydrogels for tissue regeneration [39]. 517

518

520 4. Conclusions

The effect of the inclusion of a range of salts within cystamine and ethylenediamine-crosslinked 521 HA-based hydrogels was investigated to prepare dual crosslinked bioinspired bone-like 522 nanocomposites. Specific and strong hydrogen bond interactions acting as physical crosslinks 523 were first discovered between cystamine-crosslinked HA chains and HPO₄²⁻ groups, as indicated 524 by the decreased swelling and decreased compression at the break, IR spectroscopy and DFT 525 526 calculations. The introduction of phosphorus nuclei was key to enabling this interaction, which 527 was successfully leveraged to accomplish HAp growth across the entire hydrogel structure. Gradient HAp structures were obtained and visualised by μ CT 3D reconstruction after hydrogel 528 incubation in 1.0 L c-SBF for 2 weeks. A novel method to generate dual crosslinked and 529 mineralised structures is reported that is potentially significant for hard tissue repair. 530

531

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535

536 **Declaration of competing interests**

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- 538

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- 542

543 Data availability

All the data pertaining to this study is included in the main article and supplementary material.

545 The raw/processed data is available from the corresponding authors upon reasonable request.

546

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Supporting information

Hydrogen phosphate-mediated acellular biomineralisation within a dual crosslinked hyaluronic acid hydrogel

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Figure S1. TNBS calibration curve of ethylenediamine (a) and cystamine (b)



Figure S2. Relative mass of hydrogels (n=4) C2-40 (red) and E240 (blue) after 4-week immersion in either the Na2HPO4-supplemented solution (50 mM) or deionised water.

Treating conditions		C2-40 hydrogels		E2-40 hydrogels	
		Stress at break	Strain at break	Stress at break	Strain at break
		(kPa)	(%)	(kPa)	(%)
Original		425±20 (***)	90±1.0 (***)	211±18 (****)	77±0.8 (****)
1 day	(NH ₄) ₂ SO ₄	227±48 (*)	82±0.5 (***)	146±15 (***)	77±0.9 (****)
	Na_2SO_4	199±74 (-)	80±0.8 (**)	132±29 (**)	78±0.4 (****)
	Na_2HPO_4	167±18 (*)	77±0.3 (**)	156±42 (**)	82±1.7 (****)
	CH₃COONa	100±40 (-)	80±2.1 (**)	147±22 (***)	80±1.1 (****)
	NaCl	104±27 (-)	78±0.9 (**)	98±8 (****)	76±2.0 (****)
	PBS(LONZA)	236±36 (**)	81±1.4 (**)	124±26 (**)	78±0.8 (****)
	H ₂ O	76±39	66±3.0	14.5±1.8	40±1.9
	(NH ₄) ₂ SO ₄	486±48 (****)	84±0.7 (****)	297±28 (****)	78±0.3 (****)
4 weeks	Na_2SO_4	170±32 (**)	80±0.6 (****)	338±79 (***)	79±1.3 (****)
	Na ₂ HPO ₄	126±16 (*)	54±1.5 (****)	177±14 (****)	77±0.5 (****)
	CH₃COONa	230±29 (***)	79±1.1 (****)	203±16 (****)	76±0.6 (****)
	NaCl	358±32 (****)	81±0.5 (****)	347±22 (****)	78±0.8 (****)
	PBS(LONZA)	189±33 (**)	72±1.3 (***)	206±27 (****)	76±1.1 (****)
	H ₂ O	103±3	67±1.0	27±7	46±2.4 (****)

Table S1. Compression stress and strain values at break of C2-40 and E2-40 hydrogels after synthesis('Original') and different salt treatments.



Figure S3. Compression stress at break of hydrogel C2-40 measured after synthesis ('Original') and after 1-day (a) and 4-week (b) incubation in different aqueous solutions. All the statistical analysis is presented with respect to H_2O group and labelled as *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001. All the data are presented as Mean±SD.



Figure S4. Compression measurements of hydrogel E2-40 after synthesis ('Original') and following 1-day (a, b) and 4-week (c, d) incubation in different aqueous solutions. All the statistical analysis is presented with respect to the H₂O group and labelled as **p < 0.01, ***p < 0.001, ****p < 0.0001. All the data are presented as Mean±SD.

Figure S5. SEM images of freeze-dried E2-40 networks following 4-week incubation in $(NH_4)_2SO_4$ (a), Na_2SO_4 (b), Na_2HPO_4 (c), NaAc (d), NaCl (e), PBS (f) and deionised water (g). Scale bar: 500 μ m.

Figure S6. DFT calculations of the hydrogen bond interaction between HPO₄²⁻ and hyaluronic acid crosslinked with either 1,6-Hexanediamine (H-HA), 1,4-Butanediamine (B-HA) or Ethylenediamine (E-HA). In all models, oxygen (O) atoms were presented in red, nitrogen (N) in blue, sulfur (S) in yellow, carbon (C) in grey, hydrogen (H) in white and phosphorus (P) in pink. The optimised models with lowest interaction energy are presented in a red box.

Model Name	ΔE (kcal/mol)	
C-HA1	-162.075	
C-HA2	-169.501	
*С-НАЗ	-170.751	
H-HA1	-152.556	
H-HA2	-160.341	
*H-HA3	-162.149	
B-HA1	-150.107	
B-HA2	-146.913	
*B-HA3	-167.491	
E-HA	-155.330	

 Table S2. Optimised computing results of interaction energy in selected HA models.

Figure S7. IR spectrum of freshly-synthesized C2-40 and E2-40 networks.

Figure S8. Cell Tracker Green labelled ATDC 5 cells on C2-40 surface after 2-day culture in basal medium. Optical image (left) and fluorescent image (right).

Figure S9. Laser confocal image of blank hydrogel control C2-40 in TM group following 3-day conditional culture.

Figure S10. Optical images of the wet mineralised sample C2-40 following HPO_4^{2-} treatment (side view, left) and C2-40 hydrogel control (top surface, right).

Figure S11. Macro-pattern study of mineralised sample C2-40 from μ CT scan, combining 3-dimensional image (a) and cross section of each axis (b).

Fig. S12 First and second row of **Figure 5** (in main manuscript) in higher resolution. Conditional culture of ATDC 5 cell growth. Cells adhesion study on the surface of C2-40 hydrogels in either basal medium (BM) or Na₂HPO₄-treated medium (TM) after 3 days. Scale bar: 100 μ m.

Fig. S13 Third and bottom row of **Figure 5** (in main manuscript) in higher resolution. Cells after 3-week conditional culture on C2-40 hydrogel surface (third row) and TCPs (bottom row) in either BM or TM group. Live labelling was presented in green and dead labelling was indicated in red. Scale bar: 100 μ m.