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Abstract:	<p>In biomaterials for bone tissue regeneration, ionic substitutions are a promising strategy to enhance the biological performance of calcium phosphates and composite materials. However, systematic studies have not been found on multi-substituted organic/inorganic scaffolds. In this work, highly porous composite scaffolds based on calcium phosphates substituted with Sr²⁺, Mg²⁺, Zn²⁺ and SeO₃²⁻ ions, and biodegradable polymer chitosan have been prepared by freeze-gelation technique. Scaffolds were characterized by Fourier transform infrared spectroscopy, X-ray diffraction, scanning electron microscopy, pore size distribution and porosity measurements. In vitro enzymatic degradation of scaffolds was characterized by swelling and dry weight remaining ratio measurements, molecular weight and microstructure determination. Enhanced osteogenic potential of human mesenchymal stem cells (hMSC) seeded on scaffolds has been determined by histological, immunohistochemical and RT-qPCR analysis of cultured cells in static and dynamic (U-CUP bioreactor) conditions. This work demonstrates the influence and importance of ionic substitutions on the osteogenic differentiation of hMSCs.</p>
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1 **Osteogenic differentiation of human mesenchymal stem cells on substituted**
2 **calcium phosphate/chitosan composite scaffold**

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1 **ABSTRACT**

2 In biomaterials for bone tissue regeneration, ionic substitutions are a promising strategy to
3 enhance the biological performance of calcium phosphates and composite materials.
4 However, systematic studies have not been found on multi-substituted organic/inorganic
5 scaffolds. In this work, highly porous composite scaffolds based on calcium phosphates
6 substituted with Sr^{2+} , Mg^{2+} , Zn^{2+} and SeO_3^{2-} ions, and biodegradable polymer chitosan have
7 been prepared by freeze-gelation technique. Scaffolds were characterized by Fourier
8 transform infrared spectroscopy, X-ray diffraction, scanning electron microscopy, pore size
9 distribution and porosity measurements. *In vitro* enzymatic degradation of scaffolds was
10 characterized by swelling and dry weight remaining ratio measurements, molecular weight
11 and microstructure determination. Enhanced osteogenic potential of human mesenchymal
12 stem cells (hMSC) seeded on scaffolds has been determined by histological,
13 immunohistochemical and RT-qPCR analysis of cultured cells in static and dynamic (U-CUP
14 bioreactor) conditions. This work demonstrates the influence and importance of ionic
15 substitutions on the osteogenic differentiation of hMSCs.

16 **Keywords:** chitosan, hydroxyapatite, osteogenic differentiation, perfusion-bioreactor,
17 scaffolds, ionic substitution

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1 **1.Introduction**

2 As life expectancy increases, so does the incidence of skeletal diseases resulting in
3 requirements for new and more adequate regenerative materials for bone tissue (Aubin, 2008).
4 Bone tissue is mainly composed of the calcium-deficient carbonated hydroxyapatite,
5 representing 65-70% of the matrix and an organic phase composed of collagen (mainly of
6 type I), glycoproteins, proteoglycans and sialoprotein, which comprises the remaining 25-30
7 % of the total matrix (Cardonnier, Sohier, Rosset, & Layrolle, 2011). From the biomimetic
8 scaffold design perspective, achievement of molecular, structural and biological
9 compatibility, while resembling as closely as possible the natural bone tissue should be
10 achieved for a positive regenerative response. The integration of multiple stimuli physical
11 factors (topography, porosity, pore size, stiffness, phase composition) and biochemical cues
12 (grow factors, trace elements, genes or proteins) is the way forward to the next generation of
13 biomimetic bone scaffolds. (Fernandez-Yague et al., 2015; Boanini, Gazzano, & Bigi, 2010)
14 Biomimetic scaffolds should not only be biocompatible, but also promote proliferation and
15 differentiation of progenitor cells and provide structural support until the newly-formed tissue
16 has sufficient stiffness to stimulate the progression of the regeneration (Minardi et al., 2015).

17 Naturally derived polymers such as collagen, glycoaminoglycans, gelatin, chitosan, silk,
18 fibrin, and elastin, have been widely used in a variety of tissue engineering applications. As
19 these polymers show similarity with the extracellular matrix found in human physiology, they
20 demonstrate appropriate biocompatibility for *in vivo* applications, but more importantly
21 provide a range of ligands and peptides that facilitate cell-material communication to induce
22 osteogenesis and in some cases reduce immunogenicity. (Fernandez-Yague et al., 2015;
23 Roseti, 2017; Deb, Deoghare, Borah, Barua, & Das Lala, 2018) Naturally derived polymer
24 chitosan is widely used in bone tissue engineering due to minimal inflammatory body
25 reaction, biocompatibility, biodegradability, and ability to provide a suitable environment for

1 cell growth. The $-NH_2$ groups, present in the chitosan polymer chain, are able to induce
2 osteoblast differentiation. (Cardonnier et al., 2011; Muzzarelli, 2011) However, chitosan
3 scaffolds are somewhat limited in inducing osteogenic differentiation of stem cells and usually
4 need the combination with calcium phosphate (CaP) particles, typically hydroxyapatite (HAp),
5 to mimic the organic/inorganic nature of the bone (Venkatesan & Kim, 2010).

6 Recent trends have been focused on using pharmacologicals and biologics to improve the
7 osteogenic properties of synthetic organic and inorganic biomaterials. However, using growth
8 factors is often related to ectopic or unwanted bone formation, and new trends look for
9 biomimetic approaches based on trace elements naturally present in the mineral phase of bone
10 (Bose, Fielding, Tarafder, Bandyopadhyay, 2013). The high stability and flexibility of
11 hydroxyapatite (HAp) structure allow a great variety of cationic and anionic substitutions.
12 Ions, such as Sr^{2+} , CO_3^{2-} , Zn^{2+} , Mg^{2+} , Na^+ , etc., which are present in biological apatites
13 (Boanini et al., 2010) have been found to play a vital role in stem cells proliferation and
14 differentiation, formation, growth and mineralization of extracellular matrix (ECM) (Bose et
15 al., 2013).

16 Considering all mentioned above, biomimetic multi-substituted CaPs were used as inorganic
17 phases within biopolymer chitosan to obtain highly porous scaffolds. Synergic effect of
18 substituted ions in CaPs and biomimetic composite scaffold can have significant effect on
19 differentiation of stem cells and bone regeneration. In our previous studies (Ressler et al.,
20 2020a; Ressler, Antunović, Cvetnić, Ivanković & Ivanković, 2021; Ressler, 2020) individual
21 and multiple ion substituted CaPs with various levels of Sr^{2+} , Zn^{2+} , Mg^{2+} and SeO_3^{2-} ion
22 substitution have been prepared by wet precipitation method. Also, previous studies of our
23 research group have proved non-cytotoxic and osteogenic properties of chitosan-unsubstituted
24 hydroxyapatite scaffolds (Rogina et al., 2017, Ressler et al., 2018). Composite scaffolds with
25 hydroxyapatite fraction of 30% (w/w) showed the strongest osteoinduction of mouse MC3T3-

1 E1 preosteoblasts and human mesenchymal stem cells (Rogina et al., 2017). The aim of this
2 work was to study the effect of multiple ionic substitutions of Sr^{2+} , Zn^{2+} , Mg^{2+} and SeO_3^{2-} in
3 CaPs on the osteogenic potential of chitosan-CaP scaffolds. The highly porous composite
4 scaffolds, with 30 wt% of multi-substituted CaPs, were prepared by the freeze-gelation
5 method and characterized. An extensive biological characterization in static and dynamic
6 conditions has been performed, using human mesenchymal stem cells (hMSC). To the best of
7 our knowledge, there are no data in literature regarding the effect of the studied multiple ionic
8 substitutions in CaPs on the osteogenic potential of chitosan-CaP scaffolds.

9 **2. Materials and methods**

10 *2.1. Preparation of composite scaffolds*

11 The appropriate amount of chitosan was added to 0.40 wt% acetic acid solution to obtain 1.2
12 wt% chitosan solution at ambient temperature. The CaPs substituted with various trace
13 elements (Sr^{2+} , Mg^{2+} , Zn^{2+} and SeO_3^{2-}) and precipitated from biogenic source (cuttlefish
14 bone), were previously prepared in our studies (Ressler 2020, Ressler et al., 2021a, Ressler et
15 al., 2021) and used to obtain composite scaffolds. Nominal composition and labels of
16 prepared CaPs are given in Table 1. The appropriate amounts of each CaP, alone or in a
17 mixture, were added to obtain 30 wt% of CaP in composite scaffold, based on the previous
18 study (Rogina, Rico, Gallego Ferrer, Ivanković, Ivanković, 2014). The mass fraction of each
19 CaP in the CaPs mixtures was the same (25 wt.%). The composite scaffolds are labeled as
20 shown in Table 2. The chitosan/CaP suspensions were cooled to 4 °C, set in moulds, frozen,
21 and kept at -30 °C for 8 h. Further, frozen samples were immersed into the neutralisation
22 medium of 1 M NaOH/ethanol at -30 °C for 24 h to induce gelation of chitosan. The samples
23 were rinsed in ethanol (96 wt%) at -30 °C for 24 h, washed with distilled water, frozen, and
24 lyophilized.

1 **Table 1.** Nominal composition and labels of prepared CaP samples substituted with Sr²⁺,
 2 Zn²⁺, Mg²⁺ and SeO₃²⁻ ions (mol %) used as inorganic phase in composite scaffolds.

	Sample	Sr/(Ca+Sr) (mol %)	Zn/(Ca+Zn) (mol %)	Mg/(Ca+Mg) (mol %)	Se/(P+Se) (mol %)
Synthetic source	CaP	0	0	0	0
	CaP_0	0	0	0	0
	1_Sr_CaP	1	0	0	0
	5_Sr_CaP	5	0	0	0
	1_Zn_CaP	0	1	0	0
Biomimetic source (cuttlefish bone)	5_Zn_CaP	0	5	0	0
	1_Mg_CaP	0	0	1	0
	5_Mg_CaP	0	0	5	0
	1_Se_CaP	0	0	0	1
	5_Se_CaP	0	0	0	5

3

4 **Table 2.** Labels and type of CaPs in the prepared composite scaffolds.

Precursor for CaP	CaP	Composite sample	Wt.% of CaP in composite	Wt.% of CaP in the CaP mixture	
Synthetic	CaP	SC	30	100	
	CaP_0	SC_0		100	
	1_Sr_CaP	SC_1Sr		100	
	5_Sr_CaP	SC_5Sr		100	
	1_Zn_CaP	SC_1Zn		100	
	5_Zn_CaP	SC_5Zn		100	
	5_Zn_CaP	SC_1Mg		100	
	1_Mg_CaP	SC_5Mg		100	
	1_Se_CaP	SC_1Se		100	
	5_Se_CaP	SC_5Se		100	
	Biogenic			30	100
		1_Sr_CaP			25
		1_Zn_CaP			25
1_Mg_CaP		SC_1MIX		25	
1_Se_CaP				25	
5_Sr_CaP				25	
5_Zn_CaP				25	
5_Mg_CaP		SC_5MIX		25	
5_Se_CaP			25		

5

6 2.2. Scaffolds characterization

7 The Fourier transform infrared spectra (FTIR) of composite scaffolds were recorded by
 8 attenuated total reflectance (ATR) spectrometer for solids with a diamond crystal (Bruker
 9 Vertex 70) at 20 °C over the spectral range of 4,000 – 400 cm⁻¹, with 32 scans and 4 cm⁻¹ of
 10 resolution.

1 Mineralogical composition of composite scaffolds was determined by X-ray diffraction
2 analysis (XRD) using a Shimadzu XRD-6000 diffractometer with Cu Ka radiation operated at
3 40 kV and 30 mA, in the range of $3^\circ < 2\theta < 60^\circ$ at a scan speed of $0.2^\circ/2s$. Identification of
4 crystal phases was done by International Centre for Diffraction Data (ICDD) card catalogue.
5 The morphology of composite scaffolds was imaged by the scanning electron microscope
6 TESCAN Vega3SEM Easyprobe (SEM) at electron beam energy of 10 keV. Previously to
7 imaging, samples were sputter coated with gold and palladium for 120 s. Obtained SEM
8 images and ImageJ software were used to determine the diameter of 200 pores of obtained
9 composite scaffolds. The results of pore size distribution are shown as pore density (%) of
10 each pore range in relation to the total number of measured pores.

11 The porosity of the scaffolds was determined by Archimedes' principle, immersing each
12 composite scaffold in ethanol at room temperature ($T = 22.2 \pm 0.3^\circ\text{C}$) as previously described
13 in our research (Ressler et al., 2020b, Morales-Román et al., 2019). Scaffolds porosity (%) is
14 defined as the volume fraction of pores within the scaffold; so it was calculated as the volume
15 of pores (V_p) divided by the total volume of the scaffold (V_{sc}) composed of pores and
16 composite (V_c), Eq. (1):

$$17 \text{ Porosity (\%)} = \frac{V_p}{V_{sc}} \times 100 = \frac{V_p}{V_p + V_c} \times 100$$

18 (1)

19 The previously prepared scaffolds were cut with biopsy puncher into cylindrical shape pieces
20 of 6 mm diameter (\varnothing) with uniform thickness (h) of ~ 1 mm. The dry samples
21 ($n = 5$) were initially weighted (m_d). The volume of the pores was calculated by filling them
22 with ethanol ($\rho = 0.789$ g/mL). After immersion in ethanol under vacuum atmosphere, excess
23 liquid was removed and samples were weighted again (m_e). The density of cylindrically-

1 shaped scaffold is calculated according to the Eq. (2), the volume of the scaffold (V_{SC})
2 according to Eq. (3) and the pore volume was calculated according to Eq. (4):

$$3 \quad \rho_{SC} = \frac{m_d}{\pi \times (\emptyset/2)^2 \times h} \quad (2)$$

$$4 \quad V_{SC} = \frac{m_d}{\rho_{SC}} \quad (3)$$

$$5 \quad V_p = \frac{m_e - m_d}{\rho_e} \quad (4)$$

6 2.3. *In vitro* enzymatic degradation

7 The degradation behaviour of obtained composite scaffolds (SC, SC_0, SC_1MIX,
8 SC_5MIX) was studied at two different concentrations of lysozyme (1.5 $\mu\text{g}/\text{mL}$,
9 corresponding to an activity of 164 U/mL, and 150 $\mu\text{g}/\text{mL}$, corresponding to an activity of 16
10 400 U/mL) under static physiological conditions in phosphate buffer saline solution (PBS).
11 The degradation behaviour of composite scaffolds in the PBS solution without lysozyme was
12 used as a control. Composite scaffolds ($n = 5$, $\emptyset = 6$ mm, $h \sim 1$ mm) were incubated in 5 mL
13 of PBS containing lysozyme at 37 °C during 28 days. Freshly prepared degradation medium
14 was changed every third day to maintain the activity of lysozyme and to mimic physiological
15 conditions *in vivo* (Porstmann et al., 1989). At defined time points, the degradation medium
16 was removed in order to determine the mass of swollen samples (m_s). Then, samples were
17 lyophilised (m_d) and the swelling ratio was calculated according to Eq. (5):

$$19 \quad \text{Swelling ratio}(\%) = \frac{m_s - m_d}{m_d} \times 100 \quad (5)$$

20

21 The degradation of scaffolds was determined as the ratio of remaining hydrogel weight (m_d)

1 and initial weight of the sample (m_{d0}) before enzymatic degradation Eq. (6).

2

$$3 \quad \text{Dry weight remaining ratio}(\%) = \frac{m_d}{m_{d0}} \times 100 \quad (6)$$

4

5 The influence of degradation medium on scaffolds' microstructure was analysed by SEM
6 after 28 days of degradation. Dried degraded samples were coated with a plasma of gold and
7 palladium for 120 s. The microscopic imaging was carried out by the electron microscope
8 TESCAN Vega3SEM Easyprobe at electron beam energy of 15 keV.

9

10 *2.4. Gel permeation chromatography*

11 The molecular weight distribution of chitosan before and after enzymatic degradation was
12 analysed by Gel Permeation Chromatography (GPC), at 35 °C, with a Waters Breeze GPC
13 system and a 1525 Binary HPLC pump (Waters Corporation, Milford, MA) equipped with a
14 2414 refractive index detector and four serial Ultrahydrogel columns of water (7.8 mm ID X
15 30 cm) connected in series (1,000, 500, 150 and 200) ([Sanmartín-Masiá, Poveda-Reyes and](#)
16 [Gallego Ferrer, 2017](#)). The degraded composite samples were dissolved in acetic buffer
17 (CH_3COOH 0.5 M/ CH_3COONa 0.2 M, pH = 4.5) at a concentration of 1 mg/mL. This buffer
18 was also used as the eluent phase at a flow rate of 0.5 mL/min ([Gámiz-González, 2017](#)). A
19 volume of 2 mL of sample solution was prepared and left dissolve at room temperature in
20 glass vials (covered with aluminum foils to protect the polymer from UV light). They were
21 kept under agitation for 72 h to properly dissolve. Then, the samples were filtered and injected
22 in the equipment by using 100 μL of sample per injection. A minimum of 3 replicates per
23 sample were measured. The calibration curve was prepared by using monodisperse
24 polyethylene glycol (PEG) standards of known molar mass at peak, supplied by Waters.

1 Mark-Houwink-Sakurada equation and Mark-Houwink parameters for PEG and chitosan were
2 used to calculate the molecular weight of chitosan from the value obtained in the calibration
3 curve.

4 *2.5. Biological evaluation*

5 *2.5.1. Human Mesenchymal Stem Cells Isolation and Expansion*

6 The bone marrow sample was collected during the surgery at the University Hospital of
7 Traumatology in Zagreb (Croatia) with the patient's consent and approval of the Ethics
8 Committee. The hMSCs were isolated using previously described method by Matić and
9 associates (Matić et al., 2016). Briefly, bone marrow aspirates were added to Dulbecco's
10 modified Eagle medium (DMEM) – low glucose containing 10% fetal bovine serum (FBS,
11 Gibco), 100 U/mL penicillin and 100 g/mL streptomycin (Lonza). The suspension was
12 centrifuged and pelleted cells were washed twice in PBS (Gibco). Resuspended cells were
13 strained through a cell strainer (100 µm, BD Biosciences) to remove mineral bone residuals
14 and centrifuged again. Cells were plated in 100 mm Petri dishes (Sarstedt) at a density of $1 \cdot$
15 10^8 in proliferation medium DMEM-low glucose, supplemented with 10% FBS, 100 U/mL
16 penicillin and 100 µg/mL streptomycin and 10 ng/mL human fibroblast growth factor 2
17 (FGF2, Gibco), and kept in a humidified incubator at 37 °C with a 5% CO₂. After 24 h the
18 non-adherent cells were removed, and the attached cells were grown. After 80% confluence,
19 hMSCs were detached with 0.25% trypsin/EDTA (Sigma-Aldrich) and then subcultured for
20 expansion (passage 1–5). The proliferation medium was changed every 2–3 days.

21 *2.5.2. Static 3D cell culture of Human Mesenchymal Stem Cells*

22 Composite scaffolds (SC, SC_0, SC_1MIX and SC_5MIX) were cut into cylindrical pieces of
23 6 mm diameter and ~1 mm height, sterilised in 96 % ethanol for 24 h. After sterilisation,
24 scaffolds were washed 3 times with PBS (Gibco – Thermo Fisher Scientific) and left in
25 proliferation medium DMEM-low glucose, supplemented with 10% FBS, 100 U/mL

1 penicillin and 100 $\mu\text{g}/\text{mL}$ streptomycin and 10 ng/mL human fibroblast growth factor 2
2 (FGF2, Gibco) for 24 h at 4 $^{\circ}\text{C}$. The following day, scaffolds were transported into
3 polystyrene 96-well plates with a hydrophobic surface (Corning – Sigma Aldrich). The
4 hMSCs were seeded on each scaffold in a concentration $2 \cdot 10^5$ cells/200 μL of medium per
5 well. The cell suspension was added on each scaffold and incubated for 30 min in the
6 incubator to allow cell attachment and migration inside the scaffold. Following the incubation
7 period, the medium was added to a final volume of 200 μL per well. Each experiment was
8 performed in triplicate and blanks were included as well. The cells were kept in a 5 % CO_2
9 humidified atmosphere at 37 $^{\circ}\text{C}$ for 14 and 21 days, respectively. After 14 and 21 days of cell
10 culture, the scaffolds were put in TRIzol and kept at -80°C until analyzed.

11 *2.5.3. Dynamic 3D cell culture of Human Mesenchymal Stem Cells in Perfusion Bioreactor*

12 A perfusion bioreactor (U-CUP Cellec Biotek) was used to better simulate *in vivo* conditions
13 by *in vitro* 3D culture of bone promoting scaffolds. Two samples of each scaffold (SC and
14 SC_5MIX), with a diameter of 8 mm and a height of 1 mm, were inserted into the bioreactor
15 for cell seeding with a cell suspension of $2 \cdot 10^6$ cells/scaffold in a total volume of 10 mL of
16 proliferation medium per bioreactor. The closed bioreactor chamber was placed in the
17 incubator at 37 $^{\circ}\text{C}$ and 5% CO_2 for 48 h and the flow rate for the perfusion of cells was settled
18 at 1.7 mL/min. Following the 48 h-period, the proliferation medium was replaced by
19 osteogenic induction medium containing: Minimum Essential Medium-Alpha Eagle (α -MEM,
20 Lonza), 10% FBS, 1% penicillin/streptomycin, 50 $\mu\text{g}/\text{mL}$ ascorbic acid (Sigma-Aldrich), 4
21 mmol/L β -glycerophosphate (Sigma-Aldrich) and 1×10^{-7} mol/L dexamethasone (Sigma-
22 Aldrich). The cell culture flow rate (perfusion speed) was set at 0.6 mL/min during the culture
23 and the bioreactor was placed in the incubator at 37 $^{\circ}\text{C}$ and 5% CO_2 . The osteogenic medium
24 was exchanged every 2–3 days according to the manufacturer’s instructions (Cellec Biotech).
25 After 21 days of cell culture, scaffolds were removed from the bioreactor and analyzed.

1 2.5.4 Gene expression analysis by qRT-PCR

2 The expression of genes related to osteogenesis, including alkaline phosphatase (ALP), bone
3 sialoprotein (BSP) and dentin matrix protein 1 (DMP1) were analyzed by quantitative reverse
4 transcription polymerase chain reaction (qRT-PCR). The qRT-PCR was performed to
5 evaluate the expression levels of genes of cells cultured on scaffolds (SC, SC_0, SC_1MIX,
6 SC_5MIX) after 14 and 21 days. The primer sequences used for PCR amplification are listed
7 in Table 3.

8 *Table 3.* Human primer sequences used for determination of gene expression levels by reverse
9 transcription-polymerase chain reaction analysis.

Primers	5'-	Sequences	Annealing temperature (°C)
ALP	forward	CTGTTTACATTTGGATAC	57.4
	reverse	ATGGAGACATTCTCGTTC	61.6
BSP	forward	GGAGACTTCAAATGAAGGAG	57.9
	reverse	CAGAAAGTGTGGTATTCTCAG	56.4
DMP1	forward	CAACTATGAAGATCAGCATCC	58.8
	reverse	CTCCATTCTTCAGAATCCTC	59.3

10

11 Total cellular RNA was isolated using TRIzol reagent (Invitrogen Life Technologies),
12 according to the manufacturer's instructions. Briefly, samples were washed with cold PBS
13 and homogenized in 1 mL of TRIzol (Thermo Fisher) using a mixer mill (Retsch) for 3 min at
14 30 Hz, followed by 200 μ L of chloroform addition and centrifugation for 15 min at 12,000 \times
15 G and 4 °C. The supernatant was separated, isopropanol and 0.5 μ L of glycogen were added,
16 after which the RNA was precipitated by centrifugation for 10 min at 15,000 \times G and 4 ° C.
17 The supernatant was separated and 1 mL of 75% ethanol was added to the RNA pellet, and
18 the samples were centrifuged for 5 min at 15,000 \times G and 4 ° C. The supernatant was
19 separated and the RNA pellet was air dried for ~10 min and dissolved in 30 μ L of water

1 treated with diethyl pyrocarbonate (DEPC). RNA concentration and purity were determined
2 using the Nanodrop 2000 (Thermo Scientific, USA). To remove residual genomic DNA from
3 isolated RNA, the prepared RNA suspension was treated with deoxyribonuclease I (Dnase I,
4 Invitrogen Life Technologies). Enzymatic removal of residual DNA from the sample was
5 performed using a commercial purification kit (Thermo Scientific, USA) according to the
6 manufacturer's protocol. A mixture for DNA degradation in a total volume of 20 μ L was
7 prepared. The reaction mixture contained Dnase, 10 \times buffer, DEPC-treated water and a
8 sample of isolated RNA of known concentration. The final mass of RNA in the reaction was
9 0.5 μ g. The reaction mixture was homogenized and DNA digestion was performed by
10 incubating the mixture for 60 minutes at 37 $^{\circ}$ C. Then 1.5 μ L of ethylenediaminetetraacetic
11 acid (EDTA) solution was added to the samples, and Dnase I was inactivated by incubating
12 the samples for 10 minutes at 65 $^{\circ}$ C. Dnase I treated RNA was inversely transcribed into
13 complementary DNA (cDNA) using a commercial High-Capacity cDNA Reverse
14 Transcription Kit (Applied Biosystems) according to the manufacturer's protocol. Then 1.5
15 mL of the reaction mixture was prepared and mixed with each sample (10 μ L RNA). The
16 resulting mixture was homogenized and incubated according to the protocol. Samples were
17 diluted with 5 μ L of ultrapure water and stored at -20 $^{\circ}$ C for further analysis.

18 Relative gene expression was determined by RT-qPCR using the 7500 Fast Real-Time PCR
19 System (Applied Biosystems), commercially available primers (Sigma-Aldrich) and Power
20 SYBR Green Mastermix (Applied Biosystems) were used. Each reaction consisted of a
21 duplicate in 96-well plates (ABI PRISM Optical 96-Well Plate, Applied Biosystems). The
22 PCR reaction was conducted under the following conditions: 10 min at 95 $^{\circ}$ C for 1 cycle, 15 s
23 at 95 $^{\circ}$ C, and 1 min at 58 $^{\circ}$ C for 40 cycles. The expression levels of osteogenic genes were
24 normalized to *18SRNA* as a housekeeping gene and calculated using the $\Delta\Delta$ Ct method.

25 *2.5.5. Histological Analysis*

1 After removing from static and dynamic cell culture, cultured samples after 14 and 21 days
2 were rinsed in PBS, fixed in 4% paraformaldehyde, embedded in paraffin and cross-sectioned
3 (5 μm thick) and stained with hematoxyline-eosin (H&E). All sections were deparaffinised
4 using xylene, and rehydrated through a series of ethanol washes prior to staining.

5 Detection of phosphate deposits on composite scaffolds was determined using von Kossa
6 staining. Samples were washed with distilled water and incubated in a 5% AgNO_3 solution
7 (Sigma-Aldrich) under UVB light (UVB Crosslinker, Cleaver Scientific) for 3 min. Further,
8 samples were washed with distilled water and 2% $\text{Na}_2\text{S}_2\text{O}_3$ (Sigma-Aldrich) to remove
9 unreacted AgNO_3 . All slides were dehydrated and mounted with a resinous medium
10 (Biognost). Slides were observed under a microscope Olympus BX51.

11 *2.5.6. Immunohistochemical detection of collagen type I*

12 Collagen type I (COLL I) was detected using EnVision Detection Systems Peroxidase/DAB,
13 Rabbit/Mouse (Dako/Agilent, Santa Clara, CA, USA) and anti-collagen I (Abcam,
14 Cambridge, UK), as previously described with slight modification (Rogina et al. 2017).
15 Briefly, sections were deparaffinised and rehydrated. Nonspecific binding sites were blocked
16 with 10% goat serum (Dako) in PBS for 60 min at room temperature. Sections were then
17 incubated with primary antibody (anti-collagen I, Abcam, Cambridge, UK), diluted 1:400
18 with 1% goat serum in PBS, overnight at 4 °C. After washing, the signal was detected with
19 EnVision Detection Systems Peroxidase/DAB, Rabbit/Mouse (Dako), according to the
20 manufacturer instructions. Hematoxylin was used as a counterstain. Human bone was used as
21 a positive control. Slides were observed under a microscope Olympus BX51.

22 *2.6. Statistical Analysis*

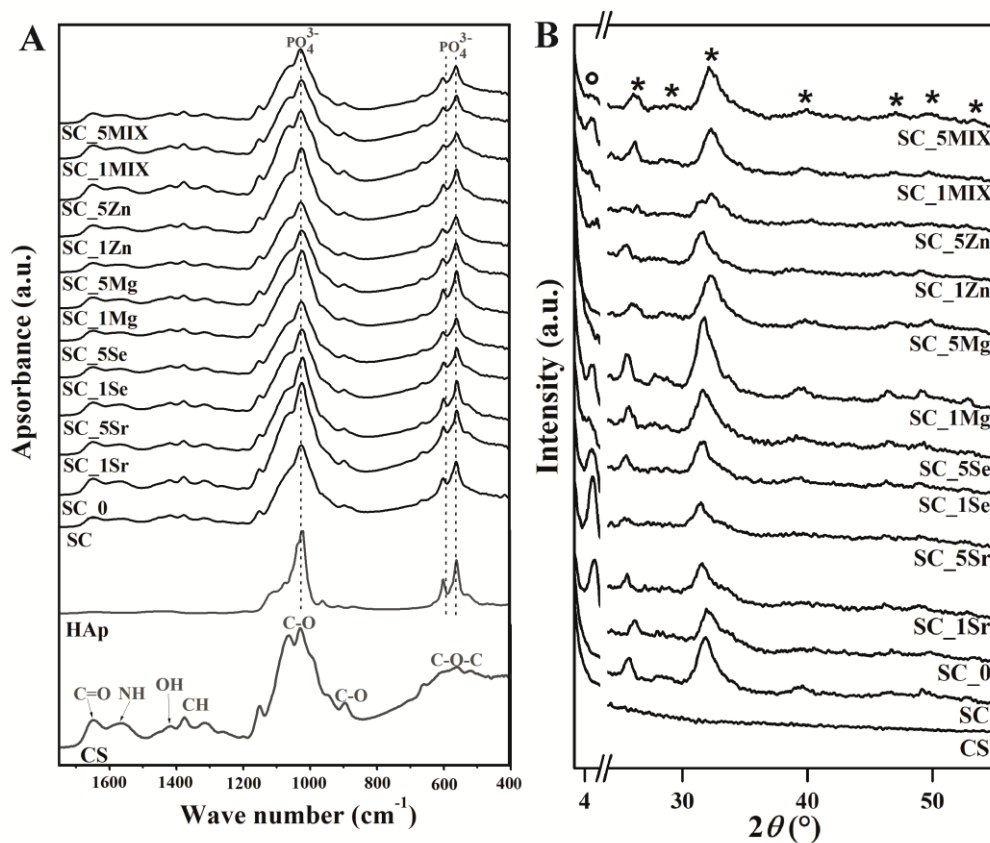
23 The swelling and dry weight remaining ratio measurements were performed in quintuplicate
24 ($n = 5$), and RT-qPCR experiments were performed in duplicate ($n = 2$). All data were
25 expressed as mean \pm standard deviation. Statistical analysis was performed using a one-way

1 ANOVA test followed by a post-hoc test to evaluate the statistical significance between
2 groups. A value of $p < 0.05$ was considered statistically significant.

3 3. Results

4 3.1. The composition identification and morphology of scaffolds

5 The characterization of prepared systems was performed by FTIR spectroscopy (Fig. 1A) and
6 XRD mineralogical analysis (Fig. 1B). The FTIR spectra for all prepared composite scaffolds
7 with different trace elements content, pure chitosan scaffold (CS) and hydroxyapatite powder
8 (HAp) have been shown in the range $400\text{--}1750\text{ cm}^{-1}$. At the wave numbers $>1550\text{ cm}^{-1}$
9 significant bands for were not detected.



10

11 *Figure 1.* FTIR spectra (A) and XRD pattern (B) of prepared composite scaffolds with
12 different substitution elements in CaP structure. Chitosan (CS) scaffold and pure

1 hydroxyapatite (HAp) were used as a control. Characteristic HAp (ICDD 9-432) diffraction
2 maxima are depicted as (*) and OCP (ICDD 11-0293) as (°).

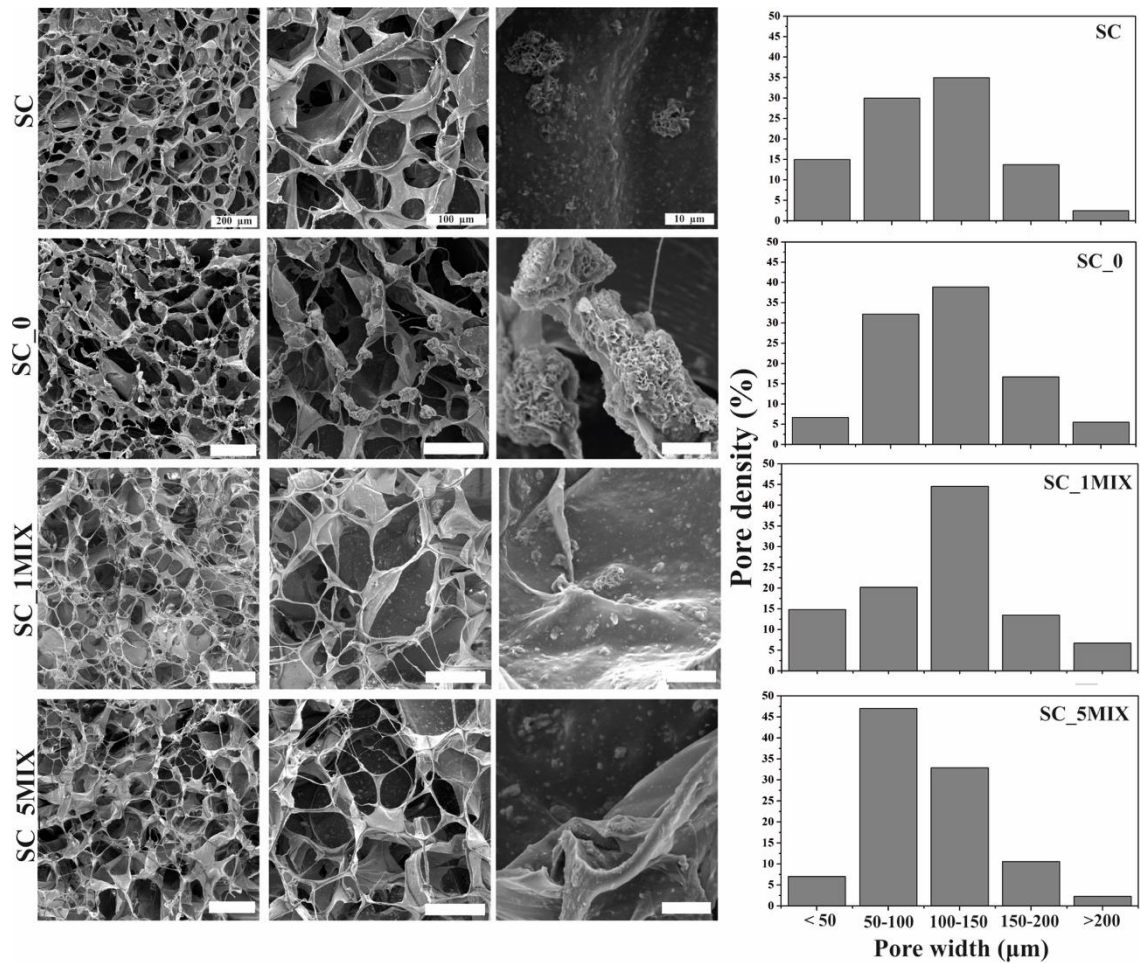
3 Absorption bands characteristic for chitosan are visible in all prepared scaffolds: at
4 1653 cm^{-1} corresponding to amid I (carbonyl band of amid), at 1567 cm^{-1} attributed to amid II
5 (amino band of amid), 1422 cm^{-1} and 1324 cm^{-1} that correspond to the vibrations of OH and
6 CH in the ring, 1378 cm^{-1} to CH_3 in amide group. Along with characteristic bands for the
7 chitosan, typical bands for PO_4^{3-} groups in HAp structure were found at 565 cm^{-1} and 600
8 cm^{-1} that can be assigned to asymmetric bending vibrations of O–P–O, and at 1028 cm^{-1}
9 attributed to the asymmetric stretching vibration of P–O.

10 The CaP powders substituted with Sr^{2+} , Mg^{2+} , Zn^{2+} and SeO_3^{2-} , obtained in our previous
11 studies from biogenic source (cuttlefish bone), have been used as the inorganic phase in
12 prepared composite scaffolds (Ressler et al., 2020a; Ressler et al., 2021; Ressler, 2020). The
13 powder samples were mainly composed of calcium-deficient HAp, octacalcium phosphate
14 (OCP, $\text{Ca}_8(\text{HPO}_4)_2(\text{PO}_4)_4 \cdot 5\text{H}_2\text{O}$) and amorphous calcium phosphate (ACP) phases.

15 The XRD analysis of composite scaffolds confirmed the presence of HAp in all prepared
16 samples as a main mineralogical phase. Contrary to expectations, the diffraction patterns of
17 SC_0, SC_1Sr, SC_5Sr, SC_5Se and SC_1MIX revealed the coexistence of HAp and OCP. A
18 minor amount of OCP phase has been detected in SC_1Se, SC_5Zn and SC_MIX composite
19 scaffolds. The calcium-deficient HAp, with a Ca/P ratio 1.50–1.67, is stable within pH range
20 6.5–9.5, while OCP, with a Ca/P ratio 1.33, is stable within pH range 5.5–7.0. The chitosan
21 and CaP suspension has been frozen and neutralized in NaOH/EtOH solution for 8 hours.
22 Even if the pH of the NaOH/EtOH solution is >10 , the portion of the OCP stayed stable
23 despite of being thermodynamically less stable at a high pH. There are several possible
24 explanations for this outcome. The OCP phase might be stabilized by substituent ions in its

1 structure and/or the neutralization time is too short for the OCP to fully transform to a
2 thermodynamically more stable phase, HAp. It has been determined that the OCP phase
3 occurs as an unstable intermediate phase during precipitation of more stable CaPs in aqueous
4 solutions. The central OCP inclusion determined in biological apatite and synthetic
5 precipitated HAp confirms obtaining more stable phases by transformation from the firstly
6 formed OCP phase as explained by Dorozkin ([Dorozhkin, 2014](#)).

7 The SEM analysis and ImageJ software were used to determine the microstructure and pore
8 size distribution of prepared composite scaffolds SC, SC_0, SC_1MIX and SC_5MIX. Fig. 2
9 shows highly porous structures with interconnected pores. The CaPs particles are
10 homogeneously dispersed in the chitosan matrix. The determined pore volume fraction was
11 74.3 ± 0.1 % in SC, 75.1 ± 0.2 % in SC_0, 75.3 ± 0.2 % SC_1MIX and 72.3 ± 0.2 % in
12 SC_5MIX scaffold. The distribution of pore size (Fig. 2) revealed no significant difference
13 between obtained samples. The pore size ranged from ~ 20 to ~ 350 μm in all obtained
14 scaffolds. The composite scaffolds SC_1Sr, SC_5Sr, SC_1Se, SC_5Se, SC_1Zn, SC_5Zn,
15 SC_1Mg and SC_5Mg have not revealed a difference in morphology compared to scaffolds
16 SC, SC_0, SC_1MIX and SC_5MIX.



1

2 *Figure 2.* Microscopic imaging (left) and pore size distribution (right) of prepared composite
 3 scaffolds (SC, SC_0, SC_1MIX, SC_5MIX). Scale bar: 200, 100 and 10 μm.

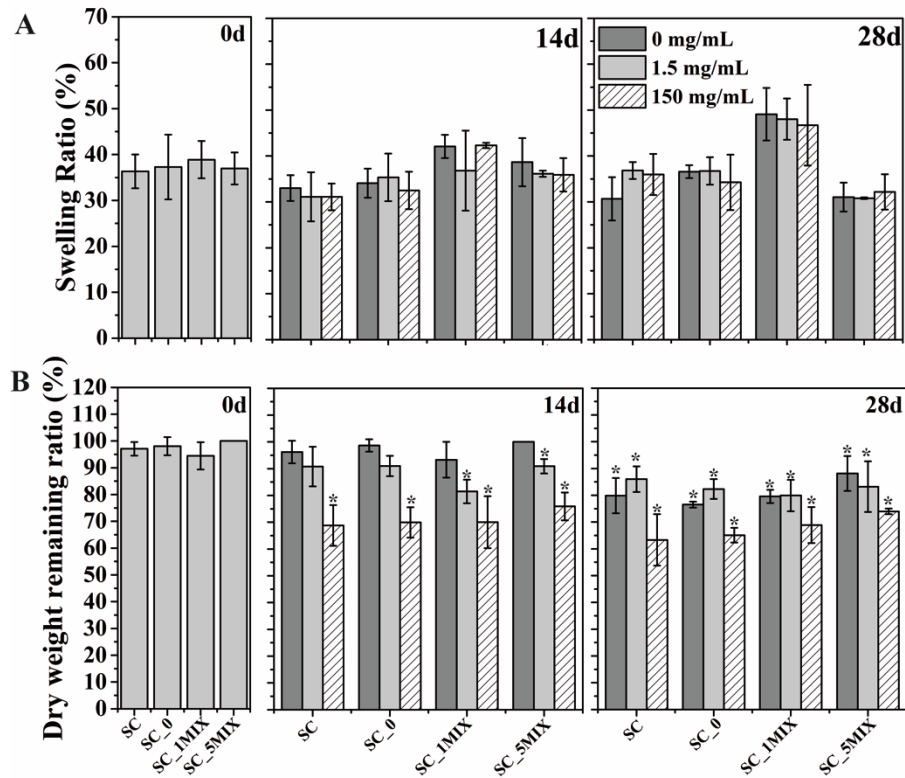
4 3.2. *In vitro* degradation testing

5 Biodegradation of porous scaffolds is a complex process, where the degradation rate depends
 6 on various scaffold characteristics as porosity, pore size, surface area, hydrophilicity, polymer
 7 structure, etc. The enzymatic hydrolytic degradation by lysozymes is the primary mechanism
 8 of chitosan degradation in the human body (Kalantari, Afifi, Jahangirian, Webster, 2019). The
 9 enzymatic degradation of SC, SC_0, SC_1MIX and SC_5MIX was studied at 37 °C as a
 10 function of time, for 28 days, by monitoring swelling behaviour, dry weight loss, weight
 11 average of molecular weight (M_w) and dispersity (D_M). To distinguish between dissolution

1 and enzymatic degradation, composite scaffolds were exposed to PBS solution without (0
2 $\mu\text{g}/\text{mL}$) lysozyme, and with (1.5 and 150 $\mu\text{g}/\text{mL}$) lysozyme, mimicking the *in vivo*
3 physiological conditions. The concentration of lysozyme in human serum is in the range 0.95–
4 2.45 $\mu\text{g}/\text{mL}$ and can increase up to 1,000- fold in the extracellular matrix (Hou, Hu, Park, &
5 Lee, 2012).

6 The results shown in Fig. 3A did not reveal a further increase of swelling ratio after initial
7 water uptake at 0 day. There is no significant difference in swelling ratios with incubation
8 time in degradation medium with 0, 1.5 and 150 $\mu\text{g}/\text{mL}$ of lysozyme. As seen from Fig. 3B, at
9 day 14 the dry weight remaining ratio of all incubated scaffolds decreases as lysozyme
10 concentration increases. After 28 days of incubation, the dry weight remaining ratio in
11 different lysozyme solutions significantly decreases for all analysed scaffolds. The changes in
12 molecular weight were monitored by GPC measurements. As shown in Fig. 4A for SC_5MIX
13 scaffold the weight average molecular weight (M_w) decreased during incubation time in all
14 investigated scaffolds, and as expected, the degradation is the most pronounced in 150 $\mu\text{g}/\text{mL}$
15 of lysozyme solution. In the PBS solution (without lysozyme) no significant change in
16 molecular weight was observed after 14 days of incubation, but, significant change was
17 observed after 28 days. In the solution with 1.5 $\mu\text{g}/\text{mL}$ of lysozyme a significant decrease of
18 M_w occur, from initial $M_{w0} \approx 180,000$ to 147,500 and 94,000, at day 14 and 28, respectively.
19 In the solution with 150 $\mu\text{g}/\text{mL}$ of lysozyme a significant decrease of M_w to 43,000 was
20 observed after 14 days. Between days 14 and 28 no significant change of M_w was observed.
21 The initial \bar{D}_M of 2.37, increased after 14 and 28 days, in 0 $\mu\text{g}/\text{mL}$ of lysozyme solution to
22 2.74 and 2.97, respectively, in 1.5 $\mu\text{g}/\text{mL}$ of lysozyme solution changed to 2.74 and 2.33,
23 respectively, and in 150 $\mu\text{g}/\text{mL}$ of lysozyme solution decrease to 1.79 and 1.90, respectively.
24 The increase of \bar{D}_M after incubation indicates that degradation products remained in the
25 matrice. The \bar{D}_M decrease after 28 days in 1.5 $\mu\text{g}/\text{mL}$ of lysozyme solution and after 14 and

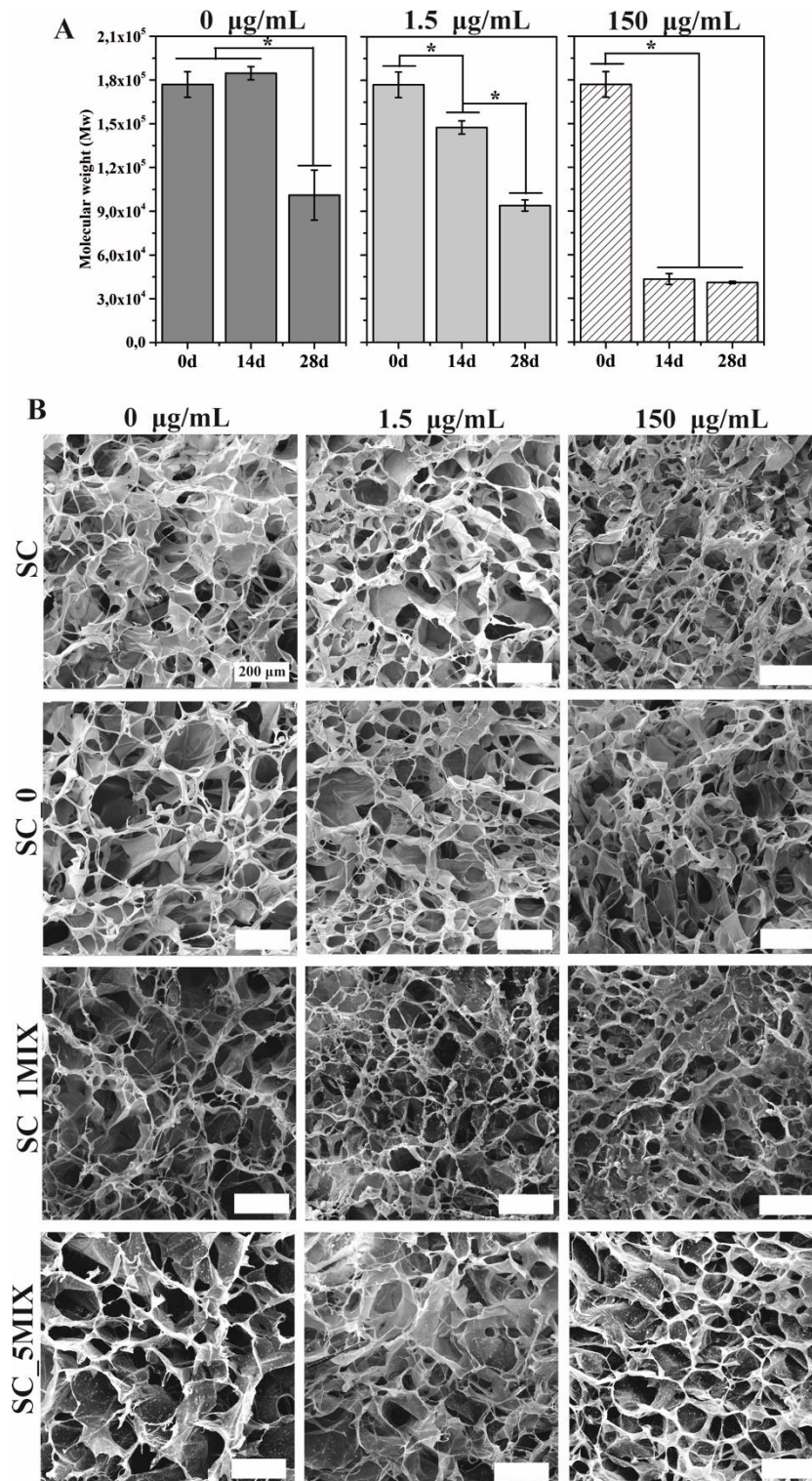
1 28 days in 150 $\mu\text{g}/\text{mL}$ of lysozyme solution indicates the release of low molecular weight
 2 degradation products into surrounding media that is related to pronounced weight loss (Ren,
 3 Yi, Wang and Ma, 2005).



4
 5 *Figure 3.* Swelling ratio (A) and dry weight remaining ratio (B) of composite scaffolds
 6 incubated in different degradation mediums at 37 °C as a function of time. Significant
 7 difference compared to scaffolds at 0 day: *(p < 0.05).

8 The microstructures of the cross-section of SC, SC_0, SC_1MIX and SC_5MIX scaffolds
 9 shown in Fig. 4B after 28 days of incubation at 37 °C did not show significant difference to
 10 initial scaffolds (Fig. 2). Additionally, there is no a significant difference among the scaffolds
 11 incubated in different concentrations of lysozyme. The highly porous structure has been
 12 retained after 28 days of enzymatic degradation which is highly important for cell migration,
 13 diffusion of metabolic waste, oxygen and nutrients. The SEM micrographs confirmed highly

- 1 stable scaffold structure and uniform degradation through the entire scaffold volume despite a
- 2 significant decrease of molecular weight in different degradation mediums.



3

4 *Figure 4.* Weight average molecular weight of SC_5MIX scaffold (A) and SEM (B)

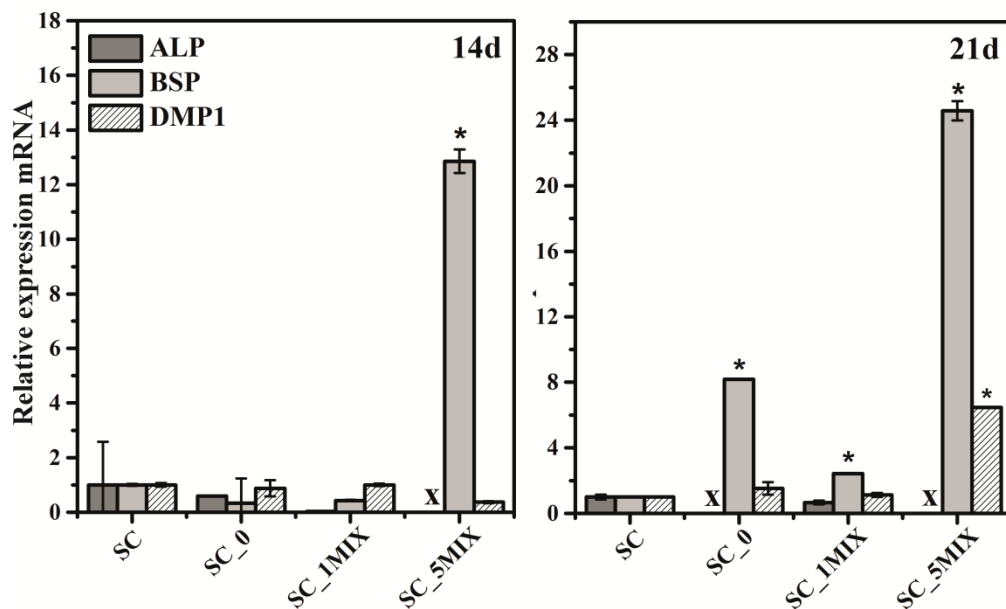
5 micrographs of SC, SC_0, SC_1MIX and SC_5MIX scaffolds incubated in different

1 degradation media at 37 °C as a function of time. Significant difference compared to scaffolds
2 at 0 day: *(p < 0.05).

3 3.3. Biological characterization

4 3.3.4. Quantitative evaluation of osteogenic differentiation in static conditions

5 Relative expression of specific osteogenic markers (ALP, BSP and DMP1) on scaffolds SC,
6 SC_0, SC_1MIX and SC_5MIX after 14 and 21 days of hMSCs culture in static conditions
7 are shown in Fig. 5. The results were normalized to the SC sample.



8

9 *Figure 5.* Relative expression of osteogenic markers (ALP, BSP and DMP1) after 14 and 21
10 days of static 3D culture. The relative gene expression was analyzed by the comparative cycle
11 threshold method ($\Delta\Delta C_t$) and the values were normalized to *18S RNA* expression. The
12 significant difference compared to non-substituted scaffold (SC): * (p < 0.05).

13

14 ALP acts as an early indicator of cellular activity and differentiation and is the first functional
15 gene expressed in the process of osteogenesis (Seibel, 2005; Golub & Boesze-Battaglia,
16 2007). The levels of ALP increase with the progress of osteoblastic differentiation, after

1 which expression of ALP decreases as mineralization progresses (Kulterer et al., 2007; Aubin,
2 2008). As seen from Fig. 5 after 14 and 21 days of cell culture low relative expression of ALP
3 has been detected in analysed samples. The analysis did not reveal any significant difference
4 in ALP expression among the composite samples. The BSP is a post-translationally modified
5 acidic phosphoprotein expressed in mineralized tissues such as bone and dentin (Gordon et
6 al., 2007). The potential to induce HAP nucleation and crystal formation indicate a potential
7 role of the protein in the early stage of mineralization. In addition, BSP can bind to HAP
8 through polyglutamic acid sequences and mediate cell attachment (Kirkham, & Cartmell,
9 2007). The BSP expression significantly increases in scaffolds SC_0 and SC_1MIX after 21
10 days of cell culture. Especially high BSP expression was determined for scaffold SC_5MIX
11 after 14 and 21 days of cell culture.

12 The DMP1 protein expression, characteristic for mineralized tissues, preferentially is
13 expressed by mature bone cells, osteocytes (Kalajzic et al., 2004). There were no significant
14 differences in DMP1 protein expression among the analysed samples, expect a significantly
15 higher expression of DMP1 after 21 days of cell culture in sample SC_5MIX.

16 *3.3.5. Qualitative evaluation of scaffolds cultured in static conditions*

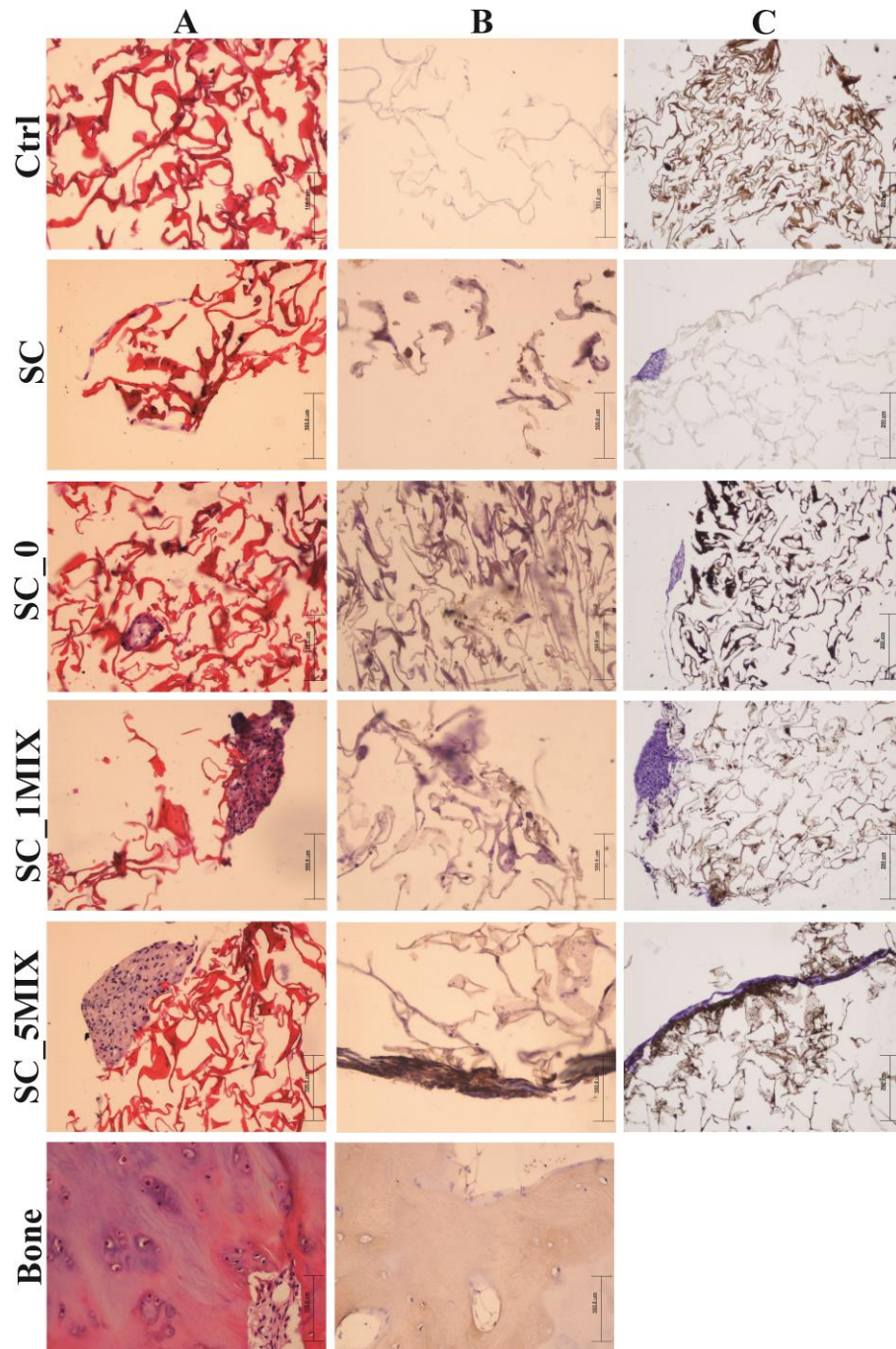
17 The stem cells can differentiate into several cell lineages such as osteoblasts, chondrocytes,
18 adipocytes, tenocytes and myoblasts with a low risk of transformation to tumor cells. The
19 MSCs are often used in evaluation of the osteogenic potential of regenerative biomaterials.
20 The stem cells can differentiate to functional matrix-synthesizing osteoblasts, while the
21 process can be subdivided into three stages: (i) proliferation, (ii) formation of ECM and
22 maturation, and (iii) ECM mineralization. During each stage, characteristic changes in gene
23 expression can be detected. (Aubin, 2008) In this study, hMSCs were used to determine the
24 osteogenic potential of prepared composite scaffolds (SC, SC_0, SC_1MIX and SC_5MIX)

1 after 21 days of cell culture. Scaffold without hMSC was used as negative control (Ctrl).
2 Human bone was used as a positive control.

3 Histological analysis of scaffolds, determined by H&E (Fig. 6A) staining, points out a
4 significant difference in the amount of newly formed tissue in the samples SC_1MIX and
5 SC_5MIX compared to the scaffolds SC_0 and SC. The scaffold SC_0 with inorganic CaP
6 phase obtained from a biogenic source showed a larger amount of formed tissue compared to
7 scaffold SC, with un-substituted HAp phase obtained from synthetic precursors. The higher
8 amount of formed tissue points the benefit of using biogenic sources in regenerative
9 biomaterials. The analysed scaffolds only differ between each other in trace element
10 composition and substitution level. From obtained results, it can be assumed that trace
11 elements play a crucial role in stem cell differentiation to osteoblasts and bone regeneration.

12 Immunohistochemical analysis of scaffolds evaluated the production of type I collagen
13 (COLL I) (Fig. 6B). The COLL I play a role in cell adhesion, proliferation and differentiation
14 of osteoblast and is considered as an early indicator of osteoblastic differentiation ([Kirkham &](#)
15 [Cartmell, 2007](#); [Kulterer et al., 2007](#)). The positive staining of COLL I on all prepared
16 scaffolds after 21 days of cell culture, indicate the early stages of osteogenesis. Along with
17 homogenous expression through the scaffold, the intense localized staining was observed in
18 the scaffold SC_5MIX.

19 Mineralized ECM in the analysed scaffolds was identified by von Kossa staining (Fig. 6C),
20 where phosphate deposits are positively stained in brown and black. After 21 days of cell
21 culture, higher intensity of brown (black) deposits in SC_0 and SC_5MIX indicate that
22 mineralization of ECM occurred. Obtained results of differentiated cells after 21 days of cell
23 culture were compared to the scaffold without seeded cells (negative Ctrl).



1

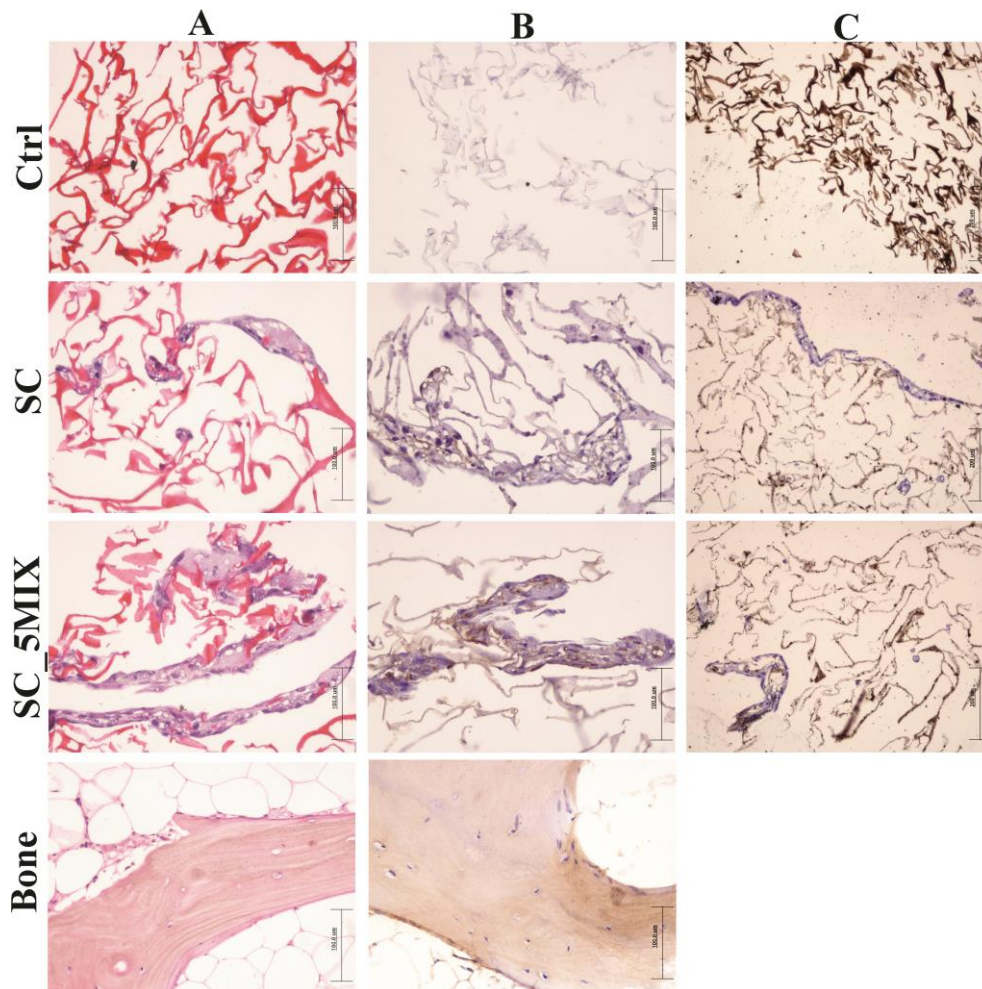
2 *Figure 6.* Detection of newly formed tissue, collagen I and mineralisation after 21 days of
 3 static 3D cell culture of hMSC on composite scaffold SC, SC_0, SC_1MIX, SC_5MIX.
 4 Hematoxylin-eosin (A) staining was used to observe newly-formed tissue,
 5 immunohistochemical (B) staining to observe collagen type I formation and von Kossa (C)
 6 staining for detection of mineralization (including hematoxylin counterstain). Scaffold

1 without hMSC was used as negative control (Ctrl). Human bone was used as a positive
2 control. Scale bar: 100 (A and B) and 200 (C) μm .

3 *3.3.6. Biological evaluation of scaffolds cultured in a perfusion bioreactor*

4 As seen before, the scaffold SC_5MIX showed the highest formation of bone tissue, COLL I
5 expression, phosphate deposition, and BSC and DMP1 protein expression. For this reason,
6 biological evaluation of the scaffold SC_5MIX, cultured in dynamic conditions in perfusion
7 bioreactor were performed.

8 In a bioreactor media flow obtained by perfusion forces provides physiochemical and
9 biomechanical stimuli for new three-dimensional bony tissue growth. The osteogenic
10 potential of the scaffold SC_5MIX was determined in the perfusion U-CUP bioreactor. H&E
11 (Fig. 7A) staining was used to observe newly-formed tissue, COLL I (Fig. 7B) staining to
12 observe collagen formation and von Kossa (Fig. 7C) staining for detection of mineralization
13 (including hematoxylin counterstain). For comparison, scaffold SC was used, to determine the
14 difference in newly-formed tissue, COLL I expression and phosphate deposition, while
15 scaffold without hMSC was used as negative control (Ctrl).



1
 2 *Figure 7.* Detection of newly formed tissue, collagen I and mineralization after 21 days of
 3 dynamic 3D cell culture of hMSC on composite scaffold SC and SC_5MIX. Hematoxylin-
 4 eosin (A) staining was used to observe newly-formed tissue, immunohistochemical (B)
 5 staining to observe collagen type I formation and von Kossa (C) staining for detection of
 6 mineralization (including hematoxylin counterstain). Scaffold without hMSC was used as a
 7 negative control (Ctrl). Human bone was used as a positive control. Scale bar: 100 (A and B)
 8 and 200 (C) μm .

9 Histological analysis of scaffolds points out a difference in the amount of newly formed tissue
 10 in the scaffold SC_5MIX compared to scaffold SC. The newly-formed tissue is uniformly
 11 distributed through the volume of the scaffold SC_5MIX. It can be assumed that obtained
 12 scaffold's composition is a suitable environment for hMSCs migration and growth after 21

1 days of cell culture. The positive staining to COLL I was determined in scaffolds SC_5MIX
2 and SC after 21 days of cell culture, indicating the early stages of osteoinduction. Along with
3 homogenous expression through the newly-formed tissue, the intense localized staining was
4 determined in scaffold SC_5MIX. After 21 days of cell culture, brown (black) deposits in
5 scaffolds SC_5MIX and SC indicate that mineralization of ECM occurred. Even the intensity
6 of the color is lower than Ctrl, the deposition (black dots) can be seen after 21 days of cell
7 culture.

8 **4. Discussion**

9 In our previous studies (Ressler et al., 2020a; Ressler et al., 2021; Ressler 2020), biomimetic
10 triphasic CaPs powders substituted with Sr^{2+} , Mg^{2+} , Zn^{2+} , and SeO_3^{2-} ions were prepared by
11 wet precipitation method, using biogenic source (cuttlefish bone) as a precursor of Ca^{2+} ions.
12 The precipitated CaPs were composed of calcium-deficient carbonated HAp, OCP and ACP.
13 Ion release study confirmed better resorbable properties of materials obtained from a biogenic
14 source and substituted with ions, while phase transformation analysis at simulated
15 physiological conditions confirmed the transformation of OCP and ACP to
16 thermodynamically more stable phase, HAp, as it occurs in natural tissue. The culture of HEK
17 293 cells indicated noncytotoxicity of prepared CaP powders substituted with Sr^{2+} , Mg^{2+} and
18 Zn^{2+} with emphasis on the cell proliferation during culture time. Additionally, the CaPs
19 powders substituted with SeO_3^{2-} ions have shown selective anticancer properties at lower
20 levels of substitution.

21 As a continuation of previous studies, highly porous composite scaffolds based on multi-
22 substituted CaPs and biopolymer chitosan have been prepared and an extensive *in vitro*
23 characterization using hMSC was performed, including enzymatic degradation during 28 days
24 at simulated physiological conditions. It was hypothesized that substituted ions Sr^{2+} , Mg^{2+} ,

1 Zn^{2+} , and SeO_3^{2-} would provide a suitable environment for hMSCs differentiation with a
2 positive influence on osteoinduction and newly-formed bone tissue.

3 Mineralogical phase composition is of high importance for suitable resorption and biological
4 performance of prepared scaffolds. Recent studies are focused on obtaining two or three CaP
5 phase systems to enhance the resorption properties of biomaterials. As explained in our
6 previous studies, foreign ions influence the phase transformation of CaPs from amorphous to
7 crystalline phases, resulting in the stabilization of metastable CaP phases. The attractive
8 aspect of OCP as a bone substitute material has been studied by Suzuki et al. (Suzuki et al.,
9 2006). It was demonstrated that the implantation of OCP markedly enhanced bone formation,
10 osteoblastic cell proliferation and differentiation compared to the implantation of Ca-deficient
11 HAp. (Suzuki et al., 2006) Obtained inorganic phase in prepared scaffolds consists of CaPs
12 characterized by small dimensions, low crystallinity and non-stoichiometric composition
13 which are characteristics of natural apatite.

14 The pore diameter, pore size distribution, pore volume, pore shape and pore wall roughness
15 are important factors for proper cell attachment, migration and tissue growth. The porosity of
16 ~75 % in prepared composite scaffolds is in the range 50-90% suitable for the efficient bone
17 regeneration process. Achieved interconnectivity between pores is essential for the diffusion
18 of essential nutrients, oxygen and extracellular fluid in and out of the cellular matrix. Both
19 macro and micro porosity play a vital role in the development of tissue inside the scaffold
20 environment. Macroporosity, having a pore size of 100-400 μm , promotes osteogenesis by
21 enhancing cell migration, cell-cell network formation, vascularization, nutrients supply, and
22 metabolic waste diffusion. Microporosity, with a pore size $< 20\mu m$, is crucial for cell seeding
23 and retention, capillaries growth, vascularization and cell-matrix interactions. (Fernandez-
24 Yague et al., 2015; Deb et al., 2018) The achieved porosity and pore size requirements of the
25 prepared scaffold should allow rapid and abundant vascularization network, essential to avoid

1 implanted-cell necrosis, formation of acellular regions and to allow new tissue formation
2 (Cardonnier et al., 2011).

3 Swelling behaviour, structural stability and degradation rate are highly important in order to
4 allow tissue formation to occur simultaneously with degradation. The scaffolds obtained from
5 natural polymers often undergo rapid degradation, difficult to control, under physiological
6 conditions. (Karageorgiou & Kaplan, 2005; Roseti et al., 2017) Chitosan has the ability to
7 swell under physiological conditions giving a microenvironment similar to natural bone tissue
8 suitable for cell attachment and growth. As explained in our previous research (Ressler et al.,
9 2018), chitosan degrades *via* hydrolysis, where polymer network breaks into smaller chains,
10 whereby the β -1-4 N-acetyl glucosamine units undergo chain scission mainly by lysozymes
11 present in body fluid. Degradation products of chitosan are non-toxic, mainly composed of
12 glucosamine and saccharide, which can then be easily excreted from the body without
13 interference with other organs (Kalantari et al., 2019). The obtained results show that
14 prepared scaffolds, based on natural polymer chitosan, are highly stable during 28 days at
15 physiological conditions, and in a much higher concentration of lysozyme, which is highly
16 important for supporting newly formed bone tissue. The reason for this outcome could be the
17 higher molecular weight of chitosan and deacetylation degree. Chitosan with a high DD tends
18 to have a slower degradation rate due to closer chain packing and hydrogen bonding
19 (Kalantari et al. 2019).

20 The MSCs are shown promising potential as a source of cell-based therapeutic strategies.
21 These cells can differentiate into osteoblasts, chondrocytes, adipocytes, tenocytes and
22 myoblasts and thus may be used in multiple types of cellular therapeutic strategies for
23 regeneration of all tissues as bone, cartilage, fat, muscle, tendons and ligaments (Aubin, 2018;
24 Kulterer et al., 2007). In bone tissue, the MSCs seem to migrate and attach to the scaffold
25 surface *in vivo* from day one after implantation, depositing bone-related proteins and creating

1 a non-collagenous matrix layer on the implant surface that regulates cell adhesion. This
2 matrix is an early-formed calcified afibrillar layer on the surface, involving poorly
3 mineralized osteoid that is rich in calcium, phosphorus, osteopontin and BSP (Mavrogenis,
4 Dimitriou, Parvizi, Babis, 2009).

5 The recent uses of growth factors have proven the advantage of manipulating synthetic
6 materials to enhance the bone regenerative process. However, the safety and stability of
7 growth factors have been questioned. In recent years, many studies have focused on the
8 biomimetic approach and incorporation of trace elements, crucial for bone function, into
9 synthetic materials (Bose et al., 2013). The CaPs used in this work are substituted with Sr^{2+} ,
10 Mg^{2+} , Zn^{2+} , and SeO_3^{2-} ions, along with Na-substitution as a result of using cuttlefish bone as
11 a precursor of Ca^{2+} ions. The biocompatibility test of ion substituted CaPs has been obtained
12 in our previous research, as the high intake of ions can have a negative side effect on cells.
13 Further, the biocompatibility test of prepared composite scaffolds confirmed biocompatibility
14 and the ability to promote cell proliferation. Significantly higher live cell (%) after 7 days of
15 culture, compared to non-substituted scaffold SC, indicate a beneficial effect of substituted
16 ions on cell proliferation (dana not shown).

17 Bonnelye et al. determined the dual mode of action of Sr^{2+} ions, where Sr^{2+} ions increased
18 expression of osteoblastic markers ALP, BSP and osteocalcin while decreasing osteoclasts
19 activity (Bonnelye, Chabadel, Salterl, Jurdic, 2008). A similar tendency was observed in *in*
20 *vivo* studies of collagen-Sr-HAp scaffold developed by Yang et al. As explained, the calcium-
21 sensing receptor (CaSR) is involved in the Sr-induced proliferation of the osteoblasts and the
22 apoptosis of osteoclasts (Yang et al., 2011). Strontium activates the CaSR in osteoblasts,
23 downstream signaling pathways, which promote osteoblast proliferation, differentiation, and
24 survival, while at the same time cause osteoclast apoptosis resulting in a decrease of bone
25 resorption. The CaSR simultaneously increases osteoprotegerin (OPG) production and

1 decreases receptor activator of nuclear factor kappa beta ligand (RANKL) expression. OPG is
2 a protein that inhibits RANKL-induced osteoclastogenesis by operating as a decoy receptor
3 for RANKL. The OPG/RANKL ratio, additionally controlled by Sr-introducing, can be a
4 powerful regulator of bone resorption and osteoclastogenesis (Bose et al., 2013).

5 In natural bone tissue, along with calcium, magnesium is the second „messenger“ for
6 regulating a wide variety of reactions involved in cell response through signaling pathways.
7 Magnesium has a key role in DNA, RNA and protein synthesis, while Mg-substituted CaPs
8 increase osteoblast cellular attachment, proliferation, and ALP production (Mammoli, 2019;
9 Bose et al., 2013). ALPs are glycosyl-phosphatidylinositol-anchored Zn^{2+} -metallated
10 glycoproteins that are released during the maturation process of the osteoblastic cell life cycle,
11 which creates an alkaline environment that favors the CaP precipitation and subsequent
12 mineralization of the ECM (Bose et al., 2013). Hwan Park et al. reported that zinc exerts
13 osteogenic effects in hMSCs by activation of Runt-related transcription factor 2 (Runx2) *via*
14 the cAMP-PKA-CREB signaling pathway. Both, ALP and Runx2 are highly important in
15 hMSC differentiation to osteoblasts, which points crucial role of zinc in bone formation. (Park
16 et al., 2018).

17 Recently, selenium attracted attention as a substituent in CaPs due to the selective effect on
18 healthy and osteosarcoma cells. Selenium induces oxidative stress in cancer cells by
19 generating reactive oxygen species and suppresses RANKL-induced osteoclastogenesis
20 through inhibition of signaling pathways while having a positive effect on healthy osteoblast
21 cells (Yang et al., 2011). In accordance with our previous study (Ressler et al., 2021) the
22 composite scaffolds, with SeO_3^{2-} substituted CaP, revealed anticancer properties on U2OS
23 cells, while at the same time enhancing the proliferation of healthy HEK 293 cells (data not
24 shown). The obtained results are in agreement with results reported by Uskoković et al.,
25 where biomaterial substituted with selenium has selective toxicity for osteosarcoma cells,

1 while not affecting the viability of non-cancerous primary cells. (Uskoković, Iyer, & Wu,
2 2017).

3 In the present work hMSCs seeded on the composite scaffolds were evaluated after 14 and 21
4 days of cell culture. The biological evaluation was carried out at static and dynamic
5 conditions using a perfusion bioreactor. The histological, immunohistochemical and RT-
6 qPCR analyses were used to determine the osteogenic properties of prepared scaffolds.
7 Detection of a higher amount of newly-formed bone tissue, ALP, BSP, COLL I, DMP1 and
8 phosphate deposits, characteristic for differentiation process, indicate good osteogenic signal
9 of the scaffolds substituted with trace elements, especially the scaffolds with the synergic
10 effect of ions, SC_1MIX and SC_5MIX. The hMSCs differentiated in osteoblasts phenotype
11 within 21 days of cell culture indicating the possible application of SC_1MIX and SC_5MIX
12 as scaffolds for bone tissue regenerative applications.

13 As the single substituent systems have demonstrated good results in recent studies, using
14 multiple substituents can be used to further increase the beneficial effects of each. A
15 combination of different trace metal ion additions in CaPs can alter the degradation kinetics,
16 the inhibitory effect on osteoclastogenesis, and the stimulatory effect on osteogenesis (Bose et
17 al., 2013). This research highlight the key role of trace elements and their synergic effect on
18 hMSC differentiation and the formation of newly-bone tissue. Trace elements have shown
19 that they have potential to replace growth factors and pharmaceologics for enhancing the
20 regenerative properties of synthetic materials without causing negative side effects. However,
21 further *in vivo* studies on animal models need to be performed to examine biodegradation and
22 osteogenesis to confirm the applicability of the SC_5MIX scaffold for bone tissue
23 engineering.

24 **5. Conclusion**

1 Present research has shown that multi-substituted scaffolds based on chitosan and CaP,
2 mainly composed of HAp, acts as three-dimensional support for hMSCs proliferation and
3 differentiation into osteoblast cells. Expression of characteristic bone genes, ALP, BSP,
4 COLL I, DMP1 and phosphate deposits, along with newly formed bone tissue, indicated that
5 ECM mineralization took place during 21 days of hMSCs culture indicating the key role of
6 Sr^{2+} , Mg^{2+} , Zn^{2+} and SeO_3^{2-} during osteogenesis. Prepared scaffolds are highly stable under
7 physiological and enzymatic conditions even the natural polymer chitosan was used.

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