3D gelatin-chitosan hybrid hydrogels combined with human platelet lysate highly support human mesenchymal stem cell proliferation and osteogenic differentiation

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Abstract
Bone marrow and adipose tissue human mesenchymal stem cells were seeded in highly performing 3D gelatin-chitosan hybrid hydrogels of varying chitosan content in the presence of human platelet lysate and evaluated for their proliferation and osteogenic differentiation. Both bone marrow and adipose tissue human mesenchymal stem cells in gelatin–chitosan hybrid hydrogel 1 (chitosan content 8.1%) or gelatin–chitosan hybrid hydrogel 2 (chitosan 14.9%) showed high levels of viability (80%–90%), and their proliferation and osteogenic differentiation was significantly higher with human platelet lysate compared to fetal bovine serum, particularly in gelatin–chitosan hybrid hydrogel 1. Mineralization was detected early, after 21 days of culture, when human platelet lysate was used in the presence of osteogenic stimuli. Proteomic characterization of human platelet lysate highlighted 59 proteins mainly involved in functions related to cell adhesion, cellular repairing mechanisms, and regulation of cell differentiation. In conclusion, the...
combination of our gelatin–chitosan hybrid hydrogels with hPL represents a promising strategy for bone regenerative medicine using human mesenchymal stem cells.

Keywords
Hybrid chitosan-gelatin hydrogel, human mesenchymal stem cells, human platelet lysate, tissue engineering, bone regeneration

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Introduction
Regenerative medicine has been receiving growing attention as treatment approach for various tissue injuries. Current tissue engineering strategies are mainly focused on the restoration of damaged tissue by using human mesenchymal stem cells (hMSCs) either alone or in combination with three-dimensional (3D) scaffolds. In particular, to restore the bone’s structural integrity, implantation of scaffolds bioengineered with hMSCs represents the most promising technique since it combines the properties of the scaffolds with the activities of MSCs.1

Scaffolds are generally 3D porous substrates designed to promote cell-biomaterial interactions, permitting sufficient transport of gases, nutrients, and regulatory factors for cell survival, proliferation, and differentiation. In addition, they can biodegrade with a controlled rate as the tissue regenerates, while triggering minimal degrees of inflammation or toxicity in vivo.2–7

For the purpose of our study, high-performance porous scaffolds were synthesized starting from gelatin (G), polyethylene glycol (PEG), and chitosan (CH) (G-CH hybrid hydrogels) to support hMSCs proliferation and osteogenic differentiation.

Hydrogels are 3D polymeric biomaterials that can be tuned to have properties similar to the natural extracellular matrix (ECM). Whereas naturally derived hydrogels are biocompatible, bioactive, and biodegradable and promote many cellular functions, synthetic hydrogels are highly reproducible, and characterized by high mechanical properties.8

G has been extensively used in the field of tissue engineering due to its biocompatibility and biodegradability.9 Similarly, CH has several beneficial properties that make it a good candidate for biomedical applications. It is biocompatible, biodegradable, non-antigenic, non-toxic, antimicrobial, and shows haemostatic activity.10 In addition, its structure mimicking natural ECM constituents and cationic surface charges are expected to promote cell attachment and growth.11

hMSCs have been demonstrated to promote tissue repair thanks to their ability to migrate to injured tissues, and their immunomodulatory and trophic effects.12 These cells can be isolated from various tissues, mainly including bone marrow (BM), adipose tissue (AT), and cord blood (CB); can be expanded in quantities that are sufficient for potential therapeutic applications; and show self-renewal capacity and ability to differentiate in vitro into osteoblasts, adipocytes, and chondroblasts.13,14

In this study, we used both BM and AT-hMSCs within the G-CH hybrid hydrogels, in the presence of either fetal bovine serum (FBS) or human platelet lysate (hPL). hPL has been recently introduced as a substitute for FBS since it allows to avoid the risks of transmitting animal diseases and potential immune responses to animal antigens and also may overcome the strict rules of regulatory authorities in charge of the approval of experimental protocols for somatic cell therapies.15–19 Furthermore, platelet derivatives are widely applied in different clinical fields, since they function as tissue sealant and delivery system for mitogenic and chemotactic growth factors (GFs). In doing so, cell proliferation, angiogenesis, and cell migration are stimulated, and tissue regeneration favored.20

Thus, in this study, we tried to establish, in vitro, a clinical grade biomedical device for potential use in bone regenerative medicine, by seeding BM-hMSCs and AT-hMSCs in G-CH hybrid hydrogels in the presence of hPL and evaluating their viability, proliferation, and osteogenic differentiation.

Materials and methods
Reagents
Type A G (pharmaceutical grade, 280 bloom, viscosity 4.30 mPs), produced from pig skin, was purchased from Italgelatine, Italy. CH (molecular weight between 50,000 and 190,000 Da and degree of deacetylation 75%–85%) was obtained from Fluka, Italy. Poly(ethylene glycol)diglycidyl ether (molecular weight 526 Da) was supplied by Sigma-Aldrich, Italy. Ethylene diamine (EDA) and acetic acid were provided by Fluka, Italy. Dulbecco’s modified Eagle’s medium (DMEM), l-glutamine, penicillin-streptomycin, and sodium pyruvate were purchased from Sigma-Aldrich, USA. Amphotericin B and minimum essential medium (MEM) non essential amino acids solution were purchased from Gibco, ThermoFisher Scientific, USA.

G-CH hybrid hydrogels synthesis. G-CH hydrogels were prepared in aqueous solution and the synthetic procedure
involved the reaction between G/CH amino-groups and the epoxy groups of functionalized PEG.

Briefly, G (6 g) was dissolved in 65 mL distilled water at 45°C under mild magnetic stirring followed by drop-wise addition of PEG (1.4 g) and EDA (70 mg). CH solution in acetic acid (2 wt%, 33 g) were added and the final reaction mixture was gently magnetically stirred at 45°C for 20 min to obtain homogeneous mixture and then poured into the glass plate for gel formation. The gels were cut into rectangular bar or dumbbell and then were frozen by dipping into liquid nitrogen bath maintained at a temperature of −196°C. The frozen samples were freeze-dried using an Edwards Modulyo freeze-drier operating under vacuum at –60°C, for sublimation of ice crystals. Finally, in order to further increase the degree of grafting, the dried samples were put into oven at 45°C for 2h under vacuum. The final products obtained after freeze drying and post-curing process maintained well their size and shape. In order to obtain proper composition, two different amounts of CH were added obtaining the final dry hydrogels, namely G-CH1 (CH content 8.1%) and G-CH2 (CH content 14.9%). Hydrogels were sterilized by gamma irradiation with Cobalt 60 gamma rays using 27–33 kGy following UNI EN ISO 11137 (Sterilization of Health Care Products).

Chemical morphological and mechanical characterization of G-CH hybrid hydrogels. The texture, morphology and porous structure of hydrogels were analyzed by scanning electron microscopy (SEM) using a LEO EVO 40 scanning electron microscope. The dry samples were cryogenically fractured. Mechanical tests were carried out by an Instron series 3366 testing machine, equipped with a 50N load cell. Tensile tests were carried out at a cross-head rate of 10 mm/min. Specimens were cut from the sample disk into a dumbbell shape. The actual width and thickness of the specimens in the narrow portion of the dumbbell were measured using an optical traveling microscope. The samples, having a gauge length of 25 mm, were clamped at a larger distance and stretched until failure. At least six specimens were tested for each hydrogel composition. Engineering stress was calculated by dividing the recorded force by the initial cross-sectional area. Engineering strain under compression was calculated as the ratio of cross-head displacement to original length/thickness. Initial elastic modulus (stiffness, E) was determined from the slope of the initial linear segment of stress–strain curves. Cyclic loading–unloading experiments were performed for up to 10 cycles with 50% maximum strain without relaxation time in order to evaluate the reversible behavior of hydrogels. Cyclic compression tests were also performed over time (1, 10 and 21 days) to monitor the mechanical integrity of the hydrogels during hydrolytic degradation.

Hydrolytic mass loss evaluation and swelling ratio. The pre-weighed ($W_i$) samples were incubated in distilled water at 37°C over a 3-week period. At regular intervals of 1, 10, and 21 days, the samples were removed, weighed ($W_w$), rinsed with fresh water, air dried followed by vacuum drying at 45°C for 4h, and finally dried mass ($W_f$) was measured. The swelling ratio (%) and mass loss (%) was calculated using following equations

\[
\text{Swelling ratio} \%(%) = \frac{(W_w - W_f)}{W_i} \times 100
\]

\[
\text{Mass loss} \%(%) = \frac{(W_i - W_f)}{W_i} \times 100
\]

hPL production. hPL for mesenchymal stem cells expansion was obtained from blood donations belonging to the Blood Bank of ASST Spedali Civili of Brescia and produced according to standardized clinical grade procedures in closed systems. From the automatic separation of whole blood (CompoMat G5, Fresenius Kabi AG, Germany), fresh frozen plasma (FFP), buffy coat (BC), and red blood cells units were obtained. Within 48 h of collection, 12 (6 + 6) group 0 BC units were pooled (BioP flex Pool, Fresenius Kabi AG, Germany), each diluted in group AB FFP (250–300 mL) and gently centrifuged (340g, 6 min, 22°C—Heraeus Cryofuge 6000i, Thermo Fisher Scientific, USA). At this point, the supernatant obtained is transferred to a satellite bag and in-line filtered for leukocytes depletion. Residual leukocyte count (<0.2/mL) was performed by flow cytometry (Kit Leucocount, Facs Canto II, Becton Dickinson, Italy). Platelet-rich plasma (PRP) obtained was transferred to a 600 mL bag (Compoplex 1F, Fresenius Kabi AG, Germany), adjusted with FFP, at a platelet concentration of 1.0 ± 0.2 × 10^6/µL (sodium heparin 40U/mL), frozen at −80°C for 24 h, and thawed at 37°C for 25 min (DH8 QuickThaw™, Helmer Inc., USA) to lyse platelets. After a second centrifugation (4000 g,
15 min, 22°C) aimed at removing platelet bodies or debris, the hPL obtained was aliquoted into 50 mL tubes (BD Falcon™ Conical Tubes, Becton Dickinson, Italy) and stored at −80°C until use. Sterility tests were done to prove hPL negativity for aerobic /anaerobic bacteria.

**Platelet poor plasma production**

Platelet-poor plasma (PPP) was produced from PRP according to standardized blood bank procedures in closed systems finally containing less than 10×10³ platelets/μL. Four units of whole blood belonging to the Blood Bank of ASST Spedali Civili of Brescia were centrifuged (3400 g, 4 min, 22°C—Heraeus Cryofuge 6000i, Thermo Fisher Scientific, USA), so that the optimal number of platelets remained in plasma (PRP). Platelets from PRP were sedimented by hard spin centrifugation (5050 g, 6 min, 22°C); the supernatant PPP was transferred to a satellite 600 mL bag (Compoflex 1F, Fresenius Kabi AG, Germany) and subsequently aliquoted into 50 mL tubes (BD Falcon™ Conical Tubes, Becton Dickinson, Italy) and stored at −80°C until use as reference background in GFs determination on hPL. All the blood units used for the production of hPL and PPP were biologically qualified according to Italian legislation complying with EU legislation.

**hPL proteomic characterization by nLC-ESI MS/MS**

Quanti-quantitative evaluation of hPL compared to PPP was achieved by nLC-ESI MS/MS shotgun proteomics. In particular, one batch of hPL and four distinct PPP samples from different donors (n=4) were processed and evaluated for proteomic characterization.

High abundant proteins, as albumin and IgG, were depleted from plasma-derived samples using ProteoPrep® Immunoaffinity Albumin and IgG Depletion Kit (PROTIA Depletion kit, Sigma-Aldrich, USA), following manufacturer’s instructions before MS analysis protocol. Both depleted and not depleted samples were processed and analyzed in MS.

In particular, a volume corresponding to 300 μg of proteins for each sample was trypsinized based on filter aided sample preparation (FASP) strategy as previously described. Digestion was performed overnight at 37°C onto ultrafiltration units (Amicon Ultra–0.5 mL 30 kDa, Millipore, USA) after reduction with 50 mM dithiothreitol and alkylation (100 mM iodoacetamide). No urea or sodium dodecyl sulfate (SDS) was added, and 50 mM ammonium bicarbonate was used as buffer in each step. About 1 μg of peptide mixtures were injected into UHPLC system (Ultimate™ 3000 RSLCnano, ThermoFisher Scientific, USA) coupled online with Impact HD™ UHR-QqToF (Bruker Daltonics, Germany) as already reported. Technical variability for label-free relative quantification was minimized injecting each sample at least twice (for each single sample of PPP) and four times (for the sample of hLP).

Protein identities were determined using an in-house Mascot search engine (version: 2.4.1), using Swissprot reference databases of *Homo sapiens* (accessed June 2018, 557,713 sequences; 200,130,199 residues) and the following parameters: enzymatic digestion by trypsin, allowing one missed cleavage; precursor mass tolerance was set as 20 ppm; fragment mass tolerance of 0.05 Da; carboxymethylation (Cys) as fixed modification. Decoy database and a false discovery rate (FDR) less than 1%, were applied. Only proteins with at least one unique peptide (p ≤ 0.05) were considered identified.

Peaks Studio 8.5 (Bioinformatics Solutions Inc., Waterloo, ON, USA) was utilized to determine the relative abundances of identified proteins. All replicates were loaded separately. The areas under the curve of the extracted ion chromatograms were measured and normalized using total ion current (TIC). As protein filter a significance of 13 (FDR with adjusted p-value of ≤ 1%) was used for identification. Only peptide features having a quality ≥2, an average area ≥104 and a charge from 1 to 10, and only proteins that had been identified with at least two unique non redundant peptides were selected for quantitation. Analysis of variance (ANOVA) test integrated into the software was chosen to evaluate the significance of the comparison (p < 0.05). A fold change of 1.5 was applied and only proteins that showed significant differences in at least 4 of 12 replicates were considered. Functional annotation was performed using String Online Tool.

**Human BM and AT mesenchymal stem cells (BM-AT-hMSCs) culture**

Human BM mesenchymal stem cells (BM-hMSCs) and human AT mesenchymal stem cells (AT-hMSCs) were purchased from PromoCell, Germany. BM-hMSCs and AT-hMSCs were expanded in the presence of a growth medium (GM) consisting of DMEM, a high glucose-based medium with 2% L-glutamine/penicillin-streptomycin/amphotericin B solution (stock solution, 200 mM L-glutamine, 10 000 U/mL penicillin, 10 mg/mL streptomycin, 250 μg/mL amphotericin B), 1 mM sodium pyruvate, MEM Non Essential Amino Acids Solution 1X and 10% FBS (referred as complete medium FBS) or 5% hPL (referred as complete medium hPL) at 37°C and 5% CO₂ in an incubator.

**BM-AT-hMSCs cell viability and cell proliferation assay**

BM-hMSCs and AT-hMSCs were detached from the flask using trypsin ethylenediaminetetraacetic acid after reaching 80% of confluence, neutralized with complete medium
FBS, centrifuged at 11000 r/min for 5 min, resuspended in the GM, and counted with a hemocytometer. A cell suspension at a cellular density of 10⁶ cells/mL (36 × 10⁵ cells/scaffold) was added to the scaffolds. Scaffolds with volume of approximately 36 mm³ (approximately dimensions: 3 mm × 3 mm × 4 mm) were used. The required amount of BM-hMSCs and AT-hMSCs cells at passage 3 was added to each scaffolds by direct cell seeding in 24 uncoated multiwell plates, then incubated at 4°C for 1 h and 37°C under 5% of CO₂ conditions for 1 h. In fact, it was previously observed that this cold static seeding method improves cell penetration generating a uniform initial cell distribution in the porous scaffolds. After this time, 1 mL of complete medium FBS or complete medium hPL were added to each scaffolds. Cell proliferation was analyzed after 2, 6, 10, 14, and 21 days. Cells viability was evaluated using a Live/Dead kit for mammalian cells (ThermoFisher, USA). The samples were washed with Dulbecco’s phosphate-buffered saline (DPBS) and incubated for 30–45 min at RT in DPBS with 2 μM of calcein AM and 4 μM of ethidium homodimer-1 (EthD-1). NucBlue® Live reagent (2 drops/mL) for nuclei staining was added to the cultures. An analysis of live (stained in green with calcein AM) and dead (stained in red with EthD-1) cells was then performed using a Zeiss Observer Z1 fluorescence microscope. Ten images were taken from three different replicates of each sample and the percentage of viable cells (% total cell number – % dead cells) was then calculated for each conditions using Image J.

Cell proliferation was determined using the Cell Counting Kit-8 (CCK-8, Sigma-Aldrich, USA) on days 2, 6, 10, 14, and 21 of cell culture, following manufacturer instructions. The formazan produced by the CCK-8 assay is shown to be proportional to the number of viable cells giving rise to a colored compound that can be measured spectrophotometrically at 450 nm. Briefly, the cell-cultured samples (three replicates) were moved to a new cell culture plate at each time points and incubated with a fresh culture medium containing the CCK-8 reagent (ratio 1:10) at 37°C for 2 h 30 min. Then, the absorbance of 100 μL of supernatant transferred to a new cell culture plate was measured at 450 nm with an Infinite 200 PRO plate reader (Tecan, Switzerland). Absorbance at 450 nm is proportional to the number of viable cells in each sample. Indeed, a standard curve was created by culturing known numbers of viable cells in order to determine the relative viable cells number proliferating in the hydrogels.

**BM-AT-hMSCs osteogenic differentiation**

A cell suspension at a cellular density of 10⁶ cells/mL (36 × 10⁵ cells/scaffold) was added to the scaffolds for osteogenic differentiation in static conditions. To induce osteogenic differentiation, cells-scaffolds constructs were cultured in 24-uncoated well plates for 3 weeks with osteogenic differentiation medium (OM), consisting of complete medium FBS or complete medium hPL supplemented with 10⁻⁷ M dexamethasone, 25 μg/mL L-ascorbic acid, and 3 mM NaH₂PO₄. All samples were analyzed in triplicates. As controls, samples were also cultured in the complete medium FBS or complete medium hPL without additional osteogenic factors for 3 weeks, to evaluate the osteogenic potential of the hydrogels alone. After 21 days of culture, scaffolds were fixed with formaldehyde 4% for 1 h at 4°C. After the fixing step, the samples were soaked overnight in 30 w/v % sucrose in DPBS, embedded in OCT compound (VWR, USA, 361603E), and frozen in liquid nitrogen. Finally, 80 μm sections were cut out with a Leica CM 1860 ultraviolet (UV) cryostat. Sections were collected and placed onto Thermo Scientific superfrost ultra plus slides (J3800AMNZ). For immunostaining, sections were encircled by immunoEdge pen, rehydrated in PBS, permeabilized with 0.1 vol % Triton X-100 in PBS IX and then blocked for non-specific binding sites by incubation with bovine serum albumin 1% in PBS (PBS-BSA 1%) for 1 h. To analyze osteogenic differentiation of cells inside the scaffolds, primary antibodies (mouse monoclonal anti-osteocalcin (OCN), Santa Cruz Biotechnologies, USA, ABOC-5021) were diluted 1:50 in PBS-BSA 1%. After 1 h of incubation at 37°C, sections were then washed in PBS and incubated with the secondary antibodies (rabbit anti-mouse Cy3, Jackson Immunoresearch, USA, 315-165-003) diluted 1:100 in PBS-BSA 1% for 1 h at RT. To analyze actin cytoskeleton of the cells in the scaffolds Alexa Fluor 488 phalloidin (Invitrogen, USA, A12379) was diluted 1:100 in PBS-BSA 1% for 30 min at RT. Finally, samples were washed three times with PBS adding 0.5% Tween-20 before mounting the coverslip with a drop of Vectashield (Vector Laboratories, USA) containing 4′,6-diamidino-2-phenylindole (DAPI) to counterstain cell nuclei (blue). Images were acquired with an inverted Zeiss Observer Z1 fluorescence microscope. The samples were analyzed in triplicates. Fluorescent intensity was analyzed using ImageJ and normalized according to the cell number.

For von Kossa staining, the sections were air-dried for 20 min and fixed in 10% formaldehyde (Kaltek, Italy) at RT for 1 h. Then, samples were rehydrated and stained with 5% aqueous silver nitrate (Sigma-Aldrich, USA) solution under UV light for 30 min, rinsed in de-ionized water, immersed in 5% sodium thiosulfate solution (Sigma-Aldrich, USA) for 5 min RT. Samples were counterstained with hematoxylin. Images were acquired using Eclipse 90i microscope (Nikon Instruments Europe BV).

**SEM and element analysis**

BM-hMSCs were seeded on the scaffolds at 10⁶ cells/mL and were fixed in paraformaldehyde 4% for 1 h at 4°C.
Table 1. Composition tensile and compressive properties of G-CH polymeric scaffolds.

<table>
<thead>
<tr>
<th>Sample</th>
<th>Composition (%)</th>
<th>Tensile properties</th>
<th>Compressive properties</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>G</td>
<td>PEG</td>
<td>CH</td>
</tr>
<tr>
<td>G-CH1</td>
<td>74.3</td>
<td>17.6</td>
<td>8.1</td>
</tr>
<tr>
<td>G-CH2</td>
<td>68.8</td>
<td>16.3</td>
<td>14.9</td>
</tr>
</tbody>
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G: gelatin; PEG: polyethylene glycol; CH: chitosan.

G-CH hybrid hydrogels

Two different polymeric scaffolds namely G-CH1 and G-CH2 having a CH content of 8.1% and 14.9 wt%, respectively (Table 1), were prepared using a green, simple and easy synthetic method under aqueous condition without using any solvents, chemical reagents such as catalysts, coupling or foaming agents.

Functionalized PEG was used as a crosslinking agent, and the cross-linking process involved mainly end epoxide groups of PEG and amino group of G and CH chains. The final well interconnected macroporous scaffold structures were obtained by ice segregation induced self-assembly, which consists in the freezing of the hydrogel into liquid nitrogen followed by freeze-drying. During freezing by liquid nitrogen, ice crystals formed, and they acted as templates to create the macroporous structure after freeze drying. Hydrogel morphology was analyzed by SEM.

The image of cross-sectional views reported in Figure 1(a) clearly showed a highly interconnected porous network morphology with macro and micro-porosity, channels and interstices of different size, well interconnected, and homogeneously distributed.

Results of tensile and compressive tests performed on wet specimens (Table 1) are compatible with stable high crosslinked hydrogels. The tensile modulus, tensile strength and elongation at break of the G-CH1 scaffold were found to be 0.68 MPa, 0.25 MPa and 29%, respectively. Increasing amount of CH into the G-CH network reduced stiffness (0.28 MPa) and strength (0.08 MPa), producing a softer material.

To investigate the mechanical stability and reversibility of the hydrogels, consecutive cyclic loading–unloading compressive tests with up to 50% of strain were done on wet cubic specimens and the value of elastic modulus and strength are reported in Table 1. During the first cycles, the hydrogels showed a Mullins effect, with the reduction of stiffness and stress at every cycle. After few cycles, the material behavior was stabilized (Figure 1(b)) and the hydrogels were able to sustain a compressive strain of 50% with full strain recovery. Both the hydrogels maintained their original shape and their load bearing capability up to such a high level of deformation. Although many microscopic flaws might have been created within the hydrogel, no macroscopic crack propagated through the sample; these broken bridges did not lead to macroscopic fracture of the hydrogel.

Controlled degradability of materials without sacrificing their structural stability and mechanical integrity is important for long-term service of hydrogels. Degradation significantly affects structural stability, swelling ratio, and mechanical integrity. Therefore, we investigated hydrogel hydrolytic degradation by measuring mass loss, swelling ratio, and mechanical stiffness/strength at predetermined time of 1, 10, and 21 days. Figure 1(c) shows the mass loss and swelling ratio as a function of degradation time over a...
Figure 1. (a) SEM micrographs showing different magnifications of G-CH1 and G-CH2. (b) Cyclic compressive stress–strain curves of G-CH1 and G-CH2 for up to 10 consecutive cycles at 50% maximum strain. (c) Swelling ratio and mass loss trends during degradation period of G-CH1 (continuous lines) and G-CH2 (dashed lines). (d) Representative cyclic stress–strain curves obtained during degradation of hydrogels G-CH1 and G-CH2. (e) Mechanical assessment of hydrogels during degradation time of 1, 10, and 21 days: variation of modulus and strength for G-CH1, and corresponding change of modulus and strength for G-CH2.
period of 3 weeks. As expected, mass loss and swelling ratio increased over the course of the experiment. Noteworthy, G, CH, and PEG as well as hydrogels originated from their grafting and crosslinking contain hydrolytically vulnerable bonds resulting in hydrolysis, breaking down the network, dissolving/leaching out the fragmented soluble fractions, and inducing mass loss, while in turn making more voids available for water and, consequently, increasing swelling ratio. Both hydrogels lost approximately 30% of their initial masses after 3 weeks. Meanwhile, swelling ratio increased by 50% and 40% of their initial values for G-CH1 and G-CH2, respectively (Figure 1(c)).

We also monitored mechanical stiffness and strength over degradation time. Representative cyclic stress–strain curves obtained during degradation of hydrogels (Figure 1(d)) demonstrated the structural and mechanical stability as well as good reversibility even after 21 days of degradation. A decreasing trend of moduli and strengths was observed over time for both hydrogels suggesting gradually breaking down of cross-linked networks and consequent mass losses. The relative change of modulus and strength over the degradation period showed that hydrogels retained about 45% of their initial stiffness and strength after 3 weeks of degradation (Figure 1(e)). In addition, hydrogels lost mass during degradation time, leaving available spaces to accommodate the growing new cells and deposition of ECM.

**Cells adhered well and maintained viability**

BM-hMSCs and AT-hMSCs were seeded in the G-CH hydrogels and after 2 and 21 days in complete medium FBS or hPL, SEM was performed to evaluate adhesion and morphology of the cells. After 2 days, adhered cells showed a spread morphology and were anchored by filopodia to the surface of the pores within the materials. Staining of nuclei (DAPI) showed uniform distribution of the cells in the hydrogels (Figure 2(a)).

After 21 days in complete medium FBS or hPL, cells viability in the scaffolds was evaluated in three replicates using a Live/Dead assay. BM-hMSCs and AT-hMSCs in the G-CH1 and G-CH2 hydrogels were viable and distributed homogenously after 21 days of culture. Specifically, cell viability within the hydrogels ranged between 75% and 90% in all conditions, independently of the presence of either hPL or FBS, of the content of CH in the scaffold, and of cell type. No significant difference of BM-hMSCs viability were noted in both the two types of hydrogels and culture media (G-CH1 + FBS, 87.79% ± 6.80 and G-CH1 + hPL, 83.18% ± 9.41, G-CH2 + FBS, 84.75% ± 5.60 and G-CH2 + hPL, 77.75% ± 7.63) and of AT-hMSCs viability in both the two types of hydrogels and culture media (G-CH1 + FBS, 79.26% ± 16.34 and G-CH1 + hPL, 85.81% ± 9.93, G-CH2 + FBS, 82.96% ± 7.85 and G-CH2 + hPL, 84.44% ± 6.31) (Figure 2(b), Supplemental Figure S1).

**G-CH1 and hPL support cell proliferation**

Cell proliferation was measured by quantifying viable cells number within the G-CH hydrogels (by the CCK8 assay) at different time points after cell seeding. We found that both BM-hMSCs and AT-hMSCs proliferated over time, reaching a maximum of cell number after 21 days of culture. In particular, G-CH1 in complete medium with hPL significantly supported enhanced cell proliferation, especially for BM-hMSCs after 10 days (G-CH1 + hPL vs G-CH2 + hPL, p = 0.0460, G-CH1 + hPL vs G-CH1 + FBS, p = 0.0099, G-CH1 + hPL vs G-CH2 + FBS, p = 0.0017), 14 days of culture (G-CH1 + hPL vs G-CH1 + FBS, p = 0.0270, G-CH1 + hPL vs G-CH2 + FBS, p < 0.0001) and 21 days (G-CH1 + hPL vs G-CH2 + hPL, p = 0.0105).

Also G-CH2 in complete medium with hPL significantly supported enhanced proliferation of BM-hMSCs (G-CH2 + hPL vs G-CH2 + FBS, p < 0.0001).

With respect to AT-hMSCs, hPL also significantly increased cell numbers after 10 days of culture (G-CH1 + hPL vs G-CH1 + FBS, p = 0.0406, G-CH1 + hPL vs G-CH2 + FBS, p = 0.0149, G-CH2 + hPL vs G-CH2 + FBS, p = 0.0312) (Figure 3(a)).

**Scaffolds and hPL contribute to osteogenic differentiation**

Then, osteogenic differentiation of BM-hMSCs and AT-hMSCs in the G-CH1 and G-CH2 hydrogels under different culturing conditions was tested. Immunofluorescence analysis for OCN was performed and the results showed that both hMSCs types used in this study could undergo differentiation to the osteogenic lineage when seeded within the hydrogel scaffolds in differentiating conditions (Figure 3(b) and (c)). Staining for OCN after 21 days of culture revealed that in the presence of FBS no differentiation occurred independently of the scaffold or cell type if no osteogenic medium was added (Figure 3(c), GM + FBS).

On the other hand, the addition of hPL significantly favored osteogenic differentiation in the absence of inducers (BM-hMSCs, G-CH1, GM + hPL vs G-CH1, GM + FBS, p = 0.0073 and G-CH2, GM + hPL vs G-CH2, GM + FBS, p = 0.0004; AT-hMSCs, G-CH1, GM + hPL vs G-CH1, GM + FBS, p < 0.0001 and G-CH2, GM + hPL vs G-CH2, GM + FBS, p = 0.0080). The effect of hPL on differentiation was significantly stronger for AT than BM derived hMSCs, where, for example, for G-CH2, hPL alone allowed for a twofold increase of OCN compared to GM with FBS (AT-hMSCs, G-CH2, GM + hPL vs G-CH2, GM + FBS, p = 0.0080), and for a threefold increase in the presence of osteogenic inducers (AT-hMSCs, G-CH2, OM + hPL vs G-CH2, OM + FBS, p = 0.0005). Moreover, for G-CH1 seeded with AT-hMSCs, the effect of hPL alone allowed for a twofold increase of OCN compared to the effect of hPL in the
The presence of osteogenic stimuli (AT-hMSCs, GM + hPL vs G-CH1, OM + hPL, p = 0.0064).

After 21 days, culture w/o osteogenic differentiation stimuli, the cells were still adherent and their spreading and migration into the inner regions of the scaffold were remarkable, because of the pore size and open porosity of both hydrogels (Supplemental Figure S2). SEM micrographs showed that BM-hMSCs adopted a more fully elongated and packed morphology in flattened sheet and deposited more ECM fibers in the pores of both the G-CH1 and G-CH2 hydrogels (Supplemental Figure S2). On the other hand, after 21 days of osteogenic differentiation, both scaffolds supported mineralization, as revealed by the presence of mineral deposits detected via SEM (Figure 4(a), Supplemental Figure S2).

**Analysis of cell mineralization**

Von Kossa staining confirmed the presence of mineralized positive area mainly localized on the scaffolds treated with OM (Figure 4(b)). SEM-EDX analysis was performed to quantify individual elements particularly focusing on calcium and phosphorus typically present in mineral deposits formed during osteogenic differentiation. EDX characterization was performed on both the G-CH hydrogels and in three distinct areas of the same sample after 21 days of osteogenic differentiation focusing on calcium and phosphorous elements. The EDX spectra associated with SEM analysis are shown in Supplemental Figure S3, from which weight and atomic ratios of elements in minerals deposited by hMSCs were determined for quantitative comparison.
Figure 3. (a) Proliferation of BM-hMSCs and AT-hMSCs cultivated in G-CH hydrogels measured by the CCK8 assay in the complete medium FBS or complete medium hPL at different time points (2, 6, 10, 14, and 21 days). Two-way ANOVA was used for statistical analysis. Data show the average values and their significant differences (*p ≤ 0.05, **p ≤ 0.01, ***p ≤ 0.001, ****p ≤ 0.0001). (b) Pictures of immunofluorescence of OCN expression by BM-hMSCs and AT-hMSCs cultured at 21 days in G-CH1 and G-CH2 hydrogels in the complete medium hPL (GM + hPL). Scale bar: 100 µm. (c) Graphs of immunofluorescence analysis of OCN expression by BM-hMSCs and AT-hMSCs cultured at 21 days in G-CH1 and G-CH2 hydrogels in the complete medium FBS (GM + FBS) or complete medium hPL (GM + hPL) or osteogenic medium FBS (OM + FBS) or osteogenic medium hPL (OM + hPL). One-way ANOVA was used for statistical analysis of the fluorescence intensity on BM-hMSCs and AT-hMSCs cultured at 21 days in G-CH1 and G-CH2 hydrogels. Data show the average values and their significant differences (*p ≤ 0.05, **p ≤ 0.01, ***p ≤ 0.001, ****p ≤ 0.0001).
EDX analysis of the constructs showed evidence of calcium phosphate crystals, a sign of mineralized matrix (Ca:P in distinct areas Supplemental Figure S3), when cells were cultured in OM with hPL. On the other hand, no calcium phosphates were observed after culture in complete medium hPL (Supplemental Figure S3). Furthermore, the Ca/P weight and atomic ratios in G-CH1 and G-CH2 were between 1.5 and 2.0 and close to that of bone, whose mineral primarily consists of Ca and P with a Ca/P ratio from 1.4–1.7.30 As shown in the table, on G-CH1 Ca/P weight and atomic ratios (1.58/1.22; 1.92/1.48; 2.15/1.66) were closer to the hydroxyapatite (HA; 2.22/1.66) and tricalcium phosphate (2.0/1.5) ones compared to G-CH2 (1.48/1.15; 1.54/1.19; 1.28/1.00) (Figure 4(c)). These results suggest that our G-CH hydrogel systems were able to mimic the physiological environment showing mineralization already detectable at day 21 using OM with hPL, but not using hPL alone.

**Proteomic characterization of hPL and PPP highlight factors involved in cell stimulating effects**

About 500 different protein IDs were identified in hPL (n = 501) and PPP (n = 491), respectively, considering both not depleted and depleted samples (Figure 5(a)). An increase in identification power was noticed in depleted plasma derivatives with respect to their non depleted counterpart, ranging from 15% to 21%.

In the depleted specimens, about 40% (169 IDs) of the proteins detected in hPL were not observed in the PPP, while in not depleted samples 36% (130 IDs) were present only in hPL (Figure 5(a)). In particular, combining the lists of protein IDs in different sample groups, 59 were uniquely present in hPL and not in common with PPP independently from the depletion process (Figure 5(b)). The related enriched biological processes and KEGG pathways, of these 59 proteins, were obtained using STRING26 and are listed in Supplemental Table 1 and in Supplemental Figure S4. Interestingly, the majority of the 59 molecules are implicated in stress response and wound healing (56% and 46% of all proteins present in the panel, respectively), platelet and cell activation (36% and 41%, respectively), cell differentiation (31%) and cell adhesion (29%) (Supplemental Table 1). It has been also observed that some of the proteins specific of hPL are involved in several pathways that are likely to be strictly linked the promotion of repair mechanisms, such as signal transduction (as RAP-1 and HIF-1 signaling pathway), cell adhesion and motility, and carbohydrate metabolisms (Supplemental Figure S4). The more critical nodes in the related networks, moreover, are represented by key proteins typical of the platelet degranulation (as ACTN1, TLN-1, PF4, VCL, MMRN1, ITGA2B, and ITGB3), neutrophil degranulation (as RAP1B), cell-matrix interaction (as PXN, ITGB1, ILX), and also osteoblast differentiation and bone development (as TPM4 and SPARC) (Supplemental Figure S4).

In addition to those proteins identified exclusively in hPL, label-free relative quantification pinpointed 7 (Supplemental Figure S5) and 23 (Supplemental Figure S6) proteins significantly altered in their abundances comparing hPL vs PPP in not depleted and in depleted samples, respectively (fold change \( \geq 1.5; p < 0.05; \geq 2 \) unique peptides; in at least 4 replicates). Three of these differences were in common and up-represented in hPL, including platelet basic protein, a factor known to be involved in the regulation of cell proliferation and observed to be more than 20 fold more concentrated in hPL vs PPP both in depleted and not depleted samples. Enrichment of the biological processes and networks that
involve de-regulated proteins in hPL compared to PPP for both depleted and not depleted samples are illustrated in Supplemental Figure S7.

Discussion

Tissue engineering using biomaterials or scaffolds as temporary matrices for hMSCs proliferation and osteochondral differentiation is a promising strategy for tissue repair of bone or cartilage. Using a freeze-drying and post-curing technique, two biocompatible CH-G hybrid hydrogels with different CH content, G-CH1 (8.1%) and G-CH2 (14.9%), were developed, with the aim to produce scaffolds with mechanical and functional properties adequate for bone repair (Figure 1).

Scaffold stiffness is a relevant property for cell proliferation, osteogenic differentiation, and for the maintenance of structural integrity during biodegradation. The hydrogels obtained using this simple and solvent/additive free preparation technique, showed improved structural and physico-mechanical properties compared to literature. They showed suitable structural stability, good swellability, porosity adequate to the cell size and high pore interconnectivity. Increasing amount of CH into the G-CH network reduced the stiffness and strength leading to a softer material (Table 1). Both hydrogels exhibited viscoelastic behavior, they maintained their original shape and showed no sign of mechanical failure under compressive loading–unloading cycles at relatively high deformation level (Figure 1(b)). During hydrolytic degradation, after 3 weeks, hydrogels retained their structural stability and elasticity, while stiffness and strength were reduced to half of their initial values (Figure 1(c)). Moreover, during degradation, both hydrogels lost approximately 30% of their initial masses after 3 weeks, while swelling ratio increased leaving spaces available to allow the growth of new cells and the deposition of ECM (Figure 1(c)). Thus, physico-mechanical properties and CH content can be modulated by this preparation process developed ad hoc to obtain high performance systems and in turn to adapt the scaffolds to the potential use in vivo.

CH is a biocompatible, biodegradable material with intrinsic antibacterial activity that we used in this study to favor adhesion, proliferation, and bone formation of BM/AT-hMSCs. Many efforts have been attempted to better support these cell functions, by adding or incorporating CH with other components, such as glycerol phosphate, poly-N-isopropylacrylamide, lactide, poly-ethylene oxide, polyethylene glycol, and poly-L-lactic acid.
or nano-particles. However, these studies did show neither a clear advantage of a specific combination of biomaterials nor a definition of the optimal concentration of CH in these hydrogel systems.

High interconnectivity and optimal porosity of both produced scaffolds allowed cell colonization of the entire structure even in static conditions without the use of a bioreactor (Figure 2). These characteristics were not fully matched in other hydrogel systems and the cell infiltration procedure was optimized using a cold static seeding method instead of the “classic” static seeding method at 37°C.

With attention to potential clinical translatable, appropriate porosity and interconnectivity are known to be important features, because they allow not only the optimal adhesion and proliferation of MSCs but also transport of nutrients and the interaction of cells with glycosaminoglycans, proteoglycans, cytokines, and GFs. It is known from laboratory practice that during the initial seeding phase, a part of the cells leaves the scaffold and they adhere to the culture well. Therefore, we moved the samples to new culture plates after a short incubation time. Consequently, proliferation results from CCK8 assay show that the number of cells proliferating after 2 days was lower compared to the number of cells added during the initial seeding phase (Figure 3(a)). However, proliferation was considerable from day 2 to day 6 of culture, in accordance with previous studies.

Interestingly, AT-hMSCs grew exponentially until reaching a plateau-phase after 10 days, which was possibly followed by cell death after 14 days. Then, cells resumed their growth and their number increased again after 21 days. This was not observed in BM-hMSCs, where an earlier onset of the osteogenic differentiation could be expected in comparison to AT-hMSCs (Figure 3(c)), limiting their proliferative potential. Hence, BM-derived cells proliferated more slowly and didn’t experience such a decrease in cell numbers after an initial exponential growth.

As confirmed by the viability assay, all the cells anchored to the surface of the hydrogels were viable and proliferating as desired. Particularly, no significant difference of BM-AT-hMSCs viability was observed in the two types of hydrogels with FBS or hPL. In all, 80%–90% of BM-AT-hMSCs were viable, suggesting that there was no significant limitation in the nutrient supply in all the conditions (Figure 2(b), Supplemental Figure S1).

The cell culture procedures were performed with FBS in comparison with hPL, and we found that hPL was not only able to stimulate cell growth and proliferation but also osteo-differentiation of BM/AT-hMSCs per se. Results from immunostaining analysis showed that BM-hMSCs and AT-hMSCs cultured in both the G-CH hybrid hydrogels were efficiently induced to differentiate toward the osteogenic lineage not only in the presence of osteogenic inducers but also using hPL alone in the absence of osteo-inductive media for both scaffolds and cell types and after 3 weeks of culture (Figure 3(b) and (c)). Interestingly, even if the differentiation levels of both cell types are similar, hPL has a stronger osteogenic effect on AT-derived cells compared to BM-derived cells. Indeed, AT-hMSCs have a lower level of osteogenic phenotype in GM with FBS compared to BM-hMSCs. This could be explained by a possible specific effect of the particular 3D environment, especially the G-CH1 hydrogel that preserves stemness of the AT-derived cells when no hPL or osteostimulants are added. Upon their addition, there is a two or even threefold increase in OCN levels. A role can also be played by the different source of these cells and their predisposition to differentiate into different lineages with different efficacy.

Our data are in line with previous literature where platelet GF cocktail was used for the ex-vivo expansion of hMSCs, showing significantly better results compared to previous strategies where a single GF was used. To better understand the positive effects of hPL through stimulation of multiple signaling pathways, we then investigated the proteomic composition of hPL by label-free nLC-MS/MS. Shotgun analysis and label-free quantification of hPL and PPP allowed to “proteomically” characterize the supplement and to highlight possible candidates responsible for the beneficial effects on the MSCs proliferation and osteo-genic differentiation. In all, 59 proteins were exclusively present in hPL and not in PPP independently from the sample processing methods (Figure 5(b)), whereas 7 and 23 proteins were observed significantly altered in their abundances when comparing hPL vs PPP in not depleted and in depleted samples, respectively (Supplemental Figure S5, Supplemental Figure S6). This specificity suggests that the list of these proteins (85 different IDs in total) could represent a possible panel of factors involved in cell stimulating effect of this supplement. In particular, CXCL7 (platelet basic protein-PBPB) a positive regulator of cell proliferation was found strongly up-represented in hPL both in depleted and not depleted samples (ratio > 20). Functional annotation tools allowed to highlight the biological processes, pathways, and networks enriched in these panels of proteins specific of hPL, not present or less concentrated in PPP (Supplemental Figure S4, Supplemental Figure S6, Supplemental Table 1). Most of these functional signatures relate to platelet degranulation, wound healing, and cell adhesion, but an involvement of other specialized functions related with the cellular repairing mechanisms and the regulation of cell differentiation was also observed, suggesting a high complexity of the molecular scenario. It also cannot be excluded that the positive proliferation and differentiation effects of hPL per se could be influenced by the interaction of hPL with the G-CH hydrogel constituents.

Indeed, we found that G-CH hydrogels, particularly G-CH1, in presence of complete medium hPL, have a great potential in terms of supporting BM-hMSC and AT-hMSC proliferation and guide them to differentiate toward the osteogenic lineage.
To further investigate the ability of our G-CH hydrogel systems to mimic the physiological environment and support bone regeneration, elemental analysis was performed and the results showed that the mineralization process was already detectable at day 21 in the presence of osteogenic medium hPL but not using hPL alone. While results from staining for the osteogenic marker OCN showed that hPL alone induces osteogenic differentiation of cells, there is no evidence of mineralization in complete medium with hPL after 21 days of culture from SEM/EDX analysis (Supplemental Figure S3). Early mineralization is only detected after adding osteogenic stimuli in the culture medium. Further analyses after four or more weeks of culture are then needed to investigate the mineralization processes supported by hPL in the absence of induction media. In fact, generally 3 weeks are necessary to lead hMSCs toward the osteogenic differentiation while almost 4 weeks or more are required for mineralization to occur.53–55

Inorganic phosphates on cell surface detected via SEM could be deemed as prime evidence for osteo-genic differentiation of hMSCs and this deposition of calcium phosphate was crucial to scaffold mineralization processes (Supplemental Figure S2). The Ca/P weight and atomic ratios supported the presence of HA, suggesting a marked ability of the G-CH hydrogels to mimic the physiological environment. Interestingly, G-CH1 characterized by less content of CH was more advantageous for cell growth and scaffold mineralization. Indeed, G-CH1 supported a significantly higher cell proliferation, particularly for BM-hMSCs (Figure 3(a)), as well as a better cell differentiation, confirmed via Von Kossa staining showing mineralization (Figure 4(b)).

In summary, our results demonstrated that both G-CH1 and G-CH2 synthetic hydrogel scaffolds are biocompatible and bioreabsorbable and their chemical, morphological, and mechanical properties are suitable to support cells growth, osteo-differentiation, and mineralization, and their potential application for bone regeneration. Moreover, we show that a cell culture procedure using hPL instead of FBS, besides avoiding the risks of transmitting animal diseases, contains crucial mitogenic and chemotactic molecules that promote cell differentiation and tissue repair. Hence, these scaffolds in combination with hPL have the potential to provide a clinical grade biomedical device. hPL has the potential to become the gold standard supplement for hMSC propagation in regenerative medicine, in addition to already commercialized products,18,19,56,57 even if a standardization of protocols has to be investigated yet.

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