

Scaffold-Based Delivery of a Clinically Relevant Anti-Angiogenic Drug Promotes the Formation of *In Vivo* Stable Cartilage

Matteo Centola, PhD,^{1,2} Franca Abbruzzese, MSc,¹ Celeste Scotti, MD,² Andrea Barbero, PhD,² Gianluca Vadalà, MD,³ Vincenzo Denaro, MD,³ Ivan Martin, PhD,² Marcella Trombetta, PhD,¹ Alberto Rainer, PhD,¹ and Anna Marsano, PhD²

Standard cartilage tissue engineering approaches, for example, matrix-induced autologous chondrocyte implantation (MACI), consist of the implantation of cell-based constructs whose survival and further development first depend on the degree of graft maturity at the time of surgery (e.g., matrix production) and, subsequently, on initial host reaction. Indeed, blood vessel ingrowth and macrophage migration within the implant may endanger graft stability of immature constructs; so, control of angiogenesis was proposed as an adjuvant of cellular therapy for the treatment of cartilage defects. In this study, we hypothesized that engineered constructs with no *in vitro* precultivation, but functionalized to block angiogenesis right on implantation, might result in better survival, as well as superior long-term cartilaginous quality. Here, we propose a clinically compatible fibrin/hyaluronan scaffold seeded with nasal chondrocytes (NC) and functionalized with an FDA-approved anti-angiogenic drug (bevacizumab; Avastin[®]), which sequesters vascular endothelial growth factor from the surrounding environment. Our results show that the sustained bevacizumab release from NC-loaded scaffolds after subcutaneous implantation in nude mice efficiently blocked host vessels ingrowth (five times lower CD31⁺ cells infiltration vs. control group, at 3 weeks after implant), and enhanced constructs survival rate (75% vs. 18% for the control, at 6 weeks after implant). *In vitro* assays, developed to elucidate the role of specific construct components in the *in vivo* remodeling, allowed to determine that fibrin degradation products enhanced the *in vitro* endothelial cell proliferation, as well as the macrophage migration; whereas the presence of bevacizumab was capable of counteracting these effects. The proposed pharmacological control of angiogenesis by a therapeutic drug released from a scaffold might enhance cartilage regeneration by MACI approaches, possibly allowing it to bypass the complex and costly phase of graft preculture to gain increased functionality.

Introduction

DAMAGED ARTICULAR CARTILAGE has a limited capacity of self-repair due to its avascular nature and low cellular mitotic activity.^{1–4} Cell-based repair techniques, such as matrix-induced autologous chondrocyte implantation, showed positive clinical outcomes, despite the formation of a fibrocartilaginous/fibrous repair tissue that is characterized by inferior mechanical properties and limited durability.⁵ Most likely, the lack of essential extracellular matrix (ECM) components, including high-molecular-weight hyaluronic acid, and other anti-angiogenic factors—such as chondromodulin, endostatin, and angiostatin⁶—exposes the not fully mature engineered tissues to an early blood vessel in-

vasion. Such a host reaction might lead to final poor cartilaginous quality⁷ and, eventually, to premature implant degradation. Thus, control of angiogenesis might be crucial for both the development and the maintenance of physiological articular cartilage.⁸ In particular, it has been demonstrated that vascular endothelial growth factor (VEGF), one of the most potent angiogenic factors, plays an essential role in the ossification process at the level of the growth plate, modulating cartilage vascularization and hypertrophy.⁹ Recent data also reveal that chondrocyte-derived VEGF promotes catabolic pathways in the osteoarthritic cartilage.¹⁰

Neo-angiogenesis is also accompanied by the massive infiltration of mononuclear cells, such as monocytes. VEGF acts as a powerful chemoattractant for monocytes,¹¹ which

¹Tissue Engineering Laboratory, Center for Integrated Research, Università Campus Bio-Medico di Roma, Rome, Italy.

²Departments of Surgery and of Biomedicine, University Hospital Basel, Basel, Switzerland.

³Area of Orthopedics and Trauma Surgery, Center for Integrated Research, Università Campus Bio-Medico di Roma, Rome, Italy.

could potentially lead to a fast macrophage-driven *in vivo* resorption of the implanted engineered cartilage. Taken together, these aspects strongly underline the importance to control angiogenesis, and, in particular, the signaling of VEGF in cartilage tissue engineering (CTE). To this extent, cell-based anti-angiogenic gene therapies for cartilage regeneration have been already successfully investigated by inducing overexpression of either endostatin^{12,13} or chondromodulin.¹⁴ In addition, overexpression of soluble VEGF receptor-1 combined with the release of growth factors belonging to the transforming growth factor beta (TGF- β) superfamily enhanced cartilage regeneration in both rat osteoarthritic¹⁰ and osteochondral defect models.¹⁵

We hypothesized that VEGF blockade by using a biomaterial-based anti-angiogenic drug release system could provide—right upon implantation—an appropriate environment for the formation of stable cartilage by freshly seeded engineered constructs. In particular, we developed a hyaluronan/fibrin-based porous scaffold which was functionalized by the incorporation of a humanized monoclonal anti-VEGF antibody (bevacizumab)¹⁶ that binds to human VEGF¹⁷ and is currently used as an anti-angiogenic therapeutic drug in the treatment of metastatic colorectal cancer, metastatic kidney cancer, and glioblastoma. The use of a drug-eluting scaffold would overcome the limitations of gene therapy in terms of a direct clinical translation.¹⁸

High-molecular-weight hyaluronan and fibrin were chosen in virtue of their biocompatibility, chondro-supportive nature,^{1,4} and extensive clinical use.^{2,19–21}

Among the promising cell sources for CTE, we opted for nasal chondrocytes (NC), as they represent one of the most interesting candidates for clinical application in virtue of (1) the relative ease and low morbidity of the harvest procedure²²; (2) a better retained capacity on cell expansion to re-differentiate and generate hyaline-like tissue²³ as compared with chondrocytes of other origin; and (3) their capacity to properly respond to mechanical forces which are typically associated with joint loading.²⁴

Despite the orthotopic model being a more clinically relevant approach, in this study, we decided to use a subcutaneous implantation in nude mice, as it represents the most efficacious model that is used for testing the intrinsic capacity of constructs to form stable cartilage tissue,⁷ being characterized by a more vascularized and hostile microenvironment and, therefore, representing a more arduous testing ground for our purposes.

Materials and Methods

All reagents were purchased from Sigma Aldrich, unless otherwise stated, and were used without further purification. Culture media and supplements were from Gibco (Invitrogen).

Bevacizumab activity and dosage

A set of preliminary experiments was performed on human umbilical vein endothelial cells (HUVEC) in order to determine the suitable bevacizumab concentration to be loaded into the scaffolds. HUVEC were cultured at a density of 5000 cells/well overnight in growth medium (GM, M199 supplemented with 20% fetal bovine serum [FBS], 100 μ g/mL endothelial cells growth supplement, 50 IU/mL heparin, 100 IU/mL penicillin, and 100 μ g/mL streptomycin). The

next day, GM was replaced with low-FBS medium (assay medium [AM], consisting of M199 with 5% FBS, 10 IU/mL heparin, and 1% penicillin/streptomycin), supplemented with recombinant human VEGF (R&D Systems) at different titers between 0 and 10 ng/mL. Bevacizumab (Avastin[®]; Roche) at different concentrations in the 1–5 μ g/mL range was added to AM.

After 2 days, HUVEC metabolic activity was measured by MTS assay (Cell Titer 96[®]; Promega).

Scaffold preparation

High-molecular-weight sodium hyaluronate (10 mg/mL) and fibrinogen (20 mg/mL) were separately dissolved in saline (0.9% w/v of NaCl) and mixed using the same volume.²⁵ Aprotinin (3000 KIU per mL of solution) and fibrin-stabilizing factor XIIIa (Abnova; 50 ng per mg fibrinogen) were added to the solution under mild stirring. Addition of bevacizumab was performed at two different concentrations with regard to the total volume, selected on the basis of the dosage experiments described earlier: 3.75 μ g/mL (hereinafter named HA-FIB-B3.75) and 5 μ g/mL (HA-FIB-B5). Samples without bevacizumab were also synthesized (HA-FIB). Polymerization of fibrinogen was achieved by the addition of a 100 IU/mL thrombin solution (0.5 IU per mg fibrinogen). Solution was transferred into a 96-well plate and incubated at 37°C for 30 min until gelation. Cylindrical porous scaffolds (6 mm diameter and 8 mm height) were obtained after freeze drying the gels.

An additional set of materials without hyaluronan, either supplemented (hereinafter FIB-B3.75) or not (FIB) with 3.75 μ g/mL bevacizumab, was synthesized in the same experimental conditions to be used as a control for further *in vitro* assays.

Degradation and water uptake assays

Weight loss of the scaffolds was monitored as a function of incubation time in cell culture medium. Scaffolds ($n=3$ for each experimental condition, weighing ~ 10 mg each) were incubated at 37°C into sealed tubes containing 1 mL of Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% FBS. Specimens were retrieved at selected time points for approximately 3 weeks, blot dried, and weighed (W_t). Medium was replaced every 2 days. Scaffold weight loss ratio was defined as $(W_0 - W_t)/W_0\%$, where W_0 is the initial wet weight and W_t is the wet weight at a given time point. Scaffold water uptake ratio was calculated as $UR = (W_{wt} - W_{d0})/W_{d0}$, where W_{d0} is the scaffold initial dry weight and W_{wt} is its wet weight after 1 h incubation in saline solution at 37°C.²⁶

Scanning electron microscopy and mercury intrusion porosimetry

Gold-sputtered scaffold specimens underwent microstructural investigation by Field Emission Scanning Electron Microscopy (FE-SEM, Leo Supra 1535; LEO Electron Microscopy). Scaffold pore size distribution was determined by mercury intrusion porosimetry (Porosimeter 2000; Carlo Erba Instruments) according to the Washburn equation:

$$p \cdot r = -2 \cdot \gamma \cdot \cos \theta,$$

where p is the applied pressure, r is the pore radius, γ is the surface tension of mercury (480 mN/m), and θ (141.3°) is the contact angle of mercury.

Compressive mechanical test

Previously, hydrated scaffolds were tested in triplicate under unconfined compression using a mechanical testing system equipped with a 10 N load cell (model 3365; Instron) at a strain rate of 0.05 min⁻¹. The compressive modulus (E^*) was calculated as the slope in the linear range of the stress-strain curve.^{26,27}

ELISA quantification of bevacizumab release

HA-FIB-B3.75 and HA-FIB-B5 constructs were incubated in DMEM at 37°C and 5% CO₂, and supernatants were withdrawn at fixed time points, for approximately 3 weeks. 96-well plates were incubated overnight at 4°C with 50 µL of supernatants, and solutions with different concentrations of bevacizumab in order to obtain the calibration curve. Unspecific protein-binding sites were saturated by 2 h incubation at room temperature (RT) with phosphate-buffered saline supplemented with 0.05% Tween 20 and 1% bovine serum albumin. After rinsing, horseradish peroxidase-conjugated anti-mouse IgG (H+L) antibody (Southern Biotech; dilution 1:3000) was added for 90 min at RT followed by o-phenylenediamine dihydrochloride substrate (Sigma Aldrich). The reaction was stopped by adding 10% sulfuric acid and, after incubation for 30 min, absorbance was read at 492 nm on a microplate reader (Infinite M200; Tecan).

Cell isolation and expansion

Biopsies were harvested from the nasal cartilage septum of four cadavers (mean age: 51 years; range 31–84 years), following informed consent by relatives and in accordance with the Local Ethical Committee. NC were isolated by digesting minced cartilage in 0.15% type II collagenase for 22 h at 37°C. NC were expanded in DMEM containing 10% FBS, 4.5 mg/mL D-glucose, 0.1 mM nonessential amino acids, 1 mM sodium pyruvate, 100 mM HEPES buffer, 100 IU/mL penicillin, 100 µg/mL streptomycin, and 0.29 mg/mL L-glutamate (complete medium) supplemented with 1 ng/mL of TGF-β1 (R&D Systems) and 5 ng/mL of fibroblast growth factor-2 (FGF-2; R&D Systems)²⁴ up to passage 3.

Cytotoxicity test

Cytotoxicity tests were performed on scaffolds seeded with NC (5 × 10³ cells/well) using Vybrant Cytotoxicity Assay Kit[®] (Life Technologies) at 4, 8, and 24 h according to the manufacturer's protocol.

HUVEC proliferation assay

Scaffolds were incubated in complete medium. Supernatants were timely collected for approximately 7 days of incubation at 37°C and 5% CO₂ and stored at -20°C until analysis.

HUVEC proliferation assays ($n=3$) were performed as previously described,²⁸ adding to VEGF-supplemented AM (0–10 ng/mL VEGF) the previously collected supernatants (mixed using the same volume). Control experiments were performed by supplementing AM with bevacizumab (1.5 or

3.75 µg/mL) or with high-molecular-weight HA (500 µg/mL). After 2 days, HUVEC metabolic activity was measured by MTS assay (Cell Titer 96[®]; Promega). Data were normalized versus proliferation of HUVEC cultured in GM.

Monocyte migration assay

Scaffolds were incubated in serum-free medium (SFM) for approximately 7 days at 37°C and 5% CO₂. Supernatants were timely collected and stored at -20°C until analysis. Peripheral blood mononuclear cells were isolated from peripheral blood of healthy donors ($n=2$) by gradient centrifugation. Monocytes were purified using the MACS CD14 isolation kit (Miltenyi Biotec), according to the manufacturer's protocol. CD14⁺ monocytes were cultured in the upper chamber of an HTS-Transwell-24-well microplate (Corning) in SFM and incubated with the previously collected supernatants loaded in the lower chamber for 20 h at 37°C and 5% CO₂.²⁹ SFM and SFM supplemented with 10 ng/mL of VEGF¹¹ were used as negative and positive controls, respectively. SFM with 10 ng/mL VEGF and bevacizumab (at the optimal stoichiometric ratio of 2.6:1 with regard to VEGF¹⁶) was used as an additional control. Cell migration was measured in terms of DNA quantification using CyQuant[®] cell proliferation assay kit (Life Technologies) and expressed as the percentage of the migrated monocytes of a given experimental condition with regard to the SFM group.

In vitro pellet culture model

Chondrogenic re-differentiation capacity of two NC primaries (at least three replicates per condition per assay) was tested in a pellet culture model.³⁰ Briefly, pellets were generated by 5 × 10⁵ cells per pellet and cultured for 2 weeks in SFM supplemented with 1% ITS+1 liquid media supplement, 1.25 mg/mL human serum albumin, 0.1 mM L-ascorbic acid 2-phosphate sesquimagnesium salt, 10⁻⁷ M dexamethasone, and 10 ng/mL TGF-β1, with or without the addition of 3.75 µg/mL bevacizumab.

In vitro culture on scaffolds

HA-FIB-B3.75, HA-FIB-B5, HA-FIB, and FIB-lyophilized scaffolds were placed in agarose-coated six-well plates and seeded with NC (two donors with at least three replicates per condition per assay) by adding 10 µL of cell suspension (at a density of 1.5 × 10⁷ cells/cm³) from the top. After an incubation of 2 h at 37°C, 3 mL of complete medium supplemented with 10 mg/mL insulin (Actrapid HM; Novo Nordisk Pharma AG), 0.1 mM ascorbic acid 2-phosphate, 10 ng/mL TGF-β3 (Novartis), and 3000 KIU/mL aprotinin^{31,32} was added, and constructs were cultured for 2 weeks by changing the medium twice a week.

Glycosaminoglycan and DNA quantification

Pellets and/or *in vitro* constructs were digested with protease K 1 mg/mL protease K in 50 mM Tris with 1 mM EDTA, 1 mM iodoacetamide, and 10 mg/mL pepstatin-A for 15 h at 56°C. Glycosaminoglycan (GAG) amount was measured spectrophotometrically using 1,9-dimethylmethylene blue chloride dye.³³ Results were normalized by DNA levels, which were assessed by CyQuant[®] cell proliferation assay kit (Molecular Probes, Invitrogen).³²

Real-time quantitative reverse-transcriptase polymerase chain reaction

Total RNA was extracted from pellets and/or *in vitro* constructs using TRIzol[®] (Life Technologies), and cDNA was generated as previously described.³⁴ TaqMan[®] Gene Expression or on-Demand assays (Life Technologies) were used on a ABI 7900 Fast Realtime PCR System (Life Technologies) for gene expression measurements of type I (Hs00164004_m1) and II (Hs00264051_m1) collagen, Sox-9 (Hs00165814_m1), VEGF (Hs00900055_m1), and chondromodulin-I (Lect-1, Hs0017087_m1), using GAPDH (Hs99999905_m1) as the housekeeping gene.

In vivo ectopic mouse model

HA-FIB-B3.75, HA-FIB-B5, and HA-FIB-lyophilized scaffolds were seeded as described earlier. After overnight incubation in DMEM containing 10% FBS, 4.5 mg/mL D-glucose, 0.1 mM nonessential amino acids, 1 mM sodium pyruvate, 100 mM HEPES buffer, 100 IU/mL penicillin, 100 µg/mL

streptomycin, and 0.29 mg/mL L glutamate, fresh NC constructs (four donors with at least three replicates per condition per assay) were implanted in the back of nude mice (CD1 nu/nu, athymic, 5-week-old females; Charles River Laboratories) in pockets between excised muscle fascia and subcutaneous tissue for 1, 3, and 6 weeks. All the animals in this study were treated according to institutional guidelines.

Histological analysis

Explanted constructs underwent fixation and routine histological processing. Stainings were performed for Safranin O and Alizarin red. Immunohistochemistry on 5-µm-thick paraffin sections was performed for mouse anti-human collagen (types I, II, and X), matrix metalloproteinase 13 (MMP-13), and bone sialoprotein (BSP) (MP Biomedicals),³⁵ using hematoxylin as a nuclear counterstain. Endogenous mouse immunoglobulins in the tissue were blocked using a M.O.M.[™] kit (Vector Laboratories) according to the manufacturer's instructions. Images were acquired using an Olympus BX-61 microscope.

TABLE 1. HISTOLOGICAL SCORING SYSTEM

Scoring categories	Score
1. Intensity of Safranin O-fast green stain³⁶	
No stain	0
Weak staining of poorly formed matrix	1
Moderately even staining	2
Even dark stain	3
2. Distance between cells/amount of matrix accumulated³⁶	
High cell density with no matrix in between (no spacing between cells)	0
High cell density with little matrix in between (cells <1 cell-size apart)	1
Moderate cell density with matrix (cells ~1 cell-size apart)	2
Low cell density (cells >1 cell-size apart) with an extensive matrix	3
3. Cell morphologies³⁶	
Condensed/necrotic/pyknotic bodies	0
Spindle/fibrous	1
Mixed spindle/fibrous with rounded chondrogenic morphology	2
Majority of rounded/chondrogenic	3
4. Uniformity of Safranin O staining³⁷	
Complete disorganization of stained areas	0
Less than 50% of Safranin O stained areas with regard to the total volume	1
50%–80% of Safranin O stained areas with regard to the total volume of the section	2
Overall homogeneity of the staining	3
5. Abnormal calcifications within the cartilage newly formed³⁷	
Strong presence of abnormal calcifications inside the neo-formed cartilage	0
Presence of some calcification spots inside the newly formed cartilage	1
Presence of very few calcification spots inside the repaired cartilage	2
Complete absence of calcifications throughout the newly formed cartilage	3
6. Vessels ingrowth (within the newly formed tissue)³⁷	
Vessels ingrowth inside the core of the repaired cartilage	0
Vessels ingrowth until the inner rim of the neo-formed cartilage (approximately two thirds of newly formed cartilage's total volume)	1
Vessels ingrowth until the outer rim of the neo-formed cartilage (approximately one third of newly formed cartilage's total volume)	2
No presence of vessels or vessels confined in the external fibrotic capsule	3
7. Final score	
Not-formed cartilage. Basically, just fibrotic tissue	0–6
Inhomogeneous and fibrocartilagineous tissue, having just a few and nonconnected areas of a good newly formed cartilage	6.1–11
Moderately good tissue, having less than 60%—with regard to the total volume—of a good newly formed cartilage	11.1–15
Very good quality of homogeneity of newly formed cartilage	15.1–18

Immunofluorescence (IF) for rat anti-mouse CD31 and F4/80 (BD) was performed on 10- μ m-thick cryosections of *in vivo* generated constructs, using DAPI (BD) as a nuclear counterstain. Images were acquired with a confocal microscope (LSM710; Zeiss).

The percentage of area infiltrated by either vessels (CD31⁺) or macrophages (F4/80⁺) was calculated on cross-sections of the implant, excluding the outer fibrotic capsule ($n=5$ per experimental group) using Image J software (NIH; Bethesda).

The quality (by Safranin O staining), calcification (by Alizarin red staining), and host vessels ingrowth (by IF for CD31) of the engineered cartilaginous tissues ($n\geq 3$ per each experimental group) were assessed blindly using an *ad hoc* scoring system, which combines the already used Bern³⁶ and ICRS II³⁷ scores (Table 1). The proposed score was validated by calculating the maximum and overall variability (σ_{max} , σ_{ov}) of the scores among five blind observers as follows:

$$\sigma = \frac{s}{\bar{x}} \cdot 100 [\%],$$

where s is the variance and \bar{x} is the average. These values were calculated to be 21.43% and 8.16% for σ_{max} and σ_{ov} , respectively.

Statistical analysis

Data are presented as means \pm standard deviation, SD ($n\geq 3$). Statistical analysis was performed using the unpaired or nonparametric t-test, taking into account the normal distribution of the collected data by means of Prism[®] software (GraphPad Software); p -values lower than 0.05 indicate statistically significant differences.

Results

Scaffold characterization

The synthetic route for the preparation of hyaluronan/fibrin-based scaffolds resulted in highly reproducible incorporation of bevacizumab that did not affect their final microstructure, pore size, and distribution (Fig. 1A, B). Scaffolds displayed a similar porosity ($70\pm 5\%$) regardless of the concentration of bevacizumab, with a similar pore size in the range of 20–220 μ m.

The compressive modulus (E^*) was similar in all the experimental groups, ranging from 8.35 ± 0.04 to 8.39 ± 0.08 kPa.

NC viability was higher than 99.2% for all scaffold compositions, demonstrating their noncytotoxicity. HA-FIB, HA-FIB-B3.75, and HA-FIB-B5 showed similar behavior in terms of weight loss and water uptake, suggesting that the addition of bevacizumab did not affect the chemo-physical properties of the final scaffolds. Scaffolds lost $\sim 50\%$ of their total weight in the first 48 h, whereas the remaining 45% was slowly degraded in the next 3 weeks of culture. Water uptake ratio was calculated to be ~ 7.6 w/w for all scaffolds.

Cumulative bevacizumab release profile was about one fourth of the initial amount during the first 48 h; while from day 4 to 10, its rate slowed down until complete elution during the third week of *in vitro* soaking (Fig. 1C).

Bevacizumab activity and dosage

HUVEC showed a VEGF dose-dependent proliferation rate, reaching a plateau at a concentration of 5 ng/mL (Fig. 1D, positive control curve). Bevacizumab, at a concentration as low as 1.5 μ g/mL, efficiently and significantly reduced HUVEC

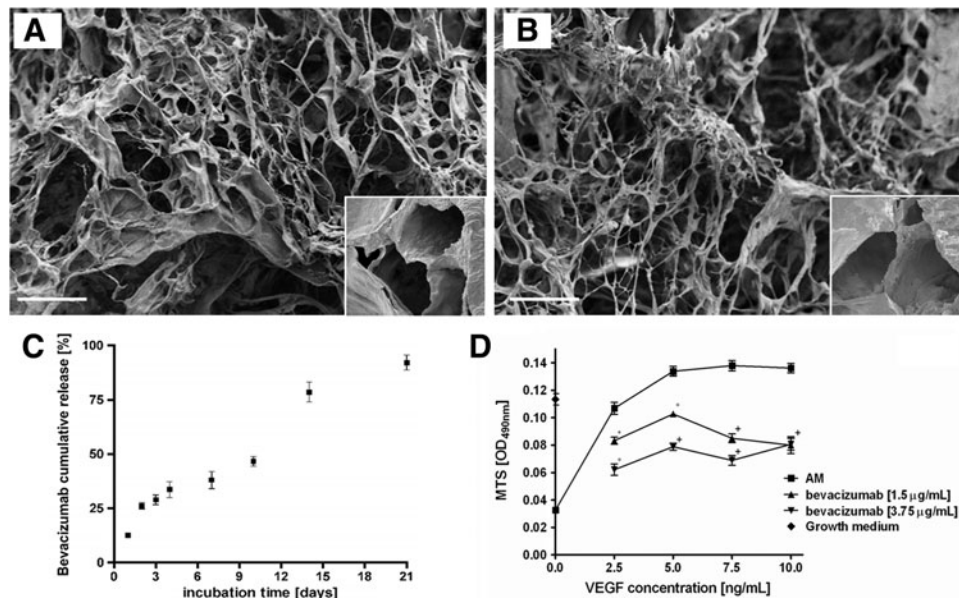


FIG. 1. Scaffold characterization. Scanning electron microscopy micrographs of scaffolds functionalized (A) or not (B) with bevacizumab. Scale bar = 100 μ m; insets show higher magnification pictures. (C) Bevacizumab cumulative release curve over 3 weeks of *in vitro* culture, expressed as % with regard to the total amount; (D) bevacizumab activity and dosage measured via MTS assay. Human umbilical vein endothelial cells (HUVEC) showed a vascular endothelial growth factor (VEGF) dose-dependent proliferation rate reaching a plateau at a concentration of 5 ng/mL (assay medium [AM] curve). Bevacizumab, at both tested concentrations (1.5 and 3.75 μ g/mL), efficiently and significantly reduced HUVEC metabolic activity for approximately 10 ng/mL of VEGF (* $p < 0.05$, ° $p < 0.01$, + $p < 0.001$).

metabolic activity for approximately 10 ng/mL of VEGF (Fig. 1D). At a higher concentration of 3.75 μg/mL, bevacizumab was more effective in blocking the VEGF-induced HUVEC proliferation. Therefore, this was selected as a suitable concentration for scaffold composition (HA-FIB-B3.75). In addition, a higher-level bevacizumab concentration of 5 μg/mL—the highest concentration not affecting the final scaffold microstructure—was chosen (HA-FIB-B5) for the *in vivo* experiments.

In vitro 3D pellet culture

Supplementation of bevacizumab in the culture medium did not affect the *in vitro* chondrogenic differentiation potential of NC. Safranin O staining showed no major differences, in terms of staining intensity and uniformity, chondrocyte morphology, and no statistically significant difference was found in the GAG content for pellets generated by NC treated or not with bevacizumab (Fig. 2A). The expression at mRNA level of Sox9, type I collagen, and VEGF was similar in the two experimental groups (Fig. 2B); whereas a slight down-regulation in type II collagen was observed. However, this effect did not alter the expression of type II collagen at protein level, as confirmed by immunohistochemistry for this marker (data not shown).

In vitro 3D culture on scaffolds

Release of bevacizumab from the scaffolds did not affect NC chondrogenic differentiation capacity, confirming the findings of the 3D pellet culture model. We, indeed, observed a quite similar cartilaginous ECM, intensively stained for GAG with chondrocyte-like cell morphology in all hyaluronan-containing scaffolds regardless of bevacizumab supplementation. Interestingly, FIB scaffolds did not result in

being chondrosupportive as those containing HA, as the deposited ECM was negatively stained for GAGs (data not shown). A comparison with nonseeded HA-FIB scaffolds enabled the exclusion of potential biases or artifacts in histological and biochemical assays.

As compared with the engineered tissues generated with FIB scaffolds, the constructs engineered with scaffolds and also containing HA deposited a significantly higher amount of GAG (Fig. 2C). The presence of HA gave rise to a significantly increased expression of genes of interest, such as type II collagen and chondromodulin (Fig. 2D). The addition of bevacizumab did not affect the expressions of cartilaginous markers.

In vivo ectopic mouse model

After subcutaneous implantation, NC-based constructs generated without bevacizumab were mostly resorbed (Fig. 3). Indeed, the HA-FIB constructs that persisted at 6 weeks *in vivo* were as low as 18.2% (2 out of 11). On the contrary, the engineered tissues generated with HA-FIB-B3.75 and HA-FIB-B5 scaffolds showed higher survival rates at 6 weeks *in vivo*, corresponding to 60% (6 out of 10) and 75% (3 out of 4), respectively. Interestingly, the percentage of not-resorbed constructs remained unaltered after 3 weeks. These data remarkably underline the role of the early blocking of angiogenesis for the successful implantation of an immature graft *in vivo*.

Despite the different survival rate, no major differences were observed in terms of cartilage quality between HA-FIB, HA-FIB-B3.75, and HA-FIB-B5 constructs once an initial critical phase of tissue formation has been overcome, demonstrating the self-sustaining capacity of the neo-formed

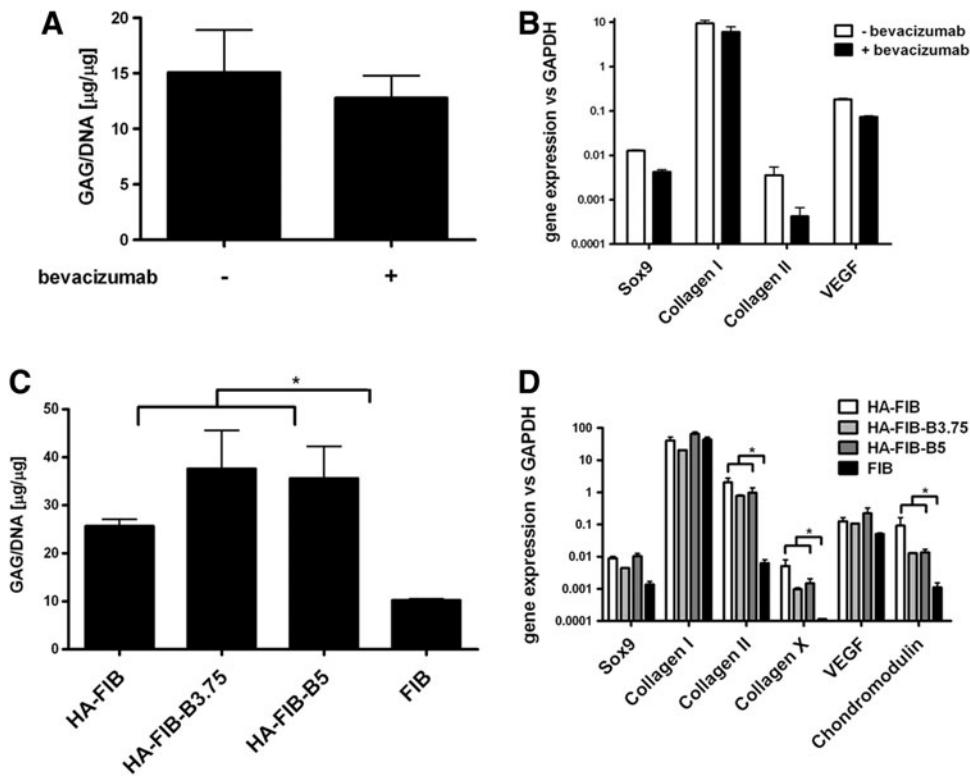


FIG. 2. *In vitro* 3D pellet and scaffold-based culture systems. (A, B) *In vitro* pellet culture. (A) Quantification of the glycosaminoglycan (GAG)/DNA ratio for pellets generated by nasal chondrocytes (NC) cultured with chondrogenic medium supplemented or not with 3.75 μg/mL of bevacizumab. (B) Analysis at mRNA level of Sox9, collagen (types I and II), and VEGF genes. (C, D) *In vitro* scaffold-based culture. (C) Quantification of the GAG/DNA ratio of cartilaginous constructs generated by NC loaded on FIB control scaffold, HA-FIB either alone, or functionalized by the supplementation of 3.75 (HA-FIB-B3.75) or 5 (HA-FIB-B5) μg/mL of bevacizumab. (D) Analysis at mRNA level was also performed for Sox9, collagen (types I, II, and X), VEGF, and chondromodulin genes. **p* < 0.01.

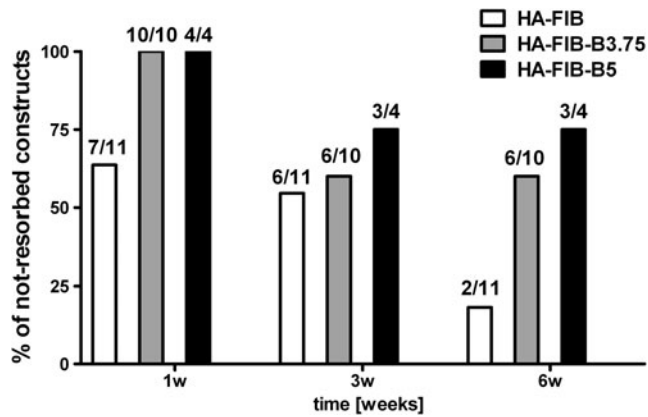


FIG. 3. *In vivo* construct degradation—percentage of the non-resorbed engineered tissues with regard to the total number of implants.

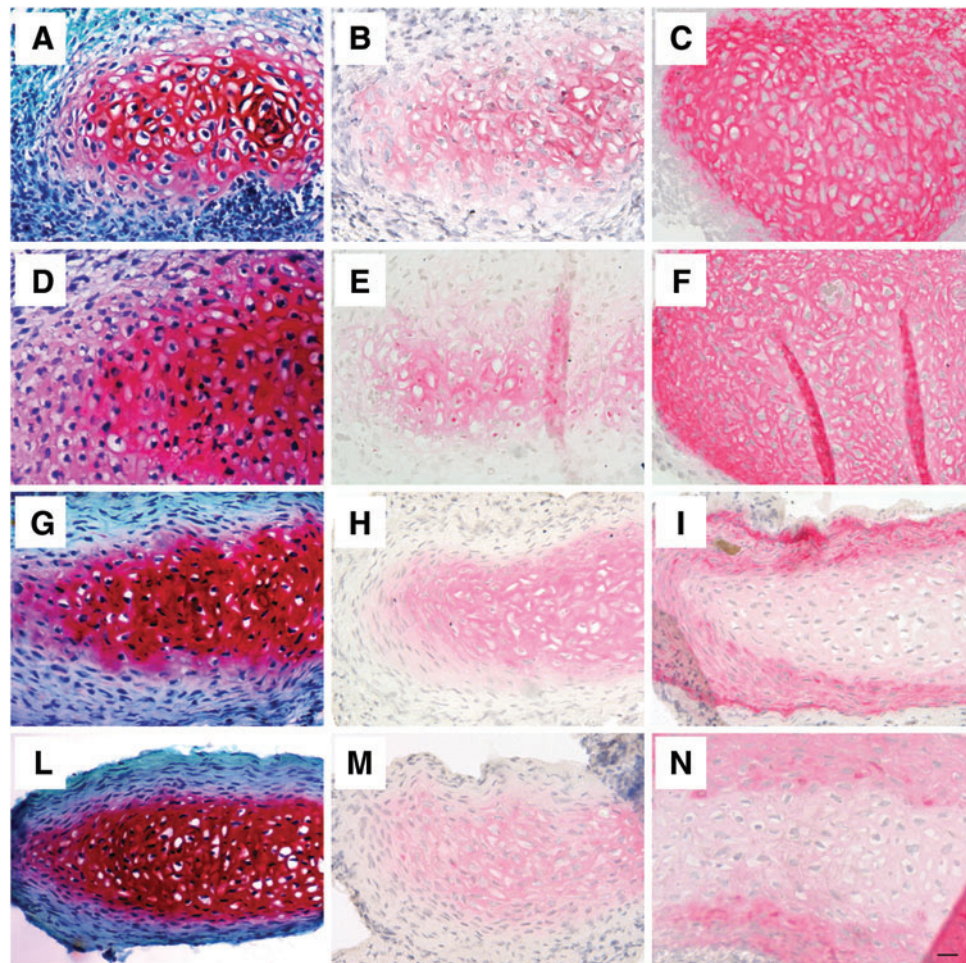
engineered cartilage. Since no remarkable differences were found between the groups at different bevacizumab concentrations (3.75 or 5 $\mu\text{g}/\text{mL}$), histological results are presented only for the HA-FIB-B3.75 group (Fig. 4). After just 1 week, we observed the formation of a cartilaginous ECM, positively stained for Safranin O and type II collagen with chondrocyte-like morphology in all the experimental condi-

tions for the not-resorbed constructs (Fig. 4A, B, D, E), although type I collagen staining was intense and uniform throughout the whole construct (Fig. 4C, F). After 6 weeks *in vivo*, the cartilaginous ECM showed an even more intense and uniform staining for GAG (Fig. 4G, L) and type II collagen (Fig. 4H, M); whereas type I collagen was present only in the outer edges of the implants (Fig. 4I, N). Cells were mainly showing a typical chondrocyte morphology, and no necrotic or pyknotic cells were found in all the experimental conditions.

The stability of the chondrogenic phenotype reached by the NC *in vivo* was assessed by a histological analysis for hypertrophic markers (such as type X collagen) and other key molecules involved in cartilage remodeling (such as MMP-13 and bonesialoprotein), which were negative in all experimental groups over time (data not shown).

At 1 week, in all the groups, no CD31⁺ cells were found inside the neo-formed cartilaginous ECM, but only in the host fibrotic capsule surrounding the implant. At 3 weeks, host vessels infiltration toward the center of the construct was reported only for the HA-FIB group, giving rise to a process eventually leading to the breakdown of the tissue-engineered cartilage at later time points (Fig. 3). On the contrary, CD31-positive vessel structures remained confined at the outer edges of the engineered tissue for both HA-FIB-B3.75 (Fig. 5B) and HA-FIB-B5 (Fig. 5C), which is

FIG. 4. *In vivo* chondrogenesis. Safranin O staining: (A) HA-FIB, 1 week, (D) HA-FIB-B3.75, 1 week, (G) HA-FIB, 6 weeks, (L) HA-FIB-B3.75 6 weeks. Immunohistochemistry for type II collagen: (B) HA-FIB, 1 week, (E) HA-FIB-B3.75, 1 week, (H) HA-FIB, 6 weeks, and (M) HA-FIB-B3.75 6 weeks. Immunohistochemistry for type I collagen: (C) HA-FIB, 1 week, (F) HA-FIB-B3.75, 1 week, (I) HA-FIB, 6 weeks, and (N) HA-FIB-B3.75, 6 weeks. Scale bar: 50 μm . Color images available online at www.liebertpub.com/tea



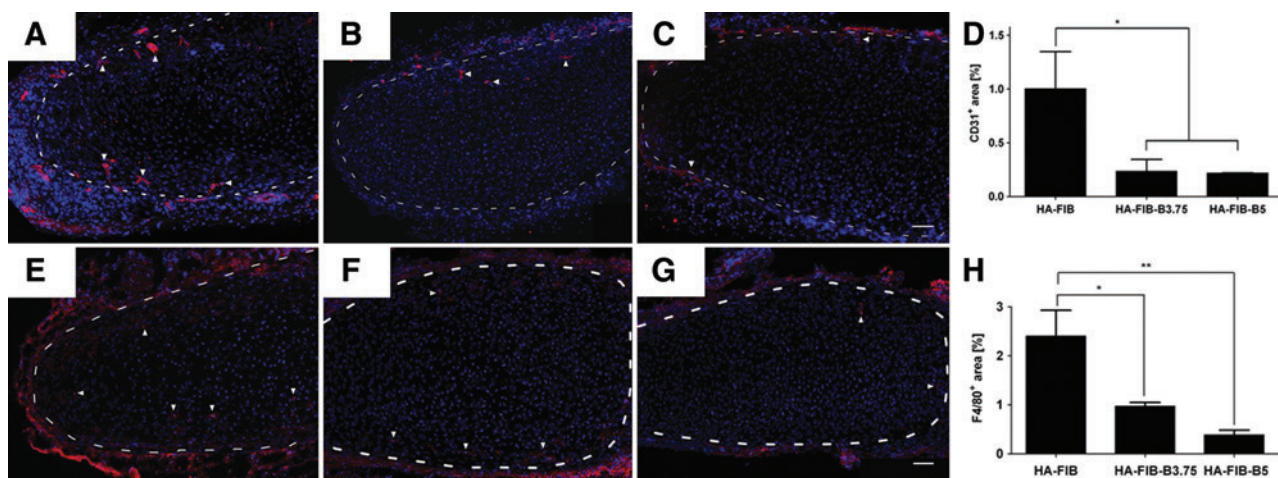


FIG. 5. *In vivo* host vessel ingrowth. Immunofluorescence (IF) for CD31 at 3 weeks' time point of (A) HA-FIB, (B) HA-FIB-B3.75, (C) HA-FIB-B5; (D) vessels quantification, calculated as the percentage of CD31⁺ area with regard to the total picture area ($n=5$ per experimental group). Inflammatory response. IF for F4/80 at 3 weeks' time point of (E) HA-FIB, (F) HA-FIB-B3.75, and (G) HA-FIB-B5 samples; (H) migrated monocytes quantification, calculated as the percentage of F4/80⁺ area with regard to the total picture area ($n=5$ per experimental group). Scale bar: 50 μm . * $p < 0.01$, ** $p < 0.001$. Color images available online at www.liebertpub.com/tea

consistent with the capacity of the eluted bevacizumab to effectively inhibit neo-angiogenesis. Quantification of the percentage of CD31⁺ areas showed a statistically significant difference ($p < 0.05$) between HA-FIB constructs and bevacizumab-containing constructs HA-FIB-B3.75 and HA-FIB-B5 (Fig. 5D).

These observations matched with the results of the histological scoring. Statistically significant differences were, indeed, observed, particularly with regard to the constructs vascularization at 3 weeks between HA-FIB and HA-FIB-B3.75 ($p < 0.05$), and at 6 weeks between HA-FIB and both HA-FIB-B3.75 ($p < 0.05$) and HA-FIB-B5 ($p < 0.001$) (Table 2).

At 3 weeks, the infiltration toward the center of the cartilaginous ECM of F4/80⁺ murine macrophages appeared to be stronger in constructs without bevacizumab as compared with both HA-FIB-B3.75 and HA-FIB-B5 (Fig. 5E–G). The quantification of F4/80⁺ cells, indeed, showed a significantly higher number of monocytes in the HA-FIB group as compared with HA-FIB-B3.75 and HA-FIB-B5 groups (Fig. 5H).

In order to elucidate the reported differences in terms of host vessels and monocytes invasion *in vivo*, all the scaffold groups were tested *in vitro* against HUVEC proliferation (Fig. 6A) and monocytes migration (Fig. 6B).

Effect of bevacizumab and scaffold degradation products on HUVEC proliferation

Figure 6A shows the results of HUVEC proliferation assay for scaffolds at different compositions, as the ratio between the given experimental condition and the GM condition (fold increase). FIB degradation products were found to strongly enhance HUVEC proliferation, with a 1.71 ± 0.09 -fold increase with regard to HUVEC cultured in GM. HA release from HA-FIB scaffolds appeared to mitigate this proliferative effect (1.02 ± 0.09 -fold vs. GM group), which was reliably due to the reported anti-angiogenic effect of high-molecular-weight hyaluronan.³⁸

Elution of bevacizumab from FIB-B3.75 scaffolds (not containing hyaluronan) induced a progressive reduction of

TABLE 2. HISTOLOGICAL SCORING EVALUATION

Sample	HA-FIB			HA-FIB-B3.75			HA-FIB-B5		
	1 week	3 week	6 week	1 week	3 week	6 week	1 week	3 week	6 week
<i>Scoring categories</i>									
1. Intensity	0.82 ± 0.98	1.82 ± 0.87	2.70 ± 0.48	0.67 ± 0.78	1.82 ± 0.60	3.00 ± 0.00	1.00 ± 1.00	1.91 ± 0.54	2.64 ± 0.50
2. Density	0.55 ± 0.69	1.36 ± 0.50	2.10 ± 0.57	1.00 ± 0.95	1.64 ± 0.47	2.27 ± 0.65	1.00 ± 0.89	1.91 ± 0.83	2.45 ± 0.69
3. Morphology	1.27 ± 1.35	1.82 ± 0.40	2.50 ± 0.53	1.08 ± 1.00	1.73 ± 0.67	2.91 ± 0.30	1.36 ± 1.29	2.00 ± 0.63	2.55 ± 0.69
4. Uniformity	0.55 ± 0.69	1.64 ± 0.50	2.20 ± 0.63	0.92 ± 1.00	1.64 ± 0.50	2.55 ± 0.69	0.91 ± 0.94	1.36 ± 0.50	2.00 ± 1.00
5. Calcifications	3.00 ± 0.00	3.00 ± 0.00	2.40 ± 0.52	3.00 ± 0.00	3.00 ± 0.00	2.91 ± 0.30	3.00 ± 0.00	3.00 ± 0.00	3.00 ± 0.00
6. Vascularization	2.67 ± 0.71	1.22 ± 0.67	1.38 ± 0.74	3.00 ± 0.00	1.78 ± 0.67 ^a	2.78 ± 0.44 ^b	2.78 ± 0.44	1.89 ± 0.60 ^a	2.67 ± 0.50 ^b
Final score	8.85 ± 2.06	10.86 ± 1.39	13.28 ± 1.43	9.76 ± 1.87	11.6 ± 1.32	16.41 ± 1.13 ^c	10.05 ± 2.13	12.07 ± 1.41 ^a	15.3 ± 1.57 ^b

Statistical differences evaluated between the scaffolds with and without bevacizumab at each time point.

^a $p < 0.05$.

^b $p < 0.01$.

^c $p < 0.001$.

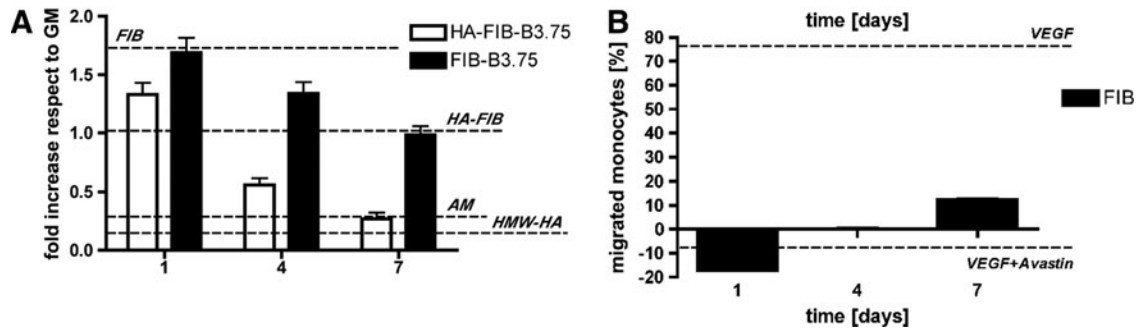


FIG. 6. Scaffold degradation products characterization. The role of each scaffold component was tested by using a HUVEC proliferation assay (A) and an *ad hoc* developed monocytes migration assay (B). (A) For HUVEC proliferation assay, high-molecular-weight HA (HMW-HA), AM, and growth medium (GM) were used as controls. HUVEC metabolic activity is presented as the ratio between the given experimental condition and the GM condition (fold increase). (B) Monocytes migration assay performed on FIB degradation products, calculated as the percentage of the migrated monocytes of a given experimental condition with regard to the serum-free medium group. Supplementation of 10 ng/mL of VEGF alone or in combination with bevacizumab (using the optimal stoichiometric ratio 1:2.6¹⁵), was used as a control.

the HUVEC metabolic activity, which was similar to the levels obtained with HA-FIB scaffolds, and, therefore, similar to the GM control. It should be noted, in HA-FIB-B3.75 scaffolds, that the synergistic effect of HA and bevacizumab contributed toward reducing the HUVEC proliferation rate since day 1. At day 7, HA and bevacizumab were able to reduce HUVEC proliferation to the threshold level represented by the negative control (AM with no VEGF).

Effect of bevacizumab and scaffold degradation products on monocyte migration

Figure 6B shows the results of monocyte migration assay for FIB scaffolds at different time points, as the percentage of the migrated monocytes of a given experimental condition with regard to the SFM group. VEGF supplement to SFM strongly increased the migration of monocytes (+78.07% ± 1.14% vs. SFM group) (Fig. 6B), as previously shown by Zentilin *et al.*,¹¹ while the addition of bevacizumab to the VEGF-containing medium completely suppressed their migration (-10% vs. SFM). It should be noted that degradation products obtained from FIB scaffolds increased the migration of monocytes to +12.28% ± 1.18% vs. the SFM group at day 7. On the contrary, no significant differences were observed between HA-FIB and SFM groups.

Discussion

Our study showed that an anti-angiogenic approach, applied to mimic the avascular nature of hyaline cartilage and to block host reaction, may support the formation of long-term, stable, and high-quality cartilaginous engineered tissue without any *in vitro* preconditioning.

Despite the promising results recently obtained in CTE field, new strategies are still warranted to overcome the intrinsic limitations to the extensive, costly, and time-consuming *in vitro* conditioning of the grafts.⁵ One possible approach relies on the direct implantation of cell-loaded 3D scaffolds, with no previous chondrogenic stimulus.³⁹ To this extent, here we proposed to streamline and overcome the *in vitro* preconditioning by using a 3D scaffold that, while supporting *in vivo* chondrogenesis, inhibits host

vessel ingrowth while preventing implant early resorption. We developed a scaffold based on well-established, FDA-approved materials for CTE, that is, fibrin and hyaluronan. In this system, fibrin provided the 3D porous structure, while hyaluronan represented the chondrogenic stimulus, which resulted in being a key requirement for inducing *in vitro* NC chondrogenic re-differentiation. The proposed scaffold displayed a strong chondrosupportive capacity particularly *in vivo*, leading to the formation of newly formed hyaline-like cartilage already after 1 week. In the last few years, several groups adopted gene therapy-based strategies to block angiogenesis for supporting cartilage regeneration. Progenitor cells were transduced to over-express autologous anti-angiogenic molecules, such as soluble Flt-1,¹⁰ endostatin,^{12,13} and chondromodulin.¹⁴ Nagai *et al.* proposed an alternative approach consisting of the systemic administration of bevacizumab (Avastin), which resulted in the augmented capacity of cartilage self-healing in rabbit osteochondral lesions, thus strengthening the key role of anti-angiogenic signaling in cartilage homeostasis.⁴⁰ The latter approach holds less concerns in terms of safety and processing time, compared with gene therapy, but still requires systemic drug administration; while, for a clinically compliant articular regeneration strategy, a local treatment is desirable. Indeed, the systemic use of bevacizumab induces severe drawbacks, such as gastrointestinal perforation, serious bleeding, in addition to inherent/acquired resistance and induction of tumor invasiveness.⁴¹

The proposed work showed that anti-angiogenesis during cartilage regeneration can be pursued using a direct and clinically oriented approach by means of time-controlled, topical administration of an anti-angiogenic drug (i.e., bevacizumab, Avastin) and by paving the way for the use of freshly isolated patients' cells with regard to the extensively expanded and genetically modified cells.

The release of bevacizumab from implanted grafts during the first 2/3 weeks appeared to effectively block human VEGF released by the implanted NC and, therefore, inhibited host vessel ingrowth,⁴² as shown by histological images, likely preserving the constructs from resorption and leading

to a long-term stability. Indeed, ~82% of scaffolds without bevacizumab (HA-FIB experimental group) completely degraded after 6 weeks *in vivo*, which was reasonably due to newly formed matrix remodeling and resorption on host vessel and monocyte invasion, which was already evident at 3 weeks on implantation. Francioli *et al.*⁴³ and Luo *et al.*⁴⁴ demonstrated how the level of maturation of cartilaginous tissues generated *in vitro* is correlated with a different profile of inflammatory chemokine production (e.g., MCP-1, IL-8), and, in particular, how the *in vitro* preconditioning can mitigate the postimplantation inflammatory reaction. In particular, Luo *et al.*⁴⁴ showed how a 2 week preculture of chondrocyte-based constructs results in a mild inflammatory response in the host tissue, with no vessels ingrowth found in the middle of the constructs and a fewer number of leukocytes in comparison with the “direct implantation group.” Therefore, we concluded that our strategy based on blocking VEGF signal by temporal release of bevacizumab on the graft implantation can streamline CTE applications (1) by overcoming the *in vitro* preconditioning phase, thus starting with freshly seeded scaffolds; (2) by limiting the inflammatory response; (3) by protecting immature cartilaginous matrix from remodeling; (4) by guaranteeing further tissue development (e.g., endogenous production of anti-angiogenic molecules that usually self-protect cartilage from macrophage invasion and modulate the postimplantation inflammatory response); and, finally, (5) by allowing long-term stability of the cell-based grafts.

In order to better study how chondrogenesis is affected by the antiangiogenic drug bevacizumab, further investigations by using an inert and not degradable scaffold are necessary. Indeed, we have reported that fibrin degradation products remarkably enhance endothelial cell proliferation, as well as monocytes migration, and that bevacizumab and high-molecular-weight HA incorporated within the scaffolds counteract that angiogenic effect. Moreover, no major differences in terms of the quality of the neo-formed engineered cartilage were found among the not resorbed constructs, regardless of the presence of bevacizumab. This aspect is probably due both to the intrinsic chondrogenic potential of the NC primary culture and to the good chondrosupportive nature of the proposed scaffold.

Future studies will also aim at investigating the chondrogenic potential of the constructs engineered by using the bevacizumab-functionalized scaffold in an orthotopic and immunocompetent model, as well as at better elucidating the complex system of host reaction and mechanical stimulations. Besides showing the indirect effect of blocking VEGF, and therefore vessel ingrowth, on monocyte infiltration, our preliminary results might also suggest a possible direct effect of bevacizumab on monocyte proliferation, differentiation, and migration. Moreover, the use of bone marrow-derived mesenchymal stromal cells (BMSCs) might be contemplated to broaden the clinical scenario and cell source choice. It has been recently shown that BMSCs undergo hypertrophy, followed by formation of micro-ossicles after ectopic transplantation in immunodeficient mice.^{45,46} Since VEGF plays an essential role in these mechanisms, future experiments will investigate whether bevacizumab-induced VEGF blockade could affect BMSC endochondral fate and thus be instrumental for CTE with BMSC.

Conclusions

Our findings suggest that blocking angiogenesis in a chondrosupportive immature graft supports the formation of a long-term stable engineered cartilage, as it effectively preserves its avascular nature and prevents its resorption. The scaffold-based approach here that is proposed to limit spatially and temporally the delivery of an anti-angiogenic drug might represent a step forward in the current CTE scenario, offering a valid alternative to conventional bio-material-induced autologous implantation techniques. The use of all FDA-approved materials for its synthesis, including the anti-angiogenic drug, and its validation with a chondrogenic and clinically relevant cell source, namely NC, are expected to allow a straightforward translation to a clinical setting.

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Disclosure Statement

There are no conflicts of interest to declare. The writing of this article was the sole responsibility of the authors.

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Address correspondence to:

Alberto Rainer, PhD

Tissue Engineering Laboratory

Center for Integrated Research

Università Campus Bio-Medico di Roma

via Álvaro del Portillo 21

Rome 00128

Italy

E-mail: a.rainer@unicampus.it

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