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Cell-collagen networks breakdown by collagen remodelling

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Abstract

Collagen model tissues are analyzed, which consist of cells embedded in a collagen matrix at different concentrations (of cells and collagen). Rheological properties are measured and complementary confocal microscopy analyses are carried out. An important feature is observed, corresponding to the breakdown of the collagen network (i.e. decrease in network elasticity) for high collagen concentrations, due to the presence of cells. Thanks to confocal microscopy, we show that cells elongate within the gel and can remodel it, this being a concentration-dependent feature. A careful analysis of the remodelling process shows that cells can attract collagen in their close neighborhood, this being an irreversible process, and that migrating cells create collagen-depleted regions behind them.

Key words: tissue; collagen; rheology; confocal; reflectance; remodelling

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1. Introduction

The rheology of biological systems is of major importance for bioengineering and materials research; it has been a source of interest for many years [21], for it concerns a wide variety of human tissues, such as muscles, arteries, connective tissues, but also new biomaterials like scaffolds to be used for restoration medicine, wound healing, and various therapies. The rheological properties of such tissues cover a wide range, going from simple elastic materials, to fluid-like ones, but in the majority of tissues, linear and nonlinear viscoelastic effects [45] as well as viscoplastic effects (gel-like) seem to be ubiquitous [49]. The essence of these properties rely on the fact that cells live in a fluid material filled with fibers (i.e. collagen, and other matrix proteins), which give rise to both viscous and elastic properties. It is the complex interplay between cell deformation, fiber reorganization, and fluid flow [37] which leads to the understanding of such viscoelastic properties and to possible modelling [41,54,55].

In order to understand the complexity of such materials, model systems are commonly studied including basic solutions with components of the extra cellular matrix (ECM) which are collagen, laminin, elastin, fibronectin, etc. or other biopolymer gels present within cells, such as actin [4,56], tubulin, fibrin [26] which are basic cell elements. The rheological properties of such systems depend on mechanical and chemical properties of the polymer network, on the ability of components to assemble and persist, and can therefore exhibit either uncrosslinked polymer features or rheological properties as in crosslinked gels, in particular different crosslinks (e.g. α -actinin, myosin II) holding the network together are shown to lead to a rich variety of such properties.

Commonly used techniques for studying biological gels range from microrheology devices (Optical tweezers [11], magnetocytometry [32], AFM [7], etc.) to conventional rheometry, compression tests [29], uniaxial tension or sometimes home-made devices [9]. Although

relaxation tests [18,35] or cycle tests [51] are commonly used, dynamic tests – in a range of angular frequencies ω - are the most versatile. Frequency-dependent viscoelastic properties (G' and G'') have been obtained previously for F-actin solutions [43] and scaling laws for the elastic modulus G' vs. concentration were found for various filamentous biopolymer networks [25], using oscillatory shearing. For example, several researchers studied the rheology of Type I collagen [29,30,48] or modified collagen [19] and found similarities with other semi-flexible polymer networks. This motivated attempts to account for such rheology using models based on polymer networks theories [31,44,45,55]; indeed the arrangement of long protein filaments into networks of different geometries can be applied to such materials.

Going towards a more sophisticated model tissue implies to embed cells into the protein network and requires particular techniques (control of temperature, percentage CO₂, pH, etc.). Under such proper conditions, it is found that unlike classical fillers (or particles) in composite materials, cells play an active role by interacting with the extra cellular matrix (or fibrillar network), leading to growth/shrinking, reorganization or proteolytic activity [10]. For example, chicken embryo fibroblasts were found to give rise to an additional contribution to the stress in fibroblast-populated collagen matrices [50]. Therefore, the stiffness of the extracellular matrix can profoundly influence cell and tissue behaviors, and conversely. Another example is the microrheometric (magnetic bead-twisting) assay used to measure changes of a cell-collagen network [33]. The authors found that cells elongate within the matrix and change the local micromechanical properties of type I collagen gel, in a cytokine-dependent manner. In this respect, the ECM is a rich signalling medium, where both biochemical and mechanical properties can rule physiological and pathological processes [46]. Therefore the ECM mechanical properties can be tuned for proper functions, thus strongly influencing the behavior of cells [14,15]. As a particular example, cellular sub-structures in myoblasts can be strongly affected by the rigidity of the underlying tissue [14].

To characterize a cellular network, one needs to measure the rheological properties, and relate these mechanical behaviors with respect to microstructure changes, using relevant techniques. The most classical ones used are SEM (Scanning electron microscopy) [1] or conventional biology tools such as histology [22,23] which require special preparations, meaning possible artefacts. Therefore such techniques are nowadays replaced by confocal microscopy (including laser reflectance for visualizing collagen fibers in the matrix) [6,16,20,24] leading to amazing observations of the scaffolds microstructure [34]. These observations have recently been coupled with micromechanical tools to analyze the stress effects on collagen microstructure [47].

Previous studies concern different cell types [17,27,53], but most of them usually focus on fibroblasts - which are major repair cells - embedded within 3D scaffolds containing matrix proteins [3,5,10,16,35,36,51]. Such model networks have been used to understand the migration/deformation of fibroblasts and the remodelling/compacting effect they can have on the ECM [12,16,17]. Collagen gels were used extensively with fibroblasts for studying the cell-matrix mechanical interactions. However, no results about the overall viscoelastic properties of such model networks using both conventional rheology and confocal microscopy have been carried out on systems with various concentrations.

Therefore, in this study, we propose to develop a 3D model network that mimics a normal tissue (e.g. connective tissue) and study its viscoelastic properties using both dynamic rheometry and confocal reflectance microscopy. To undertake such investigations, we used a collagen network at various concentrations, filled with different amounts of CHO cells (Chinese Hamster Ovary cells). It was then possible to investigate the behavior of cells in different micro-environments. Materials and methods are described in §2, whereas the model networks viscoelastic properties are investigated in §3. At the same time, we used complementary confocal scanning microscopy (combined reflectance and fluorescence) to

investigate cell shape and cell-matrix interactions. The remodelling process was studied, by measuring collagen densities in the close neighborhood of cells. In §4, we propose to combine these results and show how interactions between cells and the collagen matrix can compete and give rise to complex behaviors because of this remodelling process. Finally, conclusions are presented in §5.

2. Materials and methods

2.1 Cells, collagen, and gels

We used a CHO cell line (Chinese Hamster Ovary) - an established line of fibroblasts - previously investigated by the authors, as a suspension at high concentrations [24]. CHO cells were grown in culture medium (DMEM containing 10% Fetal Calf Serum) using standard T75 flasks under proper conditions (37°C, 5% CO₂) in a cell incubator, until they reached 70% confluence. Cells were then detached with trypsin, and suspensions at different concentrations were prepared using CHO cells and culture medium. Cell concentration was determined using a Neubauer's cell chamber, then the required concentration was adjusted.

Collagen (type I) was prepared from a rat tail collagen solution (BD Biosciences) and was mixed with culture medium. Then neutralization was carried out using NaOH (0.1M) to reach a pH of 7.4 while maintaining the temperature at 4°C. After adding cells, the system was transferred onto the rheometer at 37°C. Collagen polymerization usually takes 30 mn. Overall, the time between preparation of the gel and the beginning of the rheology experiment (including setting of the control parameters) was one hour.

Two kinds of tissues/networks were prepared in order to investigate collagen networks alone as well as cell-populated collagen matrices :

a) Collagen gels at four different concentrations (0.42-0.95-1.38-1.8 mg/mL)

b) Collagen gels including cells at different cell concentration. Three different cell concentrations were used: $0.7 \cdot 10^7$ cells/mL, $1.17 \cdot 10^7$ cells/mL and $1.8 \cdot 10^7$ cells/mL. These concentrations correspond to cell volume concentrations of 4.7%, 7.8%, 12% respectively, and were adjusted so that collagen concentrations remained the same as in a).

2.2 Rheology

Rheometrical measurements of the networks were conducted with a Bohlin Rheometer (Malvern, Gemini 150, controlled stress). The temperature was kept at 37° C (peltier system, +/- 0.1°C) which allows for collagen polymerization as stated above, and throughout the experiments. A closed chamber saturated with water was placed around the sample in order to avoid evaporation. Dynamic oscillations were performed, i.e. the sinusoidal strain $\gamma = \gamma_0 \sin(\omega t)$ was measured corresponding to the applied shear stress $\tau = \tau_0 \sin(\omega t + \phi)$, where ω is the angular frequency, and t is time. γ_0 and τ_0 are respectively the amplitude of the deformation and the amplitude of the stress. Therefore, dynamic moduli (G' , G'') can be obtained at different frequencies f (where $\omega = 2\pi f$) in the range [0.01-2Hz] by selecting the in-phase elastic deformation (G') and its out of phase viscous counterpart (G''): $G' = \tau_0 \cos\phi / \gamma_0$ and $G'' = \tau_0 \sin\phi / \gamma_0$. The viscoelastic properties (G' , G'') of our materials were measured using parallel plate geometry (20mm diameter) and a gap ranging between 500µm-800µm, since samples are of limited volumes. This is similar to the geometry previously used with CHO cell suspensions [24]. Experiments were started after gel polymerization and rheometer setup (see §2.1), which means that one hour had elapsed. The gels have a linear domain at moderate deformations γ_0 , therefore collagen samples were tested under low strains ($\gamma_0 = 0.01$) within the linear domain, in agreement with previous work [3], and similarly for cell-filled networks.

Fig. 1: Rheometrical data (G' , G'') for the four collagen gels considered (0.42-0.95-1.38-1.8 mg/mL).

Gels typically exhibit a flat plateau for $G'(f)$ – or slowly increasing $G'(f)$ – and a smaller $G''(f)$ modulus (see Fig. 1), with $G''(f)$ also slowly increasing. This is the signature of a viscoelastic behavior. In what follows, we chose to show only the dependence of G' vs. frequency f , for this is the most significant modulus, and also in order to have a limited number of curves in each graph for clarity. In most cases, the G'' modulus just followed a very similar trend as compared to G' , and was lower by half a decade (factor of 3 roughly).

2.3 Confocal Microscopy

To characterize structural changes in our model tissues, confocal microscopy was used in addition to rheometry. The idea was to analyze cell shapes as well as their behavior within the

collagen network, and compare with the rheological data. The collagen matrix organization could also be analyzed. Birefringence is indeed an intrinsic optical property of many natural ordered biopolymers, including collagen fibers [6,28]. The collagen-induced depolarisation of the linearly polarized incident light allows the imaging of nascent 3D fibers in amorphous medium with high signal-to-noise ratio in confocal reflection mode. Confocal reflection microscopy and fluorescence were proposed earlier as powerful coupled tools to study 3D cell migration [20], and were used here to examine the organization of cells within the matrix. When combined with confocal microscopy, reflection images taken at sequential focal planes along the z axis can be reconstructed in 3D for visualization and for image analysis. For an easy visualization of cells inside collagen gels, we have developed a stable CHO transfected cell line expressing GFP. Therefore reflected light from the collagen matrix was collected simultaneously with fluorescence from cells in a time-lapse mode to monitor the dynamic process of cell behavior through the 3D collagen matrix. In order to do so, collagen gels with and without cells were polymerized in a Lab-Tek chamber with coverglass and imaged using a confocal microscope (inverted Zeiss Axiovert 135M, LSM 510 model). The method was performed to visualize matrix fibers and cells using a 40X objective lens (Zeiss, C-Apochromat, water immersion). During the experiments, samples were placed in a temperature and CO₂ controlled on-stage incubator (37°C, 5% CO₂). Two lasers were coupled to the microscope: an Argon laser at 488 nm was used to visualize the GFP-labeled cells and a HeNe laser at 633 nm to visualize collagen structures.

2.4 Image processing

In order to get more precise data on cell morphology, an image processing method was used. Cells from confocal images (z-stack) were first extracted and subdivided into many building blocks using MetamorphTM. To process the data, we used the Edit3D_quant

software (kindly provided by Yves Usson, TIMC-IMAG, Grenoble). The analysis and 3D visualisation software were written in standard C language using the GL library (Silicon Graphics library) and the FORMS user interface library (courtesy of Marck Overmars, Department of Computer Science, Utrecht University, The Netherlands) [39,40,52]. The software gives the 3D-shape of the cell, its volume including the nucleus, as well as other relevant parameters (surface, major and minor axes, etc.). From this data, the cell elongation was calculated by fitting an ellipsoid around the cell and choosing the elongation ($e=M/m$) to be the ratio between major and minor axes (M and m respectively, see for example Fig. 8a-8b).

Finally, determination of the collagen densities in Fig. 10-11-12 were made thanks to the software VolocityTM (PerkinElmer Inc). Various boxes (parallelepipedic ones, cubes, or isoparametric volumes) were chosen at specific locations, and the ratio between collagen volume and box volume gave the collagen density (considered as arbitrary units).

3. Results

3.1. Rheology

The collagen solutions were tested first. The frequency dependence of G' and G'' at 1% deformation was measured for frequencies f ranging from 0.01 to 2 Hz at 37°C. The elastic modulus was always larger than the loss modulus. Both moduli increase very slowly with f and are almost parallel (see Fig. 1), a result quite similar to a previous one [19,29] obtained for collagen gels (2mg/mL) in oscillatory shear at a constant strain of 1%. The order of magnitude of the plateau modulus G_0 (i.e. the value for which G' is constant or has an inflexion point, here the value at 0.1 Hz was used) at a collagen concentration of 1.8mg/mL is of the order of 75 Pa, in good agreement with previously reported values [19,33,48,55]. The four collagen gels tested here (0.42-0.95-1.38-1.8 mg/mL) all exhibit increasing viscoelastic

moduli with collagen concentration. This can be seen in Fig. 1 and also as part of the data in Fig. 2 where the evolution of the elastic modulus G' is presented vs. frequency at four different concentrations (and three cell concentrations).

Fig. 2: Elastic modulus $G'(f)$ as a function of frequency f at four fixed collagen concentrations (0.42-0.95-1.38-1.8 mg/mL) using different cell concentrations: 0 - 0.7×10^7 - 1.17×10^7 - 1.8×10^7 cells/mL.

The cells-filled collagen networks were characterized next. Measurements of storage and loss modulus $G'(f)$ and $G''(f)$ were respectively carried out in the frequency range $0.01 \text{ Hz} < f < 2 \text{ Hz}$ at three cell concentrations ($0.7 \cdot 10^7$, $1.17 \cdot 10^7$, $1.8 \cdot 10^7$ cells/mL) for each collagen network. The results are also shown in [Fig. 2](#). We observe that for both situations, with cells and in the absence of cells, the viscoelastic moduli always increase with collagen concentration, when matrices with the same cell concentration are tested.

An interesting feature is found when solutions with different cell concentrations are analyzed, as can be seen in [Fig. 2](#).

- At small collagen concentration ($c \leq 0.95 \text{ mg/mL}$), the collagen elastic modulus G' is lower than the cell-collagen networks modulus. As more cells are included (from $0.7 \cdot 10^7$ cells/mL to $1.8 \cdot 10^7$ cells/mL), the elasticity of the network increases further.
- On the other hand, at higher collagen concentrations ($c \geq 1.38 \text{ mg/mL}$), the collagen elastic modulus G' is reduced when cells are included into the network. This is shown by the dramatic decrease of the elastic modulus G' when $0.7 \cdot 10^7$ cells/mL are included within the matrix. When more cells are added (i.e. $1.38 \cdot 10^7$ cells/mL and $1.8 \cdot 10^7$ cells/mL), the network elasticity increases again and can reach its initial level ($c=1.38 \text{ mg/mL}$) or higher ($c=1.8 \text{ mg/mL}$). This feature therefore seems to be a concentration-dependent phenomenon and will be discussed in the next part.

In [Fig. 3](#), we present the evolution of the elastic modulus G' at fixed cell densities, when varying the collagen concentration. G' increases with frequency f as usual, and always increases with collagen density.

Since the frequency dependence (slow increase of G') is in agreement with previous works, we clarify this analysis further by representing G' (more precisely its value G_0 at $f=0.1 \text{ Hz}$) as a function of concentration, in the 16 different cases studied. This is shown in [Fig. 4](#) using a log-log scale. This representation is interesting because it can allow one to determine the

exponent of the power law model. Experimental data from other works have indeed shown the concentration dependence of the plateau shear modulus to vary like $c^{2.2}$ in the case of actin gels, $c^{2.1}$ for fibrin networks [25], and c^3 for collagen networks [47]. Our data gives a dependence $G' \approx c^{2.6}$, in excellent agreement with previously reported data.

Fig. 3: Elastic modulus $G'(f)$ as a function of frequency f at four fixed cell concentrations (0.7×10^7 - 1.17×10^7 - 1.8×10^7 cells/mL) using different collagen concentrations : 0.42-0.95-1.38-1.8 mg/mL (same as Fig. 2 but shown in a different way).

The presence of cells embedded into the matrix shows a different dependence vs. collagen concentration (Fig. 4), leading to a smaller slope of order 1 (more precisely the slopes of the fits are respectively 1.11, 1.08 and 0.99 for cell concentrations respectively 0.7 - 1.17 - 1.8×10^7 cells/mL) at the three cell concentrations studied. Again the dramatic effect of cells embedded in the network is outlined by the crossover of the collagen curve with the cell-filled network data at $c > 1.38$ mg/mL.

To understand this cell effect, it seemed important to study the microstructure of the matrices, including the way cells organize within the gels, in particular through interactions with the filaments. This was possible thanks to complementary confocal microscopy data which is presented next.

Fig. 4: Concentration dependence of the shear elastic modulus G_0 (0.1 Hz) vs collagen concentration (log-log scale) with and without cells. The fits indicate power law with respective exponents 2.6 for collagen gels alone and 1.11-1.08-0.99 for cell concentrations 0.7×10^7 - 1.17×10^7 - 1.8×10^7 cells/mL.

3.2 Confocal Microscopy

3.2.1 Cells within the collagen matrix

To understand first how the matrices behave, collagen gels with the same concentrations as above (0.42-0.95-1.38-1.8 mg/mL) were prepared and analyzed using confocal reflectance microscopy. The images were taken 120 μm deep into the sample and revealed the presence of the collagen fibers. A simple visual inspection suggested that the matrices present an increase in fiber density (or decrease in mesh size) with increasing collagen concentration, as expected [28]. Fig. 5 display this clear effect and the analysis of global pixel intensity versus concentration revealed a monotonous increase (data not shown), confirming the validity of our data.

Fig. 5: Confocal reflectance images at four collagen concentrations demonstrate the architectural differences between matrices prepared from type I collagen. Bar = 40 μm for all images. (a) 0.42 mg/mL (b) 0.95 mg/mL (c) 1.38 mg/mL (d) 1.8 mg/mL.

The mechanical impact of the cells on their microenvironment (i.e. collagen fibers) was investigated next. Cells (1.8×10^7 cells/mL, i.e. 12% volume concentration) were embedded within gels (see Materials and Methods) with collagen concentrations ranging from 0.42 mg/mL to 1.8 mg/mL. This is shown in Fig. 6a-d at a low magnification first.

Fig. 6: Confocal reflectance and fluorescence microscopy for collagen gels with cells; (magnification 40x, 1 μ m z-slice, xy: 358 x 358 μ m, Zoom 0.7). (a) 0.42 mg/mL (b) 0.95 mg/mL (c) 1.38 mg/mL (d) 1.8 mg/mL.

- It seems obvious that the elongation of the cells (around 3 and 3.5) in [Fig. 6a](#) at low concentration is important. Cells adhere easily and show long protrusions. Such cells seem to deform the collagen fibers in their close neighborhood and probably also pull collagen towards them, i.e. they remodel it.

- On the other hand, cells seem to be less elongated (aspect ratio or elongation slightly higher than 1) in the next two Fig. 6b-6c. Stiffer matrices seem to lead to round cells because they might be affected by a lack of space. Still cells are able to make short protrusions.

Fig. 7: Confocal reflectance and fluorescence microscopy for collagen gels with cells; (magnification 40x, 1μm z-slice, xy: 76.8 x 76.8 μm, Zoom X3). (a) 0.42 mg/mL (b) 0.95 mg/mL (c) 1.38 mg/mL (d) 1.8 mg/mL.

- Finally, at large collagen concentration (Fig. 6d) cells seem to be stuck within the concentrated collagen. They keep a round shape (aspect ratio or elongation close to 1) similar to the ones observed previously in a dense collagen matrix 5mg/mL [23]. Interesting new

regions filled with fluid (i.e. culture medium) appear and maybe due to the interplay between cell tractions/remodelling and fluid pressure.

To quantify these observations we analyzed image stacks by selecting magnified single cells (zoom X3) embedded into collagen gels at the different concentrations (0.42-0.95-1.38-1.8 mg/mL). [Fig. 7a](#) confirms that, in a less concentrated collagen gel, cells present a large deformation/elongation, in contrast with higher collagen concentrations where cells keep their almost round shape with very small protrusions ([Fig. 7b-7c-7d](#)) corresponding to an amoeboid motion. In addition, [Fig. 7d](#) shows a high density of fibers around the cell in agreement with the fact that cells play an attractive remodelling role of the collagen. To quantify these effects more precisely, an image processing method was used. Cells were isolated from the images (see Materials and Methods) and their global parameters (positions, surface, volume, etc.) were calculated. We focused on their elongation which seems to be a relevant parameter here. [Fig. 8a](#) shows typical projected shapes of the most elongated cells found for the small collagen density (0.42mg/mL). Collecting results for a sufficient number of cells in different collagen matrices (18-16-19-16 for collagen concentration of 0.42-0.95-1.38-1.8 mg/mL respectively) leads to the chart in [Fig. 8b](#) where the number of cells vs. elongation is shown for the four collagen concentrations. Finally the data is collected in [Fig. 9](#) where cell elongation is represented against collagen concentration. It appears that the cell elongation has a mean value around 2 at 0.42 mg/mL, then decreases down to 1.2 at 0.95 mg/mL then it increases slightly again to reach a plateau of 1.5 at the higher concentrations (1.38 or 1.8 mg/mL). Probably the explanation of such a complex shape for this curve needs to be understood based on the competition between different effects. This will be discussed in the next part.

Fig. 8: (a) Examples of increasingly elongated cells in a low density collagen gel (0.42 mg/mL) obtained by surface projection using Edit3D. (b) Histograms showing cell elongation for four different collagen concentration.

Fig. 9: Cells elongation vs. collagen concentration. Error bars are standard errors of the mean

3.2.2 Detailed analysis of the remodelling process

We next consider a more careful analysis of the remodelling process, i.e. the way cells migrate through the collagen matrix and how they remodel the latter one. This was done using the Volocity softwareTM (see §2.4). Volumes can be selected and the sum of the intensities of the reflected light gives a measure of the collagen density, when referred to the global volume of the selected region. We have chosen to plot the data using arbitrary units (a.u.) and simple comparisons will be made here.

In a first experiment, we select one cell which is in motion. A small region of collagen (Fig. 10) located on the top left of the figure (see arrow) is followed in time. The collagen density measured in this region decreases as a function of time. Even after one hour, the density is reduced by a factor of two. Thus remodelling is quite fast. The overall effect is studied over 4 hours and the collagen density eventually goes down close to zero, thus

showing that cells leave empty spaces behind them, i.e. spaces filled with fluid alone (or collagen-free). This is the first effect.

Fig. 10: Evolution of collagen density in a fixed volume (white cube on the top left of the cell, indicated by the arrow), as a CHO cell migrates away from it : the collagen density clearly decreases during cell motion, thus the cell removes some collagen from this region. This effect is analyzed over a time period of 4 hours. Even after one hour, the density is decreased by a factor of 2 (from 0.8 to 0.4 in a.u.). Photographs are taken at times indicated on the graph. A large hole is left after passage of the cell. This is one way to remodel the gel.

To investigate such behaviors more closely, the collagen density was measured in the neighborhood of a CHO cell: different parallelepipedic boxes were chosen, located few microns away from each other (see Fig. 11). As a comparison, the average collagen density was measured in a similar box located nearby, with no cell present. The collagen density increases as one moves closer to the cell, in particular very close to it (1 μ m away): this is where it has a maximum, as shown in Fig. 11. As one moves away, the density is reduced quite rapidly, for example it is reduced by a factor of two only 7 μ m away (from 0.8 to 0.4 a.u.). At this distance, it becomes already close to the value of the collagen density as

measured in a nearby region (distant by two cell diameters). In this nearby region, its value is a fluctuating one (around 0.2 a.u.), in good comparison with the overall collagen density (0.19) measured for the whole gel (0.9 mg/mL in this case).

Fig. 11: Collagen density (a.u.) measured in parallelepipedic regions close to a cell and in a region with collagen alone (see insert showing the positions of two such regions, collagen is not shown for clarity). Regions are moved away from the cell or throughout the gel (case with no cell). The arrow also indicates the direction of migration of the cell. The average collagen density is also indicated to be around 0.19 (a.u.) corresponding to 0.9 mg/mL. Collagen density decreases when moving away from the cell, as opposed to the almost constant collagen density measured in the region with collagen alone. Thus the closer to the cell, the higher the collagen density: cells accumulate collagen around them.

This shows that the collagen density is larger the closer to the cell, thus cells have the ability to compact collagen around them. This was also shown in Fig. 6a, where collagen fibers are clearly visible around cells, and organize in a star-like manner.

Finally, the evolution of the collagen density was measured within isoparametric volumes ($1\mu\text{m}$ in thickness) with a shape that resembles that of a cell, i.e. the volume was drawn around the cell at a given distance from the cell edge, as shown in the insert of Fig. 12. The density of collagen, as measured with VelocityTM, again shows that the collagen density decreases as one moves away from the cell. Three measurements were made at three different times $t=0$, $t=82$ min, and $t=137$ mn. The collagen density increases close to the cell; this shows again the ability of cells to attract collagen fibers. These effects will be further discussed in the next part (§4).

Fig. 12: Evolution of the collagen density in the neighborhood of a cell. Isodistance volumes ($1\mu\text{m}$ in thickness) have been used around a cell (see an example in insert) and the corresponding volume of embedded collagen fibers was measured within volumes spaced by $1\mu\text{m}$. Measurements are made at times $t=0$, 82 min, and 137 min. The volume occupied by collagen (i.e. collagen density) always increases with time around the cell and is higher closer to the cell. This shows the ability of cells to attract collagen fibers irreversibly. This is the second way to remodel the gel.

4. Discussion

The first part of this work deals with the intrinsic properties of collagen gels. Such properties result from the interactions between two components: a network of collagen fibrils and an interstitial fluid, i.e. the culture medium. When mechanical stress is applied, the network and the medium deform together and the resistance to the interstitial flow of the solution through the network can lead to high pressure, which allows the gel to withstand compressive loads [29]. In shear, the resistance is quite important and the elastic moduli G_0 can reach rather large values, in agreement with previous works [19,29]. The ability of the network to resist such stresses is due to its particular dense crosslinked microstructure as exhibited in Fig. 5. Our data for collagen gels is also comparable (Fig. 1-2) with the results obtained in laser trap microrheology [48] where the authors obtained an increasing shear modulus G with c , the collagen concentration. The data obtained for $G'(0.1 \text{ Hz})$ vs. c exhibits a typical slope of 2.6 (see Fig. 4), in agreement with previous works on actin and collagen networks [25,47].

In the second part of this work, cells are embedded into the gel. It is shown, for the first time, using rheological measurements, that the matrix elasticity is globally reduced by a rather large amount at high values of c , this effect being due to the presence of cells (see Fig. 2-3-4). This effect, although surprising, might be due to the ability of cells to remodel the matrix, as fibroblasts were earlier shown to condense a tissue into a fibrillar structure [5], i.e. to compact it [16,17] or further colonize it [22,23]. Other living cells (like bacteria) can also modify the viscoelasticity of certain gels [8]. This effect has been considered by Leung and co-workers using a different technique [33] using bead twisting microrheology to investigate the effect of fibroblasts in a collagen matrix. The results are similar : local elasticity of the collagen matrix is higher without cells, and decreases when cells are embedded in the matrix (25K, 50K and 100K cells/mL, less than what was used in the present study). The authors also checked this

effect except for very long durations (larger than 7 days). We restrict here the analysis to short term behavior ($<24\text{h}$) because it is known that other mechanisms may take place at longer times, for example the expression of collagenase (MMP1) and gelatinase A (MMP2) leading to matrix degradation/hydrolysis [22]. Finally the modulus reduction observed at short times [33] is in agreement with our experiments.

The nonlinear effect with cell concentration found in Fig. 4 was also observed by Evans and Barocas [16]: their data shows a decrease of the Young Modulus (E) at small cell concentration, then an increase at higher cell concentration ($c=3\text{ mg/mL}$). At lower values of c ($1\text{-}2\text{ mg/mL}$), E always increases as in our case. These nonlinearities maybe due to the nonlinear properties of collagen [17], but are also due to the presence of cells, as will be discussed next. Another observed effect was the slope change of G_0 vs. collagen concentration c (Fig. 4). This slope changes drastically from a value of 2.6 to a value of 1 when cells are embedded into the gel. This effect further enhances the fact that cells play a definitive role on the organization of collagen fibers.

Let us now discuss these effects in view of the confocal microscopy analysis. The collagen networks appeared with a finer mesh as the concentration c increases, as expected (Fig. 5). Regarding cell-collagen networks, it was found (see Fig. 8-9) that cells behave differently depending on the collagen matrix concentration. Low concentrations lead to more elongated cells (Fig. 6a-7a-8a) whereas larger concentrations lead to smaller round cells (Fig. 6b-6c-6d and Fig. 7b-7c-7d); Had the concentration been smaller, then single contacts with isolated fibers would have been found [20], but this is not the case here. It has also been emphasized that new regions appear in the cell-collagen matrix (Fig. 6d) where fluid-filled spaces are present. Figure 8b and Fig. 9 quantify these effects: cell elongation (ratio of the length over width of the corresponding cell-fitting ellipse) vs. collagen concentration seems to be a decreasing function (with a possible minimum). To explain this feature, a simple analysis is

proposed based on the existence of two competing effects leading to a minimum, a phenomenon also observed for 2D cell migration [2,38] :

(a) Binding/adhesion to collagen is important for cells and is enhanced by the presence of collagen or mechanosensing (b) Size effects (related to collagen mesh size) affect cell shapes as cells cannot spread in too dense regions. The results are as found experimentally:

- At small concentrations, cells elongate thanks to space; they encounter a few fibers only, that they attract (Fig. 6a).
- At large concentrations, cells remain small because of the lack of space and do not elongate so much, because they have enough fibers around them to bind.

Most important is the effect that cells can play on the microstructure of the network, and the tensions they can generate (Fig. 6-7), which leads to remodelling of the collagen network. This has been studied thanks to the determination of collagen densities within specific volumes chosen in the neighborhood of cells.

Figure 10 showed how a cell can move through a collagen network (0.9 mg/mL) by leaving holes behind it. This was shown by a rapid decrease of the collagen density in a box close to where the cell passed. This phenomenon can occur within one hour and is quite fast.

In a second experiment, the collagen density was measured in a parallelepipedic box located close to a cell. This box had a fixed volume and was moved away from the cell. The resulting collagen density (Fig. 11) was shown to decrease rapidly away from the cell, thus showing an enhanced amount of collagen in the neighborhood of the CHO cell. This was checked with a control volume in a nearby region (away from any cell), whose density (≈ 0.2) was found close to the value far away from the cell (≈ 0.2 , at a distance of $8\mu\text{m}$). Finally this latter value was found to be in agreement with the average collagen density in the whole gel (0.19) as indicated in Fig. 11. Therefore it is shown that cells attract collagen towards them.

Finally, isovolumes (distant by $1\mu\text{m}$ each, each having a thickness of $1\mu\text{m}$) were chosen around a cell edge to measure collagen density (see Fig. 12). The result is again striking: as time goes on (during the first two hours), the collagen density increases all around the cell but even more when getting closer to the cell. This shows an irreversible attraction of collagen fibers by CHO cells. Such fibers never go back to their original positions, this corresponding to plastic deformations.

Summarizing our data, the remodelling effect is comprised of two mechanisms :

- (1) the first one shows that some cells migrate through gels and leave tunnels “behind them”;
- (2) the second one is due to the attraction of collagen by cells exerting traction forces.

These new features can share some light when compared to previous works on the compaction of gels [5,17]. It is indeed shown here that cells, either by pulling on fibers or “drilling tunnels” through gels, can remodel it and create collagen-free regions (i.e. fluid streams). Then it becomes possible for the fluid to flow outside while the network is contracted. This is a microscopic explanation of the proposed macroscopic effect [5,17].

Fig. 13: Sketch of the possible structure formed by cells surrounded by collagen network: Cells reorganize the initial collagen matrix (left) by attracting collagen thus depriving other regions from collagen, and creating less dense regions where fluid can flow. The rigidity of the network on the right is smaller.

Finally, the drastic reduction of the elastic moduli obtained previously can be explained thanks to the ability of cells to remodel the collagen network. The initial network shows strong interacting links (Fig. 13, left, and confocal image in Fig. 5) corresponding to a rigid gel. Then cells remodel it, this leading to a softer network composed of weakly connected islands drained by the interstitial fluid as sketched in Fig. 13 (right) and also confirmed using confocal microscopy (Fig. 6d) in this present work. This effect was analyzed closely thanks to collagen density measurements in the neighborhood of cells.

5. Conclusions

In this work, we focused on the mechanical impact that CHO cells (a fibroblast type) can have on their microenvironment, i.e. a collagen network, and conversely. Bulk rheology, as well as confocal reflectance/fluorescence microscopy were used to explore the complex interplay between the different network elements.

Collagen matrices were found to exhibit viscoelastic properties with an elastic modulus following a nonlinear relation with collagen concentration in agreement with previous works. Regarding cell-filled matrices, we found that cells can affect the collagen rheology dramatically. This was shown by a strong decrease of the elastic shear modulus G' of cell-collagen networks at the highest collagen concentrations used ($c > 1$ mg/mL). Confocal microscopy enabled to explain this dramatic change by revealing the cell-collagen microstructure. In particular, remodelling of the network was explained by the ability of cells to attract the collagen fibers or leave “tunnels” behind them, thus leading to new microstructures and to the subsequent breakdown of the elastic modulus.

Perspectives of this work could focus on the ability of the confocal method to capture matrix fiber heterogeneities or orientations in cell-populated collagen gels [42,47]. But they could also find interest in the combined effects of cell remodelling, adhesion, myosin II contraction

[36], as well as proteolytic degradation by MMPs [13] and growth [6,10] on longer time scales [22,23]. Finally they would be ideally suited for studying cell migration in 3D collagen matrices.

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