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A numerical model suggests the interplay between nuclear plasticity and stiffness during a perfusion assay

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Abstract

Cell deformability is a necessary condition for a cell to be able to migrate, an ability that is vital both for healthy and diseased organisms. The nucleus being the largest and stiffest organelle, it often is a barrier to cell migration. It is thus essential to characterize its mechanical behaviour. First, we numerically investigate the visco-elasto-plastic properties of the isolated nucleus during a compression test. This simulation highlights the impact of the mechanical behaviour of the nuclear lamina and the nucleoplasm on the overall plasticity. Second, a whole cell model is developed to simulate a perfusion experiment to study the possible interactions between the cytoplasm and the nucleus. We analyze and discuss the role of the lamina for a wild-type cell model, and a lamin-deficient one, in which the Young's modulus of the lamina is set to 1% of its nominal value. This simulation suggests an interplay between the cytoplasm and the nucleoplasm, especially in the lamin-deficient cell, showing the need of a stiffer nucleoplasm to maintain nuclear plasticity.

Keywords: Nucleus, Plasticity, Finite Element, Viscoelasticity, Viscoplasticity

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1. Introduction

Cell motility is a fundamental cellular mechanism involved in several biological phenomena such as bone remodelling, immune response and tumor metastasis. Tumor metastasis is estimated to be responsible for 90 % of cancer death [1]. In such cases, where cell migration through sub-nuclear pores is often necessary, the nucleus plays a critical role due to its size and mechanical properties [2]. Hence, a better understanding of the processes behind cell migration and of nuclear mechanics is primordial in order to develop new therapeutic strategies in the fight against cancer. Most publications investigate the molecular mechanisms of cancer metastasis [3], of mechanical properties of the nucleus [4] and of the mechanical coupling between the nucleus and the cell's cytoskeleton [5, 6]: the aim here is to tackle these issues through a mechanical perspective. During migration or transmigration events, the cell is capable of going through narrow constrictions down to 10% of the size of the nucleus [7]. The large size and high stiffness of the nucleus make it a major obstacle to this process, although some cells are able to overcome such difficulties [6]. Its mechanical properties mainly arise from two components: the lamina – a dense meshwork composed of A-type and B-type lamins, as well as lamin-associated proteins [8] – and the nucleoplasm. The nucleoplasm is mostly made up of chromatin surrounded by fluid and can thus be seen as a viscoelastic material. The lamina however, as a dense meshwork where fluid cannot circulate, can be seen as a solid elastic material [9]. With sheer observation of its internal organization, we can already qualitatively propose a visco-elastic model of the nucleus. Since nuclear mechanics is at stake here, numerical simulation appears as a very interesting tool to investigate the mechanical interplay between cellular components, such as the lamina and the cytoplasm, and to get new insights on some biological assumptions. As for quantitative values, many techniques are accessible to study specific features of the nucleus, as will be discussed in the following paragraphs.

1.1. From experimentation to mechanical modelling

A wide range of experimental techniques are available to investigate mechanical properties of various components of the cell and of the nucleus at various scales [10]. They include perfusion [11], micropipette aspiration - possibly coupled with relaxation experiments - [12], Atomic Force Microscopy (AFM), active or passive micro-rheology through magnetic or optical tweezers with local information [13], microplate compression [14], substrate strain, micro-needle manipulation [15], shear flow and cytoindentation [16]. The first experimental setups focused on getting the mechanical behaviour of the overall nucleus. Depending on the technique, different behaviours at various scales can be studied.

The overall nucleus. The nucleus has widely been found to behave as a viscoelastic material, all throughout different techniques that enlightened various specific aspects of its behaviour [17, 18]. The perfusion and aspiration assays give a good assessment of purely passive mechanical properties of both the cell and the nucleus since they are fast enough — cells pass through in less than a second [11, 19, 20] — to assume no cyto- or nucleoskeleton reorganization occurs. Such assays showed that nucleus was 3-4 times stiffer and twice more viscous than the cytoplasm [12, 14]. The viscoelasticity of the nucleus is mostly accepted, but a hyperelastic behaviour of the nucleus is sometimes assumed in order to fit experimental data with simulation ones and to obtain quantitative data on the mechanical parameters of the nucleus [14]. Using these global measurement techniques, the Young modulus of the cell nucleus is estimated around 1-8 kPa [14, 21, 12, 22], but is sometimes found to be much higher depending on the experimental method [23]. This illustrates one of the limits of the global techniques that raise various uncertainties due to the interaction between the nucleus and the rest of the cell. Some recent advances in AFM techniques are used to get rid of this bias by probing the nucleus more locally [22].

The nuclear envelope (NE) and the lamina. Information on the whole nucleus is essential and easier to get, but the tight interaction between mechanical forces

and gene regulation induces to look more closely and precisely at local properties of the nucleus at the scale of specific proteins such as chromatin and lamins. Studies of the nuclear envelope alone are scarce, but combined techniques of micropipette aspiration and confocal microscopy were used to characterize the nuclear envelope as purely elastic [24]. Underlying the nuclear envelope is the lamina, a stiff material ensuring the nuclear stability, sometimes described as viscoelastic, although more thorough and velocity-dependent testing would be necessary to rigorously prove the viscoelastic behaviour [25, 26]. Together, the lamina and the nuclear envelope form a very thin layer of 10-200 nm surrounding the nucleus [27]. Given the very high stiffness of the lamina, the impact of the NE, as well as lamina's viscosity can be neglected. Such stiffness protects the cell and its genetic information, but can also be a rate-limiting factor during confined migration by preventing sufficient nucleus deformation. In fact, cells have to find a good compromise between viability and motility.

The nucleoplasm. The nucleoplasm behaves as a sponge-like material that initially does not present much resistance to deformation but this resistance increases as the chromatin gets compacted [21]. Besides, chromatin exhibits a plastic behaviour, i.e. irreversible deformation, at long time-scales and after shear stresses, which decreases with up-regulation of Lamin-A [28, 29]. This suggests that the nucleoplasm sets the rheological character of the nucleus while the lamina dictates the extent of the deformation. Such plasticity is an advantage during confined migration, since the nucleus stays elongated after going through a narrowing space, making it easier for the cell to migrate once the first constriction is overcome. Most often, the lamina is considered as the main load-bearing element in the nucleus, but recent findings suggest that chromatin itself is the main structural component of the nucleus [30].

Existing computational models of the nucleus and its mechanical behaviour. The interest for cell computational models has been rising in the last two decades, as it becomes more and more obvious that mechanics plays a major role during cell migration and even in gene transcription. One strategy for representing the

cell and the nucleus is discrete modelling, just considering the cell membrane or the nuclear envelope [31], another is an energetic approach [32], and finally, the cell or nucleus can be modeled through continuum mechanics [33]. Most continuum models describe the whole cell, with or without its nucleus as a separate compartment, but fewer model focus on the isolated nucleus, as reviewed in [34] and [16]. Our approach here is to propose a model which can be employed for both the isolated nucleus and the whole cell. The nucleus, if proven to be visco-elastic, is sometimes modeled as a hyperelastic material in order to simplify the simulation and to investigate specific mechanical issues [14, 35, 32]. A visco-elastic model was later developed to simulate a micro-pipette aspiration assay [36], with the lamina and nuclear envelope taken into account. A more advanced model was proposed with the nucleus described as a poroelastic material with a plastic behaviour [37]. Interestingly, this model faithfully reproduced the irreversible deformation found in Lamin A/C deficient cells after transmigration. Although we acknowledge the validity and the interest of all these models, we observe that each one of them is designed to fit a specific experiment and can wonder whether one single model could describe several different assays. This is specifically what we aim to tackle in this article: a unified model of the whole cell that can be confronted with various experimental techniques.

1.2. *The proposed model*

Given the major role of nuclear mechanics during confined cell migration, this paper will present a two dimensions (2D) FE implementation of a cell nucleus model, representing the nuclear lamina as elastic and the nucleoplasm as visco-elasto-plastic. Even though a 3D model would be more accurate, we chose a 2D representation to facilitate the computation, since it was shown that for a cell entering a micro-channel, the model is insensitive to depth [38]. While the whole nucleus is generally described as merely viscoelastic, we decided to design a new model to be able to account for a more complex behaviour of the nucleus including plasticity. Besides, we aim at developing a model that is able to be tested in various experiment-like setups, but we focus here on modeling a

purely passive cell to fully understand the mechanics at stake without migration or skeleton reorganization. In this regard, we first build a model of an isolated nucleus that will be tested under compression mimicking the experimental setup from Caille et al. [14]. This model being thoroughly investigated, it will be implemented in a whole cell model, modeling the cytoplasm as in previous work [33]. This complete model is then tested to reproduce a perfusion experiment [11]. The mechanical parameters of the cell and its nucleus were chosen to match those of the HeLa cell, as in our previous work [33], but the versatility of our model would allow it to cover a wide range of cell types by simply adjusting the mechanical parameters.

2. Numerical simulation of a compression test of the nucleus

2.1. Nucleus geometry

The nucleus ($\Omega_{nucleus}$) has an initial circular geometry of radius $r_{nucleus} = 4 \mu m$ and is composed of the 50 nm thick lamina (Ω_{lamina}) and the nucleoplasm ($\Omega_{nucleoplasm}$) (See Figure 1). Both the lamina and the nucleoplasm have been described through characteristic functions g_l and g_{np} which are a composition of a regularized Heaviside function and a level set function (see Appendices A and B). The characteristic function representing the whole nucleus is given by $g_n(\mathbf{x}) = g_{np}(\mathbf{x}) + g_l(\mathbf{x})$, where \mathbf{x} indicates the current position of any particle of the system.

2.2. Constitutive model and mechanics of the nucleus

In most models, the nucleus is simply described as a viscoelastic material. In order to tackle its potential irreversible deformation, we chose to implement a visco-elasto-plastic material composed of the lamina, purely elastic, and the nucleoplasm, visco-elasto-plastic, as represented in the schema in Fig 2. Both nuclear components were assembled in parallel since we consider the stress from both components to add up. Indeed, the lamina being very thin, the associated mesh would need to be too small. The mechanical influence of the lamina is

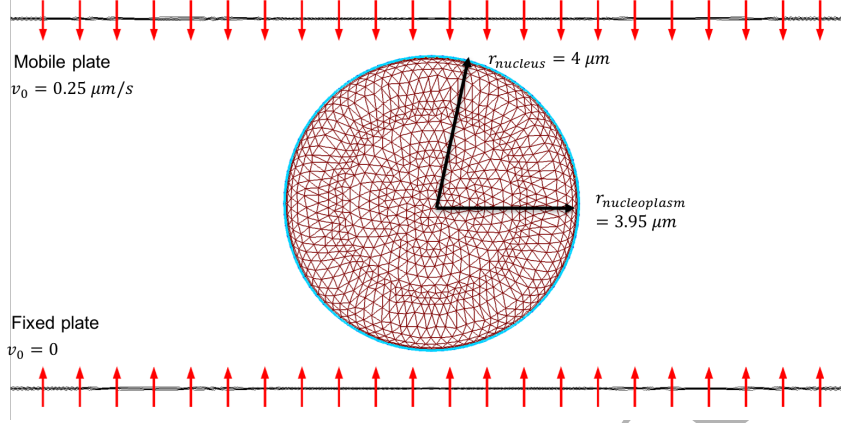


Figure 1: Geometry of the nucleus (Dark red: nucleoplasm and Light blue: lamina) and compression plates (red arrows represent the outward normal to the plates $\mathbf{n}_{\text{plate}}$) and FE mesh

thus not taken into account unless we homogenize the whole nucleus with volume ratios through the Voigt homogenization [39] (See Equation 4). An Arbitrary Lagrangian Eulerian (ALE) formulation is used in all the simulations, which allows to consider only small strains in the moving frame and Cauchy stresses [40, 41]. The total Cauchy stress $\boldsymbol{\sigma}_n$, where the subscript n stands for nucleus, and the small deformation tensor $\boldsymbol{\epsilon}_n$ are defined as

$$\begin{aligned}\boldsymbol{\sigma}_n &= (\boldsymbol{\sigma}_{np} + \boldsymbol{\sigma}_l) \\ \boldsymbol{\epsilon}_n &= \left(\frac{1}{2} (\mathbf{D}_{\mathbf{x}} \mathbf{u} + \mathbf{D}_{\mathbf{x}} \mathbf{u}^T) \right) = \boldsymbol{\epsilon}_{np} = \boldsymbol{\epsilon}_l\end{aligned}\tag{1}$$

where the subscripts np and l indicate the nucleoplasm and the lamina respectively, \mathbf{u} is the displacement, $\mathbf{D}_{\mathbf{x}} \mathbf{u}$ is the usual displacement gradient.

The nucleoplasm itself is decomposed into a visco-plastic part and a purely elastic part as follows

$$\begin{aligned}\boldsymbol{\sigma}_{np} &= \boldsymbol{\sigma}_{np,e} = \boldsymbol{\sigma}_{np,vp} \\ \boldsymbol{\epsilon}_{np} &= \boldsymbol{\epsilon}_{np,vp} + \boldsymbol{\epsilon}_{np,e}\end{aligned}\tag{2}$$

where the subscripts vp and e stand for visco-plastic and elastic, respectively.

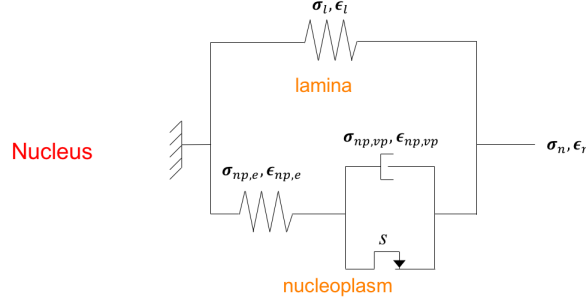


Figure 2: Rheological model of the homogenized nucleus

The constitutive equations of each part reads more specifically:

$$\begin{aligned}
 \sigma_l &= \lambda_l \text{Tr}(\epsilon_n) \mathbf{I} + 2\mu_l \epsilon_n \\
 \sigma_{np} &= \lambda_{np} \text{Tr}(\epsilon_n - \epsilon_{np,vp}) \mathbf{I} + 2\mu_{np} (\epsilon_n - \epsilon_{np,vp}) \\
 \dot{\epsilon}_{np,vp}^D &= \frac{h(\sigma_{np,VM} - s) |\sigma_{np,VM} - s|}{\eta_{np}} \frac{\sigma_{np}^D}{\|\sigma_{np}^D\|}
 \end{aligned} \tag{3}$$

where λ_l , μ_l , λ_{np} and μ_{np} are the Lamé coefficients of the lamina and the nucleoplasm, respectively and defined as $\lambda_k = \frac{E_k \nu_n}{(1+\nu_n)(1-2\nu_n)}$ and $\mu_k = \frac{E_k}{2(1+\nu_n)}$, $k = \{l, np\}$. E_l , E_{np} and ν_n are the Young moduli of the lamina and the nucleoplasm respectively and the Poisson ratio of the nucleus. Tr defines the trace operator, \mathbf{I} is the identity matrix and \mathbf{A}^D indicates the deviatoric part of the tensor \mathbf{A} defined as $\mathbf{A}^D = \mathbf{A} - \frac{1}{2} \text{Tr}(\mathbf{A}) \mathbf{I}$. We hypothesize that $\epsilon_{np,vp}$ only has a deviatoric part. We note h the regularized Heaviside function, s the plasticity threshold, $\sigma_{np,VM}$ the Von Mises stress of the nucleoplasm and η_{np} the viscosity of the nucleoplasm defined as $\eta_{np} = \tau_{np} E_{np}$, with τ_{np} its characteristic time [35].

Due to the very small thickness of the lamina (50 nm), the Young moduli were weighted with regard to the surface occupied by the lamina and the

nucleoplasm as:

$$\begin{aligned} E_{np} &= E_{np,0} \frac{A_{np,0}}{A_{n,0}} \\ E_l &= E_{l,0} \frac{A_{l,0}}{A_{n,0}} \end{aligned} \quad (4)$$

with $A_{np,0}$, $A_{n,0}$, and $A_{l,0}$ being the initial areas of the nucleoplasm, the nucleus and the lamina, respectively.

The nominal values of the mechanical parameters $E_{l,0}$, $E_{np,0}$, τ_{np} and the Poisson ratio ν_n were taken from the literature and previous works. There is no experimental data allowing us to determine the value of the plasticity threshold s , so we performed various tests in order to find a consistent value. All parameters values and references are listed in Table 1.

2.3. Compression experiment

Initially, the nucleus is placed between two rigid plates (Ω_{plate}) of length l_{plate} and height g_{plate} , the lower plate being fixed and the upper one being mobile, to simulate a compression experiment as in Caille et al. [14]. All parameters were chosen so that the compression speed is the same as in Caille et al. and the nucleus is compressed up to 70 %. Gravity is applied to the nucleus before the compression cycle begins. The upper l_{up} and lower l_{lp} plates are described by two characteristic functions $g_{up}(\mathbf{x})$ and $g_{lp}(\mathbf{x} - \mathbf{u})$, where $\mathbf{x} - \mathbf{u}$ indicates the initial position of any particle of the system (see Appendix C), in order to give the obstacles position to evaluate the contact force.

The friction force between the plate and the cell when they are in close contact is neglected. A normal force is introduced to control the contact between the nucleus and the plates, as follows

$$\mathbf{f}_{plate}(\mathbf{x} - \mathbf{u}) = \mu_{plate} g_{plate}(\mathbf{x}) \mathbf{n}_{plate} \quad (5)$$

where μ_{plate} is the penalization coefficient, \mathbf{u} is the displacement and $g_{plate}(\mathbf{x}) = g_{up}(\mathbf{x}) + g_{lp}(\mathbf{x})$ is a characteristic function that varies smoothly between 0 and 1 on a given interpenetration depth equal to $0.2 \mu m$. This depth can be adjusted

by varying the scaling parameter of the Heaviside function. Consequently, because of the property of the level set, no such force is applied when the plate and the nucleus are far from each other. When they become close enough, an eventual small overlap induces a large repulsive force cubically dependant of the overlap depth. Lastly, \mathbf{n}_{plate} is the outward normal to the plates.

As described in previous work [33], the global equilibrium of the system is expressed as

$$\rho_n \mathbf{a} = \mathbf{Div}(\boldsymbol{\sigma}_n) + \mathbf{f}_{plate} + \rho_n \mathbf{g} \quad (6)$$

where ρ_n is the nucleus density, \mathbf{a} is the acceleration and \mathbf{Div} is the usual divergence and \mathbf{g} is the gravity acceleration. \mathbf{f}_{plate} is the contact force, considered here as a localized body force in the neighbourhood of the contact with plate.

2.4. Results

The nucleus undergoes a loading/unloading cycle with T_1 , T_2 and T_3 respectively being equal to 10, 27.5 and 2 s. First, the nucleus settles down on the fixed lower plate under the action of gravity ($0 < t < T_1$). Then, the mobile upper plate goes down and compresses the nucleus up to 70% ($T_1 < t < T_1 + T_2$) and keeps the maximum compression for a time T_3 . Lastly, the upper plate goes back to its initial position as it unloads the nucleus.

In a first simulation, we set $E_{l,0}$ to 3000 Pa, $E_{np,0}$ to 25 Pa, τ_{np} to 2 s and the plasticity threshold to 4 Pa. Our model displays a non linear force-normalized deformation (with d_N the normalized vertical deformation of the nucleus) relationship during compression that shows a similar profile than those in Caille et al. [14] (Figure 3a). Additionally, a plastic behaviour is observed since the norm of the average deviatoric strain ϵ_n^D reaches a peak of 70% at $t = 30$ s as expected, but does not drop back to zero once the nucleus is unloaded (between $t = 35$ s and $t = 50$ s). In fact a residual strain of about 8% is found (Figure 3b).

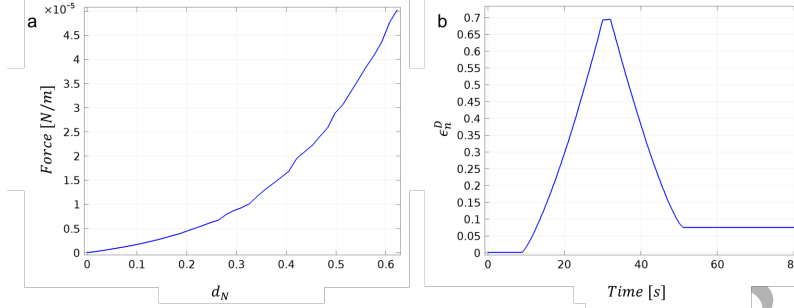


Figure 3: Simulation of compression and release of the nucleus. (a) Force-Normalized displacement curve, (b) evolution of the deviatoric strain ϵ_n^D in the nucleus with respect to time

Four test cases were implemented to study the influence of several parameters on the behaviour of the model. First, we examined the response of the nucleus model for $E_{l,0}$ equal to 100, 500, 1000, 3000 and 10000 Pa (Figure 4). When the lamina's Young modulus increases, the force needed to compress the nucleus increases as well (Figure 4a), while the plastic deviatoric strain after unloading decreases (Figure 4b). To compress the nucleus up to 50 %, the force ranges from 6 $\mu\text{N/m}$ for $E_{l,0} = 100$ Pa to 80 $\mu\text{N/m}$ for $E_{l,0} = 10000$ Pa, while the plastic deviatoric strain after unloading ranges from 30% to 2.5% (Figure 4a).

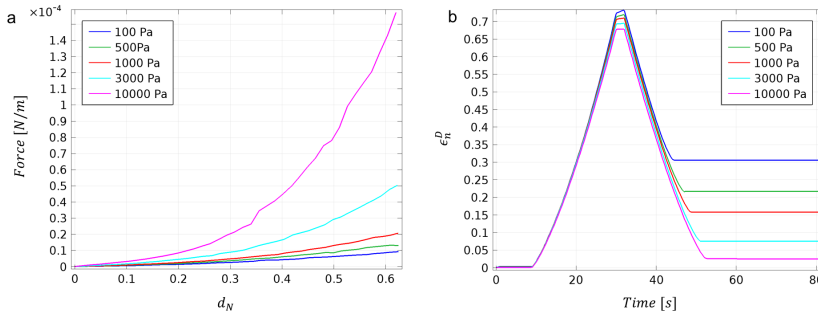


Figure 4: (a) Force-Normalized displacement curve and (b) evolution of the nuclear deviatoric strain ϵ_n^D with respect to time for various values of $E_{l,0}$

Thus, the lamina seems to be a major load-bearing element of the nucleus

and a lower Young modulus, which can be correlated with lamin-deficient nuclei, triggers higher nuclear plasticity. This result is consistent with a recent study on nuclear deformability [37] in which irreversible nuclear deformation was observed after transmigration of Lamin A/C deficient cells.

Although most reviews focus on the lamina [42, 43], the recent work of Stephens et al. highlighted the fact that chromatin may play a critical role as well [30]. The influence of the stiffness of the nucleoplasm, in which the chromatin has a major impact, was then considered with $E_{np,0}$ equal to 1, 25, 100 and 1000 Pa (Figure 5). As expected, the force required to compress the nucleus increased with the Young modulus (Figure 5a). Contrarily to the previous case, a higher nucleoplasm stiffness yielded to an increased plasticity (Figure 5b). The deviatoric strain upon unloading of the nucleus goes down at various speeds due to the definition of $\eta_{np} = \tau_{np} E_{np}$. Here, the force required for a 50 % compression ranges from 20 to 100 $\mu\text{N}/m$ (see Figure 5a). Thus, as well as the lamina, our model features the nucleoplasm as a potential load-bearing element.

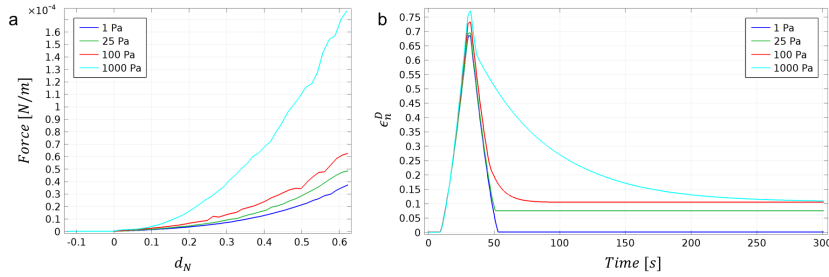


Figure 5: (a) Force-Normalized displacement curve and (b) evolution of the nuclear deviatoric strain with respect to time for various values of $E_{nucleoplasm,0}$

Finally, τ_{np} and s could also affect the overall mechanical behaviour of the nucleus. The nucleoplasm characteristic time has been found to be very disperse depending on the cell and experiment type. In this simulation, τ_{np} was set to 0.1, 1, 2, 10 and 30 s (Figure 6b). Longer characteristic times gave less plastic

behaviours down to only 2 %. As for the influence of s , with values of 0.1, 1, 2, 4, 10 and 30 Pa (Figure 6a), the last two values yielded roughly no plastic strain. It is interesting to notice that there is an increased plasticity from 0 to 5 % when the threshold increases from 0.1 to 4 Pa (see Figure 6).

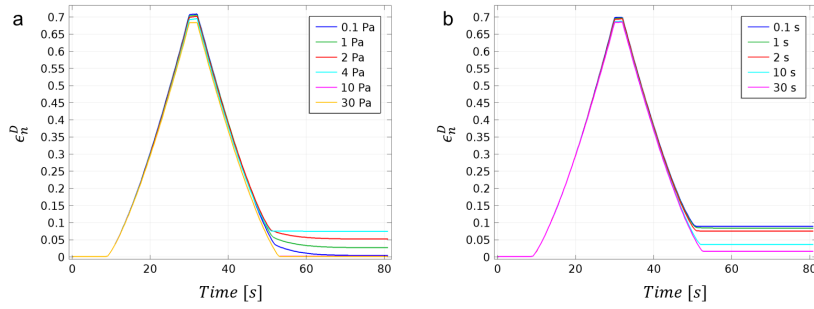


Figure 6: Deviatoric strain for various values of plasticity threshold (a) and $\tau_{nucleoplasm}$ (b)

2.5. Nuclear plasticity correlates with the lamin levels and nucleoplasm stiffness

In conclusion, our model does reproduce the visco-elastic behaviour of the cell's nucleus, as well as its plasticity upon higher strains. Depending on the values of Young's moduli of the lamina and the nucleoplasm, either of them can be the major load-bearing element of the nucleus. With respect to the experimental works in the literature, we chose to set the lamina as the major load-bearing element and the nucleoplasm as the one prone to plasticity. The aforementioned plasticity grows larger with increasing values of $E_{np,0}$ and with decreasing values of $E_{l,0}$ or τ_{np} . Smaller value of $E_{l,0}$ can be seen as the modeling equivalent of lamin-deficient nuclei, hence being in agreement with experimental results where Lamin A/C-deficient nuclei present a larger plastic deformation [37]. Now that the compression study has been completed, we aim to implement the nucleus constitutive law into a whole cellular model to simulate a perfusion test.

3. 2D model of a perfusion experiment

In this section we detail the 2D FE model developed to simulate a perfusion test. In this type of experiment, the cell passively flows through micro-channels in which a fluid flow is assured thanks to a pressure gradient [11]. Cell deformability and passing time can be studied. The objective here is to analyze the influence of the nucleus visco-elasto-plastic behavior during perfusion on the overall cellular response. Such influence will be studied more specifically on two phenotypes through variation of the mechanical properties of the lamina: a control model and a lamin-deficient model of the cell.

3.1. Cell geometry

Similarly to our previous works [33, 44], the cell (Ω_{cell}) is constituted by the surrounding actin cortex (Ω_{cortex}) and the cytosol ($\Omega_{cytosol}$), which form the cytoplasm ($\Omega_{cytoplasm}$), and by the lamina (Ω_{lamina} , Sec. 2.2) and the nucleoplasm ($\Omega_{nucleoplasm}$, Sec. 2.2), which form the nucleus ($\Omega_{nucleus}$, Sec. 2.2) (Figure 7). To describe the Ω_{cortex} and the $\Omega_{cytosol}$, we use, as we did for the lamina and the nucleoplasm (Sec. 2.1) two characteristic functions $g_{cx}(\mathbf{x})$ and $g_{cl}(\mathbf{x})$ as described in the Appendix D (Figure 7).

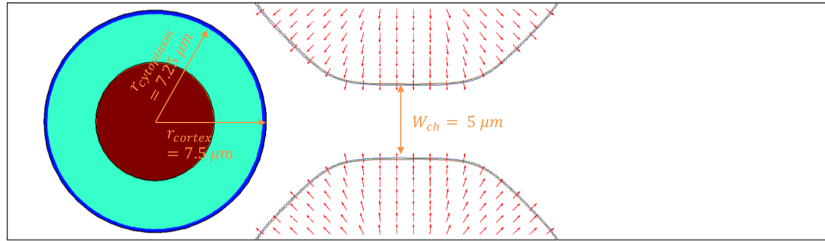


Figure 7: Cell and micro-channel geometry. In red is the nucleoplasm, orange is the lamina (too thin to be visible here), light blue is the cytosol and dark blue is the cortex. Red arrows represent the outward normal to the channel \mathbf{n}_{ch} .

3.2. Perfusion experiment

The perfusion device is modeled as a constriction channel ($\Omega_{channel}$) defined by two rigid walls. The upper (g_{uc}) and the lower (g_{lc}) walls (Figure 7) are defined by two characteristic functions (Appendix E).

We first performe a microfluidic study to obtain the velocity field inside the micro-channel. The fluid is described by the Navier-Stokes equation as follows

$$\rho_f \mathbf{a}_f = -\nabla p \mathbf{I} + \eta_f \Delta \mathbf{v}_f \quad (7)$$

where ρ_f is the volumetric mass of the fluid, \mathbf{a}_f the fluid acceleration, ∇ the gradient operator, p the pressure, η_f the fluid viscosity, Δ the Laplacian and \mathbf{v}_f the fluid velocity.

The boundary conditions include a pressure P_{in} at the left inlet and P_{out} at the right outlet. The fluid parameters are chosen in order to avoid turbulent patterns both at the entrance and the exit of the constriction (Figure 8 and Table 1). Finally, we assume that the velocity field calculated without the cell in the constriction does not vary over the perfusion assay, which constitutes a first step towards a more complex multiphysics model including fluid-structure interactions that will not be developed here.

The force \mathbf{f}_{fluid} exerted by the fluid on the cell cytoplasm is then defined as

$$\mathbf{f}_{fluid}(\mathbf{x}) = f_{fluid,0} \mathbf{v}_f \quad (8)$$

with $f_{fluid,0}$ the amplitude of the force applied on the cell, chosen so that the cell always passes through the channel in $1s \pm 0.05s$ in order to be able to compare the results (experimentally, the passing time in such devices ranges from 20 ms to a few seconds [45, 19, 20]).

Similarly to the compression test, a contact force is applied, now using the regularized characteristic function of the channel, as follows

$$\mathbf{f}_{ch}(\mathbf{x}) = \mu_{ch} g_{ch}(\mathbf{x}) \mathbf{n}_{ch} \quad (9)$$

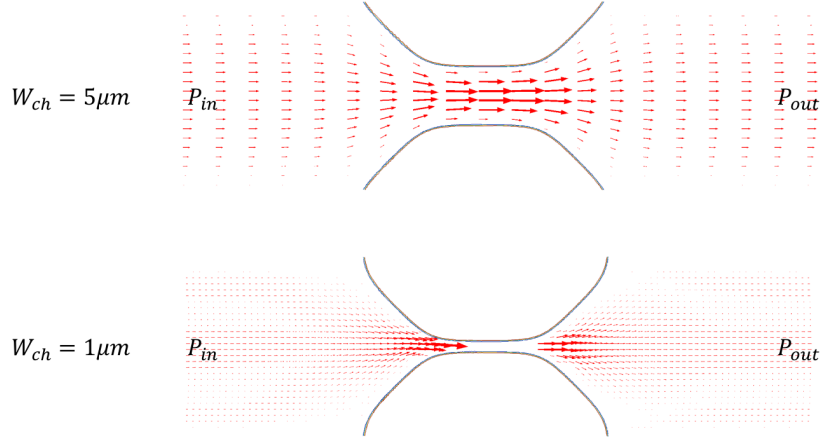


Figure 8: Profile of the fluid velocity inside the device for two sizes of constriction $W_{ch}=5$ and $1 \mu m$

where μ_{ch} is the penalization coefficient, $g_{ch} = g_{uc} + g_{lc}$ is a characteristic function as defined for g_{plate} in Section 2.3 and \mathbf{n}_{ch} is the outward normal to the channel.

3.3. Constitutive model and mechanics of the cytoplasm

As the nucleus constitutive law has been presented in Sec. 2.2, here we describe the mechanical behaviour of the other cell's components, namely the actin cortex and the cytosol. We assume that the cortex behaves as an isotropic elastic material, whereas the cytosol is described as viscoelastic, the whole cytoplasm thus being described by a generalized Maxwell model [44, 46, 33] (Figure 9). As previously developed, the cytoplasm is considered as a Voigt-homogenized material from its two constituents, namely the cortex and the cytosol. The whole cell is then a heterogeneous composed of the homogenized cytoplasm and the homogenized nucleus, such that $\boldsymbol{\sigma} = g_n \boldsymbol{\sigma}_n + g_{cp} \boldsymbol{\sigma}_{cp}$. The boundary conditions between these components are handled by the level set functions and the finite

elements. The Cauchy stress of the cytoplasm σ_{cp} is given by

$$\sigma_{cp} = g_{cp}(\mathbf{x})(\sigma_{cx} + \sigma_{cl}) \quad (10)$$

where the subscripts cp , cx and cl represent the cytoplasm, the cortex and the cytosol respectively, and g_{cp} represents the characteristic function of the cytoplasm (see Appendix D).

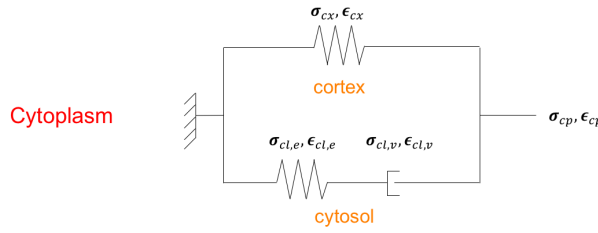


Figure 9: Rheological model of the cytoplasm [33]

The cytosol itself is decomposed into a viscous part and a purely elastic part as follows

$$\begin{aligned} \sigma_{cl} &= \sigma_{cl,e} = \sigma_{cl,v} \\ \epsilon_{cl} &= \epsilon_{cl,v} + \epsilon_{cl,e} \end{aligned} \quad (11)$$

where the subscripts v and e respectively stand for viscous and elastic.

Additionally, we have

$$\begin{aligned} \sigma_{cx} &= \lambda_{cx} \text{Tr}(\epsilon_{cp}) \mathbf{I} + 2\mu_{cx} \epsilon_{cp} \\ \sigma_{cl} &= \lambda_{cl} \text{Tr}(\epsilon_{cp} - \epsilon_{cl,v}) \mathbf{I} + 2\mu_{np}(\epsilon_{cp} - \epsilon_{cl,v}) \\ \dot{\epsilon}_{cl,v}^D &= \frac{\sigma_{cl}^D}{\eta_{cl}} \end{aligned} \quad (12)$$

where λ_{cx} , μ_{cx} , λ_{cl} and μ_{cl} are the Lamé coefficients associated to the cortex and the cytosol respectively, defined as $\lambda_k = \frac{E_k \nu_c}{(1+\nu_c)(1-2\nu_c)}$ and $\mu_k = \frac{E_k}{2(1+\nu_c)}$, $k = \{cx, cl\}$. E_{cx} , E_{cl} and ν_c are the Young moduli of the cortex and the cytosol respectively and the Poisson ratio of the cytoplasm. η_{cl} is the viscosity of the

cytosol defined as $\eta_{cl} = \tau_{cl} E_{cl}$, with τ_{cl} its characteristic time [35]. As before, we assume that $\epsilon_{cl,v}$ only has a deviatoric part.

As previously, and due to the very small thickness of the cortex (10 nm), the Young moduli were homogenized with regard to the surface occupied by the cortex and the cytosol as:

$$\begin{aligned} E_{cx} &= E_{cx,0} \frac{A_{cx,0}}{A_{cp,0}} \\ E_{cl} &= E_{cl,0} \frac{A_{cl,0}}{A_{cp,0}} \end{aligned} \quad (13)$$

where $A_{cx,0}$, $A_{cp,0}$, and $A_{cl,0}$ are the initial areas taken respectively by the cortex, the cytoplasm and the cytosol.

The nominal values of the mechanical parameters $E_{cx,0}$, $E_{cl,0}$, τ_{cl} and the Poisson ratio ν_{cp} were taken from the literature and previous work as listed in Table 1.

Similarly to the nucleus computation, an ALE formulation with updated frame and moving mesh is adopted. The contact force is also considered as a body force only applied in the possible overlapped region between the cell and the channel whenever it takes place and measured by the intersection level set of the channel and the cell and the global equilibrium of the system is expressed as

$$\rho \mathbf{a} = \text{Div}(\boldsymbol{\sigma}) + \mathbf{f}_{fluid} + \mathbf{f}_{ch} \quad (14)$$

where ρ is the cell density defined as $\rho(\mathbf{x}) = \rho_n g_n(\mathbf{x}) + \rho_{cp} g_{cp}(\mathbf{x})$, \mathbf{a} is the acceleration and $\boldsymbol{\sigma}$ is the Cauchy stress that reads $\boldsymbol{\sigma} = g_n \boldsymbol{\sigma}_n + g_{cp} \boldsymbol{\sigma}_{cp}$. \mathbf{f}_{fluid} and \mathbf{f}_{ch} respectively indicate the fluid force exerted on the cell cytoplasm and the contact force between the cell and the channel.

3.4. Results

In this study, we tackle two channel widths: $W_{ch} = 5\mu m$ (slightly sub-nuclear) and $W_{ch} = 1\mu m$ (highly sub-nuclear). The large channel corresponds to a common size for perfusion experiment [11], while the 1-micron channel was

chosen to study the behaviour of the cell under higher constriction at a size close to the 10% limit impeding cell migration completely [7]. The fluid force applied to the cell is designed so that the passing time through the constriction is of the order of 1 s, seconds, as can be found experimentally [45, 19, 20]. In each case, we are first interested in the displacement of the cell inertia center \mathbf{d}_c that decomposes as $d_{c,x}$ and $d_{c,y}$ on both directions with $d_{c,y}$ equal to 0. Then we study the evolution of the norm of the nucleus deviatoric strain ϵ_n^D , the value and distribution of the Von Mises stress, the ratio of nucleus area over cell area $\frac{A_{nuc}}{A_{cell}}$ and the vertical and horizontal component of the positive part of the resultant of the fluid force acting on the cell ($|F_{fluid,x}|$ and $|F_{fluid,y}|$ respectively), all with respect to $d_{c,x}$. To better understand the structural role of the lamina and the chromatin, we implement one model for a "wild-type" cell and another for a "lamin-deficient" one, in which $E_{l,0}$ is set to 3000 and 30 Pa, respectively.

3.4.1. 5 microns-wide constriction

We first look at the "wild-type" model in the larger channel. The cell rapidly plugs the channel ($d_{c,x} = 4\mu m$) and then requires more time until the nucleus itself clogs the channel ($d_{c,x} = 12\mu m$). Once this is accomplished, the cell goes through very rapidly (it takes here 1-2 ms, which is consistent with the results presented in [11]) and can then go back to a relaxed state (Figure 10 and Movie 1 in Supplementary Material).

The horizontal displacement of the cell center of inertia $d_{c,x}$ illustrates this phenomenon clearly (Figure 11.a). The deviatoric strain of the nucleus reaches a maximum of 35% at the center of the constriction and no plastic deformation is found in the nucleus (Figure 11.b). The Von Mises stress reaches a maximum of 5.39 Pa in the constriction, which is barely above the plasticity threshold, thus explaining the absence of plasticity (Movie 1 in Supplementary Material). Initially, the cell and the nucleus area are equal to $176 \mu m^2$ and $50 \mu m^2$ respectively. Thus, the ratio of the nucleus over the cell area thus is 28.5 % before entering the constriction. As expected, this ratio grows larger as the cell enters

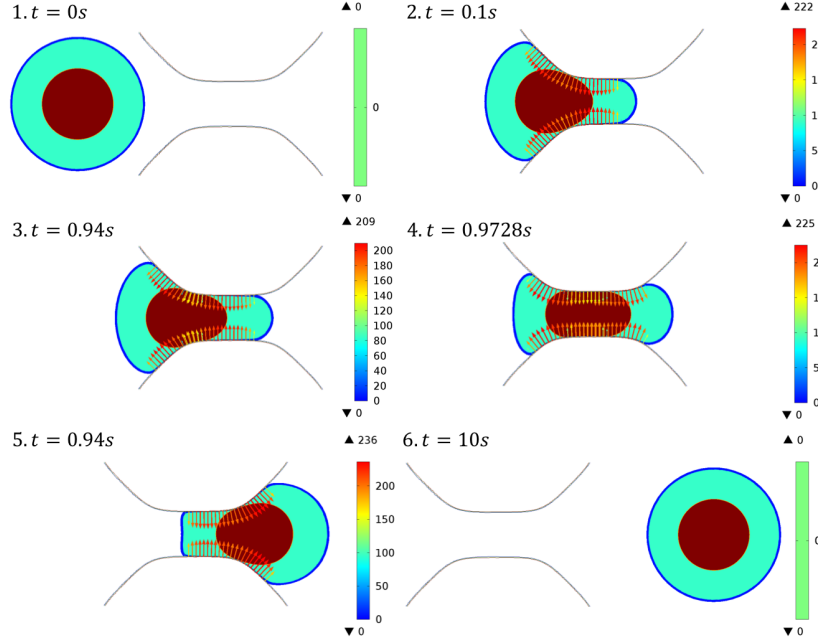


Figure 10: Simulation of the perfusion test in the 5 μm -wide micro-channel for a wild-type cell. Arrows represent the norm of the contact force (refer to the color scale for the value)

the constriction to reach a peak of 40% for $d_{c,x} = 16\mu\text{m}$ and it goes back to its initial value once the cell is completely out of the constriction after a displacement of 30 μm (Figure 11.c)). The resulting force applied on the fluid reaches 80 and 17 pN in the vertical and horizontal direction respectively (Figure 11.d). If the absolute value of the force is not specifically relevant by itself, it will be interesting to compare these value with the other test cases.

In the lamin-deficient model, the displacement of the cell inertia follows a similar curve, but the other parameters present significant differences (Figure 11). ϵ_{nuc}^D goes up to 62%, which means a 77% higher nucleus compression than for the wild-type model. The Von Mises stress reaches a maximum of 7.3 Pa in the constriction, which is slightly higher than before and should yield a higher plasticity not observed here (Movie 2 in Supplementary Material). This seemingly contradictory result will be discussed later in this section. The area

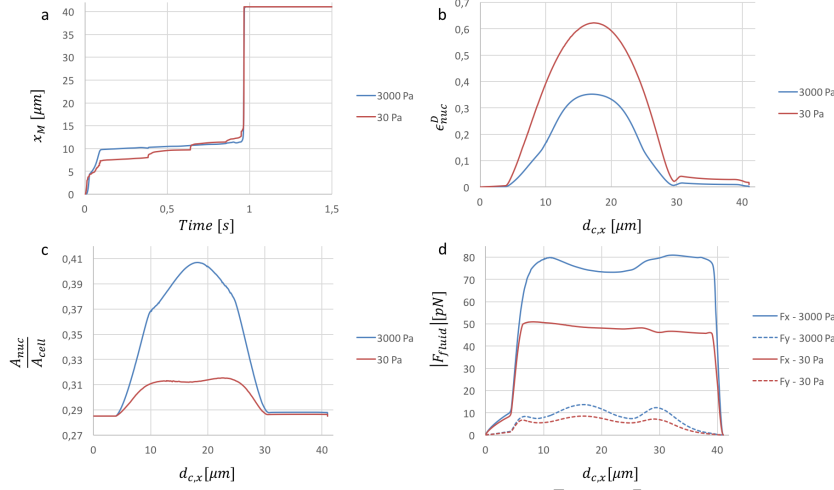


Figure 11: Comparison of the perfusion test results in the 5 μm -wide micro-channel for the wild-type (blue: $E_{l,0} = 3000 \text{ Pa}$) and the lamin-deficient (red: $E_{l,0} = 30 \text{ Pa}$) model. a) Horizontal displacement of the cell inertia center $d_{c,x}$, b) Nucleus deviatoric strain, c) Nucleus area relative to whole cell area and d) Resulting fluid force (vertical and horizontal components)

ratio curve shows two local maxima of 31 and 31.5 % for $d_{c,x} = 12$ and $24 \mu\text{m}$. Both values are around 20% lower than previously, which reinforces the earlier finding that the nucleus undergoes more deformation when the lamina does not "shield" it from stress. As it could be expected, the force required to get the nucleus through the constriction is lower now that the lamina is depleted as it reaches 51 and 9 pN in the vertical and horizontal direction respectively.

3.4.2. 1 micron-wide constriction

With the strongly sub-nuclear channel and the wild-type model, we could have expected a similar behaviour as previously, only slower. However, the cell undergoes a different path to go through the smaller constriction. Indeed, as previously, the cell rapidly clogs the constriction ($d_{c,x} = 4 \mu\text{m}$), but then, the nucleus rapidly clogs the constriction ($d_{c,x} = 12 \mu\text{m}$). Eventually, the nucleus gets gradually squeezed inside the constriction until the cell center of inertia displacement reaches $14 \mu\text{m}$, where the cell finally goes through at once (Figures

12 and 13a. and Movie 3 in Supplementary Material).

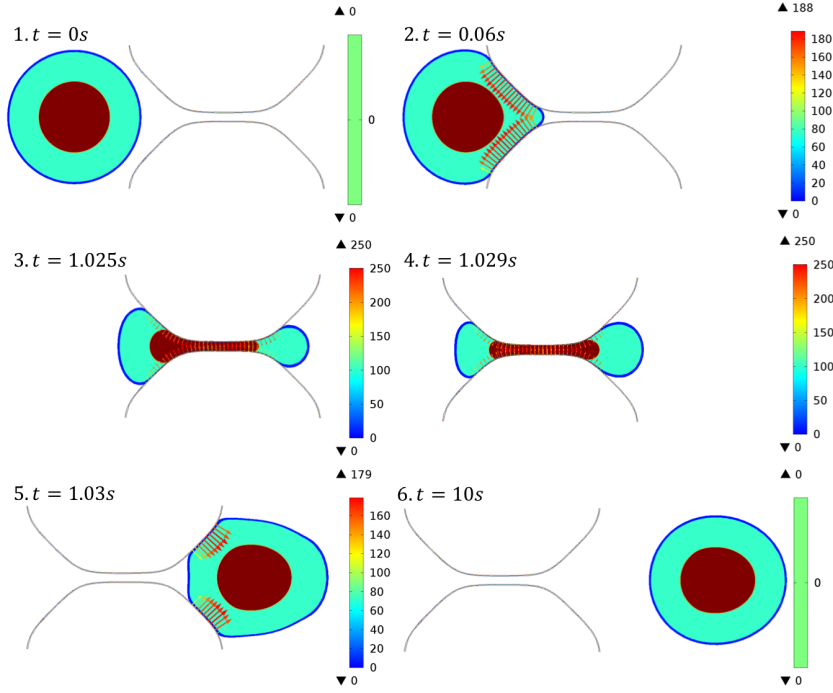


Figure 12: Simulation of the perfusion test in the 1 μm -wide micro-channel for a wild-type cell. Arrows represent the norm of the contact force (refer to the color scale for the value)

In this test case, the average nucleus deviatoric strain reaches 88% at $d_{c,x} = 18\mu\text{m}$ (Figure 13b.), which corresponds to the time when the rear of the nucleus is fully inside the constriction. Here, the nucleus presents a plastic strain of 8 % after reaching the other side of the constriction. The Von Mises stress reaches a maximum of 13.7 Pa at the center of the constrained the nucleus as it first clogs the channel and then displays a relaxation (Movie 3 in Supplementary Material). Looking at the ratio of the nucleus area over the cell area, the behaviour is significantly different from the 5 μm -wide channel since two maxima are clearly visible (Figure 13c.). The ratio reaches a maximum of 38 % at $d_{c,x} = 6.5\mu\text{m}$, corresponding to the nucleus starting to plug the channel, the front of the cytosol thus being already squeezed inside the constriction. The ratio then decreases as

the nucleus makes its way through the constriction and there is a second peak at 40% for $d_{c,x} = 28\mu m$ as the rear of the nucleus exits the constriction, before it settles down to 28.4% at the end of the perfusion test (Figure 13c.). The force needed for the cell to go through the constriction reaches 120 and 22 pN in the x and y direction respectively (Figure 13d.), which is 50% higher, for the vertical component, as in the 5- μm channel.

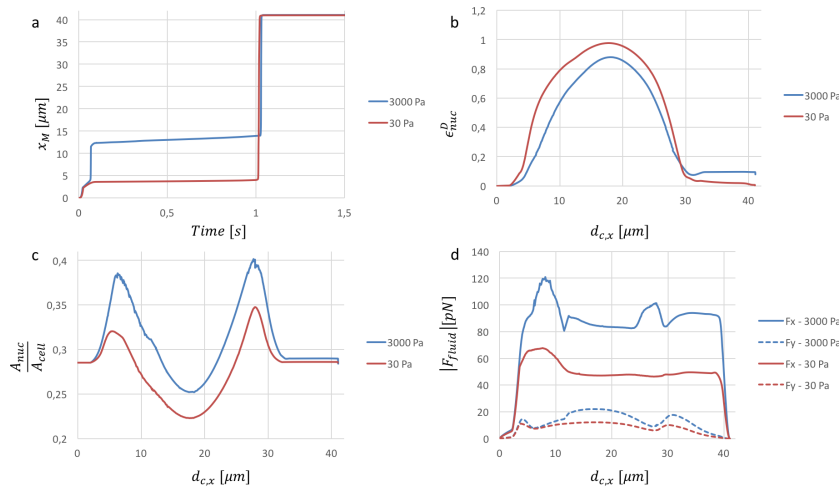


Figure 13: Comparison of the perfusion test results in the 1 μm -wide micro-channel for wild type and lamin deficient cells. a) Horizontal displacement of the cell inertia center $d_{c,x}$, b) Nucleus deviatoric strain, c) Nucleus area relative to whole cell area and d) Resulting fluid force (vertical and horizontal components)

As for the lamin-deficient model, starting in the cell inertia center displacement, differences can be spotted from the wild-type case (Figure 13). The cell first plugs the channel ($d_{c,x} = 3.5\mu m$) and, as soon as the cytoplasm tip starts getting into the constriction ($d_{c,x} = 4\mu m$), the whole cell goes through. Thus, the nucleus itself does not seem to be a barrier to transmigration anymore. The deviatoric strain reaches 98%, which is 10% higher than for the wild-type model. However, there is no plastic strain anymore compared to the wild-type, which is in contradiction with experimental results from Cao et al. [37]. This result

will be further discussed in the following section. The Von Mises stress reaches a maximum of 16.2 Pa at the center of the constrained part of the nucleus as it starts clogging the constriction and then displays a relaxation (Movie 4 in Supplementary Material). Again, this value, superior than in the wild-type cell should result in an increase of the plasticity. The area ratio between the nucleus and the cell follows the same pattern than previously although with lower values: it reaches a first maximum of 32% for $d_{c,x} = 5.5\mu m$ and second one of 35% for $d_{c,x} = 28\mu m$. Together with the values of deviatoric strain, this shows, as in the larger channel, that the nucleus gets more squeezed if no lamina protects it. The force of the fluid on the cell reaches 67 and 12 pN for its vertical and horizontal component respectively, which is almost half the force needed for the wild-type cell, once again comforting us in the idea that the nucleus is a major barrier during transmigration through small constrictions, and the lamina seems to be a crucial structural constituent of the nucleus allowing it to resist to large deformation. The decrease of plastic strain in the lamin-deficient model however raises the question of the role of the nucleoplasm as a load-bearing element of the nucleus.

3.5. The cytoplasm can "pull" on a too soft nucleus

The complete cell model with a visco-elasto-plastic nucleus was tested in 5 μm -wide and a 1 μm -wide perfusion channel both as a wild-type cell ($E_{l,0} = 3000$ Pa) and a lamin-deficient one in which the lamina was considered a hundred times weaker ($E_{l,0} = 30$ Pa). In both cases, the cell first plugs the device and then goes through very rapidly once it has progressed far enough in the constriction with a passage time of the order of 1 ms. In the larger channel, once the nucleus is in contact with both the upper and lower channels, it is sufficiently compressed for the cell to go through directly. However, in the smaller constriction, an additional step is necessary for the nucleus to fully plug the channel and only once the cell center of inertia has progressed far enough in the constriction, the whole cell can pass through.

The study of the lamin-deficient model gives very interesting results and

insights of the importance of the interaction between the cytoplasm and the nucleus. First, the lamin-deficient nucleus undergoes significantly higher deformation than the wild-type one, hence showing the importance of the lamina as a load-bearing component of the nucleus. This seems to back up studies stating that the lamina is the major structural constituent in the nucleus. However, a closer look on the behaviour of the nucleus with regards to plasticity shows that the wild-type model displays a 10% plastic strain after exiting the smaller constriction, while the lamin-deficient one does not display any irreversible deformation (a similar but less intense phenomenon is observed in the larger constriction regarding the Von Mises stress). This property of our model appears in contrast with the latest experimental results as discussed by Cao et al. [37]. Further analysis of our results revealed that the cell cytoplasm seems responsible for this loss of plasticity as it forces the lamin-deficient soft nucleus to return to its original shape. This is an exciting result that meets the recent thesis defended by Stephens et al. [30] that not only does the lamina play a role but chromatin also plays a major part as a load-bearing element in the nucleus.

4. Conclusion and Discussion

All in all, our model of an isolated nucleus does display a plastic behaviour under extensive compression, as shown experimentally [28], and shows that both the lamina and the nucleoplasm have an impact on the mechanical behaviour of the overall nucleus. A softer lamina and a stiffer nucleoplasm yield a higher irreversible deformation after unloading the nucleus. Implementing this model in a whole cell model under a perfusion test allowed us to unveil the mechanical influence of the cytoplasm on the nucleus. In the $1\ \mu\text{m}$ constriction, the plastic deformation significantly decreased with the depletion of the lamina, which we explain by the cytoplasm "pulling" on the now softer nucleus to get it back to its original shape. This is further confirmed by the fact that in the isolated nucleus model, a decreased lamina Young modulus resulted in an increased plasticity. Thus, in order to observe plastic deformation, either the nucleoplasm should

be stiffer, or the cytoplasm softer. This mechanical interplay between nuclear stiffness and its plasticity reveals the ability of numerical simulation to tackle biological issues and to shed light on different mechanisms.

Our model however presents some limitations and some improvements can be included. First of all, a 2D FE model was used, whereas a three-dimensional (3D) one would be needed in order to faithfully reproduce 3D experiments. This point is particularly sensitive looking at the compression test: a more valid reduction would be an axi-symmetric model. We still chose to develop a purely 2D model since it fits the perfusion test: the compression test model is a step towards a more complete cell model and a 2D reduction will not give faithful quantitative results, but the qualitative trends stay valid.

Moreover, we chose to cope with the thinness of the cortex and the lamina through weighted Young's moduli with respect to the relative area of each cell component. Nevertheless, a shell model would surely fit better and could provide new insights on the relation between the various cell components [47].

Lately, some very interesting models of the cell nucleus have been developed [32, 37] in which the interaction between the nucleus and the cytoplasm is represented by an active force pulling on the nucleus. Our model gives a new insight on the passive interaction between these 2 cellular compartments: the comparison between the isolated nucleus model and the complete cell model proves the major mechanical influence the cytoplasm can have on the nucleus in a passive state. This interaction would be increasingly important during an active phenomenon like transmigration.

5. Acknowledgement

This work was funded by a ministerial doctoral fellowship of the French Government.

Parameter	Description	Value (unit)	References
r_{cell}	Cell radius	$7.5 \mu m$	
r_{cortex}	Cortex radius	$7.5 \mu m$	
$r_{cytosol}$	Cytosol radius	$7.25 \mu m$	
r_{lamina}	Lamina radius	$4 \mu m$	
$r_{nucleoplasm}$	Nucleoplasm radius	$3.95 \mu m$	
t_{cortex}	Cortex thickness	250 nm	[33]
t_{lamina}	Lamina thickness	50 nm	[48, 49]
$E_{cx,0}$	Nominal cortex Young modulus	100 Pa	[33, 44]
$E_{cl,0}$	Nominal cytosol Young modulus	10 Pa	[50, 51, 33, 44]
$E_{l,0}$	Nominal lamina Young modulus	3000 Pa	[33, 44]
$E_{np,0}$	Nominal nucleoplasm Young modulus	25 Pa	[35, 36, 33, 44]
E_{cx}	Equivalent cortex Young modulus	9.08 Pa	
E_{cl}	Equivalent cytosol Young modulus	9.16 Pa	
E_l	Equivalent lamina Young modulus	74.53 Pa	
E_{np}	Equivalent nucleoplasm Young modulus	24.38 Pa	
ν_c	Cytoplasm Poisson ratio	0.3	[33]
ν_n	Nucleus Poisson ratio	0.4	[33]
τ_{cl}	Cytosol characteristic time	1 s	[52, 12]
τ_{np}	Nucleoplasm characteristic time	2 s	[18, 53, 36, 12]
s	Nucleoplasm plasticity threshold	4 Pa	
ρ_{cp}	Cytoplasm density	$1000 kg/m^3$	[54]
ρ_n	Nucleus density	$1400 kg/m^3$	[54]
$A_{cp,0}$	Initial cytoplasm area	$126 \mu m^2$	
$A_{cx,0}$	Initial cortex area	$11 \mu m^2$	
$A_{cl,0}$	Initial cytosol area	$115 \mu m^2$	

$A_{n,0}$	Initial nucleus area	$50 \mu m^2$
$A_{np,0}$	Initial cytoplasm area	$49 \mu m^2$
$A_{l,0}$	Initial cortex area	$1 \mu m^2$
T_1	First characteristic time for $y_p(t)$	10 s
T_2	Second characteristic time for $y_p(t)$	27.5 s
T_3	Third characteristic time for $y_p(t)$	2 s
v_0	Velocity of the compression plate	$0.25 \mu m/s$ [14]
$x_{p,0}$	x-coordinate of the plates	0
$y_{up,0}$	y-coordinate of the upper plate	$6 \mu m$
$y_{lp,0}$	y-coordinate of the lower plate	$6 \mu m$
l_{plate}	length of the plates	$30 \mu m$
h_{plate}	vertical width of the plates	$1 \mu m$
x_0	Geometric parameter of the perfusion constriction	$15 \mu m$
a_1	Geometric parameter of the perfusion constriction	$11 \mu m$
a_2	Geometric parameter of the perfusion constriction	$16 \mu m$
d_0	width of the constriction	$W_{ch,1} = 1 \mu m$ or $W_{ch,5} = 5 \mu m$
P_{in}	Inlet pressure in perfusion device	1 Pa
P_{out}	Outlet in perfusion device	0
ρ_f	Fluid volumetric mass in perfusion device	$1000 kg/m^3$
η_f	Fluid viscosity in perfusion device	$10^{-2} Pa.s$

Appendices

A. Characteristic functions

Throughout the paper, each cellular component is geometrically defined through a characteristic function g , which is a composition of a Heaviside function h and a spatial level set l as follows

$$g_i(\mathbf{x}) = h \circ l_i(\mathbf{x}) = \begin{cases} 1 & \text{for } l_i(\mathbf{x}) > 0 \\ 0 & \text{for } l_i(\mathbf{x}) < 0 \end{cases} \quad (\text{A.1})$$

where $i = np, l, cl, cx$. The nucleoplasm and the lamina constitute the nucleus ($g_n = g_{np} + g_l$), while the cytosol and the cortex constitute the cytoplasm ($g_c = g_{cl} + g_{cx}$).

B. Nucleus geometry

$$g_l(\mathbf{x}) = \begin{cases} 1 & \text{if } r_{nucleoplasm}^2 < \|\mathbf{x} - \mathbf{c}_{nucleus}\| < r_{lamina}^2 \\ 0 & \text{otherwise} \end{cases} \quad (\text{B.1})$$

$$g_{np}(\mathbf{x}) = \begin{cases} 1 & \text{if } \|\mathbf{x} - \mathbf{c}_{nucleus}\| < r_{nucleoplasm}^2 \\ 0 & \text{otherwise} \end{cases} \quad (\text{B.2})$$

where $\mathbf{c}_{nucleus}$ is the position of the center of the nucleus, $r_{nucleoplasm}$ and r_{lamina} are the nucleoplasm and lamina radii, respectively and \mathbf{x} is the actual spatial position.

C. Compression experiment

The compression plate are defined by the following Heaviside functions

$$g_{up}(\mathbf{x} - \mathbf{u}) = \begin{cases} 1 & \text{if } l_{up} > 0 \\ 0 & \text{otherwise} \end{cases} \quad (\text{C.1})$$

$$g_{lp}(\mathbf{x} - \mathbf{u}) = \begin{cases} 1 & \text{if } l_{lp} > 0 \\ 0 & \text{otherwise} \end{cases} \quad (\text{C.2})$$

such that $g_{plate} = g_{up} + g_{lp}$, with l_{up} and l_{lp} two level set functions expressed as

$$\begin{aligned} l_{up} &= -\left(\frac{x - x_{p,0}}{l_{plate}}\right)^8 - \left(\frac{y - y_{up,0} - y_p(t)}{g_{plate}}\right)^4 - 1 \\ l_{lp} &= -\left(\frac{x - x_{p,0}}{l_{plate}}\right)^8 - \left(\frac{y + y_{lp,0}}{g_{plate}}\right)^4 - 1 \end{aligned} \quad (\text{C.3})$$

where $(x_{p,0}, y_{lp,0})$ and $(x_{p,0}, y_{up,0})$ are coordinates of the two plates, respectively and $y_p(t)$ is the vertical displacement applied on l_{up} , the lower plate being fixed, and defined as:

$$y_p(t) = \begin{cases} -v_0(t - T_1) & \text{if } T_1 \leq t \leq T_1 + T_2 \\ -v_0 T_2 & \text{if } T_1 + T_2 \leq t \leq T_1 + T_2 + T_3 \\ -v_0(-t + T_1 + 2T_2 + T_3) & \text{if } T_1 + T_2 + T_3 \leq t \leq T_1 + 2T_2 + T_3 \end{cases} \quad (\text{C.4})$$

with v_0 the compression velocity and T_1 , T_2 and T_3 the time parameters defining the loading/unloading of the plate (see Figure C.14).

D. Cell geometry

$$g_{cx}(\mathbf{x}) = \begin{cases} 1 & \text{if } r_{cytosol}^2 < \|\mathbf{x} - \mathbf{c}_{nucleus}\| < r_{cortex}^2 \\ 0 & \text{otherwise} \end{cases} \quad (\text{D.1})$$

$$g_{cl}(\mathbf{x}) = \begin{cases} 1 & \text{if } \|\mathbf{x} - \mathbf{c}_{nucleus}\| < r_{cytosol}^2 \\ 0 & \text{otherwise} \end{cases} \quad (\text{D.2})$$

with $r_{cytosol}^2$ and r_{cortex}^2 being the cytosol and cortex radii, respectively.

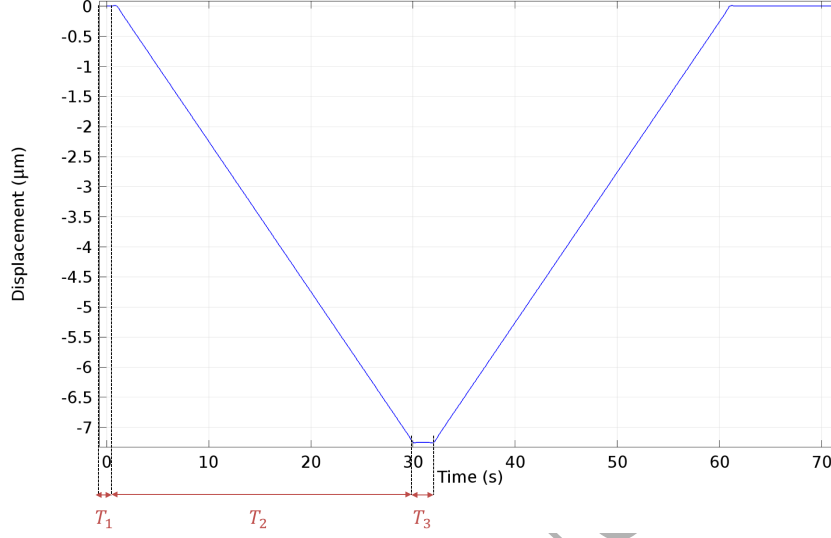


Figure C.14: Motion of the mobile upper plate

E. Perfusion experiment

E.1. Geometry

$$g_{uc}(\mathbf{x} - \mathbf{u}) = \begin{cases} 1 & \text{if } l_{uc} < 0 \\ 0 & \text{otherwise} \end{cases} \quad (\text{E.1})$$

$$g_{lc}(\mathbf{x} - \mathbf{u}) = \begin{cases} 1 & \text{if } l_{lc} < 0 \\ 0 & \text{otherwise} \end{cases} \quad (\text{E.2})$$

such the $g_{ch} = g_{uc} + g_{lc}$, where l_{uc} and l_{lc} are two level set functions defined as

$$\begin{aligned} l_{uc} &= \left(\frac{x - x_0}{a_1}\right)^{16} + \left(\frac{y - y_0}{a_1}\right)^{16} + \left(\frac{x - x_0 - y + y_0}{a_2}\right)^{16} + \left(\frac{x - x_0 + y - y_0}{a_2}\right)^{17} \\ l_{lc} &= \left(\frac{x - x_0}{a_1}\right)^{16} + \left(\frac{y + y_0}{a_1}\right)^{16} + \left(\frac{x - x_0 - y - y_0}{a_2}\right)^{16} + \left(\frac{x - x_0 + y + y_0}{a_2}\right)^{17} \end{aligned} \quad (\text{E.3})$$

where x_0 , a_1 and a_2 are geometric parameters and $y_0 = a_1 + d_0/2$, d_0 being the width of the constriction.

E.2. Parametric study

The perfusion model involves 41 parameters (see Table 1), but not all are of the same importance. We can classify them in three categories:

1. The parameters that purely define the system geometry and its dynamics (r_{cell} , r_{cortex} , $r_{cytosol}$, r_{lamina} , $r_{nucleoplasm}$, t_{cortex} , t_{lamina} , $A_{cp,0}$, $A_{cl,0}$, $A_{cx,0}$, $A_{n,0}$, $A_{np,0}$, $A_{l,0}$, T_1 , T_2 , T_3 , v_0 , $x_{p,0}$, $y_{up,0}$, $y_{lp,0}$, l_{plate} , h_{plate} , x_0 , a_1 , a_2 , d_0 , P_{in} , P_{out} , ρ_f , η_f)
2. The parameters that have been well defined in the literature (ν_c , ν_n , ρ_{cp} , ρ_n)
3. The parameters that present a high variability ($E_{cx,0}$, $E_{cl,0}$, $E_{l,0}$, $E_{np,0}$, τ_{cl} , τ_{np} , s)

We assume that the parameters of the 1st and 2nd categories do not influence the overall results of our work. Among the last category, we chose to focus on those that may play a critical role during the perfusion assay. Considering our interest in the interaction between cytosol and lamina, we chose to perform a parametric study on $E_{cl,0}$, $E_{l,0}$ and τ_{cl} . The plasticity threshold was not included since it was calibrated on the compression test and the value was then kept constant. The sensibility study was conducted on both the wild-type cell model and the lamin-deficient one. For each parameter, 3 simulations were performed using the initial value of the parameter and $\pm 10\%$. We analyzed the nuclear deviatoric strain ϵ_n^D as function of the horizontal displacement of the cell center of inertia $d_{c,x}$ and that said displacement as a function of time in order to assess the influence of cell mechanical parameters on the passage time.

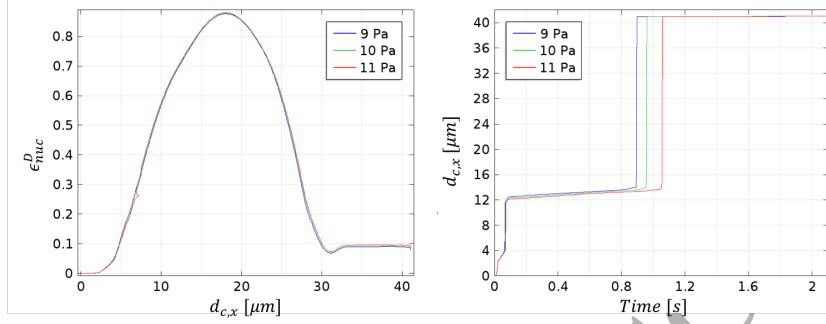


Figure E.15: Parametric study on $E_{cl,0}$. Left: Evolution of the nuclear deviatoric strain ϵ_n^D as function of the horizontal displacement of the cell center of inertia $d_{c,x}$ and Right: Horizontal displacement of the cell center of inertia as a function of time during the $1 \mu\text{m}$ -wide micro-channel perfusion simulation for the wild-type model.

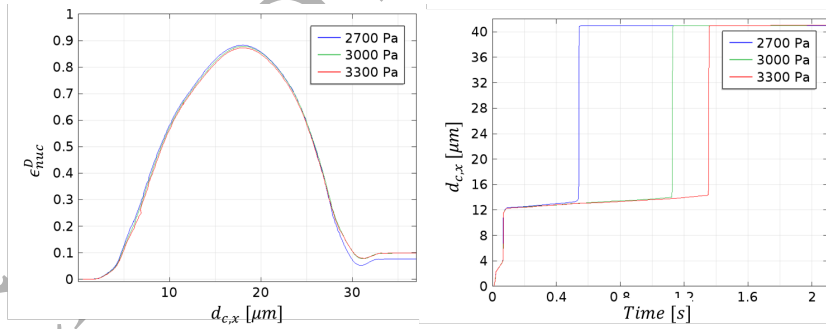


Figure E.16: Parametric study on $E_{l,0}$. Left: Evolution of the nuclear deviatoric strain ϵ_n^D as function of the horizontal displacement of the cell center of inertia $d_{c,x}$ and Right: Horizontal displacement of the cell center of inertia as a function of time during the $1 \mu\text{m}$ -wide micro-channel perfusion simulation for the wild-type model.

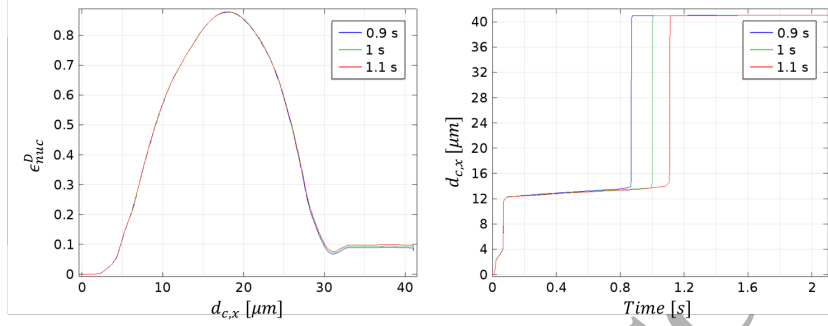


Figure E.17: Parametric study on τ_{cl} . Left: Evolution of the nuclear deviatoric strain ϵ_n^D as function of the horizontal displacement of the cell center of inertia $d_{c,x}$ and Right: Horizontal displacement of the cell center of inertia as a function of time during the 1 μm -wide micro-channel perfusion simulation for the wild-type model.

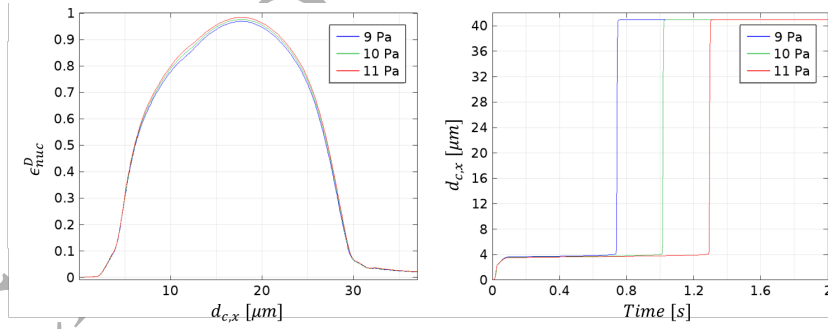


Figure E.18: Parametric study on $E_{cl,0}$. Left: Evolution of the nuclear deviatoric strain ϵ_n^D as function of the horizontal displacement of the cell center of inertia $d_{c,x}$ and Right: Horizontal displacement of the cell center of inertia as a function of time during the 1 μm -wide micro-channel perfusion simulation for the lamin-deficient model.

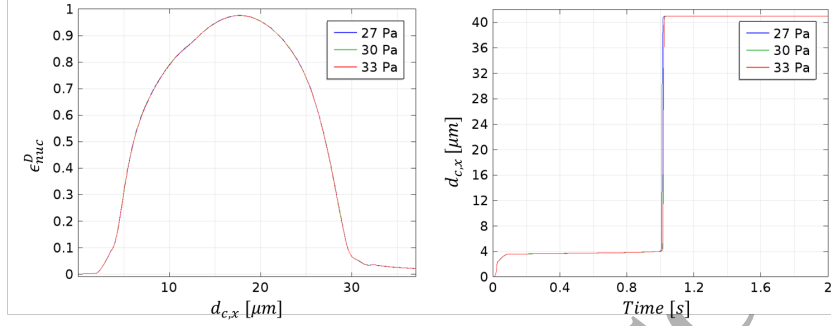


Figure E.19: Parametric study on $E_{l,0}$. Left: Evolution of the nuclear deviatoric strain ϵ_n^D as function of the horizontal displacement of the cell center of inertia $d_{c,x}$ and Right: Horizontal displacement of the cell center of inertia as a function of time during the 1 μm -wide micro-channel perfusion simulation for the lamin-deficient model.

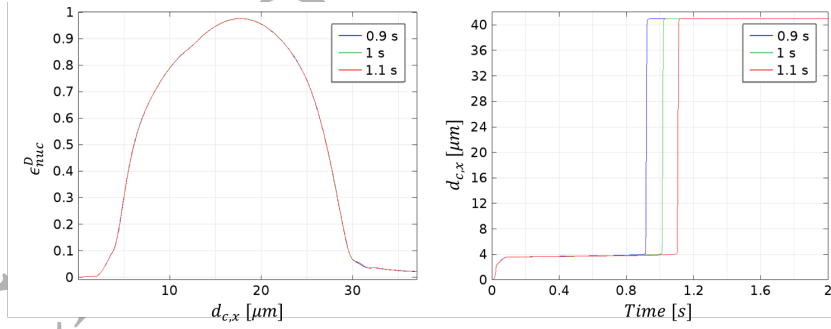


Figure E.20: Parametric study on τ_{cl} . Left: Evolution of the nuclear deviatoric strain ϵ_n^D as function of the horizontal displacement of the cell center of inertia $d_{c,x}$ and Right: Horizontal displacement of the cell center of inertia as a function of time during the 1 μm -wide micro-channel perfusion simulation for the lamin-deficient model.

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