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Pervaporation-induced drying in networks of channels of variable width

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Abstract

We study the influence of variations of cross-section on the drying dynamics by pervaporation in microfluidic channels surrounded by water-permeable polydimethylsiloxane (PDMS) walls. Single channels of linearly increasing and decreasing width display similar drying time, but different dynamics towards final drying. Single channels with stepwise increasing or decreasing width show jumps of the drying velocity when the meniscus separating the water-filled and the dry part of the channel crosses the steps. In branched networks, narrower branches tend to dry faster than wider ones. All these observations are rationalised in the framework of a model based on the solution of the diffusion equation of water through PDMS, accounting precisely for the local channel width at the meniscus location. It captures quantitatively our measurements, except the magnitude of the velocity jumps in channels with steps. Implications and perspectives of this study are discussed in the context of leaf drying by the so-called air seeding mechanism in trees.

1 Introduction

Global warming is an increasing thread for the survival of forests (Choat et al. 2018; Brodribb et al. 2020). Evapotranspiration at the surface of the leaves, the mechanism whereby sap circulates in the xylem from the roots to the leaves, puts the sap in a state of negative pressure (Stroock et al. 2014; Jensen et al. 2016), all the more that the ambient atmospheric conditions are dry and warm. Cavitation bubbles may then appear in the sap network, and subsequently invade it, which can stop the sap circulation. It is therefore of primary importance to understand this invasion mechanism in leaves, known as air seeding among the plant physiologists.

Like the vascular network of animals, leaf venation forms complex networks with different levels of hierarchy, reconnections, and different sections. The channel section tends to decrease in size after each bifurcation, when a channel divides into several channels: the section becomes smaller

Benjamin Dollet benjamin.dollet@univ-grenoble-alpes.fr and smaller until the terminal veins. It was shown the section follows Murray's law as in animal networks (McCulloh et al. 2003): the sum of conduit diameters raised to power 3 is conserved.

During evapotranspiration, water is transported from the veins to the stomata at the surface of the leaves, by diffusion through the mesophyll tissue which surround xylem sap in leaves. This transport is also called pervaporation in other contexts, such as chemical engineering; see e.g. Eijkel et al. (2005); Ziane et al. (2015); Ziemecka et al. (2015), and Bacchin et al. for a full review. To make a biomimetic system mimicking water transport through the mesophyll tissue, we use channels made of polydimethylsiloxane (PDMS), following the method by Noblin et al. (2008). This material is permeable to water vapour, like the mesophyll tissues. Among the different microfluidic or nanofluidic systems which have been used for synthetic trees or leaves, soft PDMS was preferred over the stiff hydrogels (Wheeler and Stroock 2008; Vincent et al. 2012) or nanoporous silicon (Vincent et al. 2014), which can withstand negative pressure. Here we do not focus on negative pressure and cavitation, but on latter stages when air invades the network during drying, at pressures close to atmospheric pressure. PDMS has then the advantage that it reproduces better the thinness and flexibility of real leaves.

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From a physical standpoint, to understand pervaporationinduced air invasion dynamics in biomimetic networks, we study the simplest systems, increasing the complexity by steps. We started by single channels (Dollet et al. 2019), then elementary networks Dollet et al. (2021), but still at constant section. In the current study, we release this restriction, and focus on the effect of the variation of section in single channels and simple branched networks.

2 Materials and methods

All our channels are molded in a thin PDMS layer glued on a glass plate, as follows. The glass slide was exposed for 30 s to a plasma, then the PDMS imprint was manually deposited on the plasma-activated glass surface, taking care to avoid folds and blisters of the flexible PDMS sheet. In all this study, the height h of the channel and the thickness Hof the PDMS layer (Fig. 1b) are kept as much constant as possible, within the uncertainties of the fabrication process: $h = 42 \pm 3 \ \mu \text{m}$ and $H = 88 \pm 2 \ \mu \text{m}$ (see later Table 1 for the values of h and H measured for separate designs). Hence, to study variations of the cross-section, we focus on variations of the channel width w (Fig. 1a). We study first single channels of variable width, then simple networks of channels of different widths. However, for the sake of simplicity, in the latter case, each portion of network between consecutive nodes is of constant width.

The experiments are run as follows. The channels are manually cut with a scalpel at the entrance point (the cuts are clearly visible in Fig. 2). The glass slide on which the channels are bond is plunged into a beaker of deionised water under low pressure for one hour, until the channels are completely filled. They are then placed under a constant flux of dry air, and their drying from the entrance points onwards is recorded with a CCD camera, from which the location of the meniscus (or menisci) separating the water-filled part and air-filled part of the channels is measured by image analysis. First, we study two single channels of continuously variable width between the entrance point and a terminal wall (Fig. 2a). We focus on channels of total length $L_c = 12$ mm with width varying linearly along the channel between 50 μ m and 200 μ m, either decreasing or increasing from the entrance to the terminal wall. Second, we study two single channels of total length $L_c = 10$ mm with variations of width by steps (Fig. 2b). One channel present five portions (each of length 2 mm) with widths decreasing from 175 μ m to 75 μ m, and the other has four portions (each of length 2.5 mm) with widths increasing from 75 μ m to 150 μ m.

We study next simple channel networks, and we revisit the branched networks which we have studied in Dollet et al. (2021). We focus on two configurations. First, we study single-node networks, where an entrance channel splits into two, three, four or five branches (Fig. 2c). In this paper, each of these branches has a different width. Second, we consider a "tree" with consecutive subdivisions of the channels into branches of smaller widths (Fig. 2d). The choice of the geometry of the tree results from a compromise between the presence of certain ingredients of complexity (several levels of branch divisions, branches of different lengths and widths), with yet a simple enough design to allow for simple analysis.

3 Single channels of variable width

3.1 Theory

In this Section, we model the drying dynamics by pervaporation of a single channel of varying width w(x), where *x* is the streamwise coordinate along the centreline of the channel (Fig. 1a). The channel height is denoted *h* and the PDMS thickness *H* (Fig. 1b); $\delta = H - h$ is therefore the PDMS thickness between the channel top and the outer air. If *Q* is the volumetric flux of water leaving the water-filled part of the channel, the conservation of mass writes:





Fig. 1 a Top view sketch of a channel of varying width w(x), where *x* is the streamwise coordinate along its centreline. The channel is filled with water over a length *L* along its centreline. The origin x = 0 is located at the closed end of the channel, and the water occupies the

region $0 \le x \le L$; the meniscus separating the water-filled part of the channel and its air-filled part is thus located at x = L. **b** Transverse side view of the channel of width w(x) and height *h* between a glass slide and a PDMS sheet of thickness *H*

Fig. 2 Snapshots of the channel designs studied in this paper. a Single channels of linearly varying width. Notice their slight curvature, due to a slight distortion of the flexible PDMS layer when it is bond to glass; this has no noticeable influence on the dynamics. b Single channels with widths varying by steps, which are marked by circles outside the channels. c Singlenode networks with varying number of branches (the results with the network with four branches will not be presented here). All entrance channels have a width 100 μ m and a length $L_e = 3$ mm. All branches have a length $L_b = 4$ mm. **d** "Tree" design. Branches have a length of 2, 1 or 0.5 mm, and their width decrease at each bifurcation: 175 μ m for the entrance channel, then 150, 125, 100 and 75 µm



$$-Q = \frac{d}{dt} \left[h \int_0^{L(t)} w(x) dx \right] = h \dot{L} w(x = L), \qquad (1) \qquad Q_\ell = \int_0^L q_\ell(w(x)) dx, \qquad (2)$$

where the term in brackets is the volume of water inside the channel at time *t*, *L* being the length of the water-filled part (henceforth called water length) and $|\dot{L}|$ the velocity of the meniscus. It is worth noting that at a given flux *Q*, the larger the width w(x = L) at the location of the meniscus, the slower the meniscus.

There are two contributions to the flux $Q = Q_{\ell} + Q_g$: one by diffusion from the water-PDMS interface (Q_{ℓ}) , and an evaporative flux Q_g at the meniscus. We restrict our analysis to a slowly varying width: $|dw/dx| \ll 1$. This hypothesis is suitable for slender channels, such as those encountered in leaves. For a slowly varying width, the streamwise dependence of the concentration field of water into PDMS is much weaker than the spanwise dependence, and we may neglect the width variation to estimate the diffusive flux. Hence: where $q_{\ell}(w)$ is the diffusive flux per unit length of a channel of constant width *w*. Inserting in (1), we get:

$$h\dot{L}w(x=L) = -\int_{0}^{L} q_{\ell}(w(x))dx - Q_{g}.$$
 (3)

In Dollet et al. (2019), we have derived the following analytical prediction of the flux q_{ℓ} :

$$q_{\ell} = D_P \bar{c}_P^{\text{sat}} (1 - \text{RH}) \tilde{q}_{\ell}, \tag{4}$$

with $D_p = 10^{-9} \text{ m}^2/\text{s}$ the diffusivity of water in PDMS (Watson and Baron 1996), $\bar{c}_p^{\text{sat}} = MC_p^{\text{sat}}/\rho$ the mass fraction of water in PDMS at saturation (with M = 0.018 kg/mol the molar mass of water, $\rho = 10^3 \text{kg/m}^3$ its density, and $C_p^{\text{sat}} = 40 \text{ mol/m}^3$ the saturation concentration of water in PDMS (Randall and Doyle 2005), this gives $\bar{c}_p^{\text{sat}} = 7.2 \times 10^{-4}$), RH the relative humidity of the outer air

(henceforth set to zero, consistently with our experiments) and \tilde{q}_{ℓ} a dimensionless shape factor coming from the resolution of a diffusion problem:

$$\tilde{q}_{\ell} = \frac{w}{\delta} + \xi, \tag{5}$$

with ξ another dimensionless shape factor quantifying diffusion between the channel side walls and the outer air; under reasonable geometrical assumptions quantified in Dollet et al. (2019) and holding in our experiments, it equals:

$$\xi = \frac{2}{\pi} \left[\ln \frac{(H+\delta)h}{\delta^2} + \frac{H}{\delta} \ln \frac{H+\delta}{h} \right].$$

We also use the prediction of Q_g made in Dollet et al. (2019) with the local value of the thickness at the meniscus location, simply denoted w in the remainder of this Section. To be rigorous, the assumption $|dw/dx| \ll 1$ is not sufficient to make such an approximation, and should be replaced by the more stringent condition $|dw/dx| \ll \epsilon^{1/2}$ where $\epsilon = D_P/\alpha D_a$ is a small parameter (with $\alpha = 0.03$ the Henry constant quantifying the water affinity in PDMS (Harley et al. 2012) and $D_a = 2 \times 10^{-5}$ m²/s the diffusivity of water vapour in air, hence $\epsilon = 2 \times 10^{-3}$), because the value of Q_g depends on the concentration field of water at distances of order $\epsilon^{-1/2}w$ away from the meniscus (Dollet et al. 2019). Under this approximation, the following prediction holds:

$$Q_g = \sqrt{\alpha D_a D_P h w \tilde{q}_\ell}.$$
(6)

3.2 Experiments

3.2.1 Channels with linearly variable width

We now test our theory on the two channels with linearly variable width. Figure 3a shows the time evolution of the water length in these two channels. It is a decreasing function of time, consistently with the drying process. More interestingly, both curves are convex, but the convexity is larger for the channel with increasing width compared with the one with decreasing width. The meniscus in the first channel is faster at the beginning of the drying process, but slower at the end, and the total drying time is almost similar for both channels.

Qualitatively, this behaviour highlights the influence of the local value of the channel width at the meniscus location in Eq. (1): the narrower the channel, the faster the meniscus at given flux Q. Quantitatively, we can apply the model of Sect. 3.1 for a width profile of the form:

$$w(x) = w_0 + (w_1 - w_0) \frac{L_c - L - x}{L_c},$$
(7)

with L_c the channel length, w_0 the width at the channel entrance and w_1 the width at its closed end. Inserting (7) in (2) and using (4) and (5), we get:

$$Q_{\ell} = D_P \bar{c}_P^{\text{sat}} \left[\left(\frac{w_1}{\delta} + \xi \right) L - \frac{w_1 - w_0}{2\delta L_c} L^2 \right]$$

Using the expression (6) of Q_g , we get after some algebra:





Fig.3 Drying dynamics in channels of width varying linearly between 50 and 200 μ m, for a decrease (\circ) and an increase (\Box) of the width over a length of 12 mm. **a** Time evolution of the water length, and **b** velocity of the meniscus as a function of the water length. In **b**,

the plain curves are fits by Eq. (8) with D_1 and D_2 as fitting parameters. In **a**, the plain curves result from numerical integration of the fitting curves of **b**. The dotted curves show the case of a channel of constant width 125 μ m

$$\begin{bmatrix} 1 - \left(1 - \frac{w_0}{w_1}\right)\bar{L}\end{bmatrix}\frac{d\bar{L}}{dt} \\ = -\frac{D_1}{hw_1}\left[\left(\frac{w_1}{\delta} + \xi\right)\bar{L} + \frac{w_1 - w_0}{2\delta}\bar{L}^2\right] - \frac{D_2}{L_c\sqrt{h\delta}} \\ \times \sqrt{\left(1 + \frac{w_0 - w_1}{w_1}\bar{L}\right)\left(1 + \frac{w_0 - w_1}{w_1}\bar{L} + \frac{\delta}{w_1}\xi\right)},$$
(8)

with the dimensionless quantity: $\overline{L} = L/L_c$, and two physicochemical parameters having the dimension of a diffusivity:

$$D_1 = D_P \bar{c}_P^{\text{sat}}, \qquad D_2 = \sqrt{\alpha D_a D_P}.$$
(9)

Eq. (8) provides an explicit relationship between the instantaneous velocity of the meniscus and its location. Therefore, to compare our data and our model, we plot the velocity of the meniscus as a function of the water length in Fig. 3b, and we fit the curves by (8) with D_1 and D_2 as fitting parameters. We anticipate that these two parameters will serve as fitting parameters for each set of experiments to be discussed throughout this study, and we refer to Sect. 5 for a discussion of their values.

Figure 3 shows that the fit is excellent. Only small discrepancies are visible for largest values of L, which corresponds to cases where the meniscus is close to the open end of the channel and may be influenced by the uncontrolled opening of the channel. Furthermore, if we fit independently the data for each channel, the values of the fitting parameters are in close agreement (see later Table 1). This shows the accuracy of our model to explain data in channels of slowly varying width.

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3.2.2 Channels with stepwise varying width

We proceed with two channels of stepwise varying length. Figure 4a shows the time evolution of the water length in these two channels. It is once again a decreasing function of time. However, the novel feature of the curves is the occurrence of finite changes of velocity at each step, which shows more clearly when we plot the velocity of the meniscus as a function of the water length (Fig. 4b). The velocity jump is positive, respectively negative, at the steps where the width decreases, respectively increases.

Qualitatively, as discussed in Sect. 3.2.1, this behaviour is consistent with the influence of the local value of the channel width at the meniscus location in (1). We can compare the experimental data with our model, bearing in mind that strictly speaking, the latter does not apply to stepwise variations in width, because it relies on an hypothesis of slowly varying width. We assume that the meniscus is in a portion of width w_0 and at a distance $L_0(t)$ from the upcoming step, and that the water-filled part of the channel spans N full portions of length L_i and constant width w_i (for *i* ranging from 1 to N). The total water length is then $L(t) = L_0(t) + \sum_{i=1}^{N} L_i$, and the total diffusive flux (2) writes:

$$Q_{\ell} = q_{\ell}(w_0)L_0 + \sum_{i=1}^{N} q_{\ell}(w_i)L_i = D_P \bar{c}_P^{\text{sat}}$$
$$\times \left[\left(\frac{w_0}{\delta} + \xi \right) L_0 + \sum_{i=1}^{N} \left(\frac{w_i}{\delta} + \xi \right) L_i \right],$$

where (4) and (5) have been used. Inserting into (3), we get:





Fig. 4 Drying dynamics in channels of stepwise varying width. **a** Time evolution of the water length. Here, we use all available data points, which are close enough to constitute a seemingly continuous curve. The top curve corresponds to a decreasing width, and the bottom one to an increasing width. Successive portions of constant

width are alternatively figured in plain and dotted lines. **b** Velocity of the meniscus as a function of the water length in channel of stepwise increasing (triangles) or decreasing (circles) width. The lines are fits by Eq. (10) with D_1 and D_2 as fitting parameters

$$\dot{L} = -\frac{L + L_{\rm sup}}{\tau},\tag{10}$$

with:

$$\tau = \frac{1}{D_1 A},\tag{11}$$

and:

$$L_{\rm sup} = \frac{D_2}{D_1 \sqrt{A}} - \sum_{i=1}^{N} \frac{w_i - w_0}{w_0 + \xi \delta} L_i,$$

where A is a channel-dependent geometrical parameter:

$$A = \frac{1}{hw_0} \left(\frac{w_0}{\delta} + \xi \right),\tag{12}$$

and D_1 and D_2 are given by (9).

We fit the data of Fig. 4b by Eq. (10) with D_1 and D_2 as fitting parameters. Contrary to the channels with linearly variable width, the agreement between the model and the data is not very good; in particular, the model predicts jumps in velocity which are much smaller in amplitude than what is experimentally observed. Possible reasons for this discrepancy are discussed in Sect. 5.2.

4 Branched channels with different widths

To extend further our study, we now consider branched channels, yet without reconnections, and with different widths. For simplicity, we study only channels where individual branches have constant widths.

4.1 Theory

Let us first consider a meniscus in the entrance channel. It is in contact with a water region which spans part of the entrance channel and all branches. All these contribute to the diffusive flux Q_{ℓ} introduced in Sect. 3.1. Hence, Eq. (3) is replaced by:

$$h\dot{L}w_m = -\int q_\ell(w(s))\mathrm{d}s - Q_g,\tag{13}$$

where *L* is the total length of the water region, w_m is the value of the width at the location of the meniscus, and the integral on the right-hand side spans all the water region, with *s* a curvilinear coordinate along the centreline of that region.

Let us specify the analysis to the case relevant to the experiments, of N branches of length L_i and constant width w_i (for *i* ranging from 1 to N) extending from the first node. We also consider that the entrance channel has

a constant width w_0 , and that the water inside this channel spans a length $L_0(t)$. The total water length is then $L(t) = L_0(t) + \sum_{i=1}^{N} L_i$, and (13) becomes:

$$h\dot{L}_0 w_0 = -q_\ell(w_0)L_0 - \sum_{i=1}^N q_\ell(w_i)L_i - Q_g.$$
(14)

Proceeding in a similar fashion as in Sect. 3.2.2, we obtain that the water length should obey Eq. (10), where the sum extends over all the branches ahead of the meniscus. The dynamics in intermediate channels also follows (14) and thus (10), provided w_0 is taken as the local width at the meniscus location, and the sum runs over all water-filled branches ahead of the meniscus. The dynamics in terminal branches is obtained by suppressing the sum from (10). In the current case where all branches have a constant width, the dynamics is therefore of the form $\dot{L}_0 = -(L_0 + L_{sup})/\tau$, where the characteristic time scale τ depends only on the local width (where the meniscus is located), while the length scale L_{sup} depends on the whole network ahead of the meniscus.

We proved in Dollet et al. (2021) that this analysis is true as long as all branches are slender (of length much larger than the transverse dimensions) and well separated (of mutual distances much larger than the transverse dimensions), in order to neglect diffusive coupling between channels separated by a small distance Noblin et al. 2008.

4.2 Experiments

4.2.1 Single-node networks

Figure 5a, c and e shows the distance travelled by the menisci as a function of time in the single-node networks with respectively two, three and five branches. In the entrance channel, the meniscus velocity increases as the number of branches increases, and there is a velocity jump as the meniscus in the entrance channel splits into several menisci in the branches. This result was already found for singlenode networks with uniform width, and can be ascribed to a jump of connected water length when meniscus splitting occurs at the node (Dollet et al. 2021). The new feature here is that the dynamics of each branch depends on its width: the smaller the latter, the larger the meniscus velocity and the shorter the drying time. Qualitatively, this is a signature of the fact that the typical drying time τ is an increasing function of the width. This can be seen from Eqs. (11) and (12), which show that such a dependence on width arises from the interplay between the diffusion between the channel top wall and the outer air [giving the term w/δ in (12)], and the diffusion between the channel side walls and the outer air [giving the term ξ in (12)]. We showed in Dollet et al. (2019) that this interplay is responsible for the faster drying of narrow

Fig. 5 Drying dynamics in single-node networks. From bottom to top, networks with two, three and five branches. Left column: temporal evolution of the distance travelled by the menisci from the entrance point. Right column: velocity as a function of the water length. Insets in panels **d** and **f** are zooms on the terminal branches. Symbols: entrance channel (0), branches of width 75 (∇), 100 $(\triangle), 125 (\triangleright), 150 (\triangleleft)$ and 175 μ m (\Box). The lines are fits by (10), with D_1 and D_2 as fitting parameters



channels because they are more prone to lose water from their side walls, compared to wider channels.

4.2.2 Tree network

To test more quantitatively the predictions of Sect. 4.1, we plot the velocity of the meniscus as a function of the water length in the right panel of Fig. 5, and we fit the curves by (10). For each single-node network, we impose that D_1 and D_2 be the same for the entrance channel and all the branches. The right panel of Fig. 5 shows that the fit is satisfactory; in particular, the terminal branches are correctly described, also the experimental data show some concavity, which may be ascribed to evaporative coupling effects discussed in Dollet et al. (2021). The entrance channels are also well fitted, with the exception of the five-branch network where the experimental slope is too large. We now turn to the tree network of Fig. 2d. It comprises 27 branches, but some of them share the same dimensions and "order" in the tree hierarchy, and it is found that they then follow a similar dynamics. We gather such branches into 9 different groups, represented by different symbols in Fig. 6a.

Figure 6b shows the distance travelled by the menisci as a function of time in the tree, for all branches. Like for the single-node networks, velocity jumps occur at each node. The new feature here is that at given depth in the tree network, the velocity is an increasing function of the water length ahead of the considered meniscus: the shorter the branches, the slower the dynamics (see inset of Fig. 6b). **Fig. 6** Drying dynamics in a tree. **a** Symbols used for the different branches. **b** Temporal evolution of the distance travelled by the menisci from the entrance point. The inset is a zoom on the terminal branches, to help distinguishing their drying dynamics. **c** Velocity as a function of the water length. The lines are fits by (10), with D_1 and D_2 as fitting parameters. The inset is a zoom on the bottom left part of the main plot



Table 1 Values of the fitting parameters D_1 and D_2 , and of the dimensions *h* and *H*, for all experiments

Experiment	Figure	$\frac{D_1}{(\mu m^2/s)}$	D_2 (×10 ⁴ µm ² /s)	<i>h</i> (μm)	Η (μm)
Linearly decreasing width	3	0.410	2.53	39.3	89.3
Linearly increasing width		0.394	2.42		
Stepwise decreasing width	4	0.184	2.83	45.6	84.5
Stepwise increasing width		0.540	0.79		
2-branch network	5a,b	0.389	1.83		
3-branch network	5c,d	0.395	1.62	42.2	87.4
5-branch network	5e,f	0.437	1.32		
tree	6	0.265	1.57	41.4	89.2

Here again, we plot the velocity of the meniscus as a function of the water length (Fig. 6c), and we fit the curves by (10). We find once again a good agreement between the experiments and the data, which suggests that our physical modelling remains adapted to the case of more complex branched networks.

5 Discussion

5.1 Values of the fitting parameters

In each experiment, the two physicochemical parameters $D_1 = D_p C_p^{\text{sat}}$ and $D_2 = \sqrt{\alpha D_a D_p}$ have been used as fitting parameters. Their values are reported in Table 1. They show some dispersion, especially for the channels with stepwise varying width, for which it has been shown that the model is inaccurate. The other experiments yield $D_1 = 0.4 \ \mu \text{m}^2/\text{s}$ and $D_2 = 2 \times 10^4 \ \mu \text{m}^2/\text{s}$. Taking values from the literature: $D_p = 10^{-9} \ \text{m}^2/\text{s}$ (Watson and Baron 1996), $D_a = 2 \times 10^{-5} \ \text{m}^2/\text{s}$, $\alpha = 0.03$ (Harley et al. 2012) and $\overline{c}_p^{\text{sat}} = M C_p^{\text{sat}} / \rho = 7.2 \times 10^{-4}$ [using in particular $C_p^{\text{sat}} = 40 \ \text{mol/m}^{-3}$ (Randall and Doyle 2005)], we compute $D_1 = 0.7 \ \mu \text{m}^2/\text{s}$ and $D_2 = 2 \times 10^4 \ \mu \text{m}^2/\text{s}$. Bearing in mind that the values of the physicochemical parameters D_p , α and C_p^{sat} are known with relatively poor precision, we can conclude that the values of our fitting parameters are in good agreement with the reported values of D_p , α and C_p^{sat}

5.2 Limitations of the current model and perspectives

We have compared all our experiments with a model based on the hypothesis of slowly varying width. This enables to consider that water diffusion through PDMS acts mainly in the transverse direction. We have shown that such a model captures well our experimental observations, except the case of channels with stepwise varying width.

Obviously, in such a case, the hypothesis of slowly varying width is violated at the location of the steps. To go beyond and extend the model, the full 3D diffusion of water through PDMS should be solved, at least in the vicinity of the steps. However, this requires time-consuming numerical simulations, beyond the scope of this study.

Moreover, an additional ingredient may become important in the case of steps: when passing an abrupt variation in channel section, the meniscus curvature necessarily varies at given contact angle, to match the geometry of the channel walls. In other contexts, such an effect is important for droplet breakup in microfluidic T-junctions (Ody et al. 2007) and in soap films and foams in porous media (Xu and Rossen 2003; Cox et al. 2004; Géraud et al. 2017). This induces variations of capillary pressure, hence of pressure in the liquid, which may deform the channel, an effect hitherto disregarded in the model. This might explain why velocity jumps in Fig. 4 are much more marked in experiments that in the model. Interestingly, in the context of tree physiology, such effects are relevant in the xylem network of trees (Tyree and Sperry 1989; Choat et al. 2003), where xylem vessels are separated by pits of nanometric dimensions where menisci between embolism and liquid can get trapped while undergoing huge variations of capillary pressure. In an upcoming study, we will revisit this effect by mimicking pits in microfluidic experimental micromodels, to clarify the couplings between drying, meniscus dynamics, capillary pressure variations and channel deformability.

6 Conclusions

Our study revealed that drying in networks with channels of variable sections are well described by a generalised version of our previously published model, consisting in the integration of the section-dependent diffusive flux over the length of the channels (Eq. 3). This model fits very well with experiments for linearly varying section, stepwise-decreasing width and division in branches with different sections. The drying dynamics is always smooth, without any arrest, just velocity changes. However, studies on real leaves (Ponomarenko et al. 2014; Brodribb et al. 2016) showed a stop-and-go dynamics with arrests and sudden jumps. This peculiar dynamics may be the consequence of the fact that conduits are not fully continuous in plants, but are made of collection of separate conduits connected by pits. Future work will focus on the effect of tiny constrictions in drying dynamics.

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Declaration

Conflict of interest The authors declare that they have no conflict of interest.

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