

# A linear-elastic-nonlinear-swelling theory for hydrogels. Part 2. Displacement formulation

## Electronic supplementary material

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### 1. Experimental details

To carry out the motivating experiments for the drying cylinders, cubes of commercially-available poly-(acrylamide-potassium-acrylate) copolymer (*Deco Cubes*<sup>TM</sup>, JRM Chemical Inc, 4881 NEO Parkway, Cleveland, Ohio 44128, USA) were placed in deionised water and left to swell until they reached a steady size. These were then removed, and placed in a dish of deionised water, which was kept topped up through the process of the experiment such that it never dried out fully. No quantitative measurements were taken at any point, and the experiments were used solely to motivate the subsequent theoretical analysis that followed, with the key qualitative features (curved top and bottom interfaces) apparent.

### 2. A numerical scheme for solving equation (5.36)

Since the size of the domain on which this differential equation is to be solved changes in time, introduce the scaled variable  $Y = Z/\mathcal{H}(T)$  and instead solve on  $Y \in [0, 1]$ . Then

$$\frac{\partial}{\partial T} \rightarrow \frac{\partial}{\partial T} - \frac{Y}{\mathcal{H}} \frac{\partial \mathcal{H}}{\partial T} \frac{\partial}{\partial Y} \quad \text{and} \quad \frac{\partial}{\partial Z} \rightarrow \frac{1}{\mathcal{H}} \frac{\partial}{\partial Y}, \quad (2.1)$$

and equation (5.36) thus becomes, after rearrangement,

$$\begin{aligned} \frac{\partial \Phi}{\partial T} - \Phi^{1/3} \frac{\partial \Phi}{\partial Y} \int_0^Y \frac{\partial \Phi^{1/3}}{\partial T} dY' - \frac{\Phi^{1/3}}{\mathcal{H}} \frac{\partial \mathcal{H}}{\partial T} \frac{\partial \Phi}{\partial Y} \int_0^Y \Phi^{1/3} dY' = \\ \frac{\Phi}{\mathcal{H}^2} \frac{\partial}{\partial Y} \left[ f(\Phi) \frac{\partial \Phi}{\partial Y} \right] - \frac{1}{\mathcal{H}} \left[ \frac{2f(\Phi)}{3\mathcal{H}} \frac{\partial \Phi}{\partial Y} + Y \left( \phi^{2/3} - 1 \right) \frac{\partial \mathcal{H}}{\partial T} \right] \frac{\partial \Phi}{\partial Y} + 2\Phi^{4/3} U_s, \end{aligned} \quad (2.2)$$

with the height  $\mathcal{H}(T)$  set as

$$\mathcal{H}(T) = \left( \int_0^1 \Phi^{1/3} dY' \right)^{-1}, \quad (2.3)$$

and the Neumann boundary condition replaced by

$$f(\Phi) \frac{\partial \Phi}{\partial Y} = \mathcal{H} U_t \quad \text{at } Y = 1. \quad (2.4)$$

Now, discretising the domain  $[0, 1]$ , and sampling at  $Y = 0, \Delta, \dots, N\Delta$ , let  $U$  be the  $(N+1)$ -dimensional vector with  $i^{\text{th}}$  component

$$U_i = \Phi((i-1)\Delta, T), \quad (2.5)$$

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for  $\Delta = 1/N$ . Similarly define  $\mathbf{V}$  to have  $i^{\text{th}}$  component

$$V_i = \Phi((i-1)\Delta, T)^{-2/3} \left. \frac{\partial \Phi}{\partial T} \right|_{Z=(i-1)\Delta}. \quad (2.6)$$

Also introduce the diagonal matrices  $\mathbf{U}$  with entries  $U_i^{2/3}$  and  $\mathbf{D}$  with entries  $D_i = \Phi^{1/3} \partial \Phi / \partial Y$  evaluated at  $Y = (i-1)\Delta$ ,

$$\mathbf{U} = \text{diag}(U_1^{2/3}, U_2^{2/3}, \dots, U_{N+1}^{2/3}) \quad \text{and} \quad \mathbf{D} = \text{diag}(D_1, D_2, \dots, D_{N+1}). \quad (2.7)$$

Then, the time derivative  $\partial \Phi / \partial T$  in equation (2.2) can be replaced by  $\mathbf{UV}$ . It is also now possible to discretise the integrals, noticing that

$$\int_0^Y \Phi^{-2/3} \frac{\partial \Phi}{\partial T} dY \rightarrow \frac{1}{2N} \underbrace{\begin{pmatrix} 0 & 0 & \dots & 0 \\ 1 & 0 & \dots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ 1 & 1 & \dots & 0 \\ 1 & 1 & \dots & 1 \end{pmatrix}}_{(N+1) \times N} \underbrace{\begin{pmatrix} 1 & 1 & 0 & \dots & 0 & 0 \\ 0 & 1 & 1 & \dots & 0 & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & \dots & 1 & 1 & 0 \\ 0 & 0 & \dots & 0 & 1 & 1 \end{pmatrix}}_{N \times (N+1)} \mathbf{V}, \quad (2.8)$$

where the first matrix represents the integral and the second matrix product gives the values of  $\Phi^{-2/3} \partial \Phi / \partial T$  in the centres of each grid space. This can be more concisely written  $\mathbf{MV}$ , for

$$\mathbf{M} = \frac{1}{2N} \begin{pmatrix} 0 & 0 & 0 & 0 & \dots & 0 & 0 \\ 1 & 1 & 0 & 0 & \dots & 0 & 0 \\ 1 & 2 & 1 & 0 & \dots & 0 & 0 \\ \vdots & \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 1 & 2 & 2 & 2 & \dots & 2 & 1 \end{pmatrix}, \quad (2.9)$$

an  $(N+1) \times (N+1)$  matrix. Then, equation (2.2) can be rewritten in discretised form with

$$\left( \mathbf{U} - \frac{2}{3} \mathbf{DM} \right) \mathbf{V} = \mathbf{F} \left( \mathbf{U}, \mathcal{H}, \dot{\mathcal{H}}; \mathcal{M}, \mathcal{Q} \right), \quad (2.10)$$

where spatial derivatives are approximated as finite differences and  $\mathbf{F}$  represents the right-hand side terms. Then,

$$\frac{\partial \mathbf{U}}{\partial T} = \left[ \left( \mathbf{U} - \frac{2}{3} \mathbf{DM} \right)^{-1} \mathbf{F} \left( \mathbf{U}, \mathcal{H}, \dot{\mathcal{H}}; \mathcal{M}, \mathcal{Q} \right) \right] \mathbf{U}. \quad (2.11)$$

This is then solved with a predictor-corrector method. Let  $\mathbf{U}^{(n)} = \mathbf{U}(n\Delta T)$ , and use the same superscript notation for other quantities evaluated at  $T = n\Delta T$ , then

$$\begin{aligned} \mathbf{U}^{(n+\frac{1}{2})} &= \mathbf{U}^{(n)} + \frac{\Delta T}{2} \left[ \left( \mathbf{U}^{(n)} - \frac{2}{3} \mathbf{DM}^{(n)} \right)^{-1} \mathbf{F} \left( \mathbf{U}^{(n)}, \mathcal{H}^{(n)}, \dot{\mathcal{H}}^{(n)}; \mathcal{M}, \mathcal{Q} \right) \right] \mathbf{U}^{(n)} \quad \text{and} \\ \mathbf{U}^{(n+1)} &= \mathbf{U}^{(n)} + \Delta T \left[ \left( \mathbf{U}^{(n+\frac{1}{2})} - \frac{2}{3} \mathbf{DM}^{(n+\frac{1}{2})} \right)^{-1} \mathbf{F} \left( \mathbf{U}^{(n+\frac{1}{2})}, \mathcal{H}^{(n+\frac{1}{2})}, \dot{\mathcal{H}}^{(n+\frac{1}{2})}; \mathcal{M}, \mathcal{Q} \right) \right] \mathbf{U}^{(n+\frac{1}{2})}. \end{aligned} \quad (2.12)$$

We impose the Dirichlet boundary condition on the base of the gel by requiring  $U_1^{(n)} = 1$  and the

evaporative Neumann condition at the top by requiring

$$\frac{U_{N+1}^{(n)} - U_N^{(n)}}{\Delta} = \frac{\mathcal{H}(T)U_t}{1 + (4M/3) \left( U_{N+1}^{(n)} \right)^{1/3}}. \quad (2.13)$$

This system was solved using MATLAB for all of the plots and numerical results in this paper, carrying out the matrix inversion using `mldivide`. An example function which solves this equation is shown below for illustrative purposes.

### 3. MATLAB code

```

1 function [phiReturns, dPhiReturns, hReturns] = numericalDry(M,
    Ut, Us, times, N, dt)
2     %NUMERICALDRY Numerically solve the governing equation for
    the polymer fraction
3     % field when drying of a cylinder from the top and sides.
    Returns phiReturns, the polymer
4     % fraction field; dPhiReturns, the Y-derivative of the
    polymer fraction field; hReturns,
5     % the height of the cylinder \mathcal{H}(T) -- all at the
    specified timesteps.
6
7     % M: the material parameter M
8     % Ut: the non-dimensional flux Ut from the top
9     % Us: the non-dimensional flux Us from the sides
10
11    % times: the times to save values at
12
13    % N: the number of spatial grid steps
14    % dt: the timestep
15
16    % ---
17
18    % Number of steps to save at
19    savedSteps = max(size(times, 2), size(times, 1)); % allow
    for row/column vectors for time
20
21    % *** Variables to return
22    phiReturns = NaN*ones(N+1, savedSteps);
23    dPhiReturns = NaN*ones(N+1, savedSteps);
24    hReturns = NaN*ones(1, savedSteps);
25
26    t = 0;
27    currentIndex = 1;
28
29    % *** Key variables used at each timestep
30    p = ones(N+1, 1); % polymer fraction, evaluated at edges of
    cells

```

```

31  pMids = ones(N, 1); % polymer fraction, evaluated at
    middles of cells
32
33  dpMids = zeros(N, 1); % dPhi/dY, evaluated at middles of
    cells
34  dp = zeros(N+1, 1); % dPhi/dY, evaluated at edges of cells
35  dp(end) = Ut/(1+4*M/3); % (straight away, set evaporation
    flux from top)
36
37  h = 1; % scaled height of gel
38  hDot = 0; % dH/dT
39
40  % *** Construct the matrix M
41  MBase = zeros(N+1, N+1);
42  for k = 2 : N+1
43      MBase(k, 1) = 1/(2*N);
44      if(k~=2)
45          for j = 2:(k-1)
46              MBase(k, j) = 1/N;
47          end
48      end
49      MBase(k, k) = 1/(2*N);
50  end
51
52  % *** Loop until we've reached the final timestep
53  while (currentIndex <= savedSteps)
54      holdP = p; % introduce a holding variable for polymer
    fraction to allow the predictor-corrector steps
55
56      % Step forward by dt/2 and correct to improve stability
57      for timestep = [dt/2 dt]
58          % *** Diffusive bracket term
59          bktMids = (1 + (4*M/3)*pMids.^(-2/3)).*dpMids;
60          baseBkt = (1+(4*M/3)*p(1).^(-2/3))*dp(1);
61          endBkt = Ut*h;
62
63          % d[]/dY
64          dBkt = N*[2*(bktMids(1)-baseBkt); bktMids(2:end)-
    bktMids(1:end-1); 2*(endBkt - bktMids(end))];
65
66          % prefactor is the matrix U-(2/3)DM
67          prefactor = diag(holdP.^(2/3)) - (2/3)*diag(dp.*
    holdP.^(1/3))*MBase;
68
69          % *** Advective terms on the right-hand side
70          advect = (2/(3*h^2))*(1+(4*M/3)*holdP.^(-2/3)).*dp
    .^2 + ...
71          (hDot/h)*linspace(0, 1, N+1)'.*(holdP.^(2/3)-1)
    .*dp - ...
72          (hDot/h)*MBase*holdP.^(1/3);

```

```

73
74     % The right-hand side
75     rhs = holdP.*dBkt/h^2 + 2*holdP.^(4/3)*Us - advect;
76
77     % Calculate d(Phi)/dT using mldivide
78     dpdt = (prefactor\rhs)./holdP.^(-2/3);
79
80     % *** TIMESTEP
81     holdP = p + timestep*dpdt;
82     holdP(1) = 1; % Dirichlet BC on base
83
84     % Get midpoint phis
85     pMids = (holdP(1:end-1)+holdP(2:end))/2;
86
87     % Calculate d(Phi)/dY, including applying Neumann
88     % BC on top
89     dp = N*[2*(pMids(1)-holdP(1)); pMids(2:end)-pMids
90         (1:end-1); h*Ut/(N*(1+(4*M/3)*holdP(end)^(-2/3))
91         )];
92     dpMids = (dp(1:end-1)+dp(2:end))/2;
93
94     % Calculate H and Hdot
95     hNew = N/sum(pMids.^(1/3));
96     hDot = (hNew-h)/dt;
97     h = hNew;
98 end
99
100 % Catch numerical errors
101 if(isnan(holdP))
102     break;
103 end
104
105 p = holdP;
106
107 % Save the values if we're at a reporting step
108 if(t >= times(currentIndex))
109     phiReturns(:, currentIndex) = p;
110     dPhiReturns(:, currentIndex) = dp;
111     hReturns(currentIndex) = h;
112
113     % Increment next counter
114     currentIndex = currentIndex + 1;
115 end
116
117 t = t+dt;
118 end
119
120 end

```