

A systematically reduced model for organoid expansion

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Industrially Focused Mathematical Modelling

1. INTRODUCTION

Organoids are:

- three-dimensional multi-cellular structures;
- grown *in vitro* to **recapitulate** *in vivo* micro–anatomy;
- derived from patient tissue and retain many of their features (e.g. structure, pathology, heterogeneous cellular composition [1]);
- a more realistic model of *in vivo* cells than flat 2D cell line cultures;
- suitable for use in large-throughput drug screens;
- labour-intensive and time-consuming to grow.

Cellesce utilise bioreactor technology to grow organoids at scale. Key priorities are ensuring **reproducibility** of organoid output and **uniformity** of organoid size, through bioreactor design and identifying suitable operating conditions. Achieving these goals is time-consuming via experimental work alone. Through the development of a mechanistic mathematical model, we are able to provide quantitative predictions of fluid flow and metabolite concentrations throughout the bioreactor.

Key Question: How do the operating conditions affect the nutrient delivery to and waste removal from the organoids?

The key experimental control is the **flow rate**. We consider the effect of the flow velocity on the metabolite concentrations within the bioreactor and the extent to which the flow facilitates nutrient delivery and waste removal.

Organoids are cultured within the 'CXP1' bioreactor (see Figure 1) using the following protocol:

- Organoids are seeded as single cells in a thin layer of hydrogel, which acts as a porous scaffold;
- Nutrient-rich culture media is pumped across the top of the hydrogel;
- The organoids are grown for approximately 7 days, until they are around $40-80\mu m$ in diameter and contain about 50 cells;



Organoids are extracted from the hydrogel and tested for quantity, size, and viability.



Figure 1 – (Top left) Example of colorectal organoid⁴, stained for nuclear (blue) and cytoskeletal (red) markers for imaging. Scale 50µm. (Top right) Schematic of 'CXP1' bioreactor [2]. (Bottom) Twodimensional reduction of bioreactor, with arrow showing direction of media flow. Blue is media, yellow is hydrogel, grey is organoid biomass.

2. Model development



14)

boundaries, we impose no flux.

3.MODEL REDUCTION

We derive reduced models of the full system, Eqs. (1)-(4), for metabolite concentrations $c(t,x) = c_M = c_H$ and $w(t,x) = w_M = w_H$ by considering the system in the limit $\epsilon \to 0$ and obtaining the leading-order behaviour of the system by averaging in z, where \bar{u} is the depth-averaged flow velocity and we define the parameters $\theta = h_H/(h_M - h_H)$, and $D_C = (D_{CM} + \theta D_{CH})$ and $D_W = (D_{WM} + \theta D_{WH})$:

Longwave approximation:
$$\frac{[u]}{L} \sim \nu N_0 \sim \frac{D_{ij}}{L^2}$$

 $(1+\theta)\frac{\partial c}{\partial t} + \bar{u}\frac{\partial c}{\partial x} = D_C \frac{\partial^2 c}{\partial x^2} - \theta \nu N_0 c e^{pt}$, (7)
 $(1+\theta)\frac{\partial w}{\partial t} + \bar{u}\frac{\partial w}{\partial x} = D_W \frac{\partial^2 w}{\partial x^2} + 2\theta \nu N_0 c e^{pt}$, (8)
with boundary and initial conditions:
 $\bar{u}c - D_C \frac{\partial c}{\partial x} = \bar{u}, \ \bar{u}w - D_W \frac{\partial w}{\partial x} = 0 \text{ at } x = 0$, (9)
 $\frac{\partial c}{\partial x} = 0, \ \frac{\partial w}{\partial x} = 0 \text{ at } x = L$, (10)
 $c = \frac{c-\infty}{1+\theta}, \ w = 0 \text{ at } t = 0$. (11)
A,
 a_{1}
 a_{2}
 a_{2}
 a_{3}
 B_{1}
 a_{3}
 a_{3}
Sublimit model: $\frac{[u]}{L} \sim \nu N_0 \gg \frac{D_{ij}}{L^2}$
 $(1+\theta)\frac{\partial c}{\partial t} + \bar{u}\frac{\partial c}{\partial x} = -\theta \nu N_0 c e^{pt}$, (12)
 $(1+\theta)\frac{\partial w}{\partial t} + \bar{u}\frac{\partial w}{\partial x} = 2\theta \nu N_0 c e^{pt}$, (13)
with boundary and initial conditions:
 $c = c_{-\infty}, \ w = 0 \text{ at } x = 0$, (14)
 $c = \frac{c_{-\infty}}{1+\theta}, \ w = 0 \text{ at } t = 0$. (15)



4. EFFECT OF FLOW RATE FOR A SPECIFIED ORGANOID LINE

We characterise a cell line via the cell proliferation and glucose uptake rates. We consider the following metrics to optimise the CXP1 operating parameters, e.g. peak media velocity, for a specified cell line.



Figure 4 – Results showing how glucose (A, B, C) and lactate (D, E, F) concentrations change over time during a typical experiment. (\mathbf{A}, \mathbf{D}) Results from z-averaged full model; (\mathbf{B}, \mathbf{E}) longwave approximation; (\mathbf{C}, \mathbf{F}) sublimit of longwave approximation. Parameter values given in Eq. (16).

$c_{-\infty} = 0.36 \mathrm{mol}\mathrm{m}^{-2},$	$p = 3.9 \times 10^{-6} \mathrm{s}^{-1},$	$[u] = 10^{-6} \mathrm{m s^{-1}},$	$N_0 = 2.7 - 4 \times 10^{10} \text{cell m}^{-2},$	
$L = 9 \times 10^{-2} \mathrm{m},$	$h_H = 1 \times 10^{-3} \mathrm{m},$	$h_M = 3 \times 10^{-3} \mathrm{m},$	$\nu = 9.4 \times 10^{-17} \mathrm{m}^2 \mathrm{cell}^{-1} \mathrm{s}^{-1},$	(16)
$D_{CH} = D_{CM} = 6 \times 10^{-2}$	$^{10} {\rm m}^2 {\rm s}^{-1}, \qquad D_{WH}$	$= 1.2 \times 10^{-9} \mathrm{m}^2 \mathrm{s}^{-1},$	$D_{WM} = 1.4 \times 10^{-9} \mathrm{m}^2 \mathrm{s}^{-1}.$	

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denoted by $\epsilon = h_M / L \ll 1$.

5. Key points

• Presented an unsteady, two-dimensional model for metabolite transport within the CXP1 bioreactor. • Derived two reduced models by exploiting extreme spatial and temporal parameter ratios in system. • Provided framework for improving organoid viability through varying bioreactor operating conditions.

REFERENCES

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