

MATHEMATICAL MODELING AND OPTIMIZATION IN CRYOBIOLOGY

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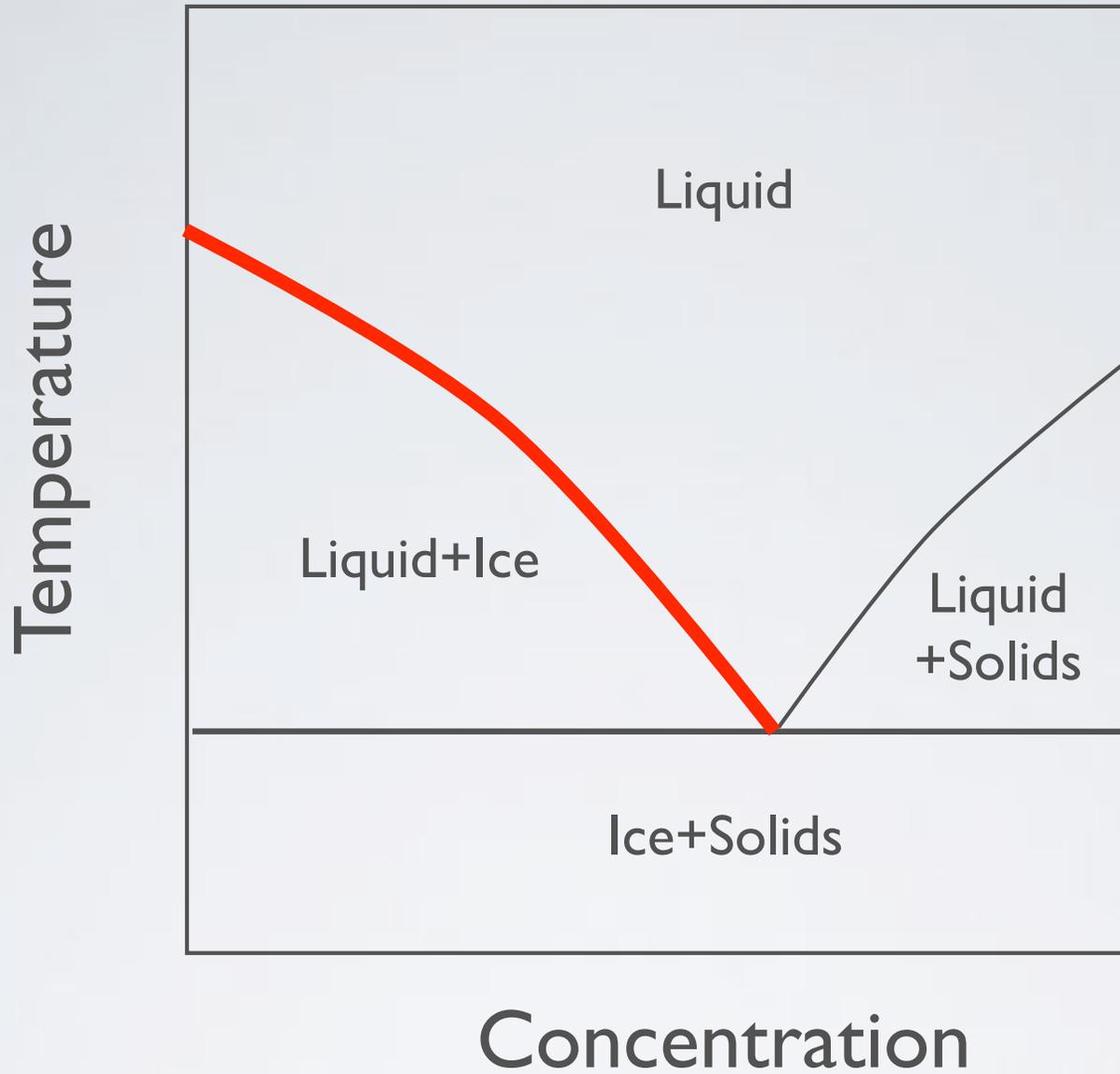
OUTLINE

- Brief introduction to principles of Cryobiology
- Model development at three length scales
- Optimal control
- Current and future directions

WHY FREEZE BIOSPECIMENS?

- Colder temperatures mean longer storage: at least 100 years in LN
- Banking, distribution and testing of cells and tissues, maybe organs in the future
- Worldwide initiatives to preserve genetic samples
 - Millennium Seed Bank, Svalbard Seed Bank, UK Biobank (0.5M samples)
 - JAX Sperm bank (>10000 strains)
 - NCRR, MMRRRC, MRRRC
 - NCI-Office of Biorepositories and Biospecimine Research
- Kill unwanted cells and tissues in living systems





HOW CELL FREEZING WORKS



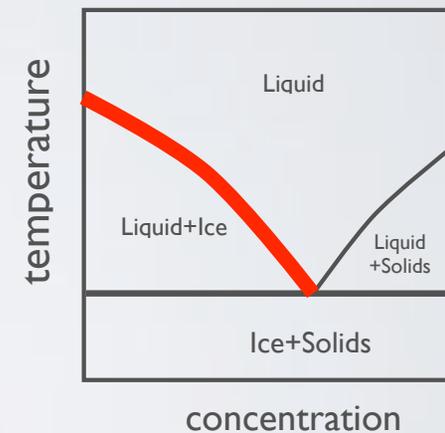
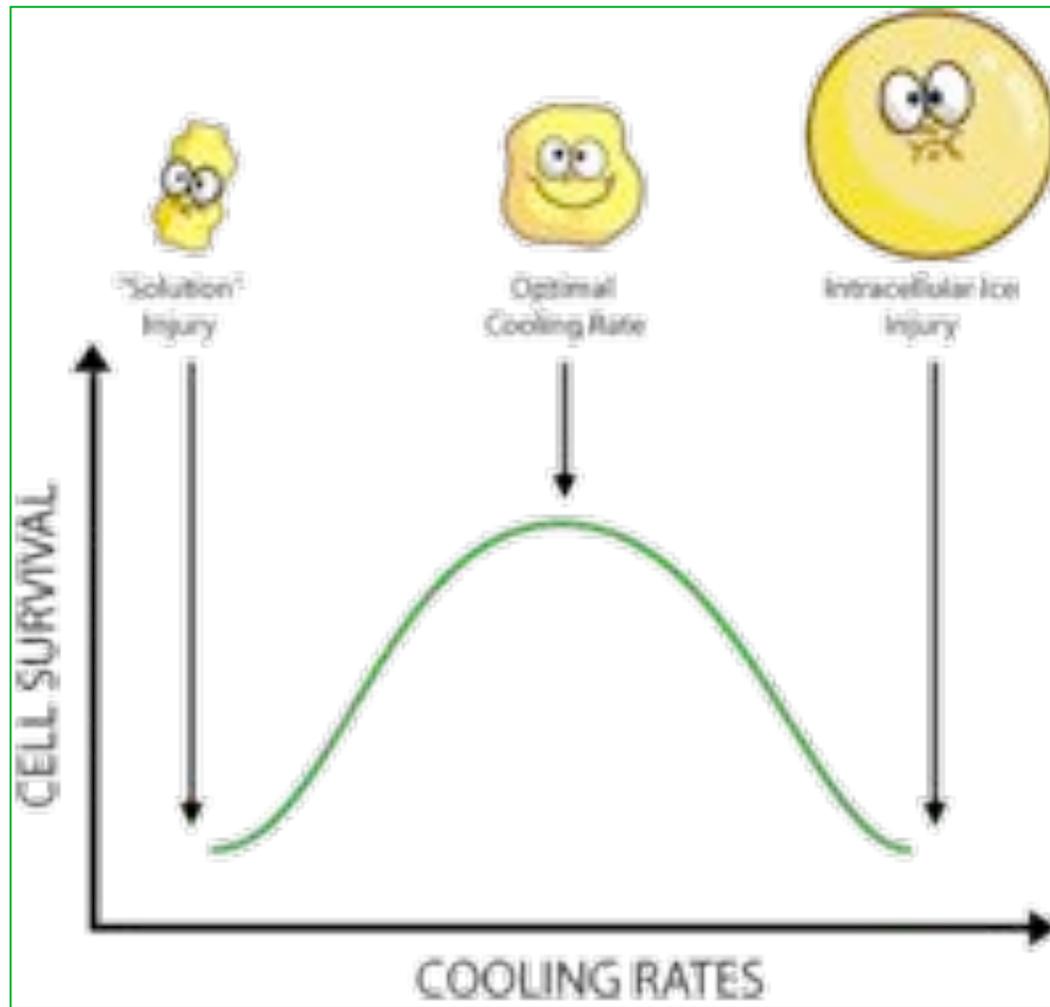
HOW CELL FREEZING WORKS





Salt Alone	Salt	CPA
1%	1%	10%
10%	1.5%	15%
30%	3%	30%
-	4%	40%

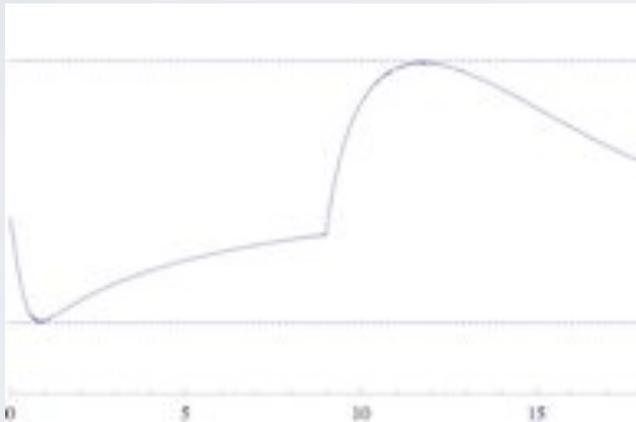
- To reduce the effects of high salt concentrations and to aid in “glass formation” we add *cryoprotective agents* (CPAs)



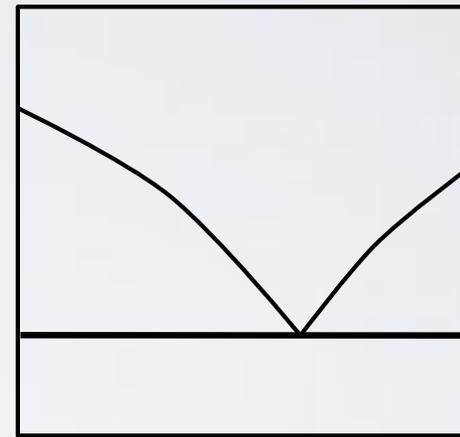
THE TWO FACTOR HYPOTHESIS

CRITICAL CRYOBIOLOGICAL QUANTITIES

Concentration



Heat



Above 0°C these quantities govern osmotically induced damage

Below 0°C these quantities govern the likelihood of intracellular ice

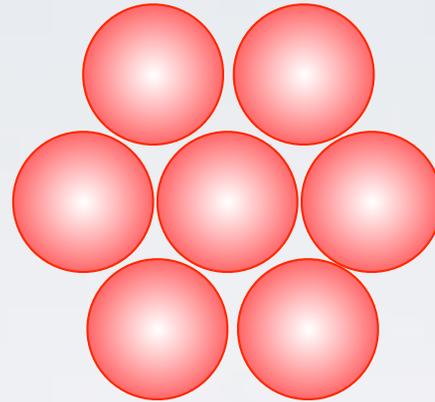


TRANSPORT PROBLEMS

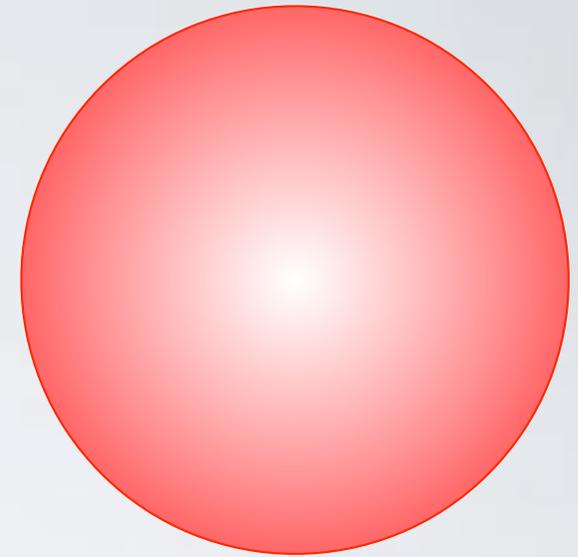
Model Selection



Single Cell
Suspensions



Multi Cell
Tissues



Larger Tissues
and Organs

Mass

ODE
System

Hybrid ODE/PDE
System

PDE
System

**Heat/
Solidi-
fication**

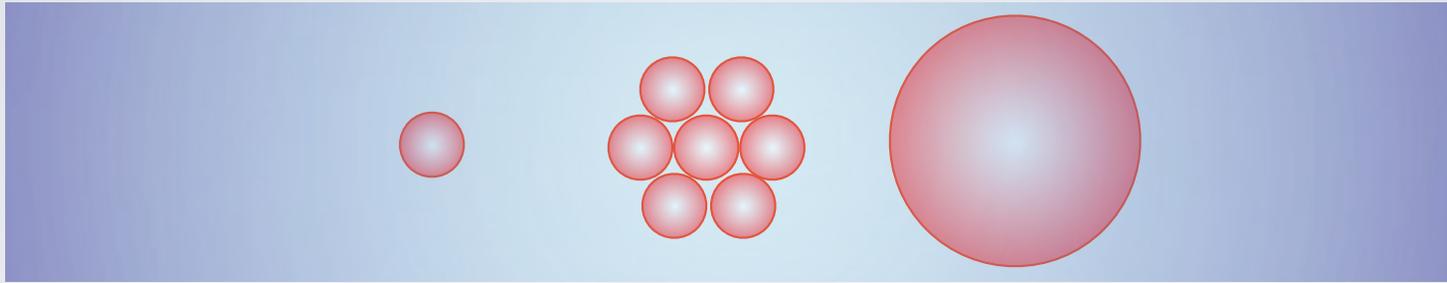
Stochastic
ODE

Large Monte
Carlo System

Nonlinear heat &
stefan problem

TRANSPORT PROBLEMS

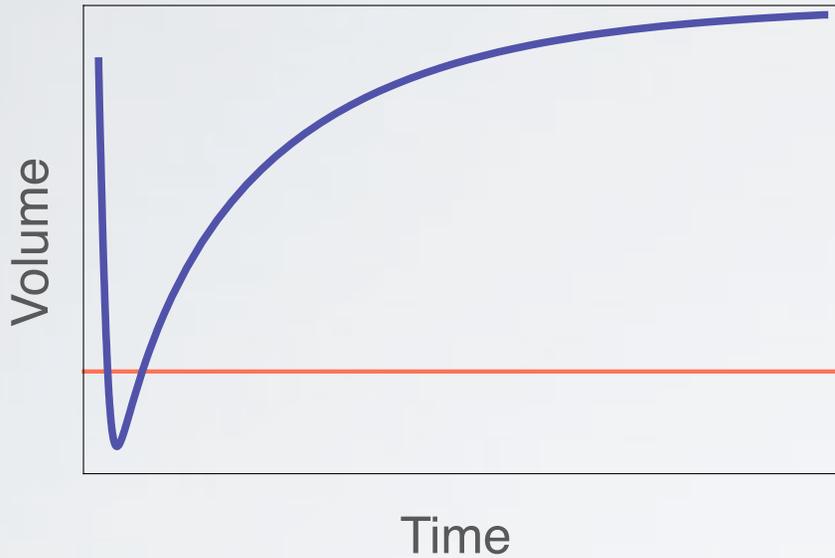
Model Selection



All models in cryobiology are coupled systems!

THE SINGLE CELL PROBLEM





$$\dot{n}_1 = P_1(\mu_1^{ext} - \mu_1^{int})$$
$$\dot{n}_2 = P_2(\mu_2^{ext} - \mu_2^{int})$$

MASS TRANSFER



MASS TRANSFER

THE CHOICE OF μ

$$\Phi(T, P, N) = N_1 \mu_0 + \sum_{i=2}^n N_i kT \ln \frac{N_i}{e N_1} + \sum_{i=2}^n N_i \psi_i + \frac{1}{2N_1} \sum_{i,j=2}^n \beta_{ij} N_i N_j$$

Differentiating with respect to N_1 or N_i and setting $\beta_{ij}/kT = (B_i + B_j)$

$$\mu_1 = \mu_0 - kT \left(\sum_{i=2}^n m_i + \frac{1}{2} \sum_{i,j=2}^n (B_i + B_j) m_i m_j \right)$$

$$\mu_i = kT \left(\ln m_i + \psi_i^* + \sum_{j=1}^n (B_i + B_j) m_j \right).$$

Specific Model: set $B_i = 0$ and $M_i \approx x_2/x_1$.

$$\dot{x}_1 = \frac{x_{np}}{x_1} + \sum_{j=2}^k \frac{x_j}{x_1} - \sum_{i=1}^n M_i,$$

$$\dot{x}_2 = b_2 \left(M_2 - \frac{x_2}{x_1} \right),$$

\vdots

$$\dot{x}_n = b_n \left(M_n - \frac{x_n}{x_1} \right),$$

Cellular Quantities

x_1 = Water Volume

$x_{2,\dots,n}$ = Moles of permeating solute

x_{np} = Moles of nonpermeating solute

$b_{2,\dots,n}$ = Relative permeability

Extracellular Quantities

M_1 = Nonpermeating solute molality

$M_{2,\dots,n}$ = Permeating solute molality

\bar{M}_i = Maximal i th solute molality

SINGLE CELLS

Specific Model

$$\dot{x}_1 = \frac{1}{x_1} \left(x_{\text{np}} + \sum_{j=2}^k x_j - \sum_{i=1}^n M_i x_1 \right),$$

$$\dot{x}_2 = \frac{b_2}{x_1} (M_2 x_1 - x_2),$$

⋮

$$\dot{x}_n = \frac{b_n}{x_1} (M_n x_1 - x_n),$$

Cellular Quantities

x_1 = Water Volume

$x_{2,\dots,n}$ = Moles of permeating solute

x_{np} = Moles of nonpermeating solute

$b_{2,\dots,n}$ = Relative permeability

Extracellular Quantities

M_1 = Nonpermeating solute molality

$M_{2,\dots,n}$ = Permeating solute molality

\bar{M}_i = Maximal i th solute molality

We have a system of the form $\dot{x}(t) = \lambda(x(t))f(x(t))$, where $\lambda(x(t)) = 1/x_1(t)$ is a positive scalar function. In this case, we can define an invertible transformation

$$q(\tau) = \int_0^\tau \frac{1}{\lambda(x(s))} ds = \int_0^\tau x_1(s) ds$$

and a new system $w'(\tau) = f(w(\tau))$ such that

$$w(\tau) = x(q(\tau))$$

meaning that we may, without any penalty, linearize the system by removing the $1/x_1$ term.



$$x'_1(\tau) = x_{np} + \sum_{j=2}^n x_j - \sum_{i=1}^n M_i(\tau)x_1,$$

$$x'_2(\tau) = b_2 (M_2(\tau)x_1 - x_2),$$

$$\vdots$$

$$x'_n(\tau) = b_n (M_n(\tau)x_1 - x_n).$$

or

$$x' = f(x, M) := A(M)x + x_{np}e_1,$$

where

$$A(M) = \begin{pmatrix} -\sum_{i=1}^n M_i & 1 & 1 & \dots & 1 \\ b_2 M_2(t) & -b_2 & 0 & \dots & 0 \\ b_3 M_3(t) & 0 & -b_3 & \dots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ b_n M_n(t) & 0 & 0 & \dots & -b_n \end{pmatrix}.$$

Define

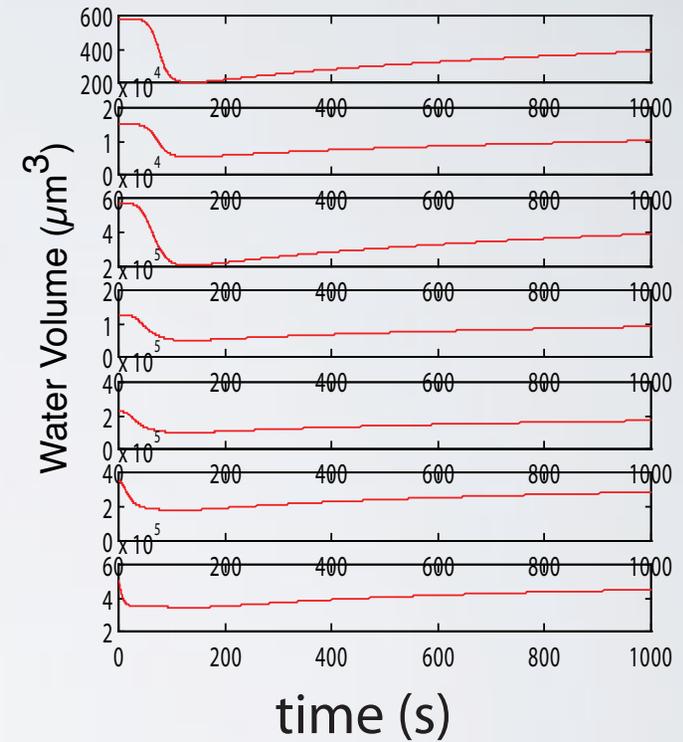
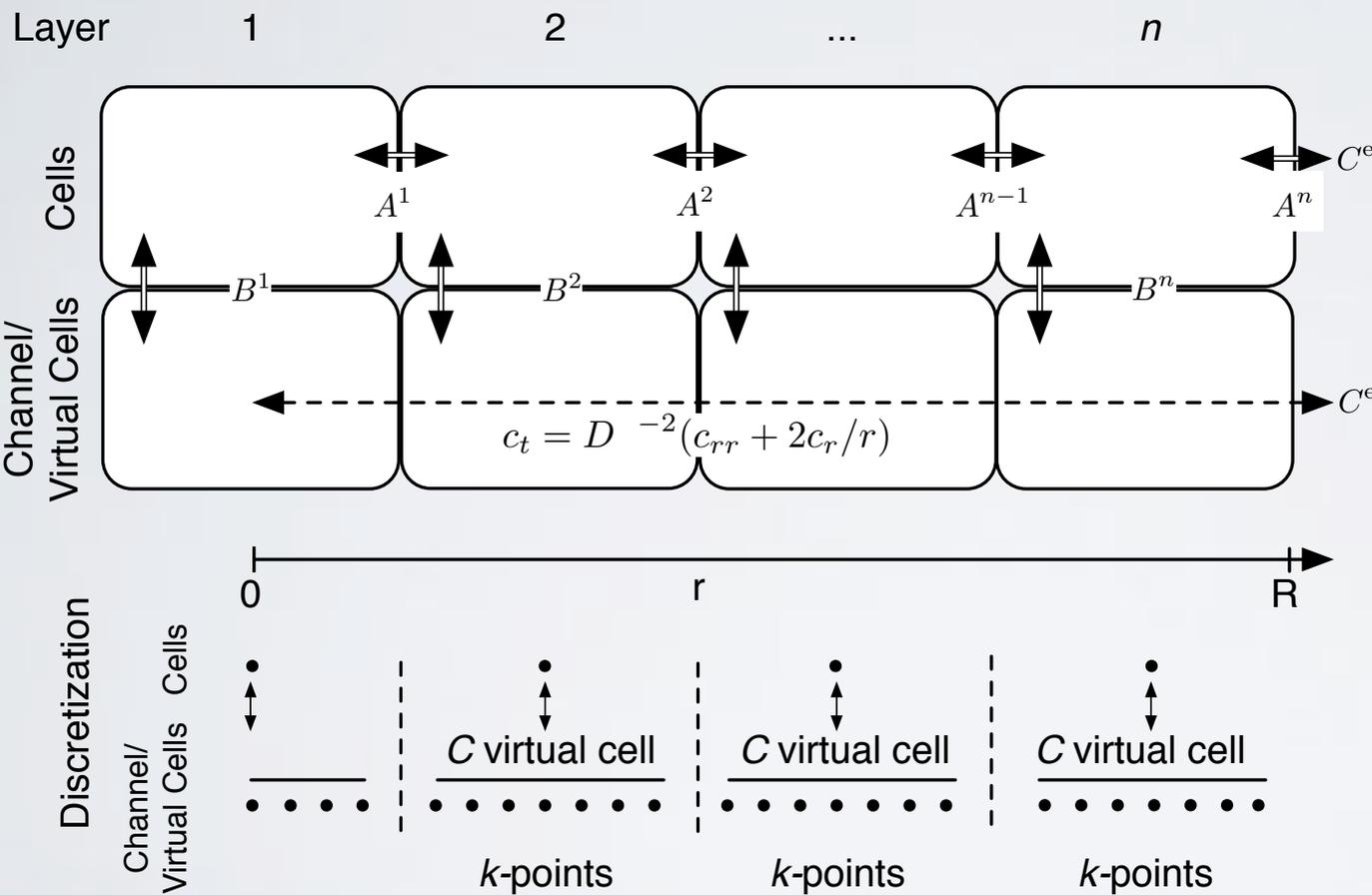
$$D := \text{diag}(1, (b_2 M_2)^{-1/2}, \dots, (b_n M_n)^{-1/2})$$

Then:

$$DA(M)D^{-1} = \begin{pmatrix} -\sum_i M_i & \sqrt{b_2 M_2} & \sqrt{b_3 M_3} & \dots & \sqrt{b_n M_n} \\ \frac{\sqrt{b_2 M_2}}{\sqrt{b_2 M_2}} & -b_2 & 0 & \dots & 0 \\ \frac{\sqrt{b_3 M_3}}{\sqrt{b_3 M_3}} & 0 & -b_3 & \dots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ \frac{\sqrt{b_n M_n}}{\sqrt{b_n M_n}} & 0 & \dots & 0 & -b_n \end{pmatrix}$$

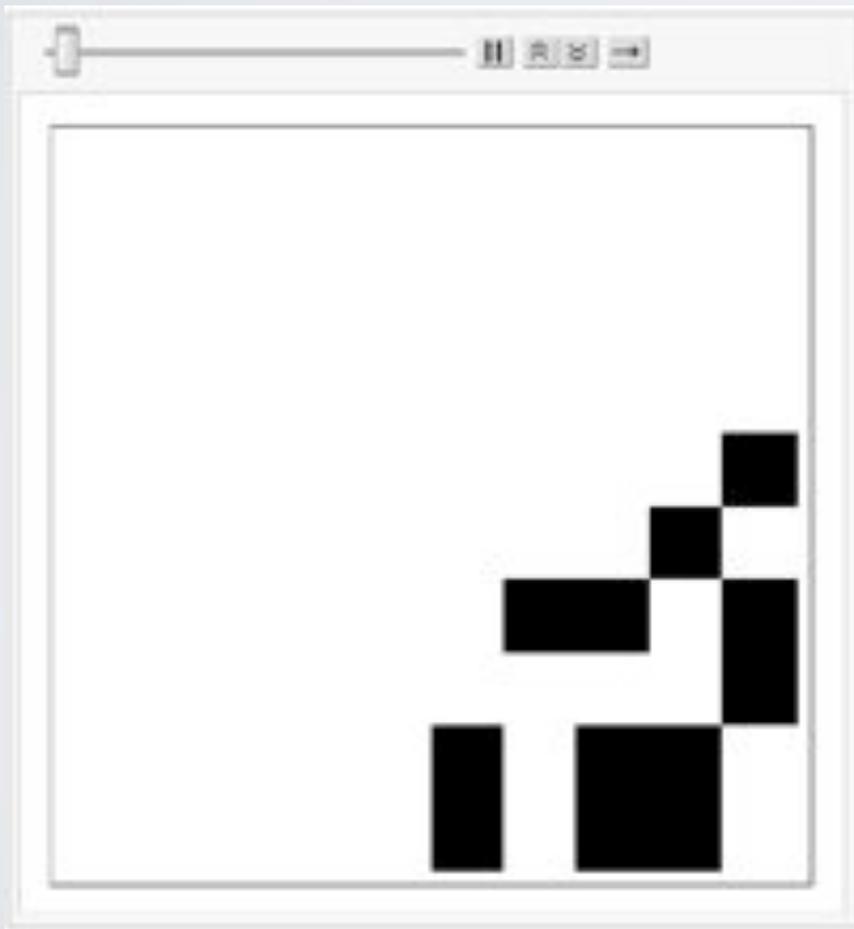
is symmetric, negative definite, and our original n-dimensional nonlinear system is globally asymptotically stable.

MASS TRANSPORT IN SMALL TISSUES



SOLIDIFICATION DURING COOLING, SMALL TISSUES:

Monte Carlo Simulation of IIF



$$p_j(\delta\tau) \approx p_j^i + p_j^p$$
$$\approx (1 + k_j\alpha)\delta t$$

p_j is the probability of ice

p_j^i is the probability of ice forming spontaneously

p_j^p is the probability of ice propagating from neighbors

k_j is the number of icy neighbors

MASS TRANSFER IN LARGE TISSUES

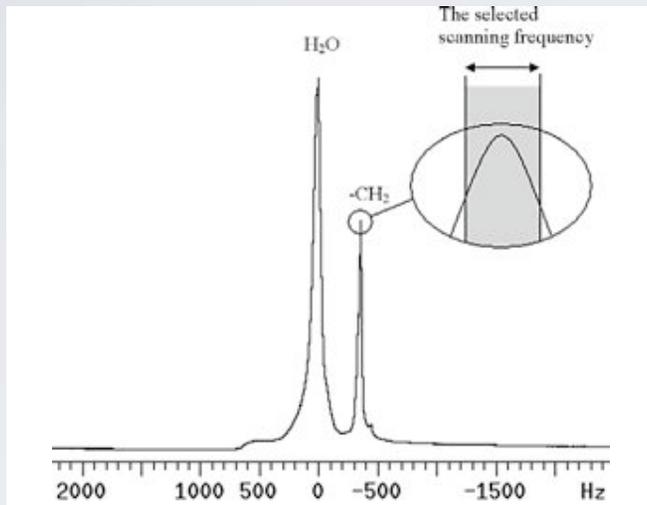


Fig. 2. The proton MR spectrum at 7 T from the sample holder loaded with two ovaries and 40% (w/w) EG solution. The excitation frequency was centered at the resonance frequency for the -CH₂ group in EG molecules.

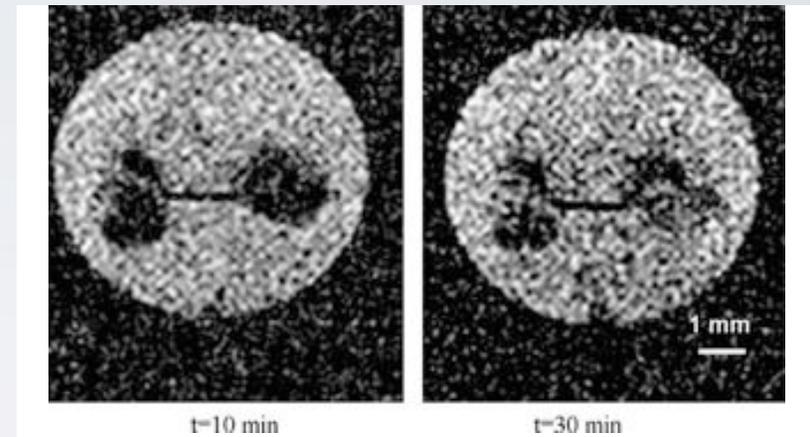


Fig. 5. Two sample MR images, with water signal saturated, showing the increasing EG concentration in ovaries during perfusion.

$$\frac{\partial c}{\partial t} = \nabla \cdot (D \nabla c)$$

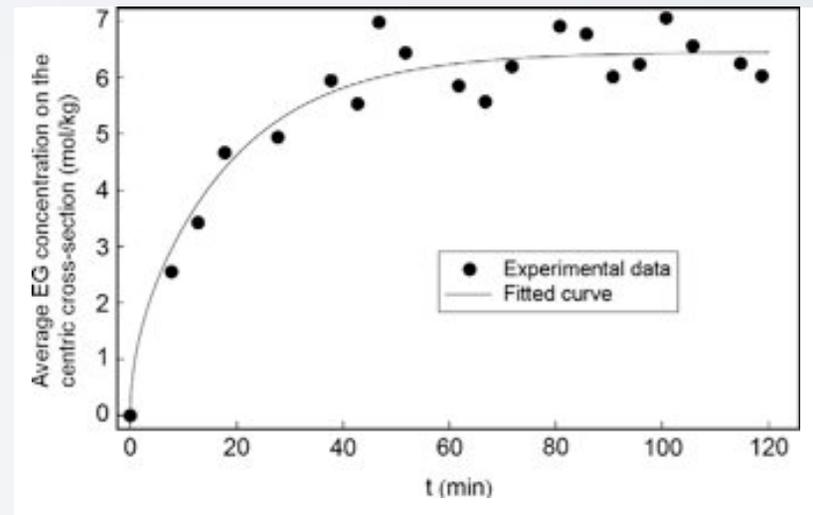
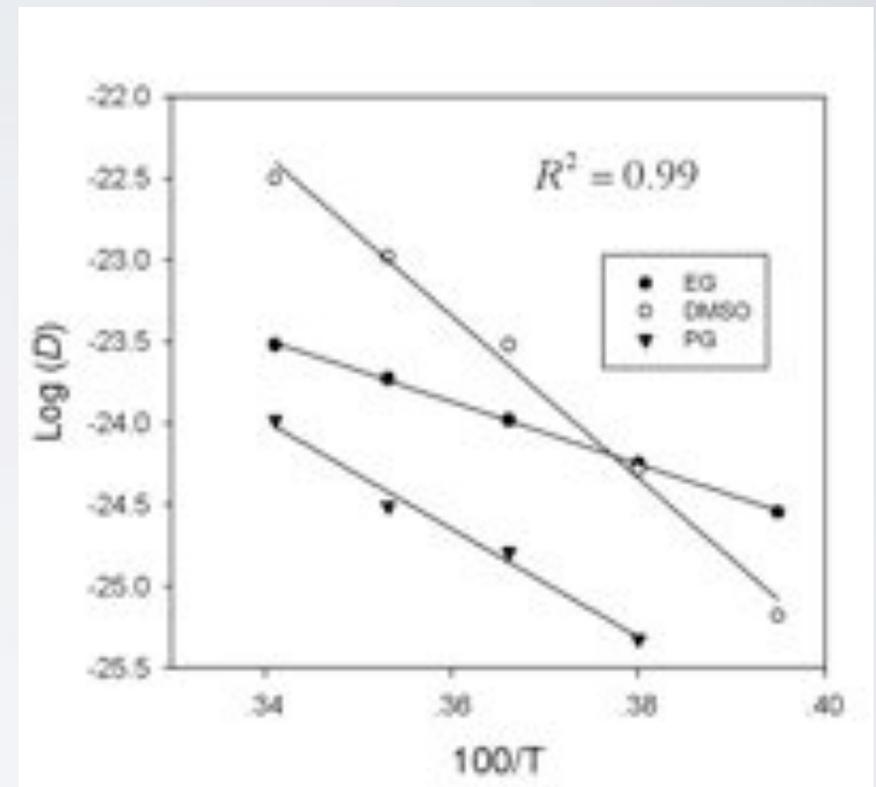
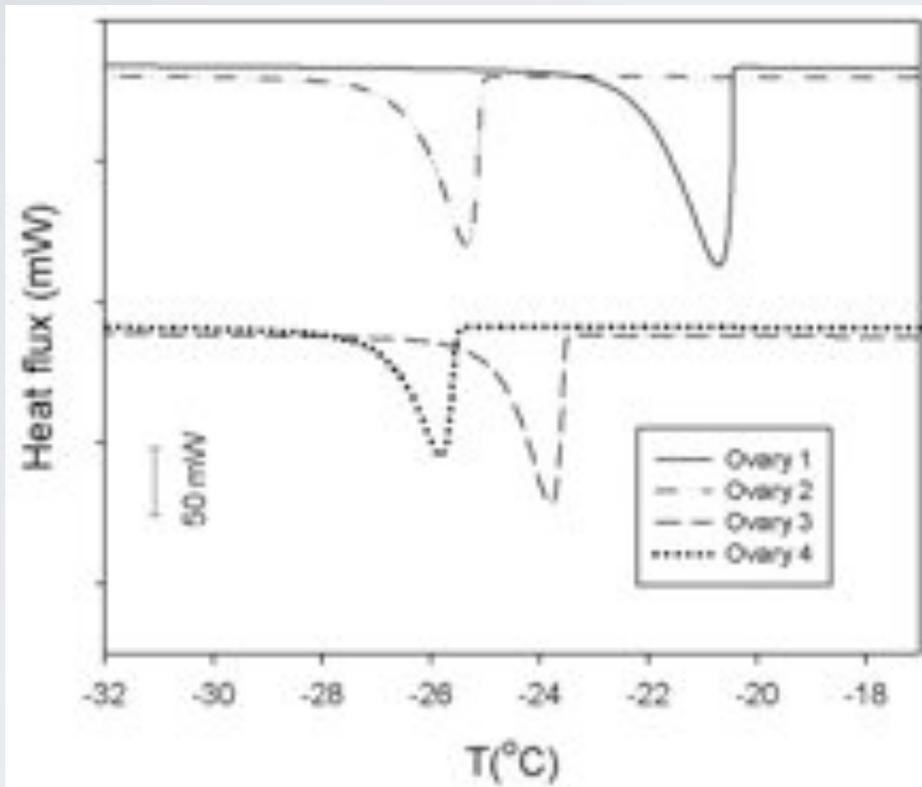


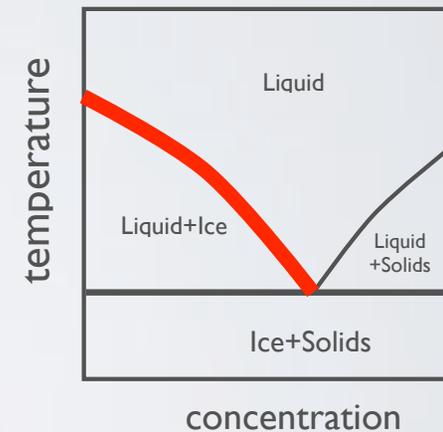
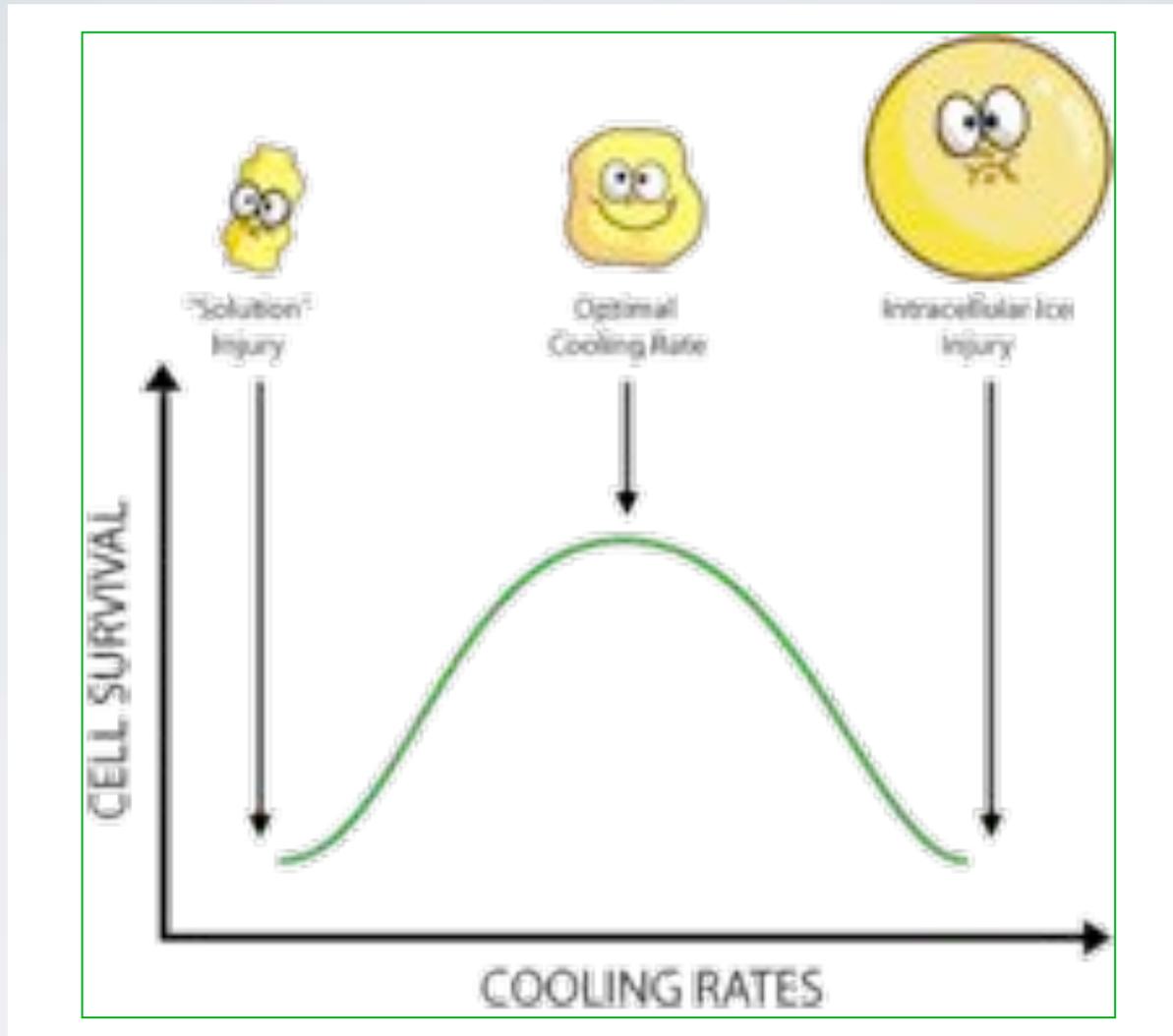
Fig. 6. The experimental data with their fitted curve for the average EG concentration change on the centric cross-section of an ovary with 1.1 mm as its identical radius.

MASS TRANSFER IN LARGE TISSUES

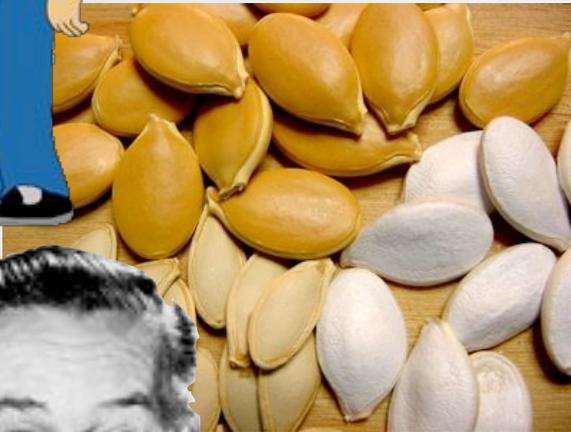
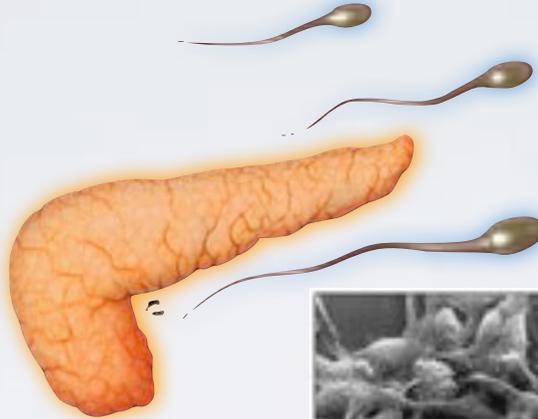
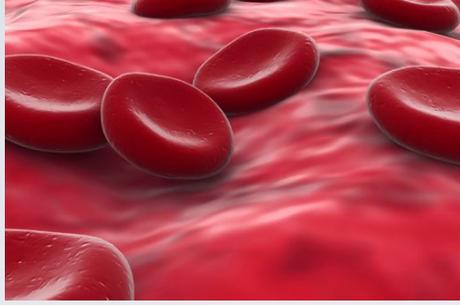


$$\frac{\partial c}{\partial t} = \nabla \cdot (D(T) \nabla c)$$

$$D(T) = \exp(-E_a/RT)$$

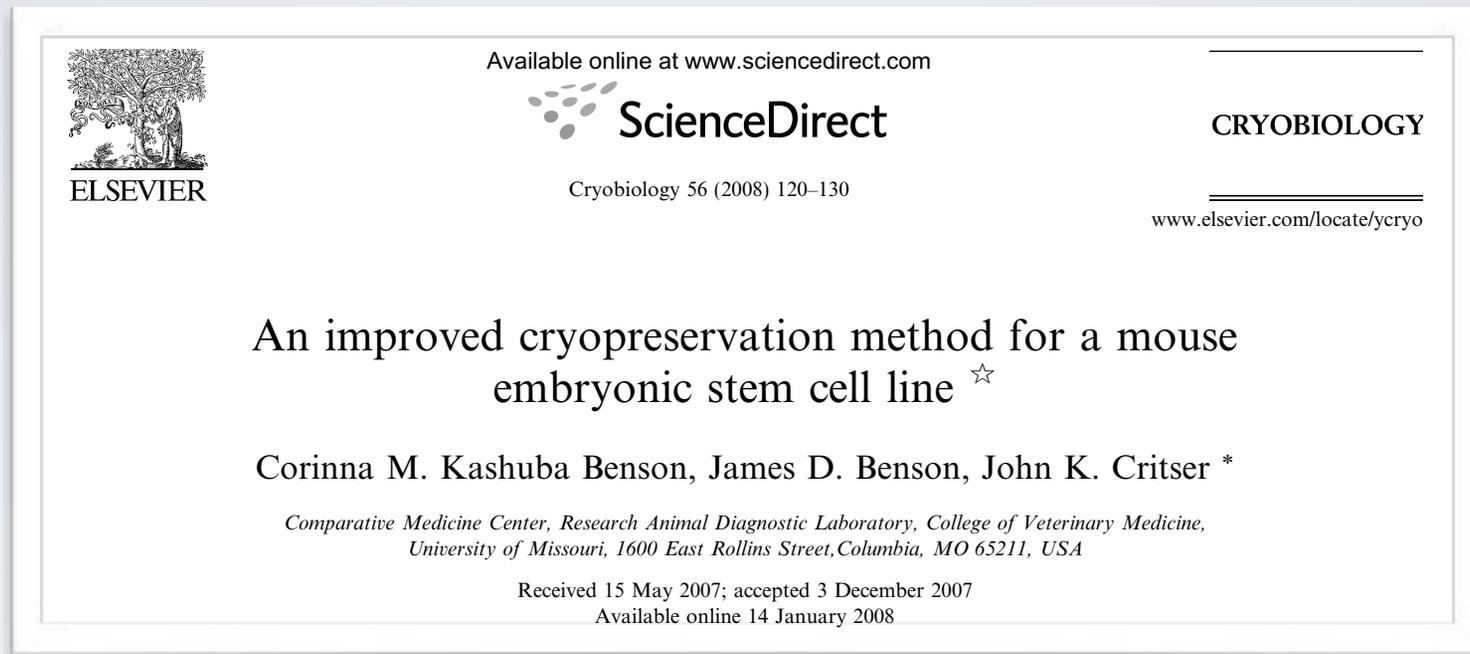


WHAT CAN WE CONCLUDE FROM THE ABOVE MODELS?



HEAT AND MASS
TRANSFER LIMIT THE SIZE
OF FREEZABLE TISSUE!

DOES MODELING IN CRYOBIOLOGY WORK?

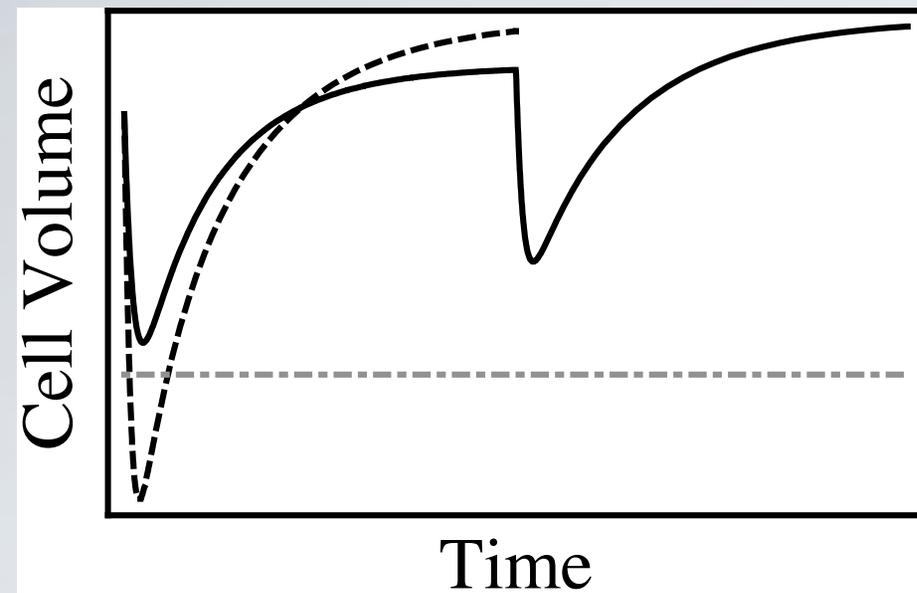


- Previous best protocol: 31% recovery
- “Optimally” defined new best protocol: 64% recovery

OPTIMAL CONTROL IN CRYOBIOLOGY:

- control quantity to minimize cost J (e.g. time, energy, stress, P_{IIF} or combinations.)
- subject to exact and inequality constraints:
 - exact constraints: governing physical system, (e.g. 2P model, heat equation, diffusion, etc).
 - inequality constraints: state or control constraints, (e.g. cell volume > 0).





$\min_{M \in \mathcal{A}} s_f$ subject to

$$\dot{w}_1 = \frac{x_{np}}{w_1} + \frac{w_2}{w_1} - M_1 - M_2,$$

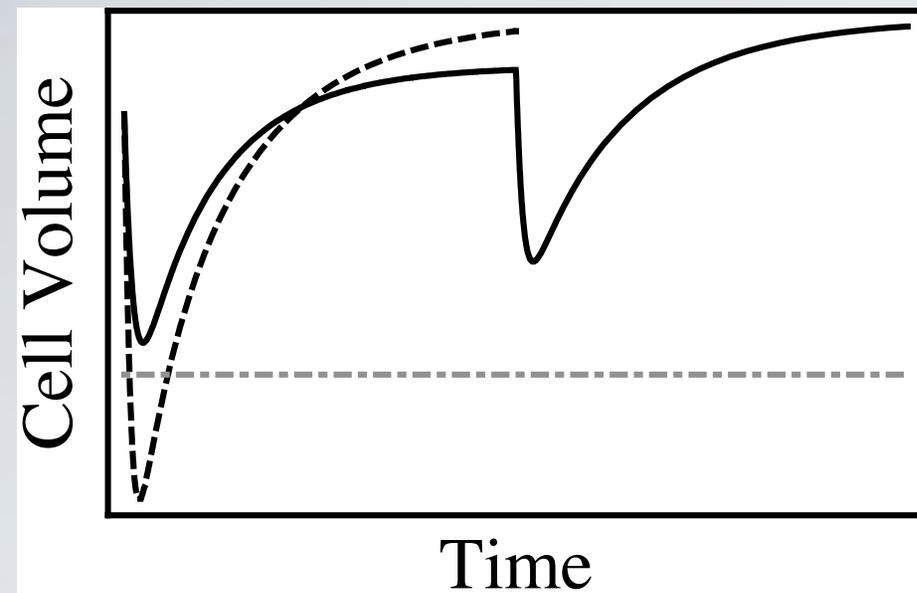
$$\dot{w}_2 = b_2 \left(M_2 - \frac{w_2}{w_1} \right),$$

and

$$w_1 + \gamma w_2 - k^* \leq 0,$$

$$k_* - w_1 - \gamma w_2 \leq 0.$$

FIRST CONTROL PROBLEM



$$\min_{M \in \mathcal{A}} s_f = q(t_f) = \int_0^{t_f} x_1(\tau) d\tau$$

subject to

$$\dot{x}_1 = -(M_1 + M_2)x_1 + x_2 + x_{np}$$

$$\dot{x}_2 = bM_2x_1 - bx_2$$

and

$$x_1 + \gamma x_2 - k^* \leq 0,$$

$$k_* - x_1 - \gamma x_2 \leq 0.$$

Bilinear state equation (in controls and state) give:

Existence ✓

Controllability ✓

OPTIMAL CONTROL

$$H(x^*, p^*, M^*) = \max_{M \in CP} (A(M)x + x_1 e_1) \cdot p - x_1$$

$$= \max_{M \in CP} \left(-M_1 x_1 p_1 + x_1 \sum_{i=2}^n M_i (b_i p_i - p_1) \right)$$

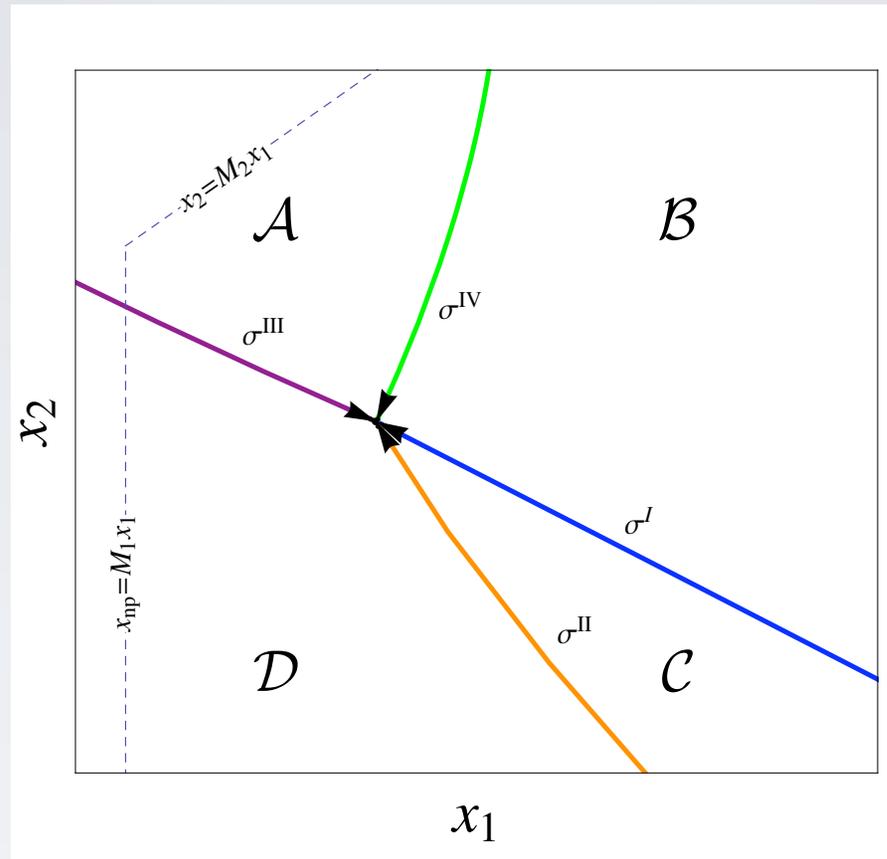
+ terms with no M)

$$M_1(t) = \begin{cases} 0, & p_1 > 0 \\ \bar{M}_1, & p_1 \leq 0 \end{cases}$$

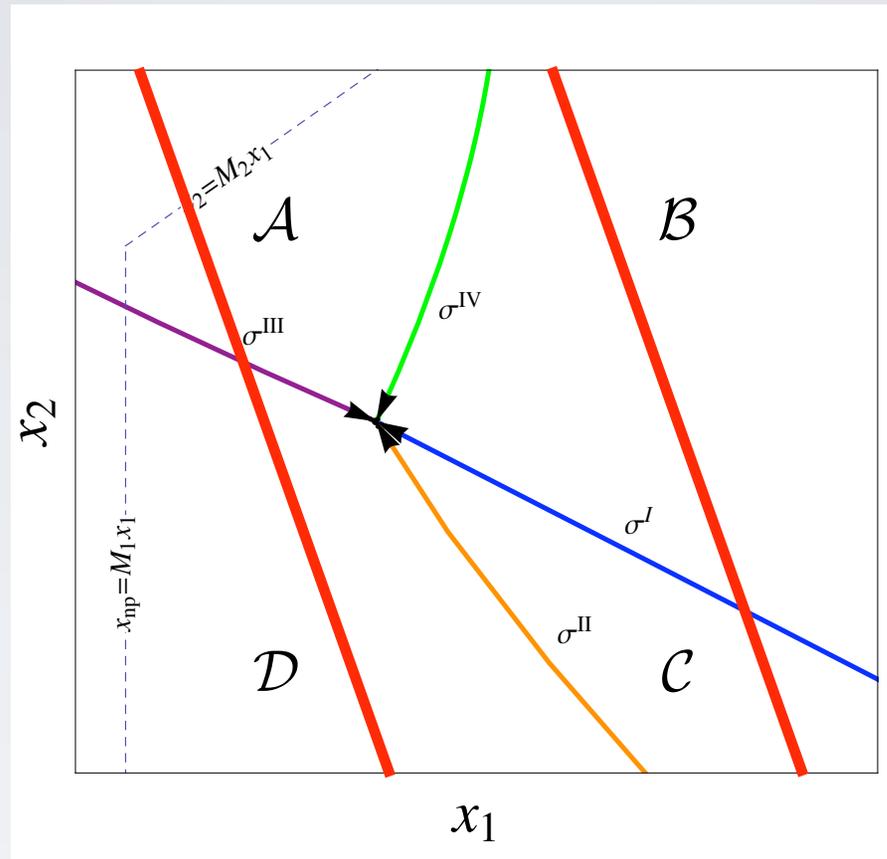
$$M_i(t) = \begin{cases} 0, & b_i p_i - p_1 < 0 \\ \bar{M}_i, & b_i p_i - p_1 \geq 0 \end{cases} \cdot$$



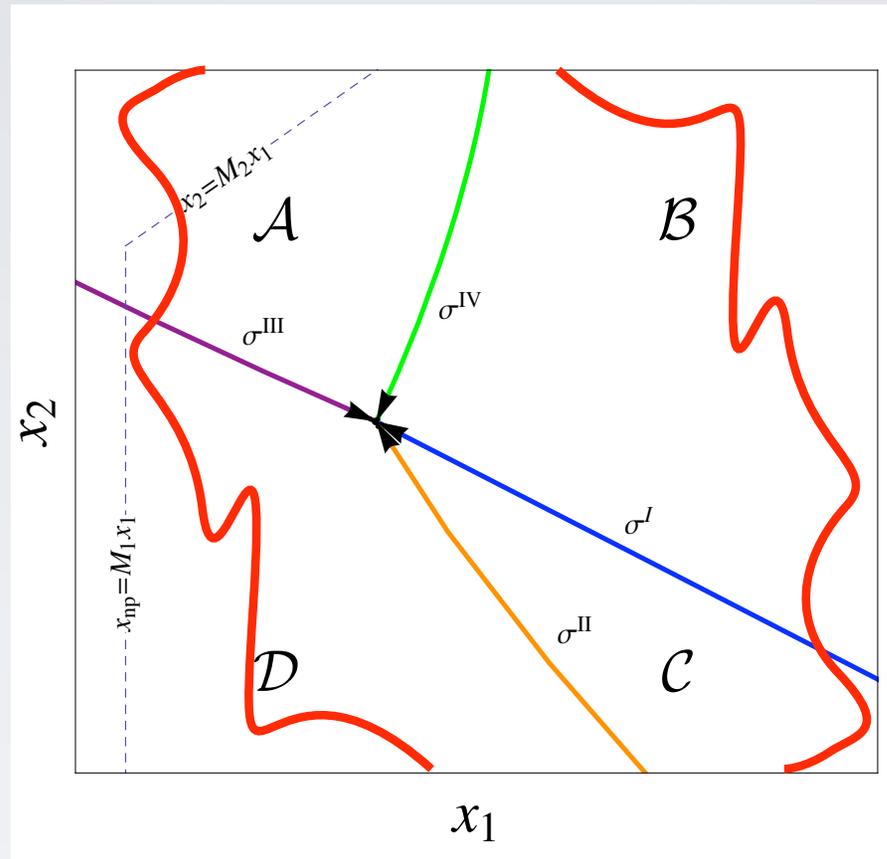
Optimal controls maximize the Hamiltonian



WHY GEOMETRIC OPTIMAL CONTROL?

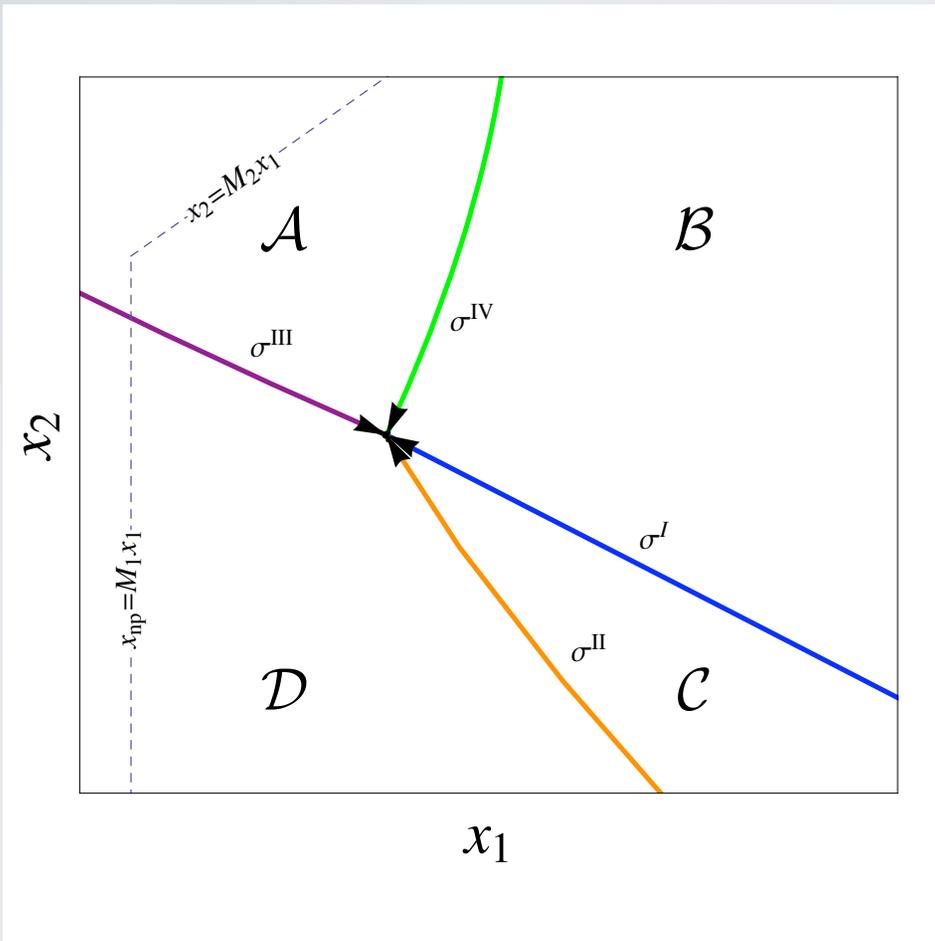


WHY GEOMETRIC OPTIMAL CONTROL?



WHY GEOMETRIC OPTIMAL
CONTROL?

Boltayanskii sufficiency theorem: a “regular, distinguished” trajectory defined by a state dependent control function $v(x)$ is optimal.



Region	Control Scheme	M_1	M_2
σ^I	M^I	\bar{M}_1	\bar{M}_2
C, D, σ^{II}	M^{II}	0	\bar{M}_2
σ^{III}	M^{III}	0	0
A, B, σ^{IV}	M^{IV}	\bar{M}_1	0

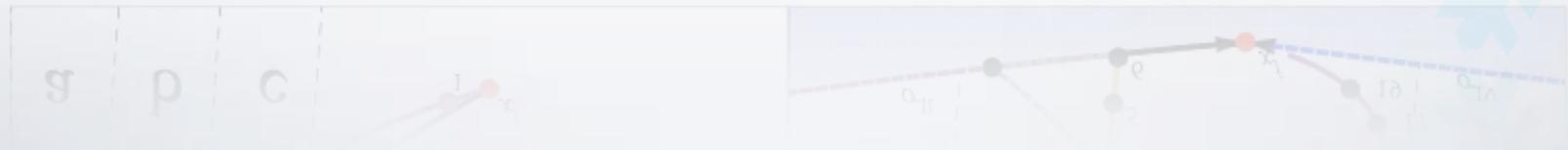
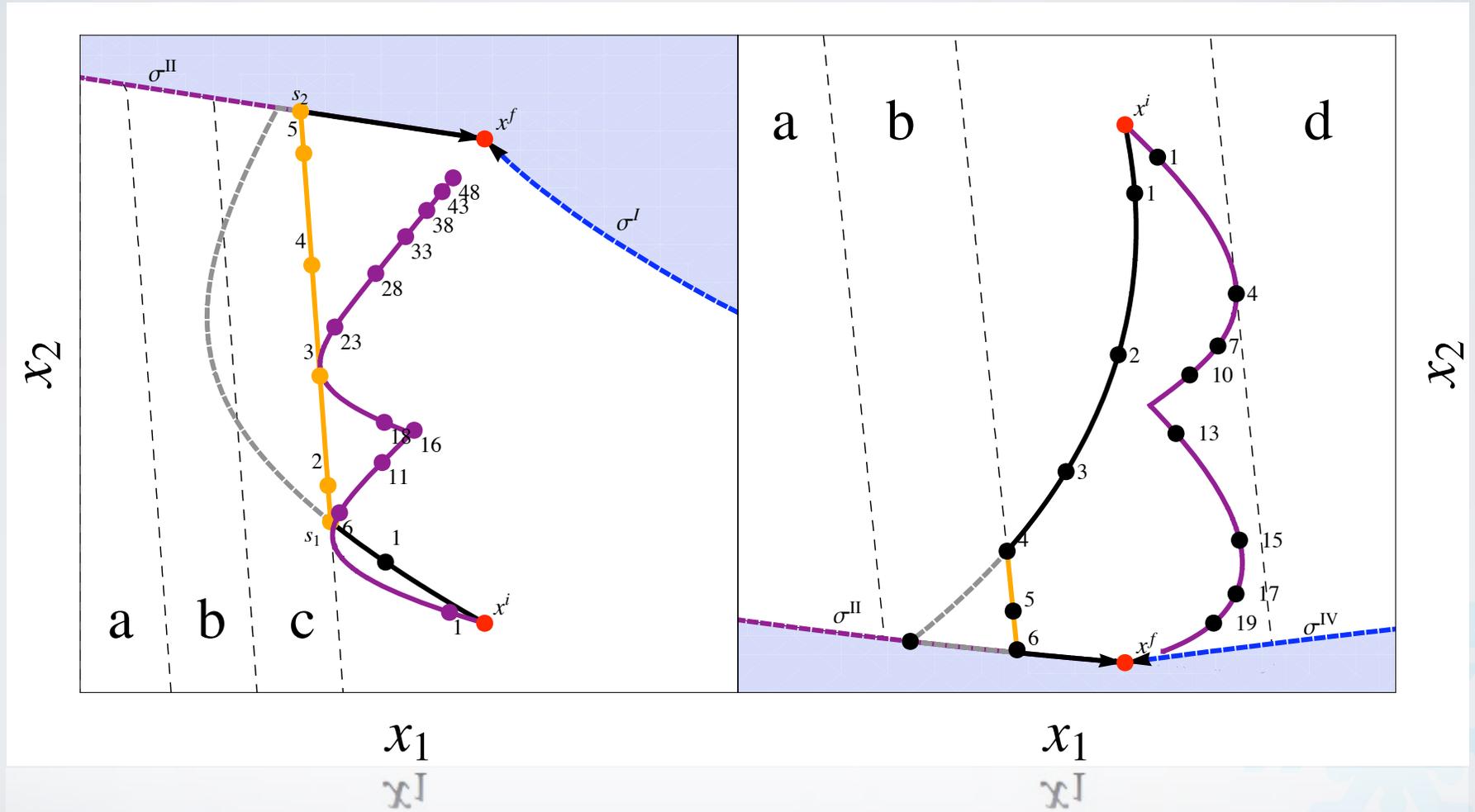
SUFFICIENCY



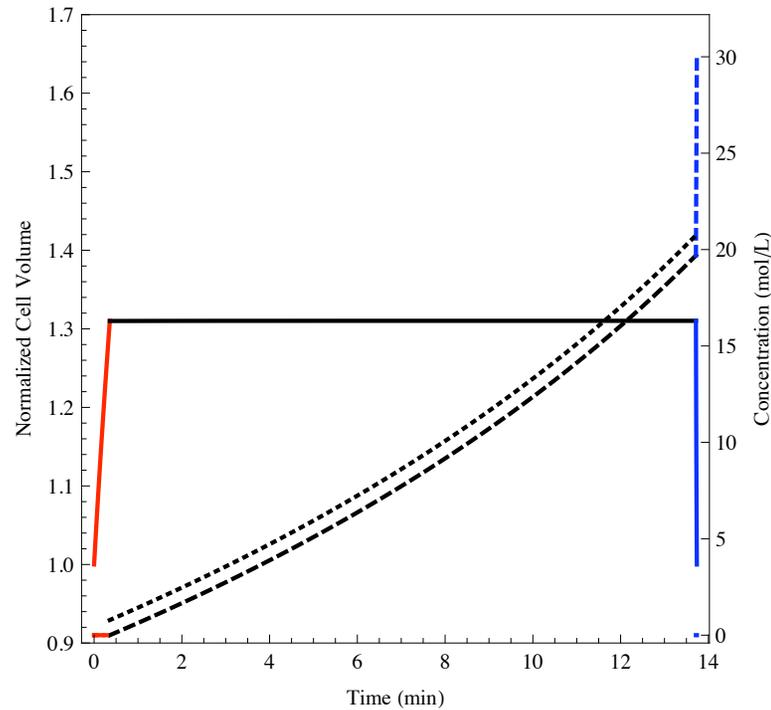
RESULTS

CPA Addition

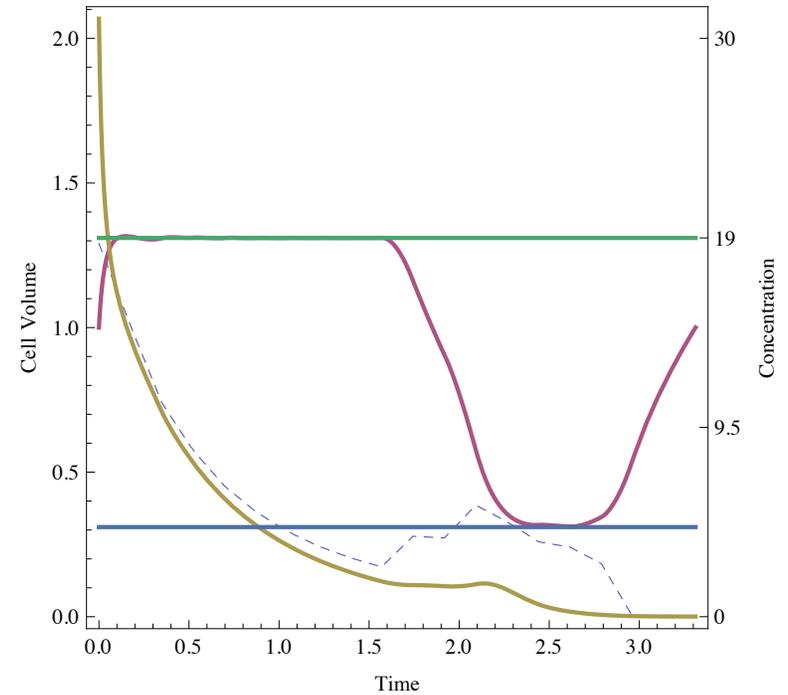
CPA Removal



Addition



Removal



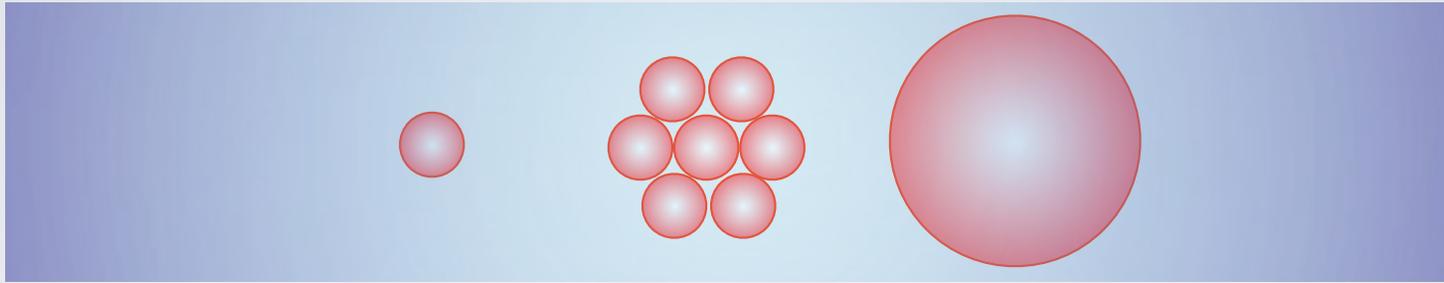
Cost function:

$$J = \int_0^T C_{\text{cell}}(t)^2 dt$$

Solved with a direct method: parametrize system with piecewise linear controls, minimize constrained system with a truncated-Newton approach to the augmented Lagrangian

OPTIMAL CONTROL

System Coupling



All models in cryobiology are coupled systems!

Before cooling

During cooling

**Mass &
Heat**

Parabolic
System

Thermal
convection/mushy
layers/etc...



EXTENSION TO TISSUES, SYSTEMS



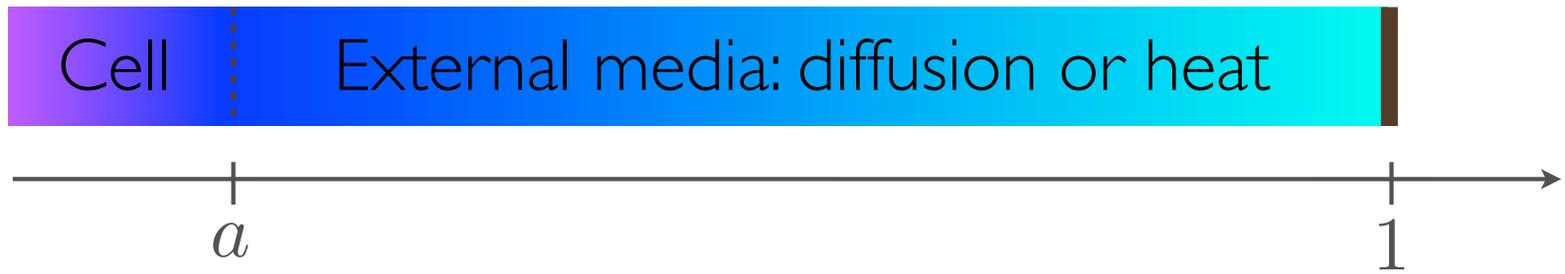
- Find $c^e(t)$ such that $f(t)$ approximates the desired independent control $u_D(t)$, known a priori
- Can use “inverse problem” techniques to solve analytically
- This gives a tool to develop numerical schemes for completely novel optimal control problems



*A Carasso, *SIAM J App. Anal.* 198



In Media	$c_t = D \left(c_{rr} + \frac{2}{r} c_r \right) \quad (r, t) \in (a, 1) \times [0, \infty),$ $\frac{\partial c}{\partial r} = k(c^c - c), \quad (r, t) \in \{a\} \times [0, \infty),$ $c = c^e, \quad (r, t) \in \{1\} \times [0, \infty),$ $c = c_0, \quad (r, t) \in [0, 1] \times \{a\},$
In Cell	$\dot{x} = h(x, c(a, t))$ $c^c = \mathcal{M}c(a, t)$ $\mathcal{M} = M_1 M_2, \quad M_1 : (x, y) \rightarrow x/y,$ $d(M_2 c(a, t))/dt = h(M_2 c(a, t), c(a, t))$



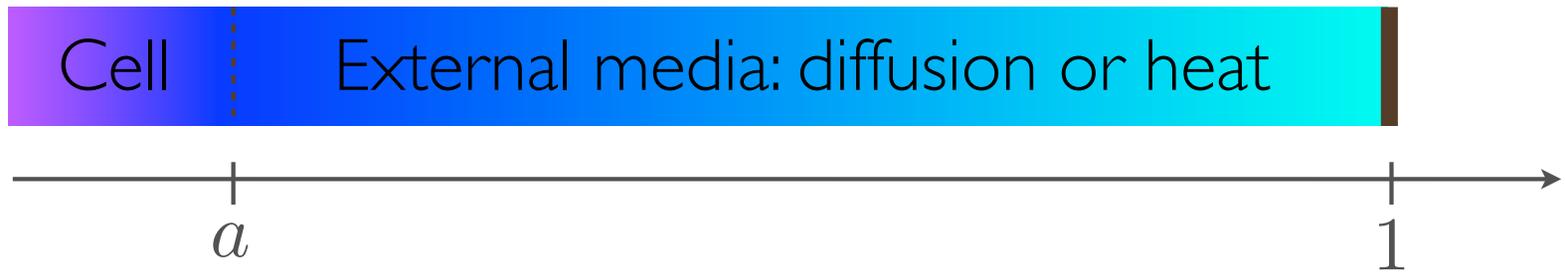
After Laplace transform, we may solve for

$$\bar{f}(s) = \bar{h}_1(s)\bar{c}^e(s) + \bar{h}_2(s)\bar{c}^c(s)$$

where $\bar{h}_1(s)$, $\bar{h}_2(s)$ are modified spherical Bessel functions*. Thus,

$$\begin{aligned} f(t) &= \int_0^t c^e(\tau)h_1(t-\tau) + c^c(\tau)h_2(t-\tau) d\tau, \\ &:= K_1c^e + K_2c^c. \end{aligned}$$

Define $K = K_1[I - K_2M_2]^{-1}$.



Lemma: \mathcal{M} , M_2 exist and are bounded.

Lemma*: K_1 and K_2 are compact linear operators with zero spectral radius and unbounded inverse.

Formally:

$$\begin{aligned} c^e &= K_1^{-1} K_2 c^c + K_1^{-1} f \\ &= K_1^{-1} (K_2 \mathcal{M} + I) f \end{aligned}$$

which exists for f with sufficient decay.

*A Carasso, *SIAM J App. Anal.* 1982

Problem

Define $J_1(v) = \{T : |(M_2 v)(T) - x^d| = 0\}$.

Define the cost

$$J(c^e) := T + \epsilon_1 |(M_2 K c^e)(T) - x^d|^2 + \epsilon_2 \|c^e\|^2.$$

Find $\min_{c^e \in \mathcal{A}} J(c^e)$ subject to above PDE-ODE system and with state constraints $\Gamma \cdot x \leq 0, \Gamma \in \mathbb{R}^2$.

Proposition: Let $\epsilon_2 = 0$. Then there exists an ϵ_1 such that

$$\begin{aligned} c_j^e(t) &= K_1^{-1} (K_2 M + I) f \\ &:= K^{-1} f \\ &= \operatorname{argmin} J(c^e) \end{aligned}$$

From above, we recall that f has step changes, and thus the frequency spectrum will not exponentially decay.

Theorem: The PDE-ODE system has no exact optimal controls.

We must use approximate controls.

Define

$$J_2(v) := \|c(a, t) - Kv\|^2 + \left(\frac{\epsilon}{M}\right)^2 \|v\|^2.$$

Theorem: The unique minimizer of J_2 is

$$v(t) = \frac{1}{\sqrt{2\pi}} \int_{-\infty}^{\infty} e^{i\xi t} \frac{\hat{h}_1^{-1}(\xi) \left(\hat{h}_2(\xi) \widehat{M_2 v}(\xi) + \hat{v}(\xi) \right)}{1 + \omega^2 \hat{h}_1^{-1}(\xi) \left(\hat{h}_2(\xi) M_2 \hat{v}(\xi) \right)} d\xi$$

Pf: Solve the overdetermined system

$$Kv = c^p, \quad \omega v := \frac{\epsilon}{M} v = 0,$$

in the frequency domain and take inverse FT.

Define

$$J(c^e, t) := t + \epsilon_1 |(M_2 K c^e)(t) - x^d|^2 + \epsilon_2 \|c^e\|^2.$$

Theorem: Fix $t^f = J_1(f)$. Then there exist ϵ_1 and ϵ_2 and $\omega(\epsilon_1, \epsilon_2, M_2, t^f)$

$$K_\omega^{-1} f = \operatorname{argmin} J(c^e, t^f).$$

Pf:

$$\begin{aligned} |(M_2 K c^e)(t) - x^d|^2 &= |(M_2 K c^e)(t) - (M_2 K K^{-1} f)(t)|^2, \\ &= |(M_2 K c^e)(t) - (M_2 f)(t)|^2, \end{aligned}$$

Since T is fixed, and M_2 and K are bounded, there exists $\epsilon_3 > 0$ (depending on T, M_2), such that

$$\int_0^T |M_2 K c^e(t) - M_2 f(t)|^2 dt \geq \epsilon_3 \|K c^e - f\|_{L^2}^2,$$

and

$$J(c^e, t^f) - t^f \geq \epsilon_1 \epsilon_3 (\|K c^e - f\|^2 + \omega_2 \|c^e\|^2) > 0.$$

$$\begin{aligned} \text{Thus, } \operatorname{argmin} J(c^e, t^f) &= \operatorname{argmin} \|K c^e - f\|^2 + \omega_2 \|c^e\|^2 \\ &= K_{\omega_2}^{-1} f. \end{aligned}$$

Now note that

$$\|M_2\|^2 \|Kc^e - f\|_{L^2}^2 \geq |(M_2Kc^e)(t) - (M_2f)(t)|^2,$$

and thus

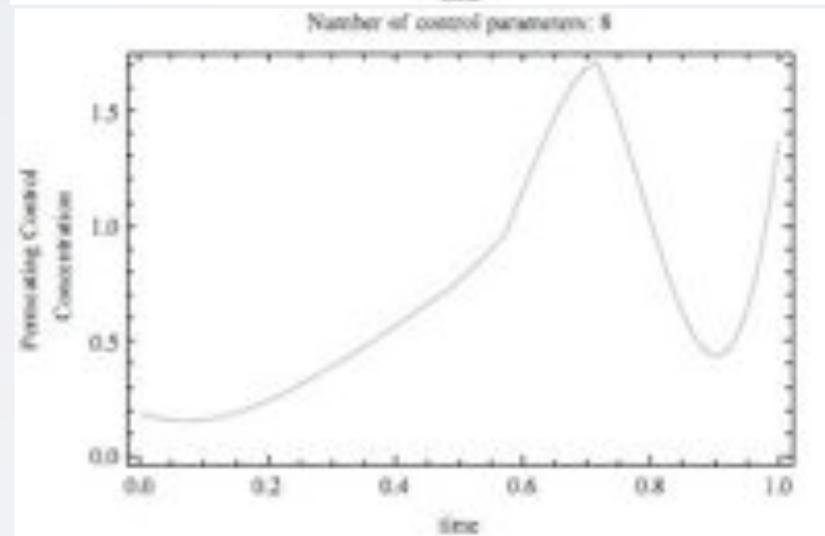
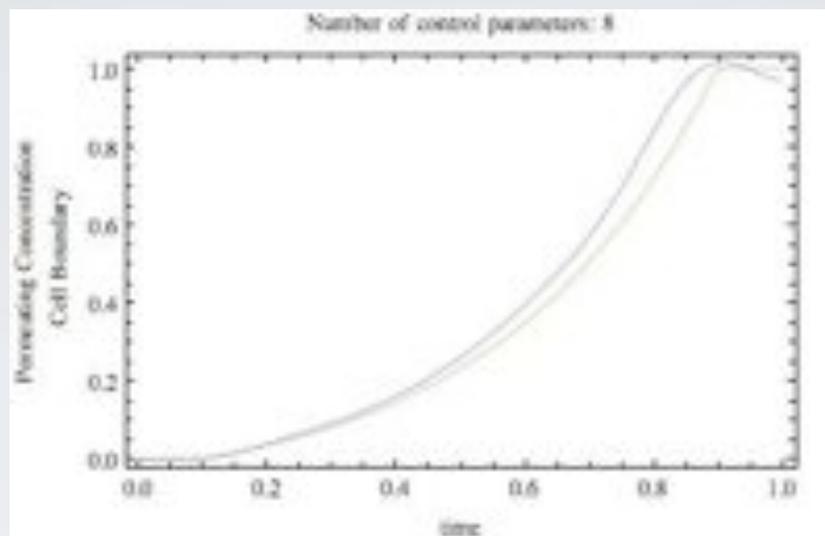
$$\begin{aligned} \|M_2\|^2 \|(KK_\omega^{-1} - I)\|^2 \|f\|_{L^2}^2 &\geq \|M_2\|^2 \|(KK_\omega^{-1} - I)f\|_{L^2}^2 \\ &\geq \|M_2\|^2 \|KK_\omega^{-1}f - f\|_{L^2}^2 \\ &\geq |(M_2KK_\omega^{-1})(t) - (M_2f)(t)|^2, \end{aligned}$$

and with $\|M_2\| \|f\| < k < \infty$, given $\delta(\omega) > 0$ there exists an $\omega > 0$ such that

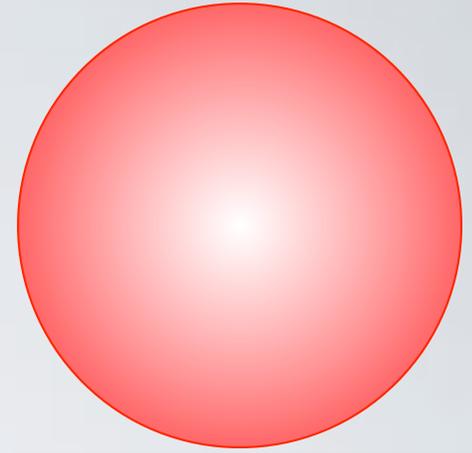
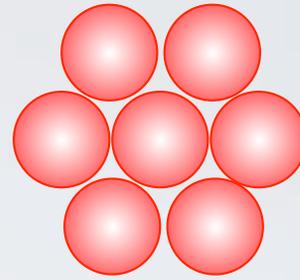
$$\begin{aligned} \delta &\geq \|KK_\omega^{-1} - I\| \\ &\geq k^{-2} |(M_2KK_\omega^{-1})(t) - (M_2f)(t)|^2. \end{aligned}$$

NUMERICS

- PHAML: hp-adaptive multilevel elliptic solver
- Implicit-Filtering minimization algorithm: adaptive secant approximation to gradient



Model scaling shows where future work lies:



MODELS	Mass	ODE System ✓	Hybrid ODE/PDE System ✓	PDE System ✓
	Heat	Stochastic ODE ✓	Large Monte Carlo System ✓	nonlinear heat equation
CONTROLS	Mass	ODE System ✓	Hybrid ODE/PDE System	PDE System
	Heat	Stochastic ODE	Large Monte Carlo System	nonlinear heat equation

CURRENT AND FUTURE PROBLEMS

- Develop cost functions for entire cryo-protocol
- Extend 'inverse' approach to 2D and 3D systems
- Model multiphase ternary solidification and interaction with biomaterials
- Iterative optimization of freezing protocols
- Optimal design of counter-current dialysis devices



QUESTIONS?

