

From Cardiac Cells to Genetic Regulatory Networks

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Joint work with

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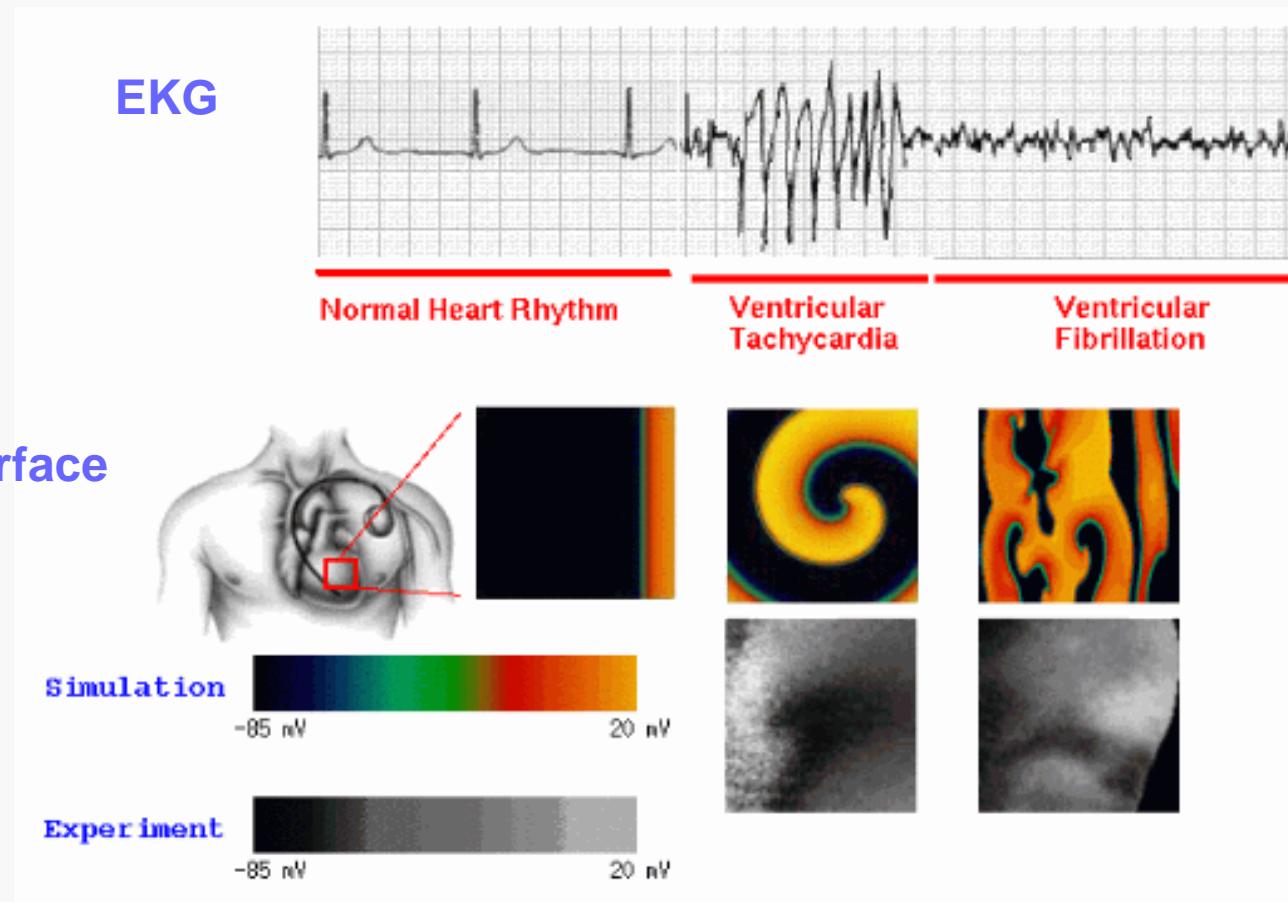


Overview

- **Background**
 - Cardiac Cells, Action Potential, Restitution
- **Biological Switching**
- **Minimal Model**
 - Resistor Model
 - Sigmoid Closure and Conductance Model
- **Piecewise Multi Affine Minimal Model**
 - Optimal Polygonal Approximation
 - Model Comparison
- **Parameter Identification**
 - RoverGene
- **Conclusion**

Background

Emergent Behavior in Heart Cells



Arrhythmia afflicts more than 3 million Americans alone

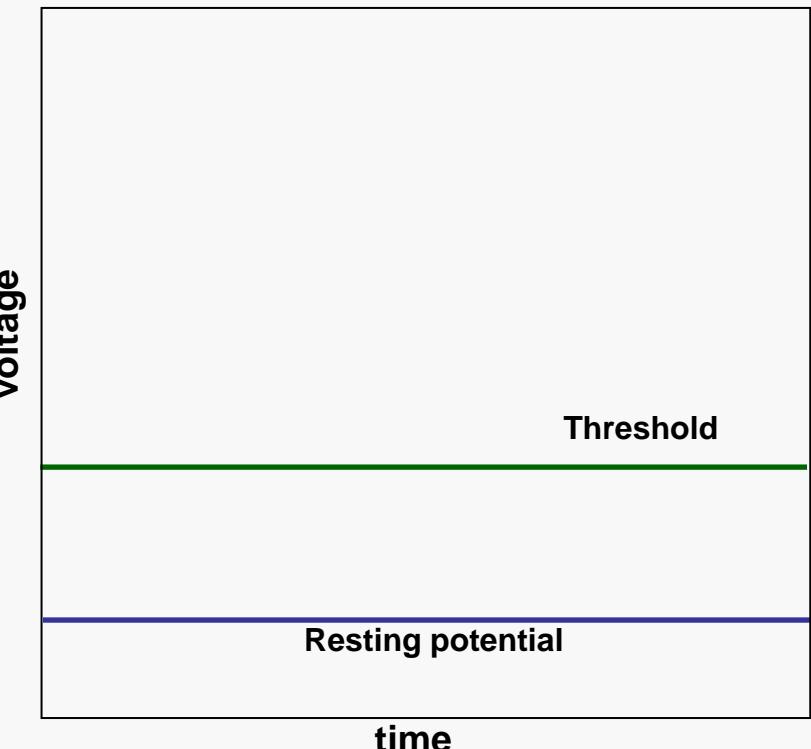


Single Cell Reaction: Action Potential

Membrane's AP depends on:

- **Stimulus (voltage or current):**
 - External / Neighboring cells
- **Cell's state**

Schematic Action Potential



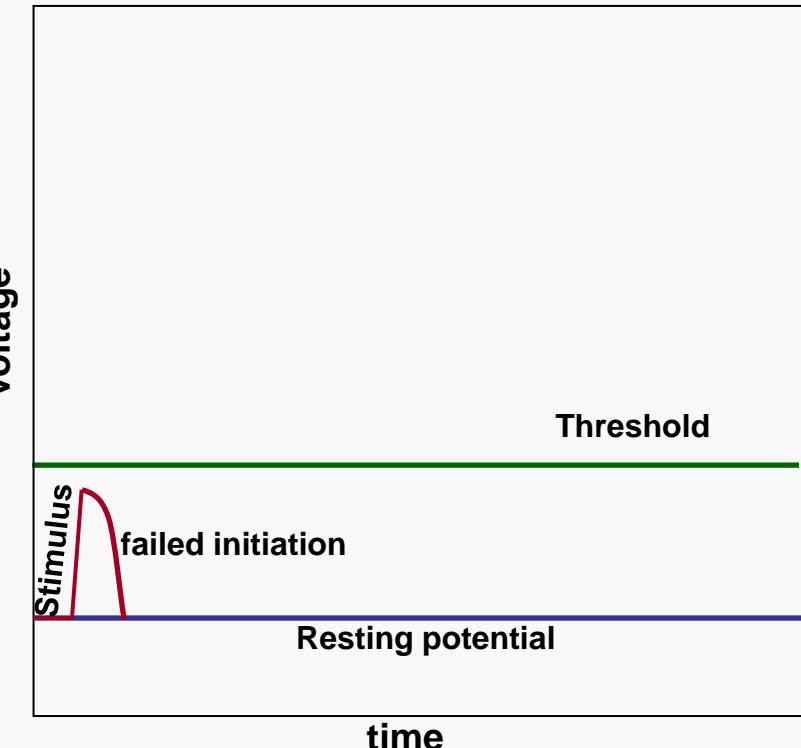


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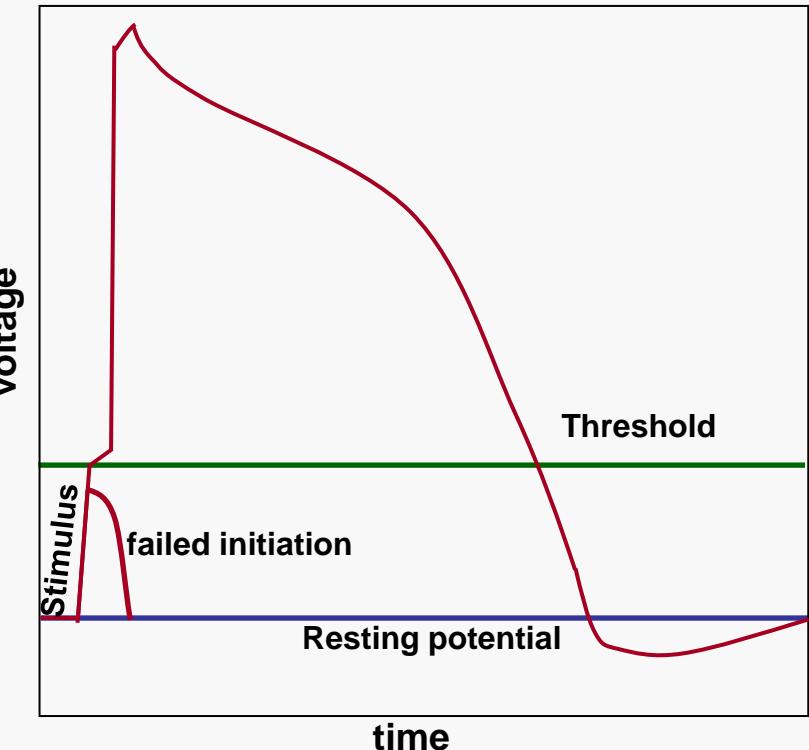


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Single Cell Reaction: Action Potential

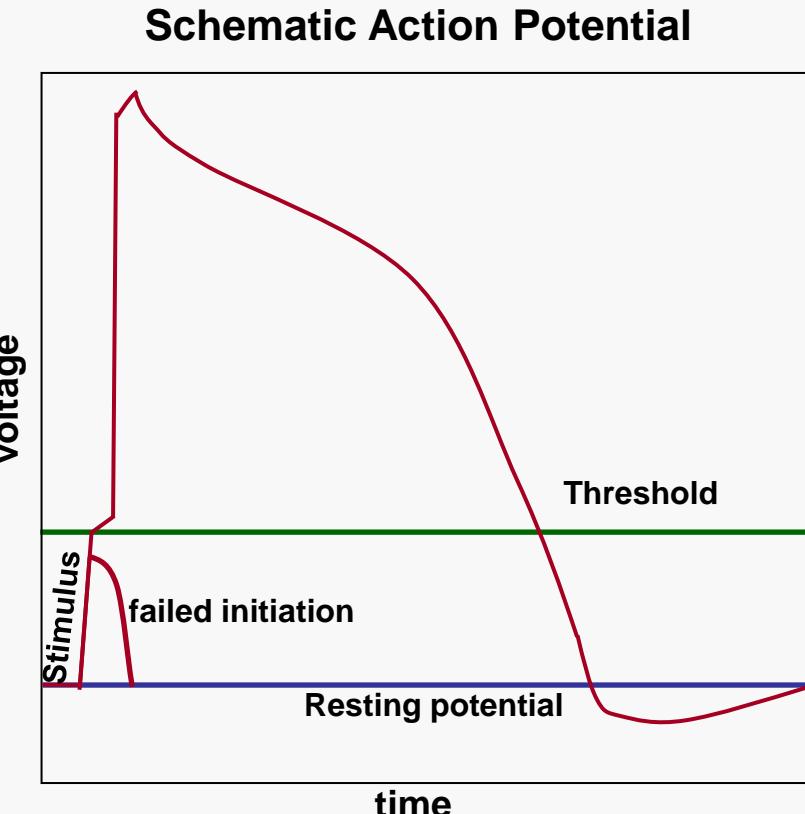
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AP has nonlinear behavior!

- Reaction diffusion system:

$$\frac{\partial \mathbf{u}}{\partial t} = R(\mathbf{u}) + \nabla(D\nabla\mathbf{u})$$





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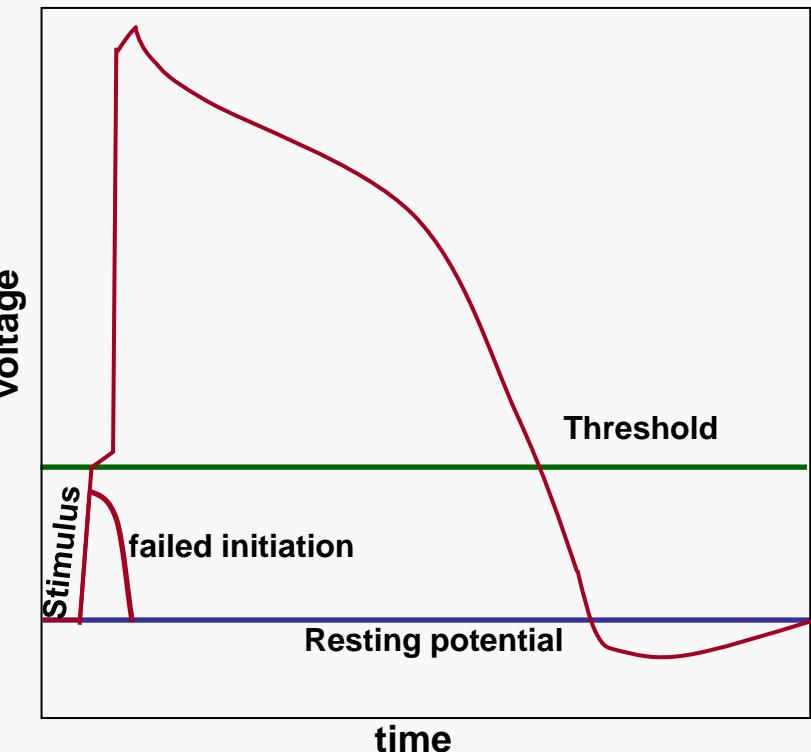
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Schematic Action Potential





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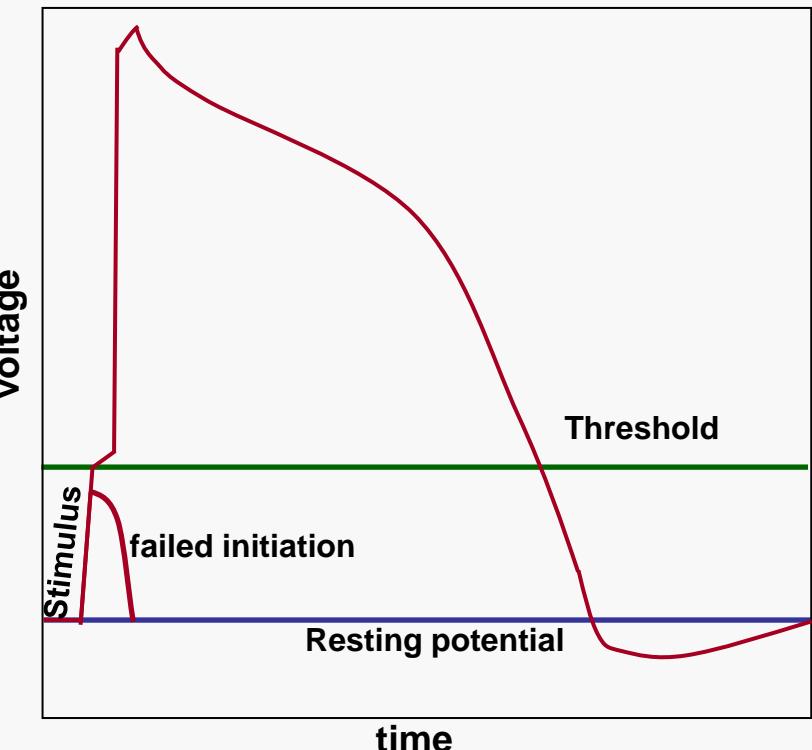
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Schematic Action Potential





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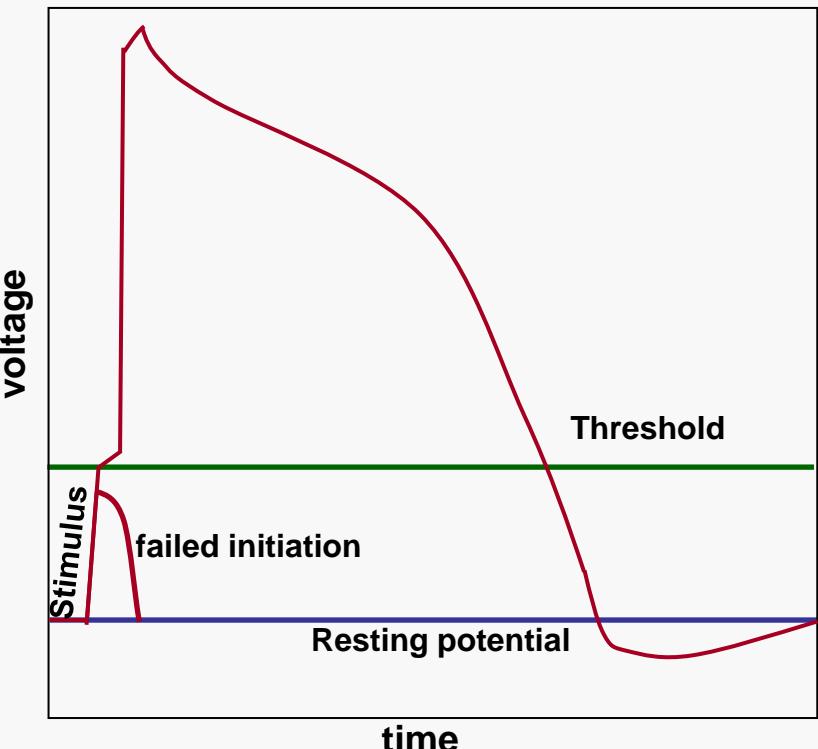
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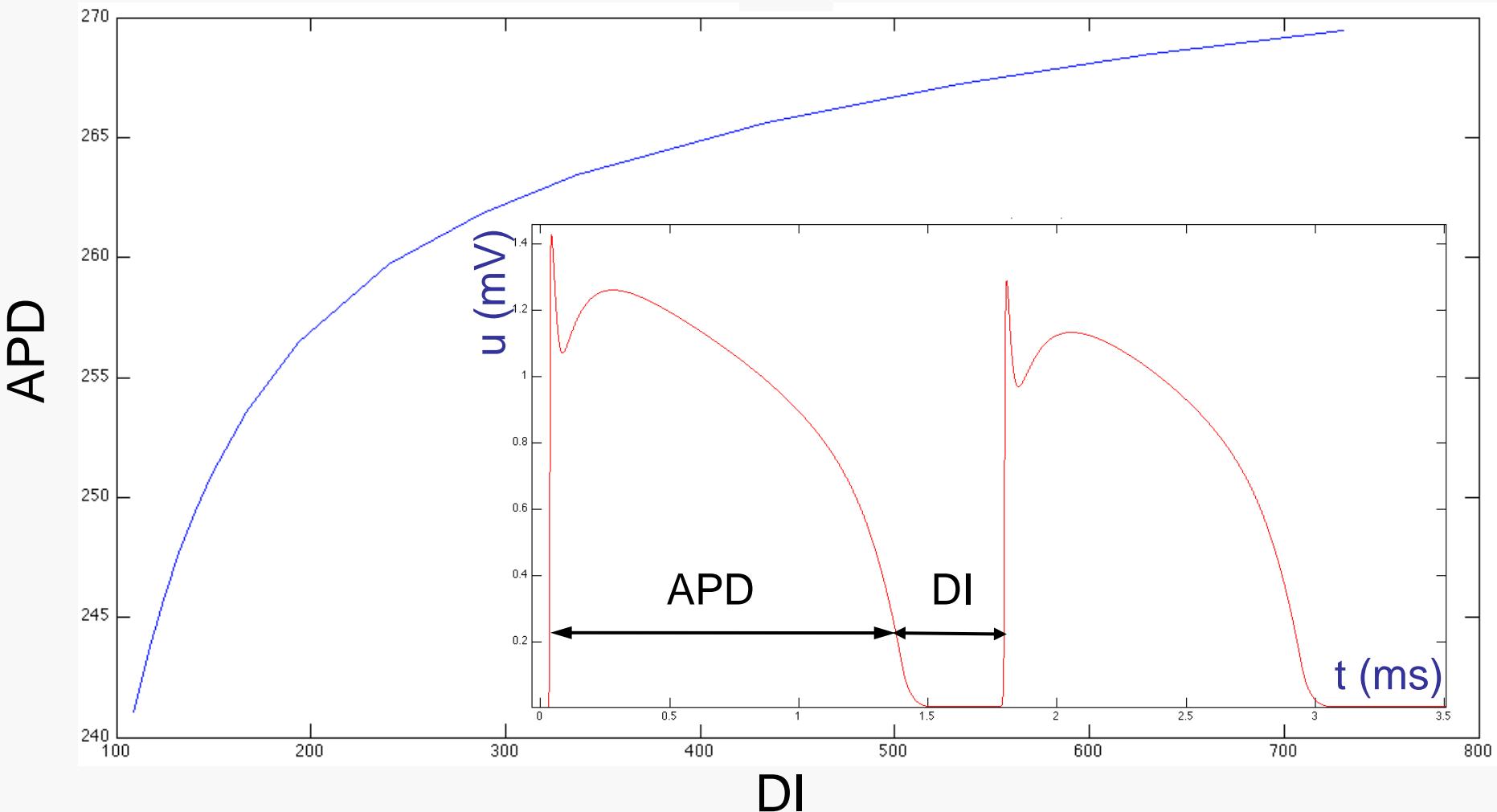


Schematic Action Potential





Frequency Response





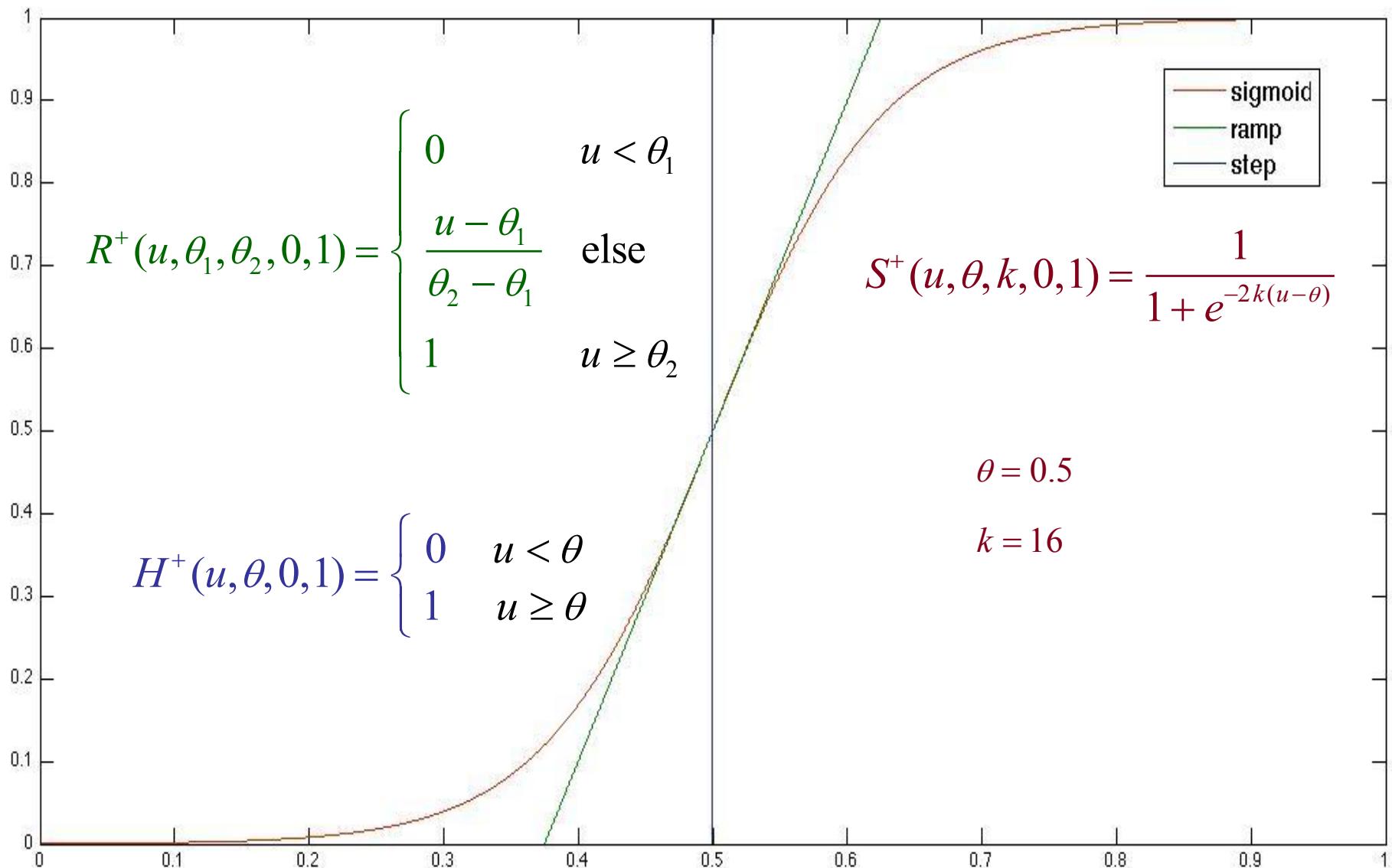
Existing Models

- **Detailed ionic models:**
 - **Luo and Rudi: 14 variables**
 - **Tusher, Noble² and Panfilov: 17 variables**
 - **Priebe and Beuckelman: 22 variables**
 - **Iyer, Mazhari and Winslow: 67 variables**
- **Approximate models:**
 - **Cornell: 3 or 4 variables**
 - **SUNYSB: 2 or 3 variable**

Biological Switching



Biological Switching





Threshold-Based Switching Functions

- Arithmetic Generalization of Boolean predicates $u \leq \theta$:

- Step: $H^+(u, \theta, 0, 1), H(u, \theta, 0, 1) = 1 - H^+(u, \theta, 0, 1)$
- Sigmoid: $S^+(u, \theta, k, 0, 1), S(u, \theta, k, 0, 1) = 1 - S^+(u, \theta, k, 0, 1)$
- Ramp: $R^+(u, \theta_1, \theta_2, 0, 1), R(u, \theta_1, \theta_2, 0, 1) = 1 - R^+(u, \theta_1, \theta_2, 0, 1)$

- Boolean algebra generalizes to probability algebra:

$$\sim(u \leq \theta) : H(u, \theta, 0, 1) = 1 - H^+(u, \theta, 0, 1)$$

$$(u \leq \theta_1) \& (v \leq \theta_2) : H^+(u, \theta_1, 0, 1) * H^+(v, \theta_2, 0, 1)$$

$$(u \leq \theta_1) \mid (u \leq \theta_2) : H^+(u, \theta_1, 0, 1) + H^+(v, \theta_2, 0, 1) - H^+(u, \theta_1, 0, 1) * H^+(v, \theta_2, 0, 1)$$

- Generalization: $H^\pm(u, \theta, u_m, u_M), S^\pm(u, \theta, k, u_m, u_M), R^\pm(u, \theta_1, \theta_2, u_m, u_M)$

$$S^\pm(u, \theta, k, u_m, u_M) = u_m + (u_M - u_m)S^+(u, k, \theta)$$



Gene Regulatory Networks (GRN)

- GRNs have the following general form:

$$\dot{x}_i = \sum_{m=1}^{m_i} \prod_{n=1}^{n_m} a_{mn} s^\pm(x_{mn}, \theta_{mn}, k_{mn}, u_{mn}, v_{mn}) - b_i x_i$$

where:

a_{mn} : are activation / inhibition constants

b_i : are decay constants

$s^\pm(..)$: are possibly complemented sigmoidal functions

- Note: steps and ramps are sigmoid approximations

Minimal Model



Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) - (J_{fi} + J_{si} + J_{so})$$

$$J_{fi} = -H^+(u, \theta_v, 0, 1) (u - \theta_v)(u_u - u)v / \tau_{fi}$$

$$J_{si} = -H^+(u, \theta_w, 0, 1) ws / \tau_{si}$$

$$J_{so} = H^-(u, \theta_w, 0, 1) u / \tau_o + H^+(u, \theta_w, 0, 1) / \tau_{so}$$

$$\dot{v} = H^-(u, \theta_v, 0, 1) (v_\infty - v) / \tau_v^- - H^+(u, \theta_v, 0, 1)v / \tau_v^+$$

$$\dot{w} = H^-(u, \theta_w, 0, 1)(w_\infty - w) / \tau_w^- - H^+(u, \theta_w, 0, 1)w / \tau_w^+$$

$$\dot{s} = (S^+(u, u_s, k_s, 0, 1) - s) / \tau_s$$

$$J_{fi} = -H(u - \theta_v)(u - \theta_v)(u_u - u)v / \tau_{fi}$$



Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) - (J_{fi} + J_{si} + J_{so})$$

voltage

$$= -H^+(u, \theta_v, 0, 1) (u - \theta_v)(u_u - u)v / \tau_{fi}$$

voltage

$$= -H^+(u, \theta_w, 0, 1) ws / \tau_{si}$$

J_{so}

$$= H^-(u, \theta_w, 0, 1) u / \tau_o + H^+(u, \theta_w, 0, 1) / \tau_{so}$$

$$\dot{v} = H^-(u, \theta_v, 0, 1) (v_\infty - v) / \tau_v^- - H^+(u, \theta_v, 0, 1) v / \tau_v^+$$

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Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) - (J_{fi} + J_{si} + J_{so})$$

$$J_{fi} = -H^+(u, \theta_v, 0, 1)(u - \theta_v)(u_u - u)v / \tau_{fi}$$
$$J_{si} = H^-(u, \theta_w, 0, 1)w / \tau_{si}$$

$$J_{so} = H^-(u, \theta_w, 0, 1)u / \tau_o + H^+(u, \theta_w, 0, 1) / \tau_{so}$$

$$\dot{v} = H^-(u, \theta_v, 0, 1)(v_\infty - v) / \tau_v^- - H^+(u, \theta_v, 0, 1)v / \tau_v^+$$

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Cornell's Minimal Resistor Model

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$$J_{si} = -H^+(u, \theta_w, 0, 1)w / \tau_o$$

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Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) - (J_{fi} + J_{si} + J_{so})$$

$$J_{fi} = -H^+(u, \theta_v, 0, 1)(u - \theta_v)v / \tau_{fi}$$

$$J_{si} = -H^+(u, \theta_w, 0)$$

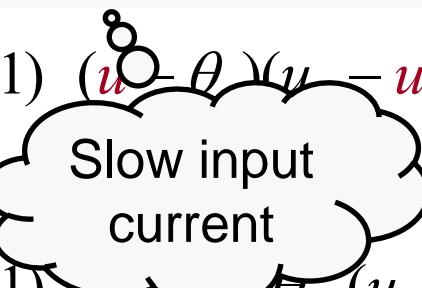
$$J_{so} = H^-(u, \theta_w, 0, 1)H(u, \theta_w, 0, 1) / \tau_{so}$$

$$\dot{v} = H^-(u, \theta_v, 0, 1)(v_\infty - v) / \tau_v^- - H^+(u, \theta_v, 0, 1)v / \tau_v^+$$

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Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) - (J_{fi} + J_{si} + J_{so})$$

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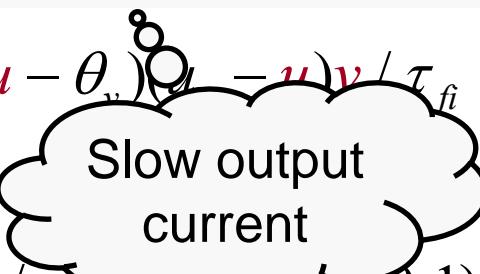
$$J_{so} = H^-(u, \theta_w, 0, 1) u / \tau_o - H^-(u, \theta_w, 0, 1) / \tau_{so}$$

$$\dot{v} = H^-(u, \theta_v, 0, 1) (v_\infty - v) / \tau_v^- - H^+(u, \theta_v, 0, 1) v / \tau_v^+$$

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Cornell's Minimal Resistor Model


$$\dot{u} = \nabla(D\nabla u) - (J_{fi} + J_{si})$$
$$J_{fi} = -H^+(u, \theta_v, 0, 1) (u - \theta_v)(u_u - u)v / \tau_{fi}$$
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Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) - (J_f)$$

Heaviside
(step)

$$J_{fi} = -H^+(u, \theta_v, 0, 1) (u - \theta_v)(u_u - u)v / \tau_{fi}$$

$$J_{si} = -H^+(u, \theta_w, 0, 1) ws / \tau_{si}$$

$$J_{so} = H^-(u, \theta_w, 0, 1) u / \tau_o + H^+(u, \theta_w, 0, 1) / \tau_{so}$$

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Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) - (J_{fi} + J_{si} + J_{so})$$

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Fast input
Gate



Cornell's Minimal Resistor Model

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Resistance
Time Cst



Cornell's Minimal Resistor Model

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$$J_{fi} = -H(u - \theta_v)(u - \theta_v)(u_u - u)v / \tau_{fi}$$

Piecewise
Nonlinear



Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) \quad \text{Slow Input Gate}$$

$$J_{fi} = -H^+(u, \theta_v, 0, 1)(u_u - u)v / \tau_{fi}$$

$$J_{si} = -H^+(u, \theta_w, 0, 1)ws / \tau_{si}$$

$$J_{so} = H^-(u, \theta_w, 0, 1)u / \tau_o + H^+(u, \theta_w, 0, 1)/\tau_{so}$$

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$$J_{fi} = -H(u - \theta_v)(u - \theta_v)(u_u - u)v / \tau_{fi}$$



Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) - (J_{fi} + J_{si} + \text{Slow Output Gate})$$
$$J_{fi} = -H^+(u, \theta_v, 0, 1) (u - \theta_u) v / \tau_{fi}$$
$$J_{si} = -H^+(u, \theta_w, 0, 1) ws / \tau_{si}$$
$$J_{so} = H^-(u, \theta_w, 0, 1) u / \tau_o + H^+(u, \theta_w, 0, 1) / \tau_{so}$$
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Cornell's Minimal Resistor Model

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$$J_{si} = -H^+(u, \theta_w, 0, 1) ws / \tau_{si}$$

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$$J_{fi} = -H(u - \theta_v)(u - \theta_v)(u_u - u)v / \tau_{fi}$$

Piecewise
Bilinear



Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) - (J_{fi} + J_{si} + J_{so})$$

$$J_{fi} = -H^+(u, \theta_v, 0, 1) (u - \theta_v)(u_u - u)v / \tau_{fi}$$

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Piecewise
Nonlinear



Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) - (J_{fi} + J_{si} + J_{so})$$

$$J_{fi} = -H^+(u, \theta_v, 0, 1) (u - \theta_v)(u_u - u)v / \tau_{fi}$$

$$J_{si} = -H^+(u, \theta_w, 0, 1) ws / \tau_{si}$$

$$J_{so} = H^-(u, \theta_w, 0, 1) u / \tau_o + H^+(u, \theta_w, 0, 1) / \tau_{so}$$

$$\dot{v} = H^-(u, \theta_v, 0, 1) (v_\infty - v) / \tau_v^- - H^+(u, \theta_v, 0, 1) v / \tau_v^+$$

$$\dot{w} = H^-(u, \theta_w, 0, 1) (w_\infty - w) / \tau_w^- - H^+(u, \theta_w, 0, 1) w / \tau_w^+$$

$$\dot{s} = (S^+(u, u_s, k_s, 0, 1) - s) / \tau_s$$

$$J_{fi} = -H(u - \theta_v)(u - \theta_v)(u_u - u)v / \tau_{fi}$$



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$$J_{fi} = -H(u - \theta_v)(u - \theta_v)(u_u - u)v / \tau_{fi}$$

Piecewise
Linear



Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) - (J_{fi} + J_{si} + J_{so})$$

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$$J_{so} = H^-(u, \theta_w, 0, 1) u / \tau_o + H^+(u, \theta_w, 0, 1) / \tau_{so}$$

$$\dot{v} = H^-(u - v) / \tau_v^- - H^+(u, \theta_v, 0, 1)v / \tau_v^+$$

$$ws = H^-(u - w) / \tau_w^- / \tau_w^+$$

$$s = (S^+(u, u_s, k_s, 0, 1) - s) / \tau_s^+$$

$$J_{fi} = -H(u - \theta_v)(u - \theta_v)(u_u - u)v / \tau_{fi}$$



Cornell's Minimal Resistor Model

$$\dot{u} = \nabla(D\nabla u) - (J_{fi} + J_{si} + J_{so})$$

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Voltage-controlled resistances

$$\tau_v^- = H^+(u, \theta_v^-, \tau_{v1}^-, \tau_{v2}^-)$$

$$\tau_o^- = H^-(u, \theta_v^-, \tau_{o2}^-, \tau_{o1}^-)$$

$$\tau_s^- = H^+(u, \theta_w, \tau_{s1}^-, \tau_{s2}^-)$$

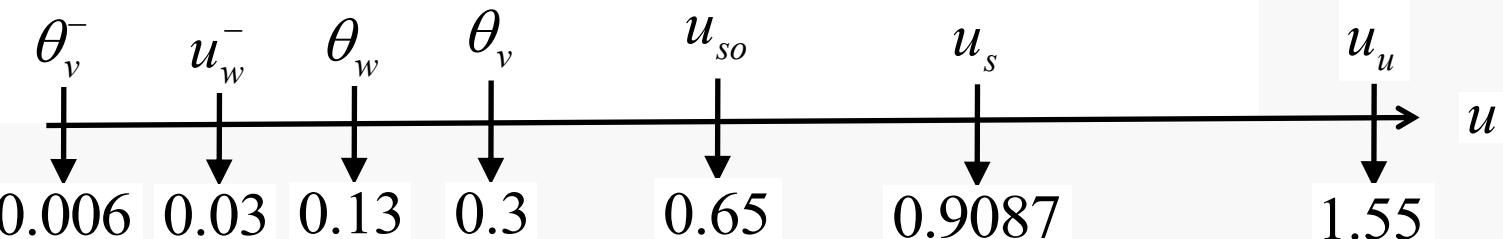
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$$\tau_{so}^- = S^+(u, k_{so}^-, u_{so}^-, \tau_{so1}^-, \tau_{so2}^-)$$

•••
Piecewise
Constant

$$v_\infty = h^-(u, \theta_v^-, 0, 1)$$

$$w_\infty = h^-(u, \theta_v^-, 0, 1) (1 - u / \tau_{w\infty}) + h^+(u, \theta_v^-, 0, w_\infty^*)$$





Voltage-controlled resistances

$$\tau_v^- = H^+(u, \theta_v^-, \tau_{v1}^-, \tau_{v2}^-)$$

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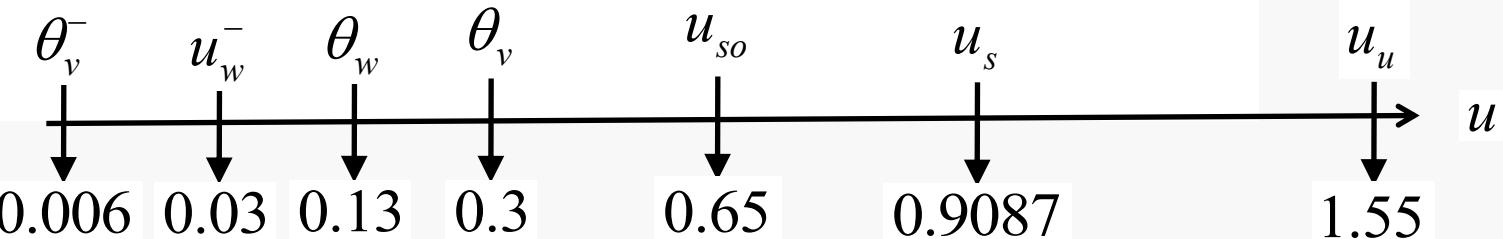
$$\tau_{so}^- = S^+(u, k_{so}^-, u_{so}^-, \tau_{so1}^-, \tau_{so2}^-)$$

Piecewise Constant

Sigmoidal

$$v_\infty = h^-(u, \theta_v^-, 0, 1)$$

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Voltage-controlled resistances

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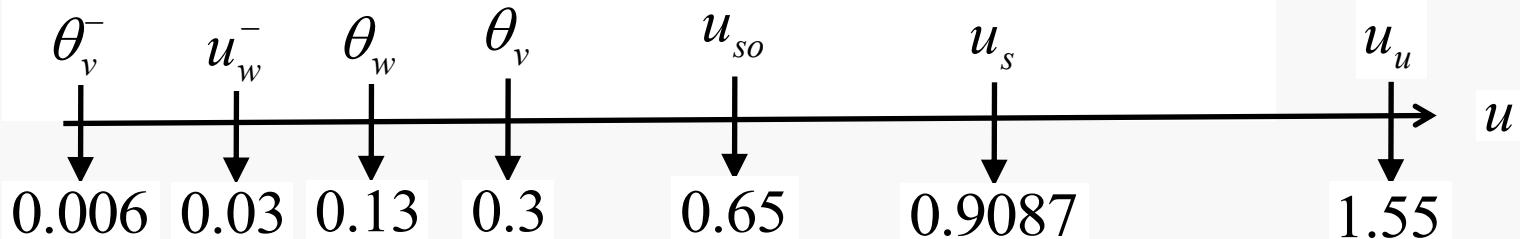
Piecewise Constant

Sigmoidal

Piecewise Linear

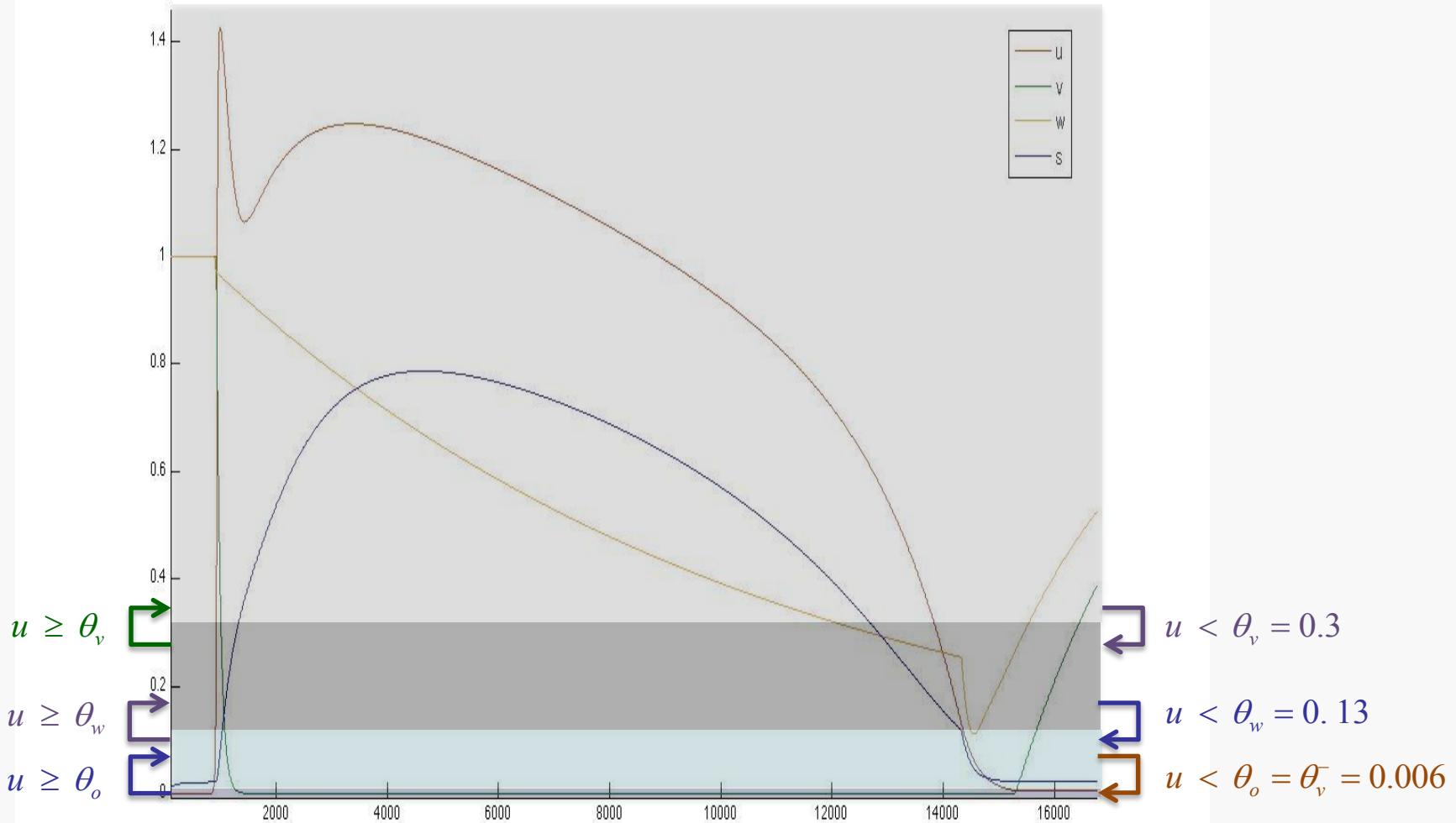
$$v_\infty = h^-(u, \theta_v^-, 0, 1)$$

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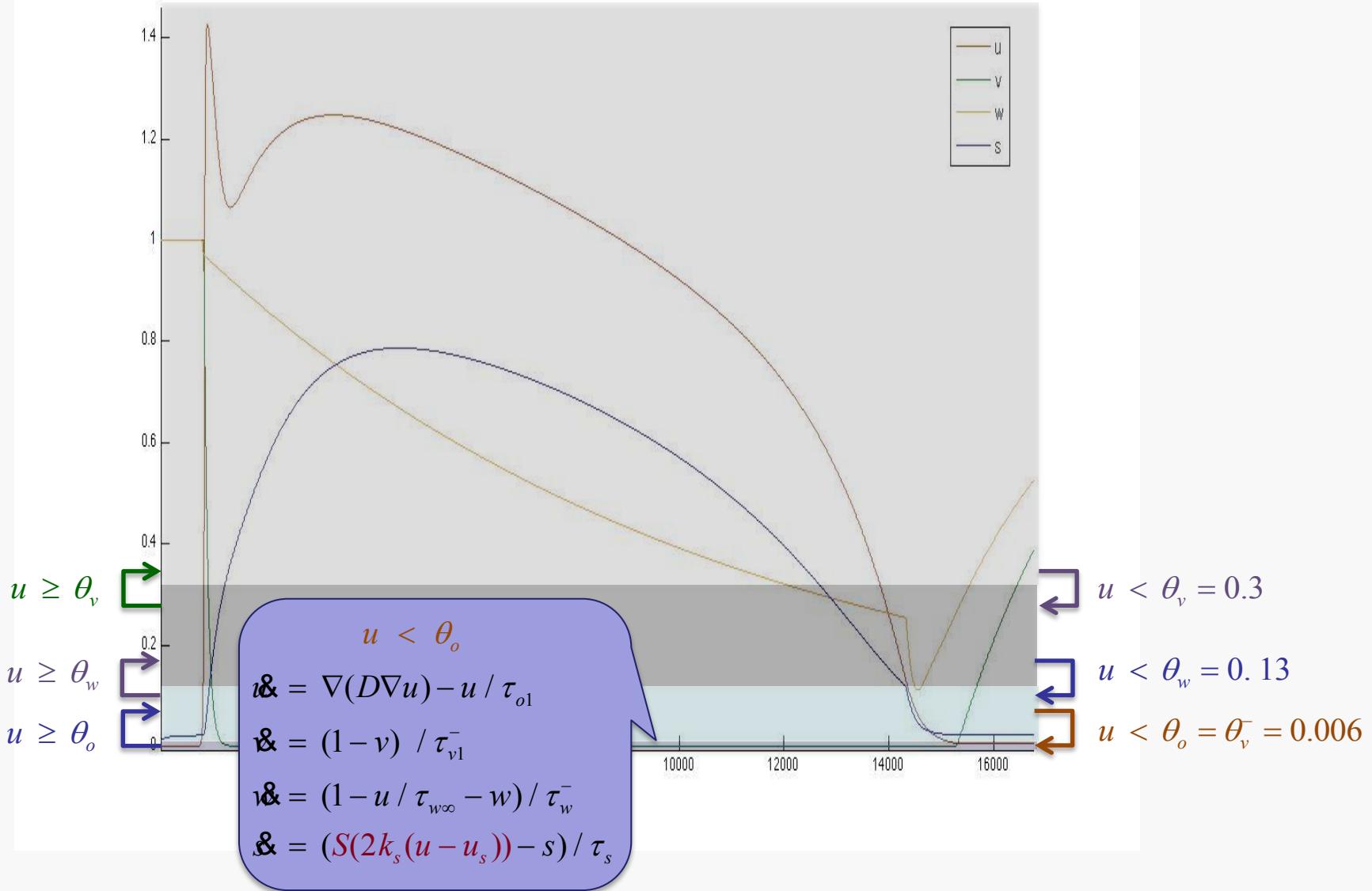




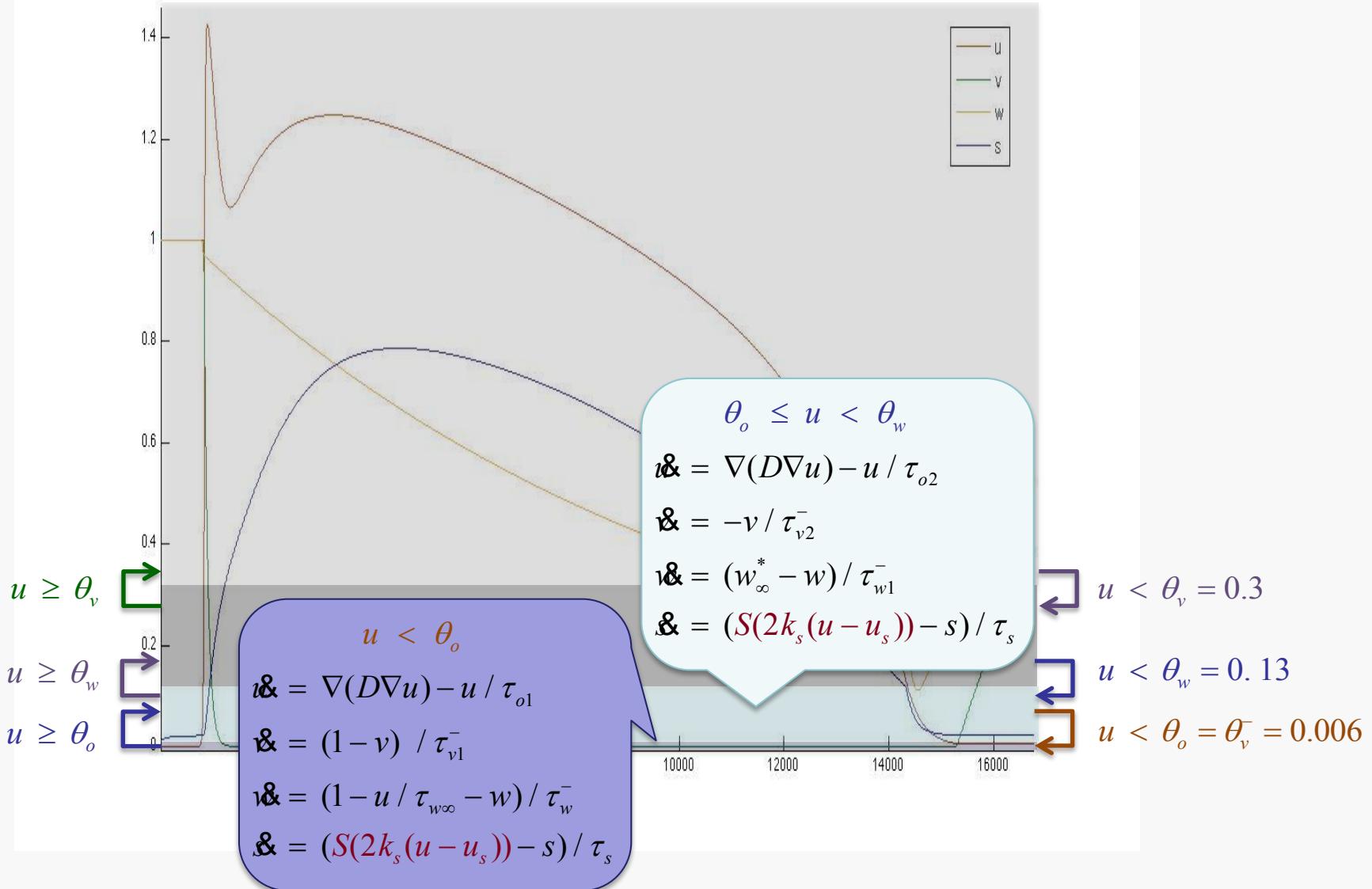
Cornell's Minimal Resistance Model



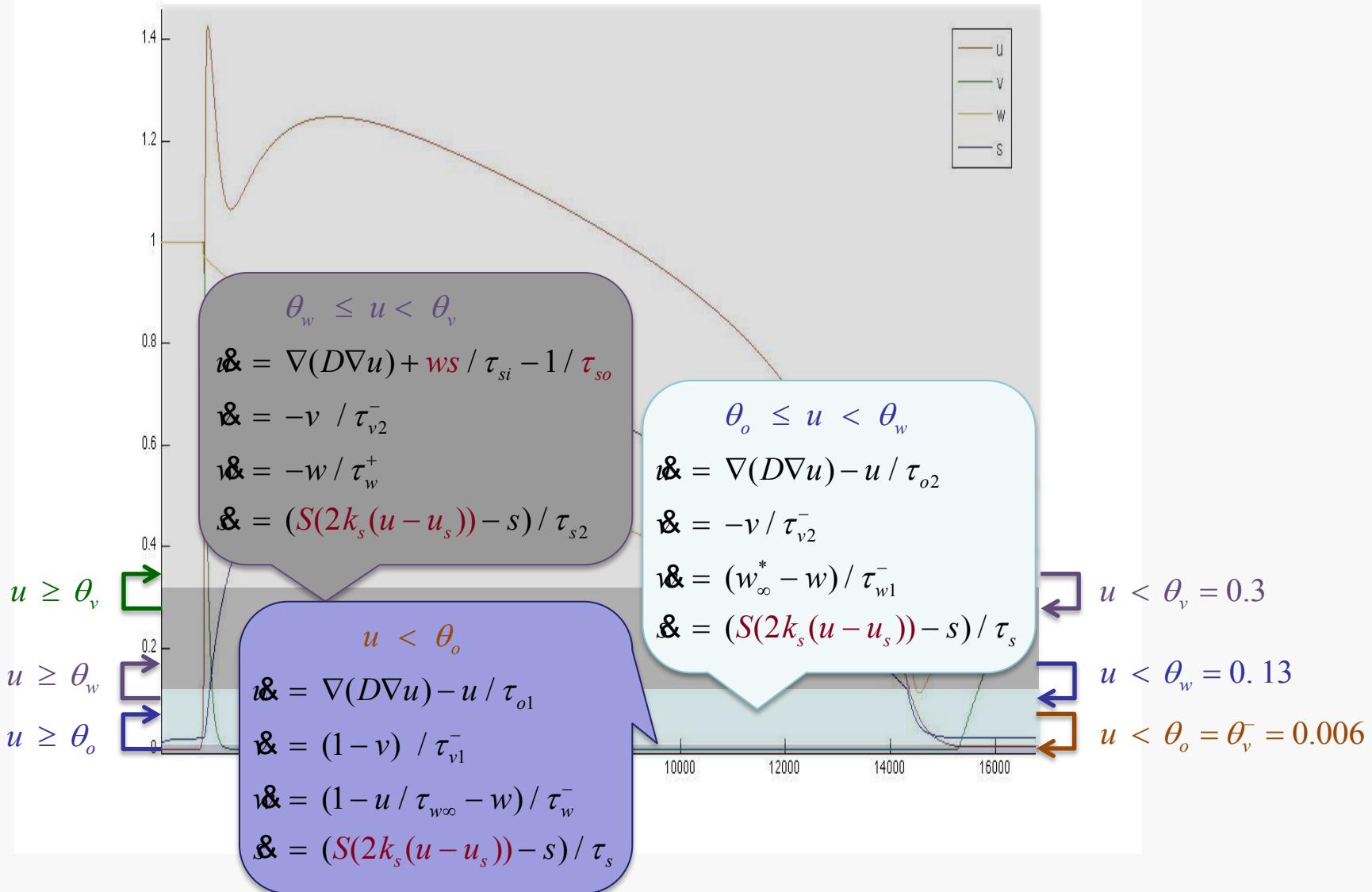
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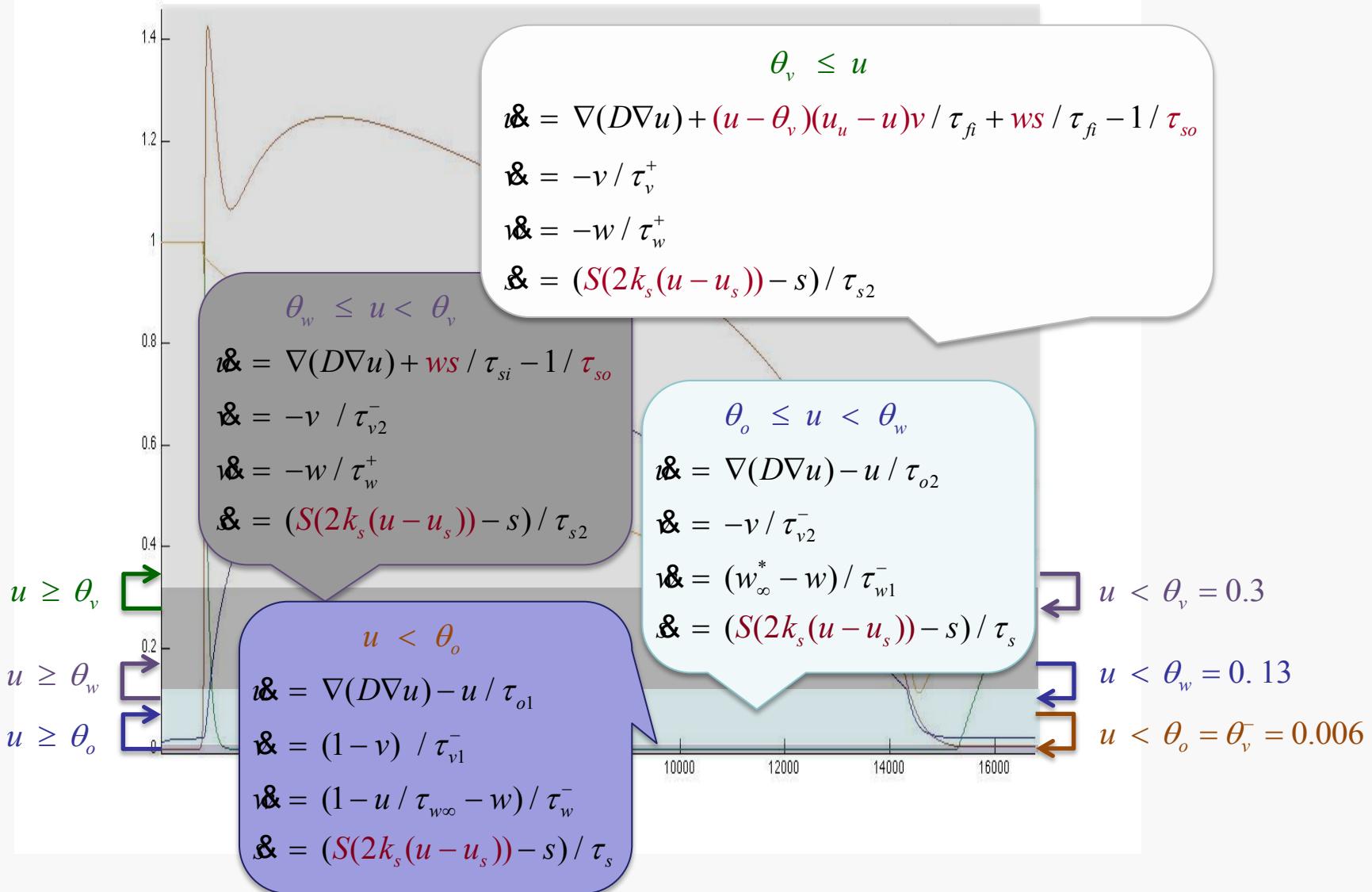
Cornell's Minimal Resistance Model



Cornell's Minimal Resistance Model



Cornell's Minimal Resistance Model





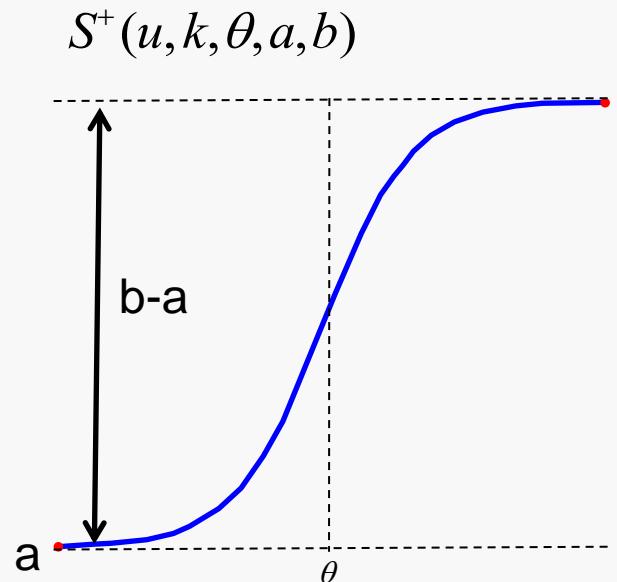
Sigmoid Closure

For $ab > 0$, scaled sigmoids are closed under multiplicative inverses (division):

$$S^+(u, k, \theta, a, b)^{-1} = S^-(u, k, \theta + \ln(a/b)/2k, b^-, a^-)$$

Proof

$$\begin{aligned} S^+(u, k, \theta, a, b)^{-1} &= \frac{1}{a + \frac{b-a}{1+e^{-2k(u-\theta)}}} = \frac{1+e^{-2k(u-\theta)}}{b+ae^{-2k(u-\theta)}} = \\ &= \frac{1}{a} \times \frac{a-b+b+ae^{-2k(u-\theta)}}{b+ae^{-2k(u-\theta)}} = \frac{1}{a} - \frac{\frac{1}{a}-\frac{1}{b}}{1+\frac{a}{b}e^{-2k(u-\theta)}} = \\ &= \frac{1}{a} - \frac{\frac{1}{a}-\frac{1}{b}}{1+e^{-2k(u-(\theta+\frac{\ln a-\ln b}{2k}))}} = S^-(u, k, \theta + \frac{\ln \frac{a}{b}}{2k}, \frac{1}{b}, \frac{1}{a}) \end{aligned}$$





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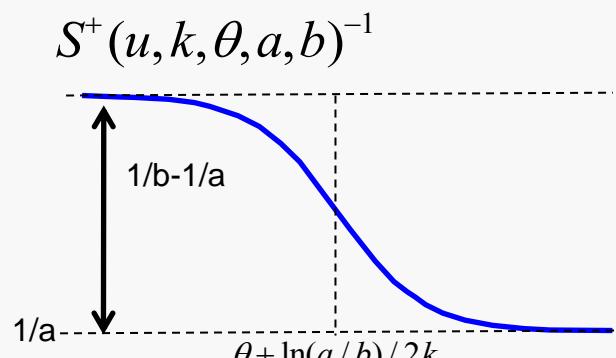
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Resistances vs Conductances

Removing Divisions using Sigmoid Closure

$$\tau_w^- = S^-(u, k_w^-, u_w^-, \tau_{w1}^-, \tau_{w2}^-) \quad g_w^- = 1 / \tau_w^- = S^+(u, k_w^-, u'_w^-, g_{w1}^-, g_{w2}^-)$$

$$\tau_{so}^- = S^-(u, k_{so}^-, u_{so}^-, \tau_{s01}^-, \tau_{s02}^-) \quad g_{so}^- = 1 / \tau_{s0}^- = S^+(u, k_{so}^-, u'_{so}^-, g_{s01}^-, g_{s02}^-)$$

Removing Divisions using $H^+(u, \theta, a, b)^{-1} = H^-(u, \theta, b^{-1}, a^{-1})$

$$\tau_v^- = H^+(u, \theta_v^-, \tau_{v1}^-, \tau_{v2}^-)$$

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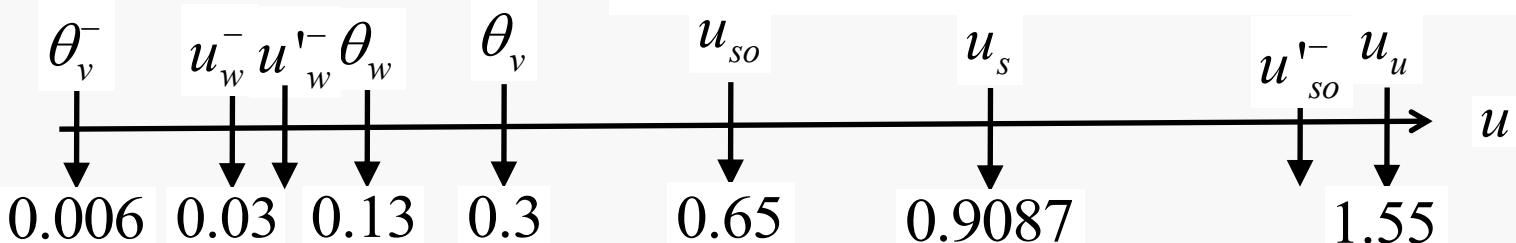
$$g_o^- = 1 / \tau_o^- = H^+(u, \theta_v^-, g_{o1}^-, g_{o2}^-)$$

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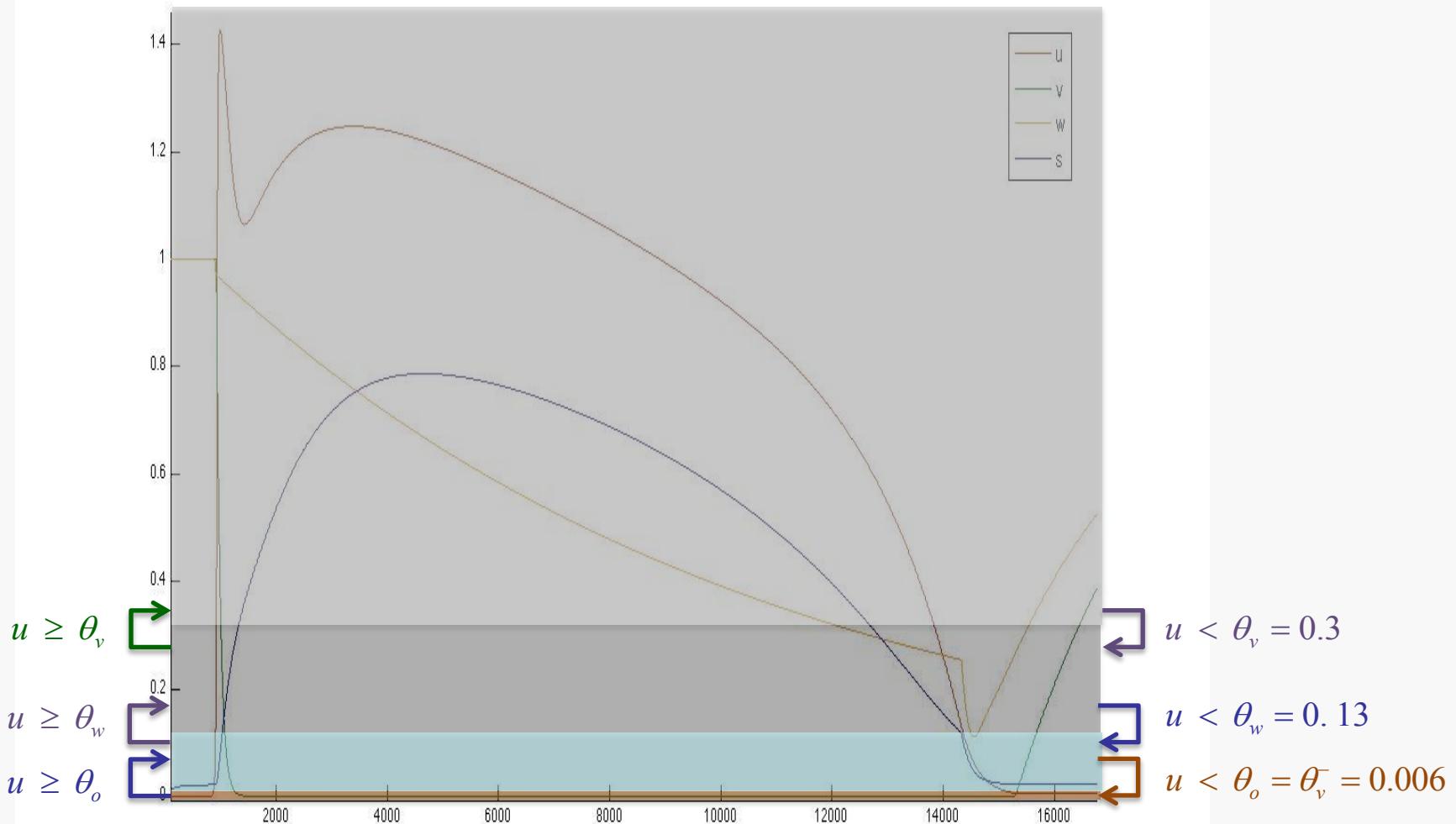
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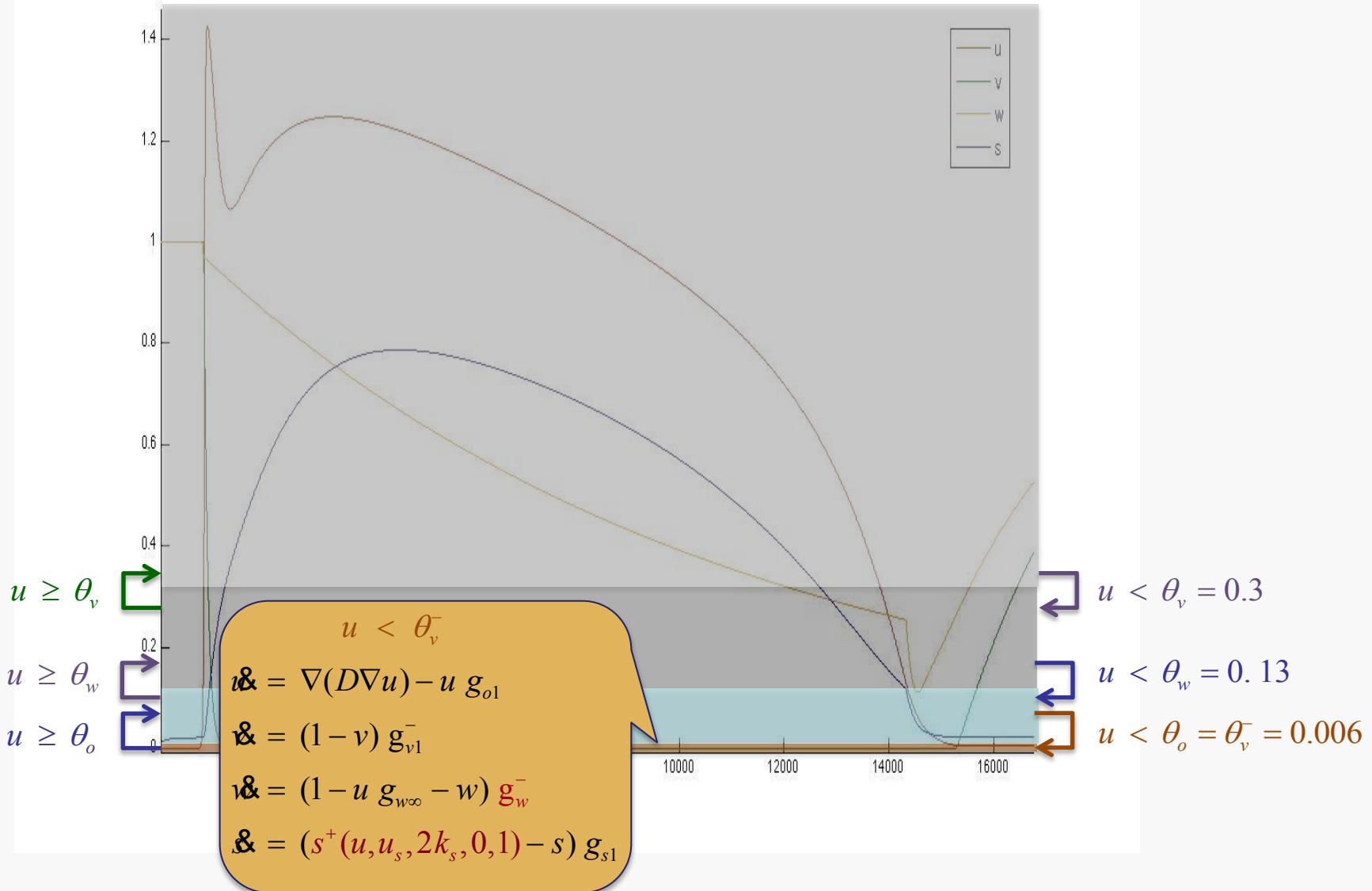




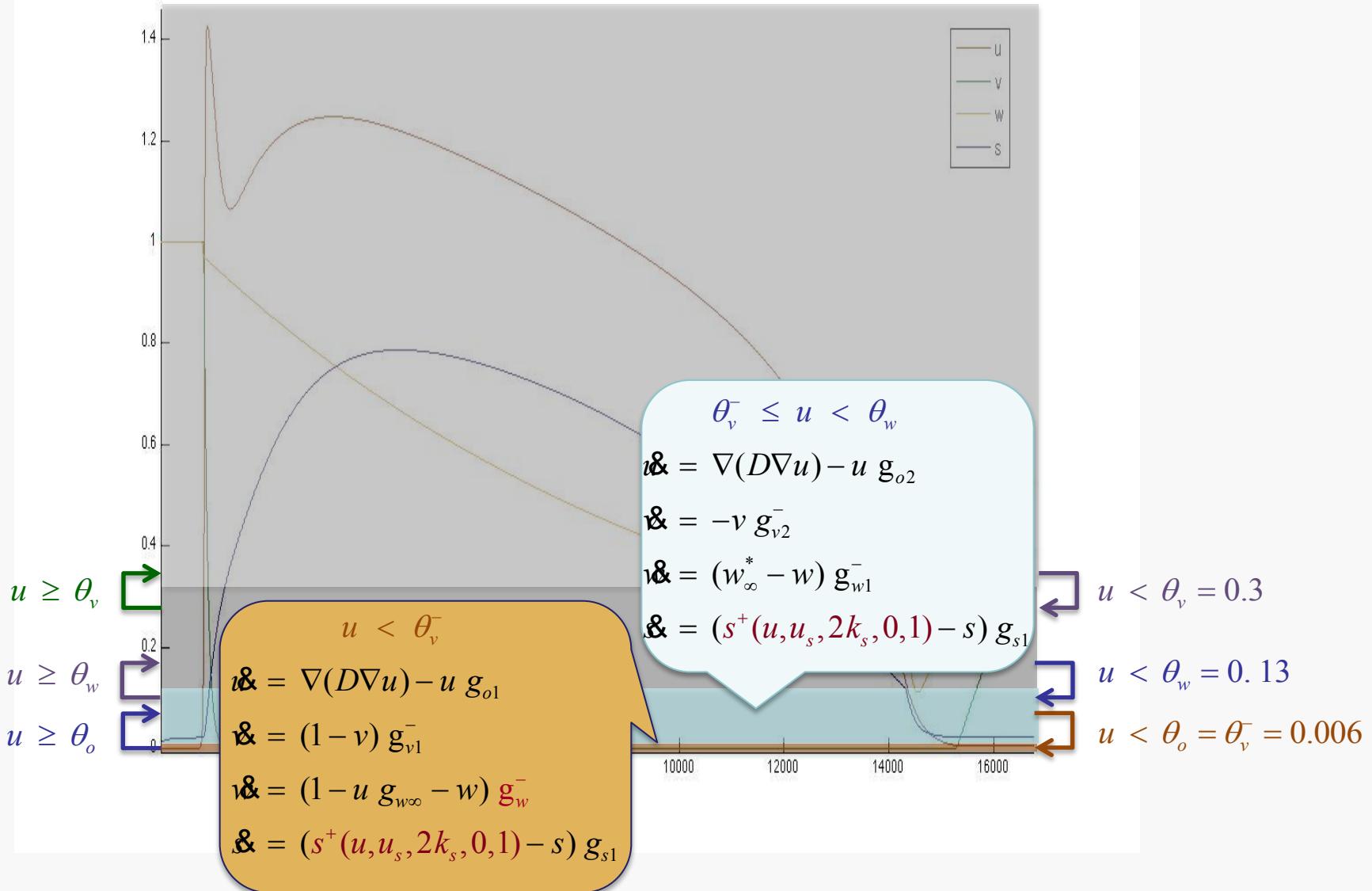
Conductances Minimal Model



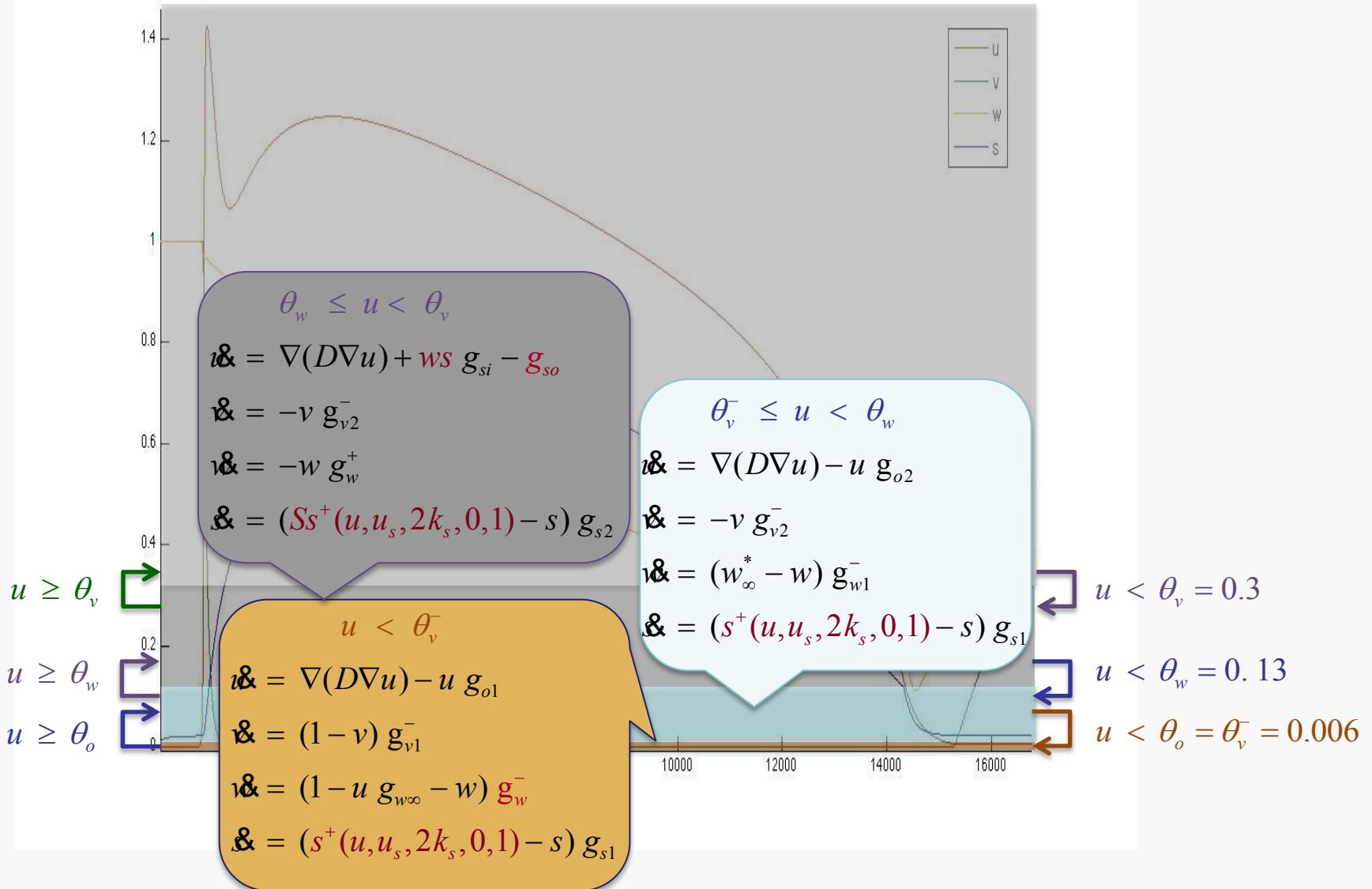
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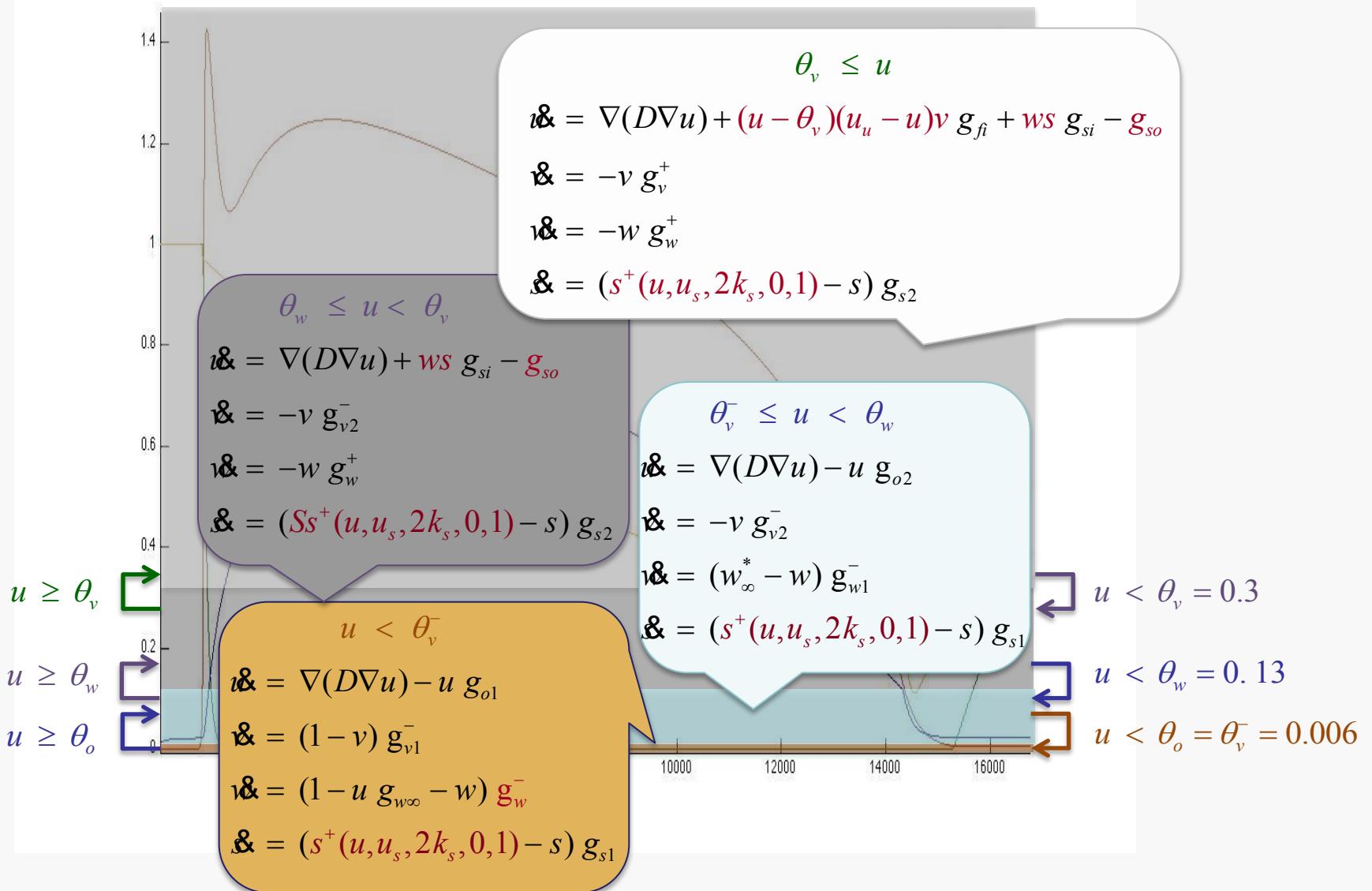
Conductances Minimal Model

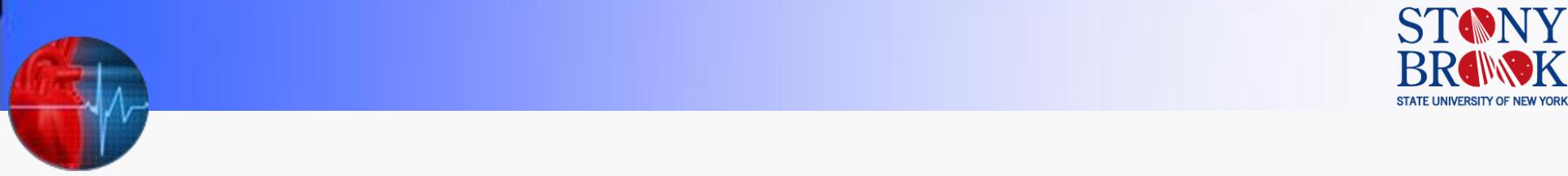


Conductances Minimal Model



Conductances Minimal Model





Piecewise Multi Affine Model

Our goals

- **Derive a Piecewise Multi Affine model:**
 - This should facilitate analysis
 - We want to improve the computational efficiency
- **Identify the parameters based on:**
 - Data generated by a detailed ionic model
 - Experimental, in-vivo data

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Optimal Polygonal Approximation

Problem to solve:

Given a nonlinear curve and the desired number of the segments return the optimal polygonal approximation:

Example:

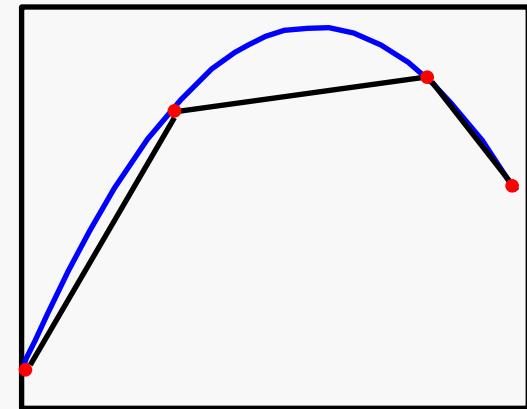
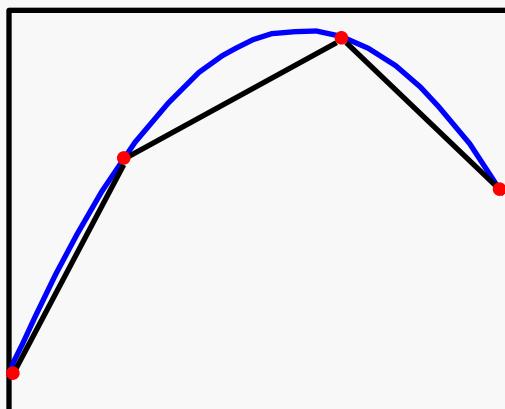
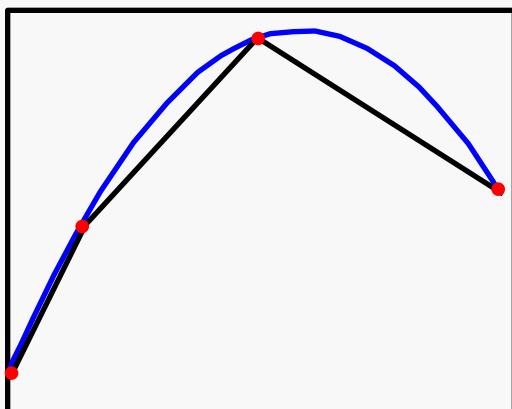


Optimal Polygonal Approximation

Problem to solve:

Given a nonlinear curve and the desired number of the segments return the optimal polygonal approximation:

Example: What is the optimal polygonal approximation of the blue curve with 3 segments ?



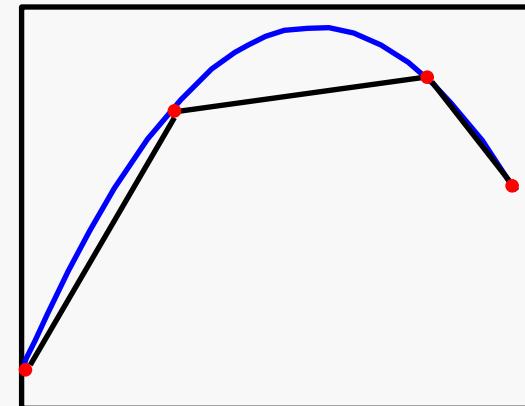
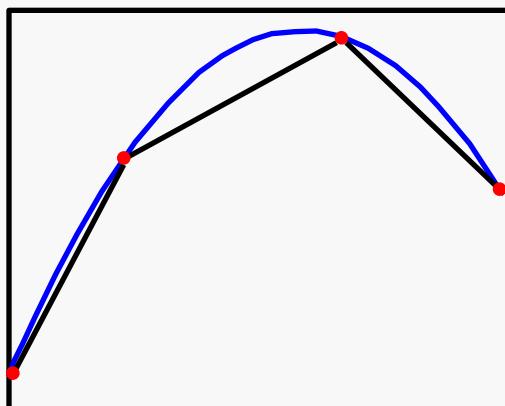
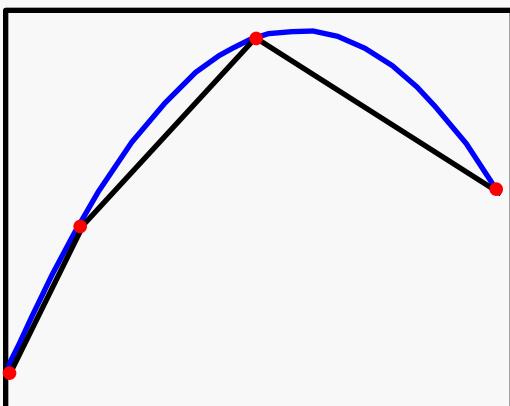


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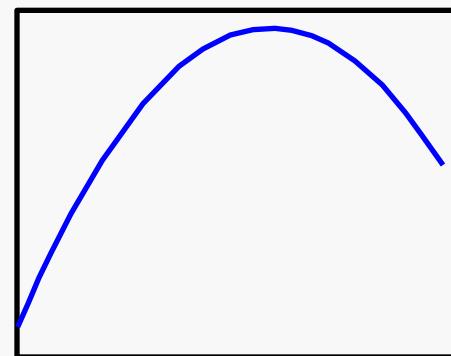


Dynamic Programming Algorithm
with Complexity $O(P^2S)$

P the number of points of the curve

S the number of segments

Marc Salotti, *An efficient algorithm for the optimal polygonal approximation of digitized curves*, Pattern Recognition Letters 22 (2001), Pag 215-221



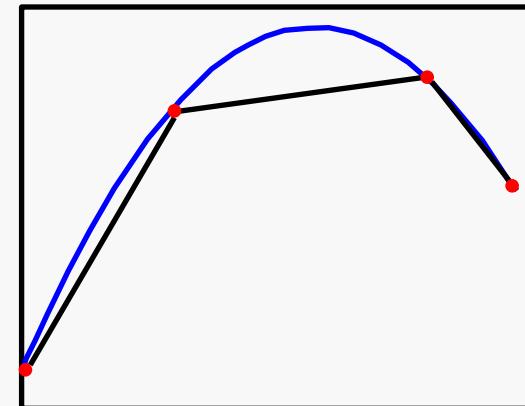
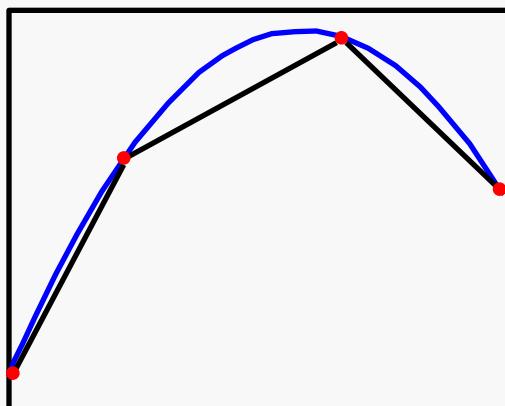
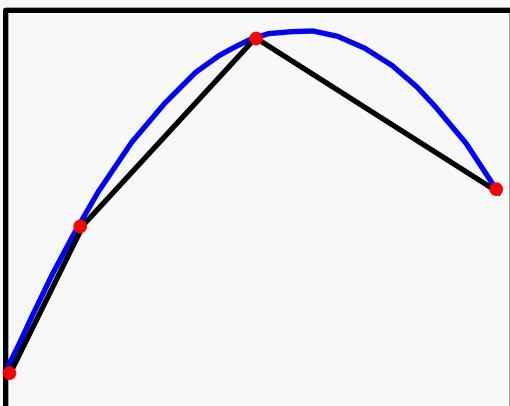


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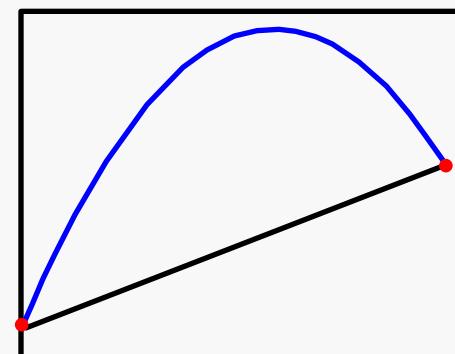


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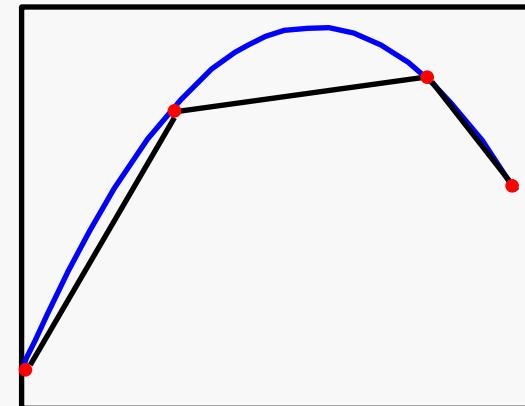
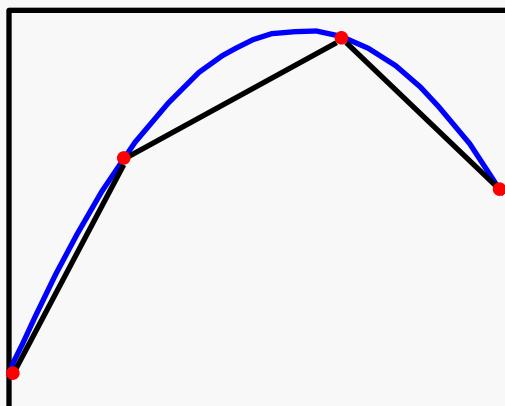
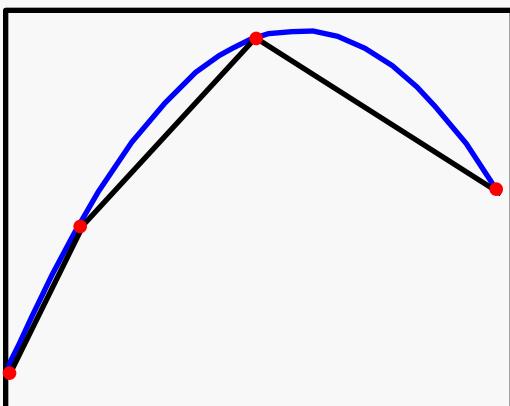


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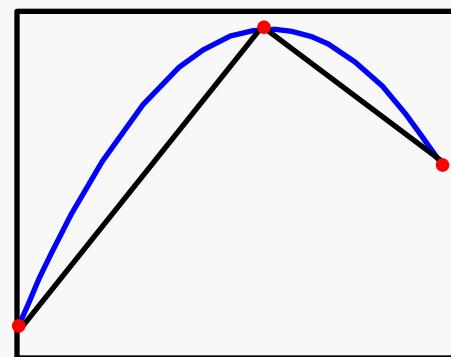


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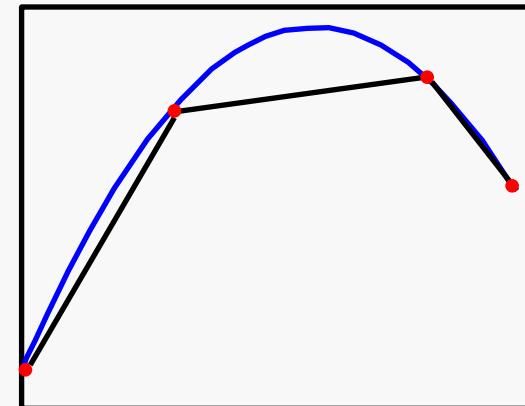
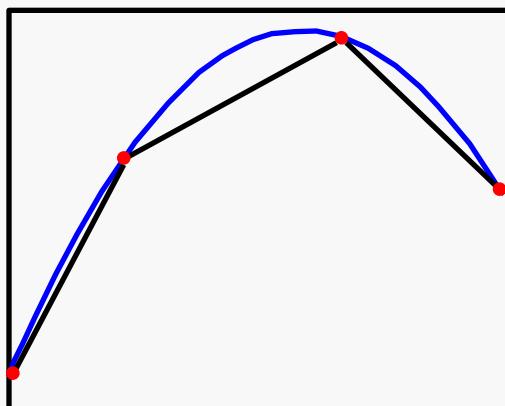
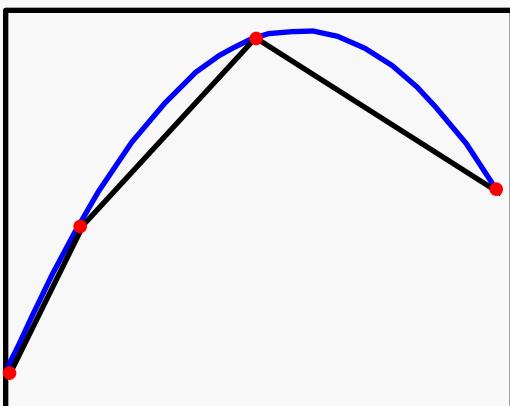


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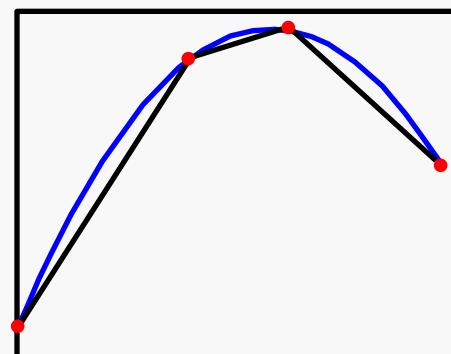


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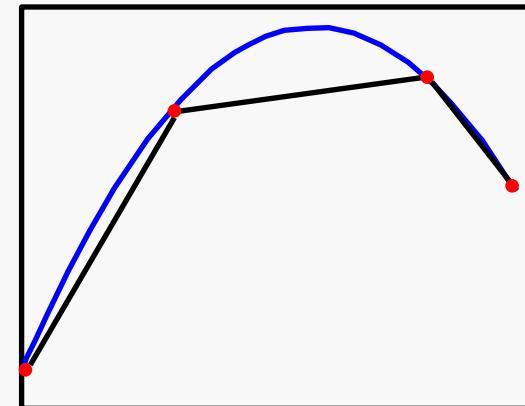
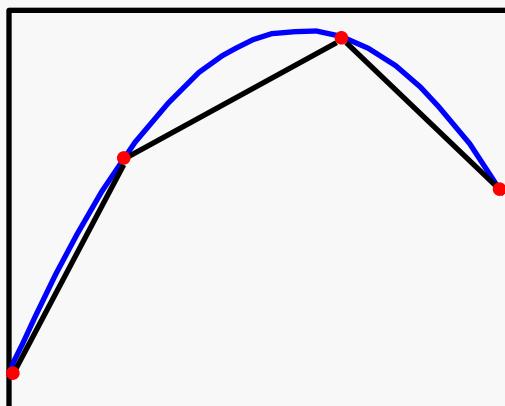
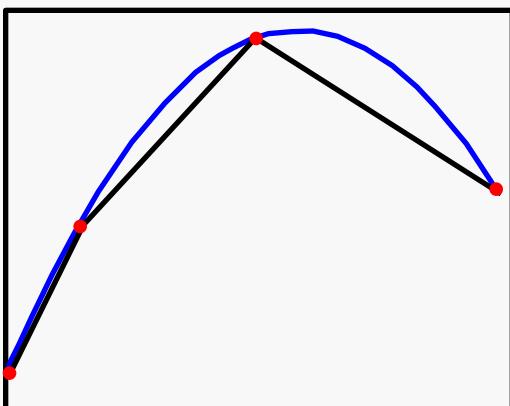


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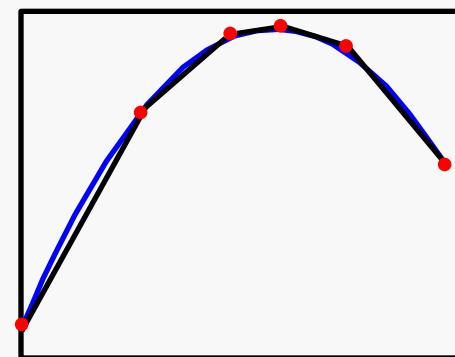


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Marc Salotti, *An efficient algorithm for the optimal polygonal approximation of digitized curves*, Pattern Recognition Letters 22 (2001), Pag 215-221



Global Optimal Polygonal Approximation

Our problem:

Given a set of nonlinear curves and the desired number of the segments return the optimal polygonal approximation:

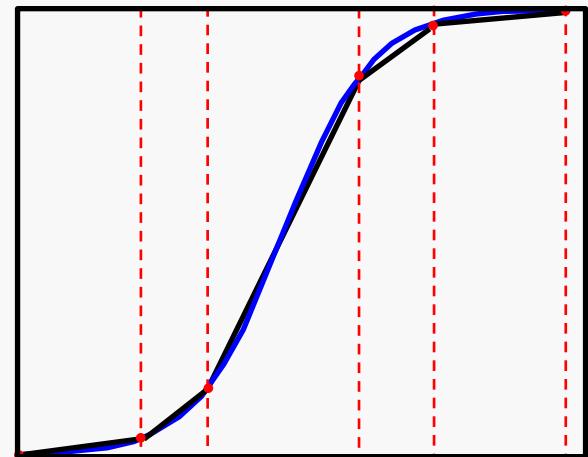
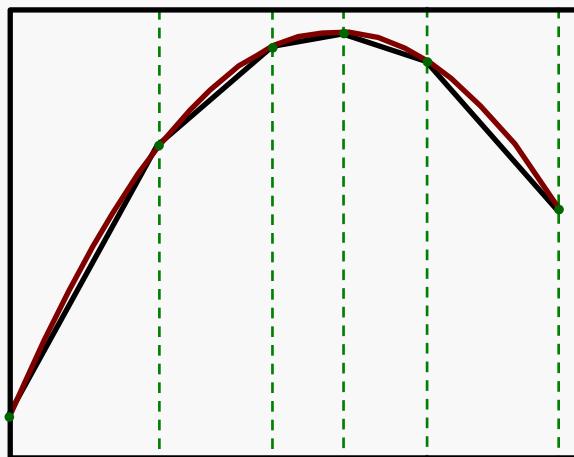


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Given a set of nonlinear curves and the desired number of the segments return the optimal polygonal approximation:

Example: What is the optimal polygonal approximation of the blue and the red curve with 5 segments ?



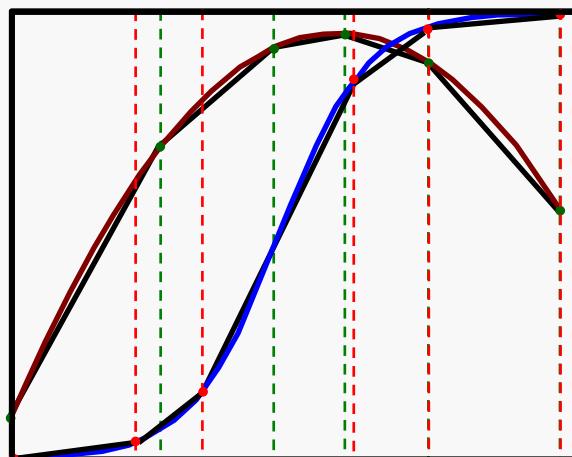


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But combining the two we obtain 8 segments and not 5 segments

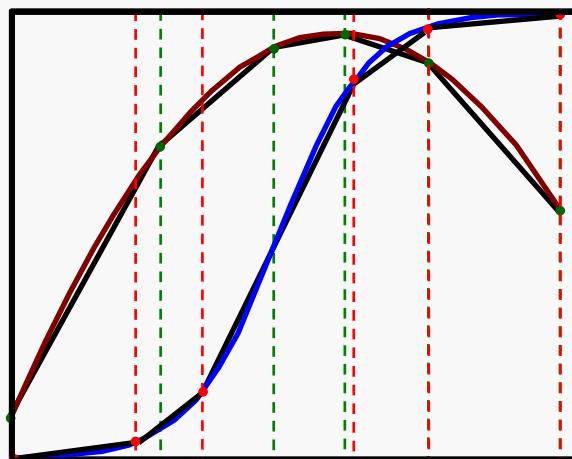


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Our solution: We modify the optimal polygonal approximation algorithm to perform the linearization on a set of curves trying to minimize the maximum error.

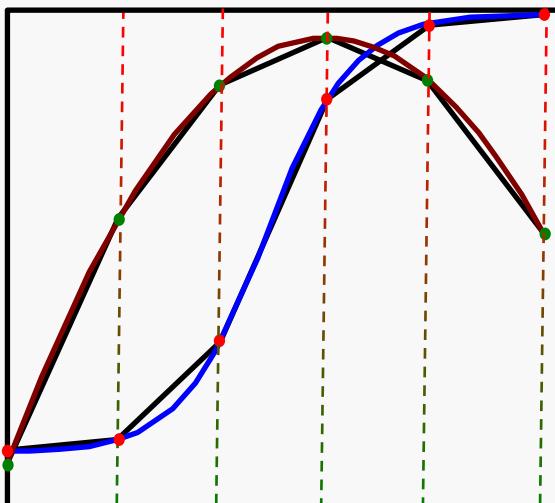


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Deriving the Piecewise Multi Affine Model

$$\begin{aligned} & (\theta_v < u \leq u_u) \\ \dot{u} &= \nabla(D\nabla u) + (\mathbf{u} - \theta_v)(u_u - u)v g_{fi} + ws g_{si} - g_{so} \\ \dot{v} &= -v g_v^+ \\ \dot{w} &= -w g_w^+ \\ \dot{s} &= S^+(u, k_s, u_s) g_{s2} - s g_{s2} \end{aligned}$$

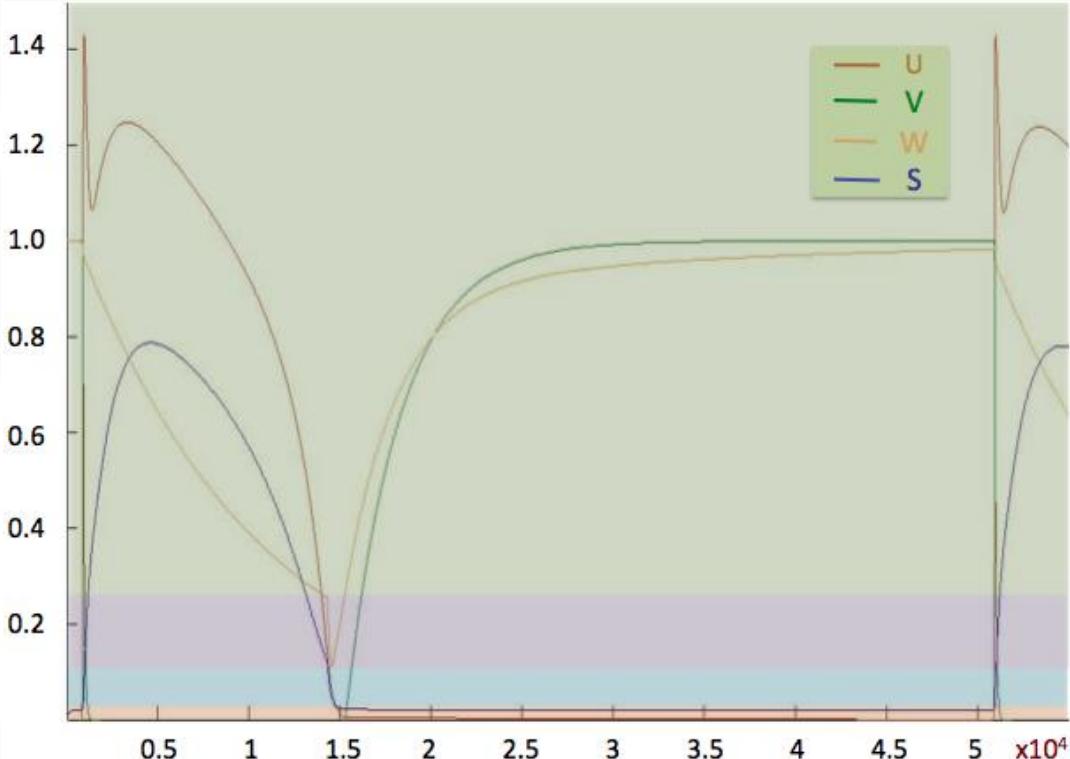
$$\begin{cases} u < \theta_v \\ u \geq \theta_v \end{cases}$$

$$\begin{aligned} & \theta_w \leq u < \theta_v \\ \dot{u} &= \nabla(D\nabla u) + ws g_{si} - g_{so} \\ \dot{v} &= -v g_v^- \\ \dot{w} &= -w g_w^+ \\ \dot{s} &= S^+(u, k_s, u_s) g_{s2} - s g_{s2} \end{aligned}$$

$$\begin{aligned} & \theta_v^- \leq u < \theta_w \\ \dot{u} &= \nabla(D\nabla u) - u g_{o2} \\ \dot{v} &= -v g_{v2}^- \\ \dot{w} &= (w_\infty - w) g_w^- \\ \dot{s} &= S^+(u, k_s, u_s) g_{s1} - s g_{s1} \end{aligned}$$

$$\begin{cases} u < \theta_w \\ u \geq \theta_w \end{cases}$$

$$\begin{aligned} & 0 \leq u < \theta_v^- \\ \dot{u} &= \nabla(D\nabla u) - u g_{o1} \\ \dot{v} &= (1-v) g_{v1}^- \\ \dot{w} &= (1-u g_{wo} - w) g_w^- \\ \dot{s} &= S^+(u, k_s, u_s) g_{s1} - s g_{s1} \end{aligned}$$





Deriving the Piecewise Multi Affine Model

$$\begin{aligned} & (\theta_v < u \leq u_u) \\ \dot{u} &= \nabla(D\nabla u) + (\mathbf{u} - \theta_v)(u_u - u)v g_{fi} + ws g_{si} - g_{so} \\ \dot{v} &= -v g_v^+ \\ \dot{w} &= -w g_w^+ \\ \dot{s} &= S^+(u, k_s, u_s) g_{s2} - s g_{s2} \end{aligned}$$

$\xrightarrow{u < \theta_v}$ $\xleftarrow{u \geq \theta_v}$

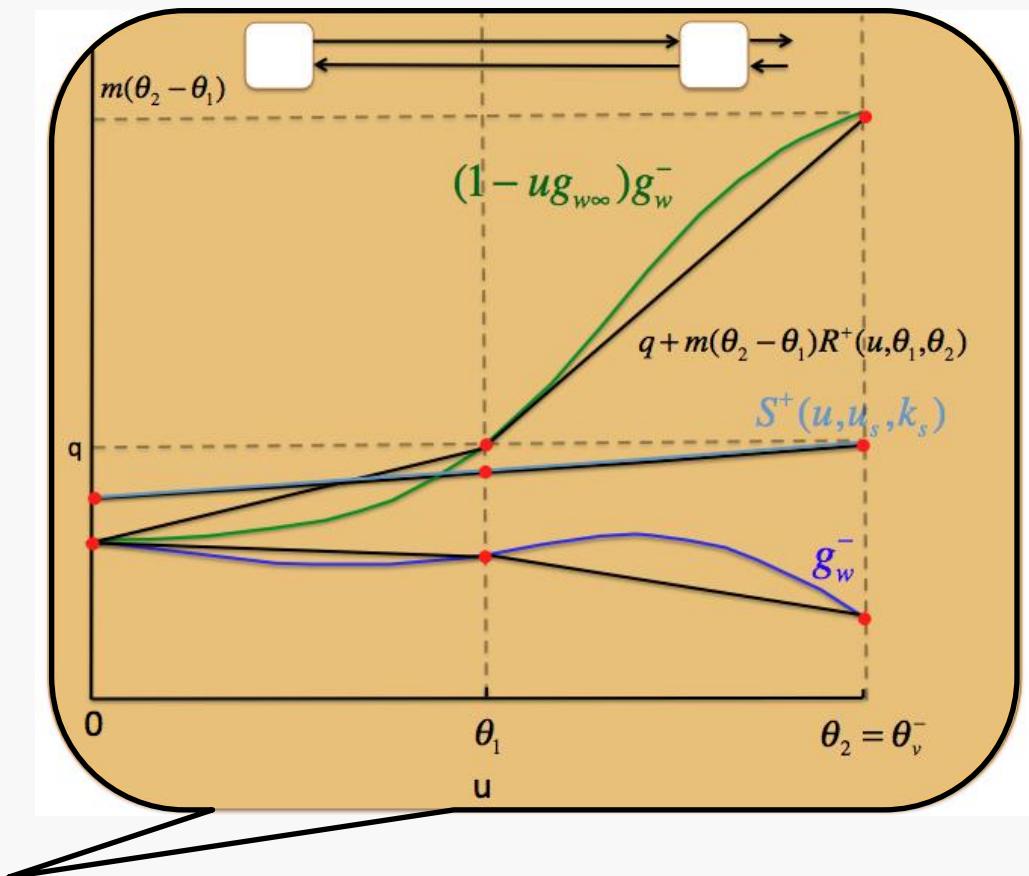
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Deriving the Piecewise Multi Affine Model

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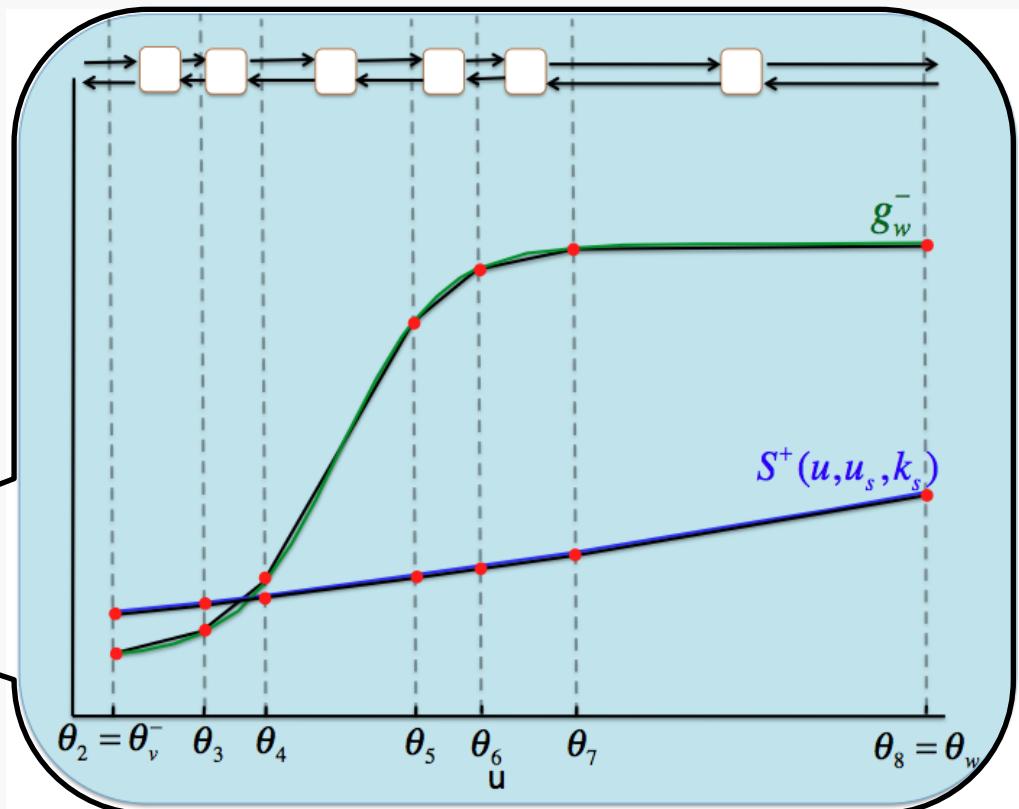
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$\xrightarrow{u \geq \theta_v^-}$



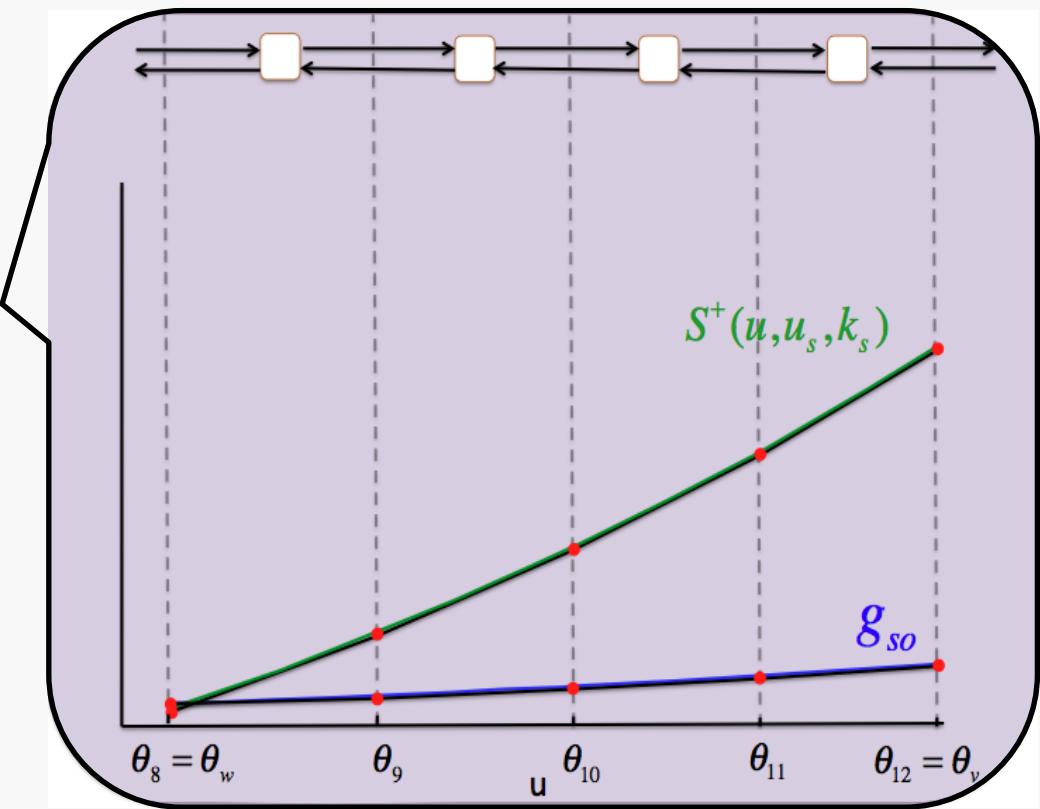


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Diagram showing piecewise transitions:

- From $u < \theta_v$ to $\theta_w \leq u < \theta_v$:
 $\dot{u} = \nabla(D\nabla u) + ws g_{si} - g_{so}$
 $\dot{v} = -v g_{v2}^-$
 $\dot{w} = -w g_w^+$
 $\dot{s} = S^+(u, k_s, u_s) g_{s2} - s g_{s2}$
- From $u \geq \theta_v$ to $\theta_v^- \leq u < \theta_w$:
 $\dot{u} = \nabla(D\nabla u) - u g_{o2}$
 $\dot{v} = -v g_{v2}^-$
 $\dot{w} = (w_\infty - w) g_w^-$
 $\dot{s} = S^+(u, k_s, u_s) g_{s1} - s g_{s1}$
- From $u < \theta_v^-$ to $0 \leq u < \theta_v^-$:
 $\dot{u} = \nabla(D\nabla u) - u g_{o1}$
 $\dot{v} = (1-v) g_{v1}^-$
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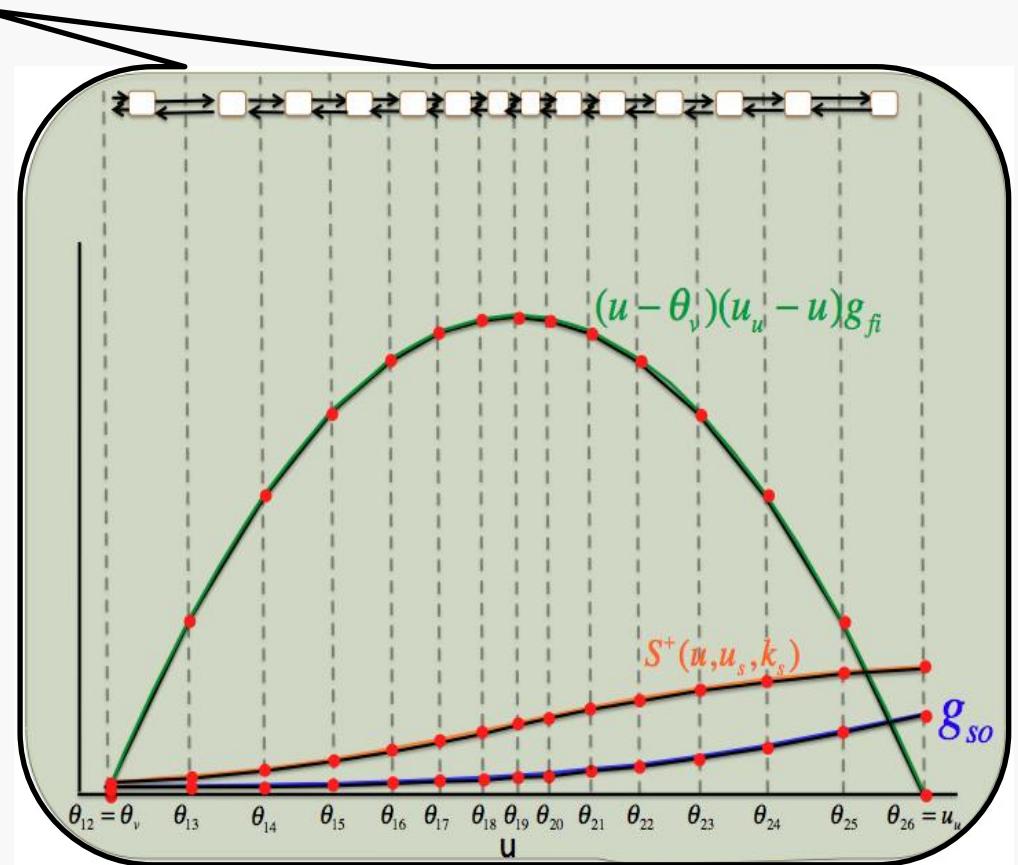


Deriving the Piecewise Multi Affine Model

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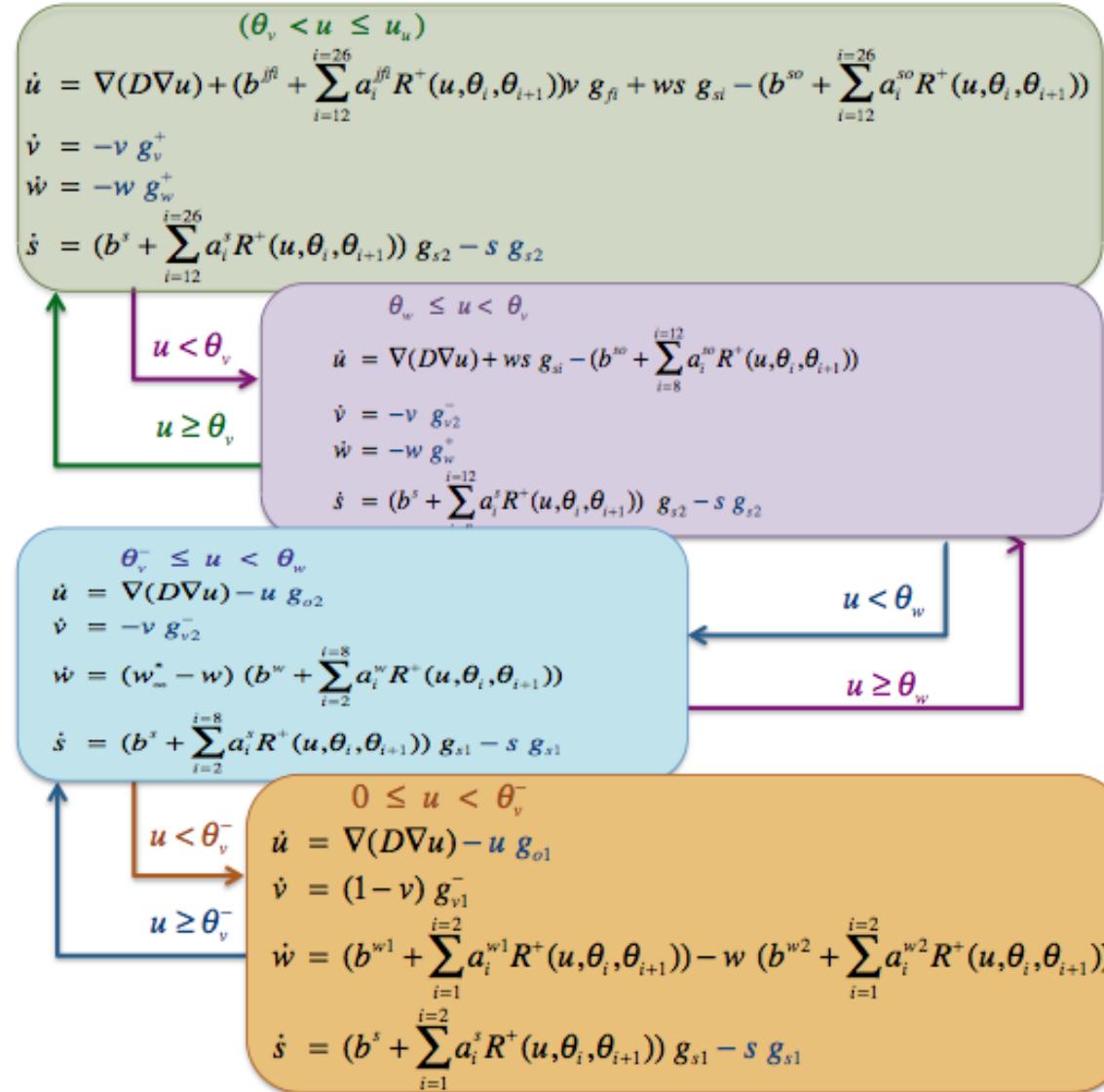
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- From $0 \leq u < \theta_v^-$ back to $u < \theta_v$:
 $\dot{u} = \nabla(D\nabla u) - u g_{o1}$
 $\dot{v} = (1-v) g_{v1}^-$
 $\dot{w} = (1-u g_{w\infty} - w) g_w^-$
 $\dot{s} = S^+(u, k_s, u_s) g_{s1} - s g_{s1}$



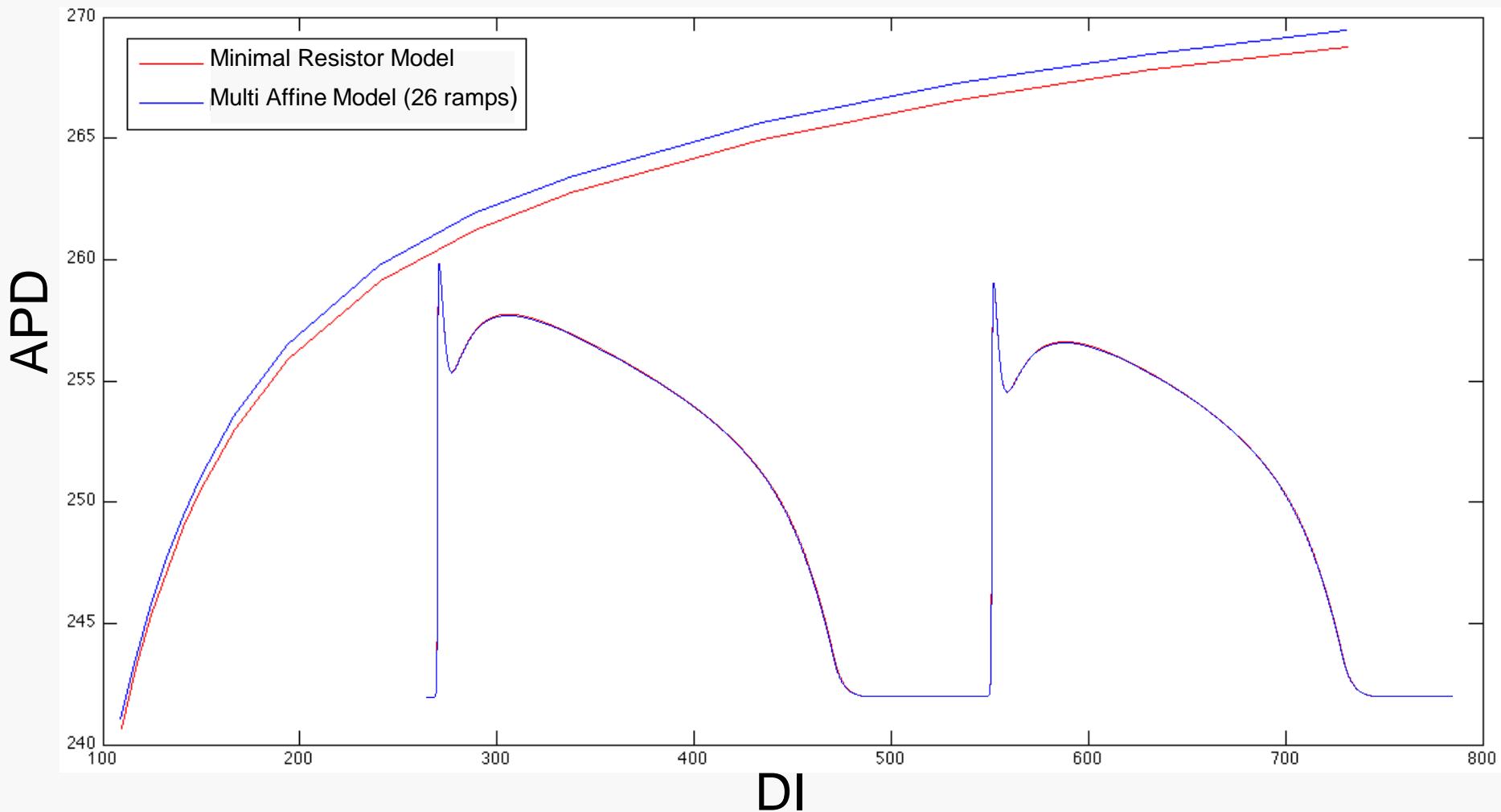


Deriving the Piecewise Multi Affine Model



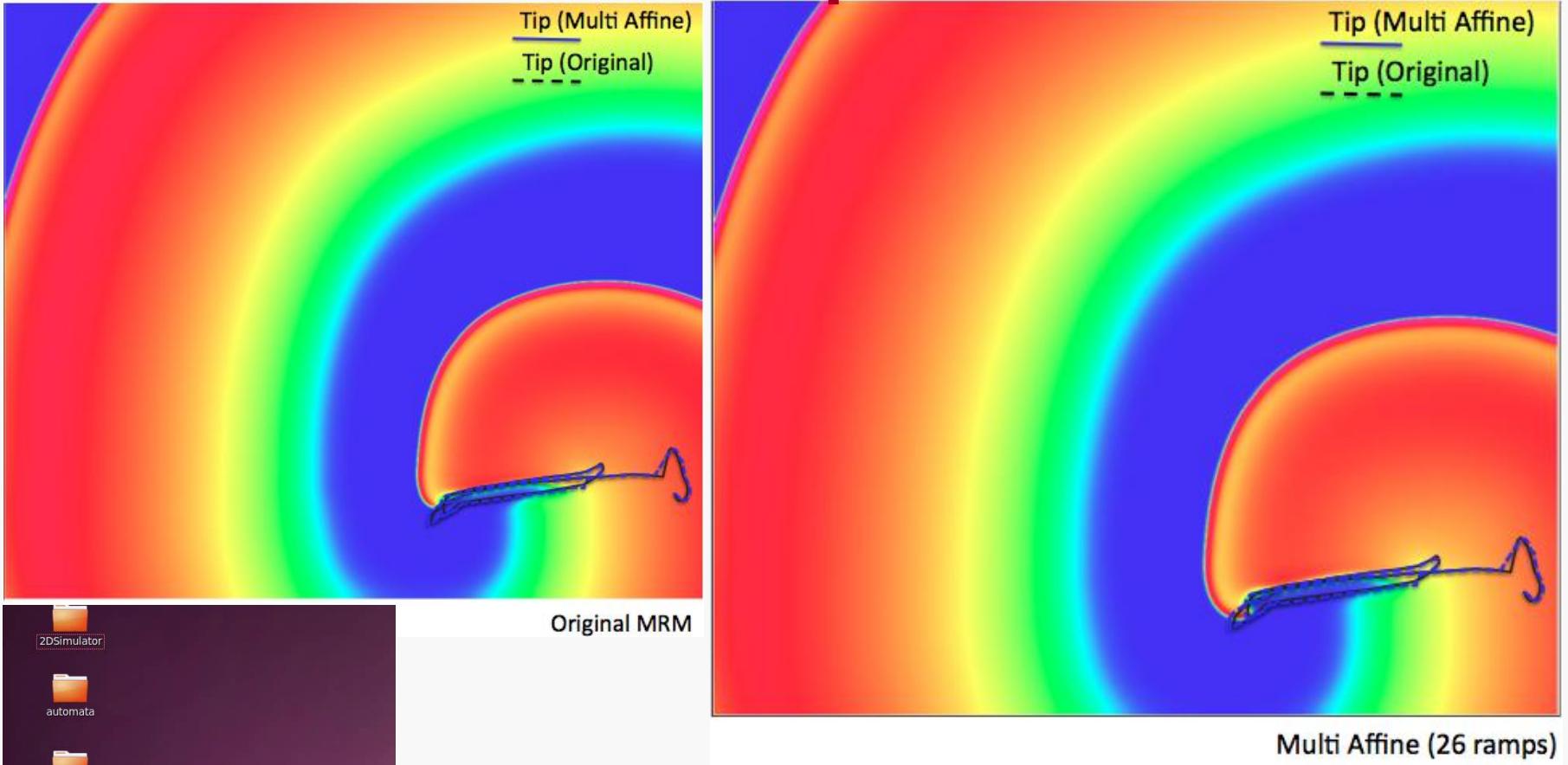


1D Cable Comparison





2D Comparison



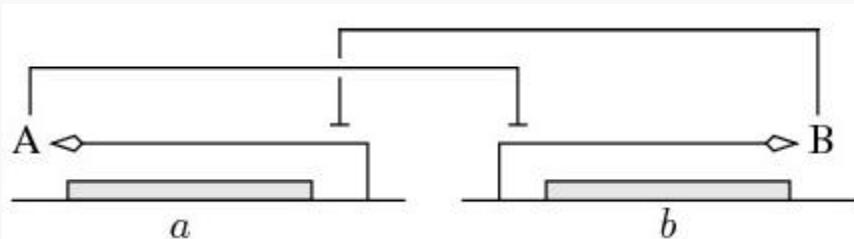
1024x1024 cells 100000 iterations 556 sec. vs. 810 sec.
(GPU Tesla C1060)

Simulation of the Multi Affine Model is
computational efficient (1.43 x)

Parameter Identification



Genetic Regulatory Networks

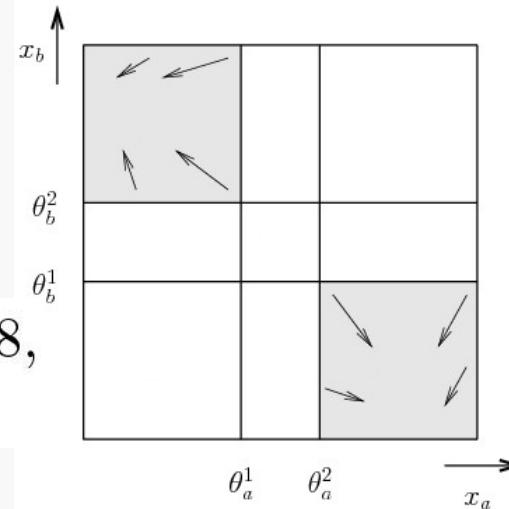
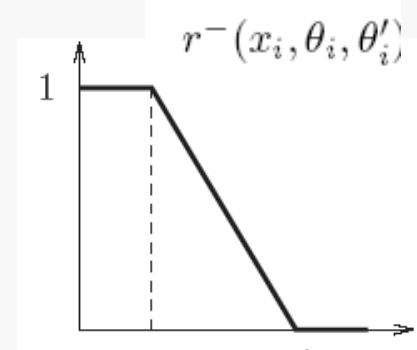


$$\begin{aligned}\dot{x}_a &= \kappa_a r^-(x_b, \theta_b^1, \theta_b^2) - \gamma_a x_a \\ \dot{x}_b &= \kappa_b r^-(x_a, \theta_a^1, \theta_a^2) - \gamma_b x_b\end{aligned}$$

G. Batt, C. Belta and R. Weiss (2008) **Temporal logic analysis of gene networks under parameter uncertainty**

cross-inhibition network

x : protein concentration
 $\theta_a^1, \theta_a^2, \theta_b^1, \theta_b^2$
threshold concentration
 $k_a, k_b, \gamma_a, \gamma_b$: rate parameters



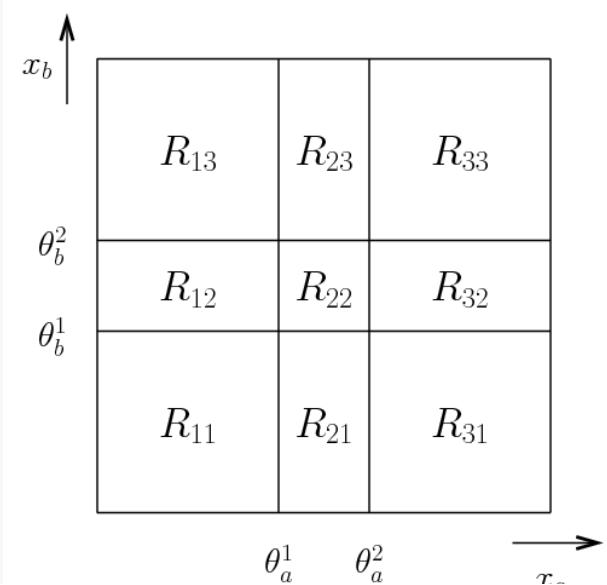
- Find parameters such that network is bistable

$$p = (\kappa_a, \kappa_b) \in \mathcal{P} = [0, 40] \times [0, 20] \quad \begin{aligned}\gamma_a &= 1, \gamma_b = 2, \theta_a^1 = 8, \\ \theta_b^1 &= 8, \theta_a^2 = \theta_b^2 = 12\end{aligned}$$

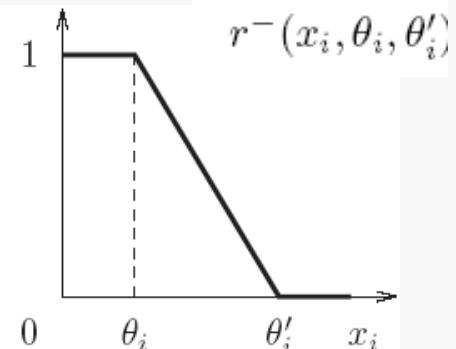
Genetic Regulatory Networks

- Partition of the state space: rectangles

$$R \in \mathcal{R}$$



$$\begin{aligned}\dot{x}_a &= \kappa_a r^-(x_b, \theta_b^1, \theta_b^2) - \gamma_a x_a \\ \dot{x}_b &= \kappa_b r^-(x_a, \theta_a^1, \theta_a^2) - \gamma_b x_b\end{aligned}$$



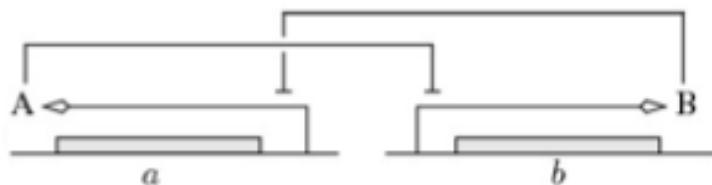
- Differential equation models $\dot{x} = f(x, p)$, with
 - f is **piecewise-monomial** (PMA) function of state **variables** x
 - f is **affine** function of rate **parameters** p k_a, k_b
- (monomial functions: products of different state variables allowed)



Specifications of dynamical properties

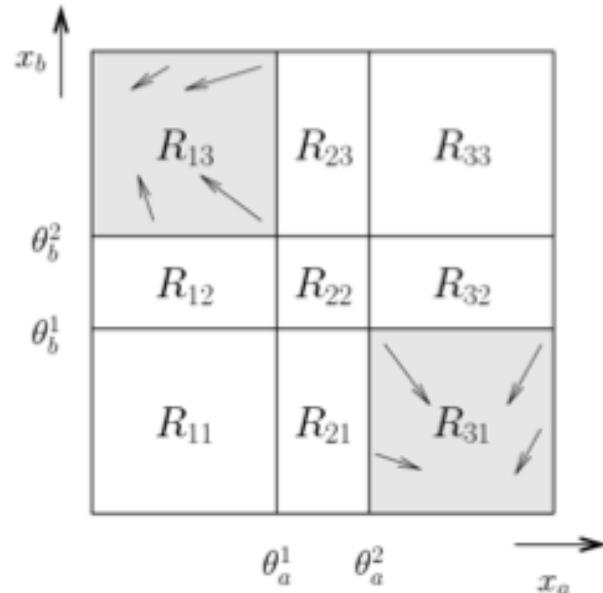
❖ Dynamical properties expressed in temporal logic (LTL)

- set of atomic proposition Π : $x_i < \lambda_i, x_i > \lambda_i$
- usual logical operators $\neg\phi, \phi_1 \wedge \phi_2, \phi_1 \vee \phi_2, \phi_1 \rightarrow \phi_2, \dots$
- temporal operators $X\phi, F\phi, G\phi, \phi_1 U \phi_2, \dots$



bistability property:

$$\begin{aligned}\phi_1 = & (x_a < \theta_a^1 \wedge x_b > \theta_b^2 \rightarrow G(x_a < \theta_a^1 \wedge x_b > \theta_b^2)) \\ & \wedge (x_b < \theta_b^1 \wedge x_a > \theta_a^2 \rightarrow G(x_b < \theta_b^1 \wedge x_a > \theta_a^2))\end{aligned}$$

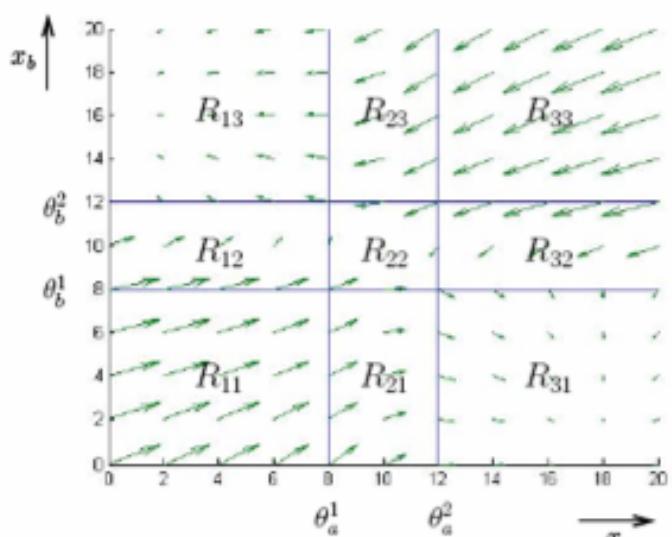


How to define that the system satisfies an LTL property ?

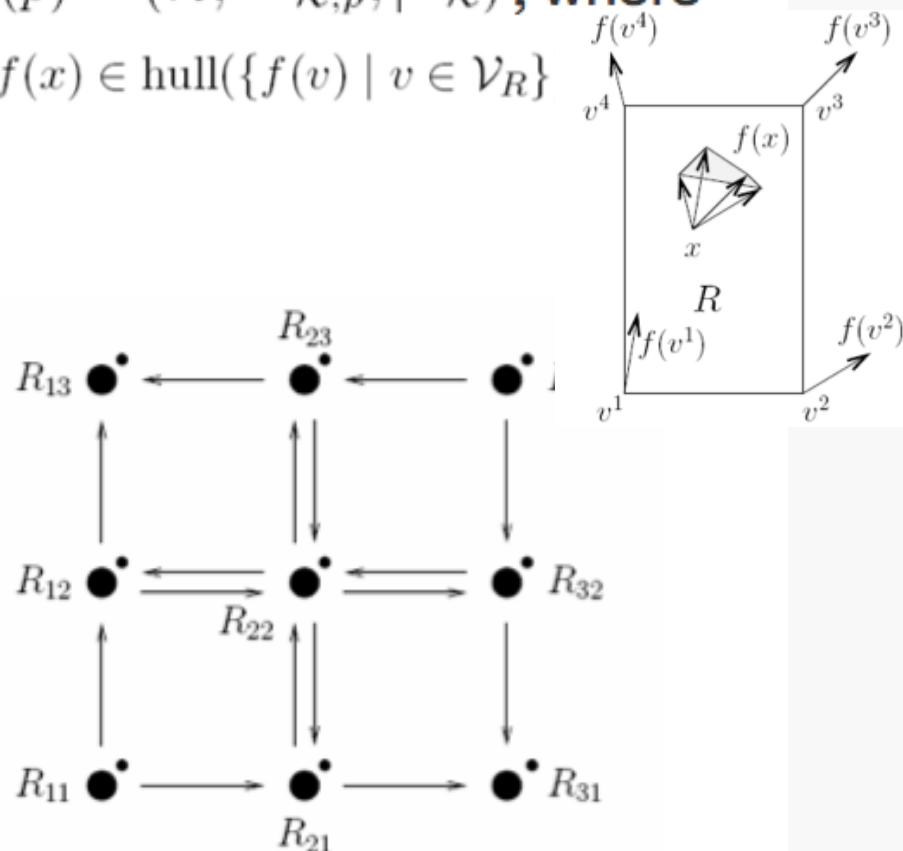


❖ Discrete transition system, $T_{\mathcal{R}}(p) = (\mathcal{R}, \rightarrow_{\mathcal{R},p}, \models_{\mathcal{R}})$, where

- \mathcal{R} finite set of rectangles $\forall x \in R, f(x) \in \text{hull}(\{f(v) \mid v \in V_R\})$
- $\rightarrow_{\mathcal{R},p}$ quotient transition relation
- $\models_{\mathcal{R}}$ quotient satisfaction relation



$$\begin{aligned} R_{11} &\rightarrow_{\mathcal{R},p} R_{21}, \quad R_{21} \rightarrow_{\mathcal{R},p} R_{31} \\ R_{11} &\rightarrow_{\mathcal{R},p} R_{11}, \end{aligned}$$



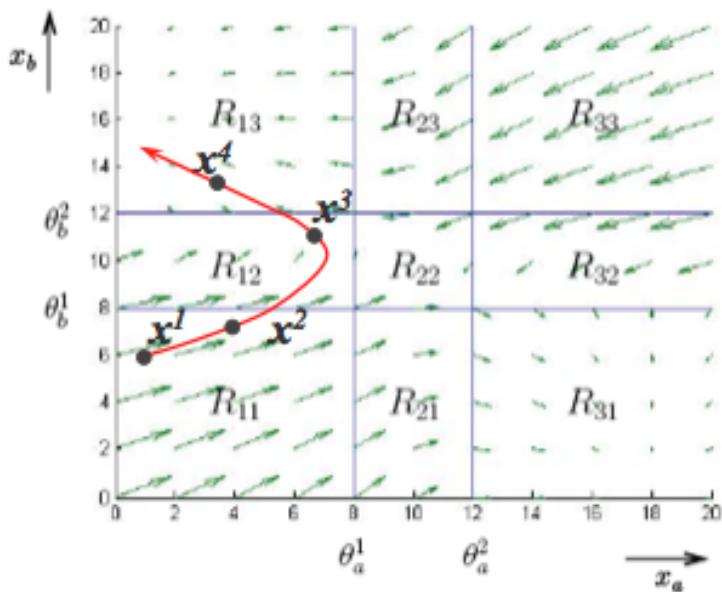
$$R_{11} \models_{\mathcal{R}} x_a < \theta_a^1, \quad R_{11} \models_{\mathcal{R}} x_b < \theta_b^1$$



Embedding transition system

❖ **PMA system**, $\Sigma = (f, \Pi)$ associated with **embedding transition system**, $T_{\mathcal{X}}(p) = (\mathcal{X}_{\mathcal{R}}, \rightarrow_{\mathcal{X}, p}, \models_{\mathcal{X}})$, where

- $\mathcal{X}_{\mathcal{R}}$ continuous state space
- $\rightarrow_{\mathcal{X}, p}$ transition relation
- $\models_{\mathcal{X}}$ satisfaction relation



$$x^1 \rightarrow_{\mathcal{X}, p} x^2, \quad x^1 \rightarrow_{\mathcal{X}, p} x^3,$$

$$x^2 \rightarrow_{\mathcal{X}, p} x^3, \quad x^3 \rightarrow_{\mathcal{X}, p} x^4$$

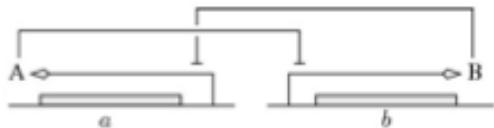
$$x^1 \models_{\mathcal{X}} x_a < \theta_a^1, \quad x^1 \models_{\mathcal{X}} x_b < \theta_b^1,$$

$$x^4 \models_{\mathcal{X}} x_a < \theta_a^1, \quad x^4 \models_{\mathcal{X}} x_b > \theta_b^1$$



Iterative exploration of parameter space

gene network



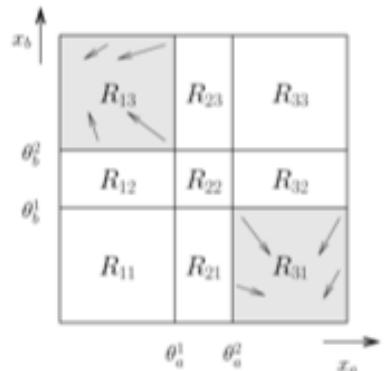
PMA model

$$\begin{aligned}\dot{x}_a &= \kappa_a r^-(x_b, \theta_b^1, \theta_b^2) - \gamma_a x_a \\ \dot{x}_b &= \kappa_b r^-(x_a, \theta_a^1, \theta_a^2) - \gamma_b x_b\end{aligned}$$

intervals for
uncertain parameters

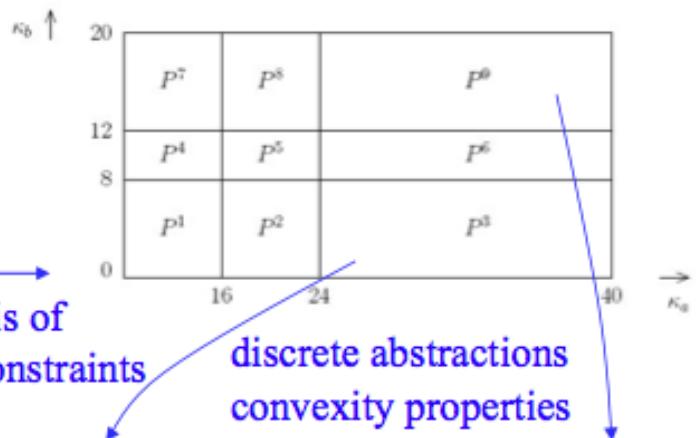
$$\begin{aligned}\kappa_a &\in [0, 40], \kappa_b \in [0, 20] \\ \gamma_a &= 1, \gamma_b = 2, \theta_a^1 = 8, \\ \theta_b^1 &= 8, \theta_a^2 = \theta_b^2 = 12\end{aligned}$$

specifications

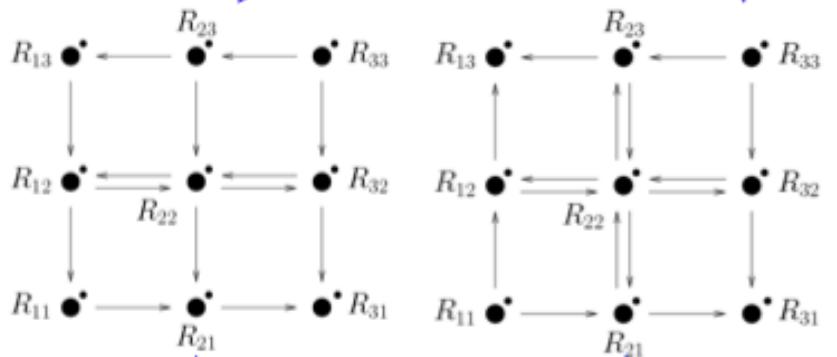


$$\begin{aligned}(x_a < \theta_a^1 \wedge x_b > \theta_b^2 \rightarrow G(x_a < \theta_a^1 \wedge x_b > \theta_b^2)) \\ \wedge (x_b < \theta_b^1 \wedge x_a > \theta_a^2 \rightarrow G(x_b < \theta_b^1 \wedge x_a > \theta_a^2))\end{aligned}$$

synthesis of
parameter constraints



discrete abstractions
convexity properties



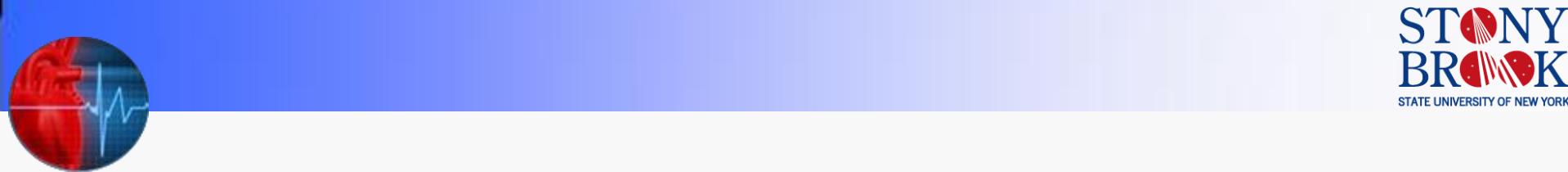
model checking

$$T_{\mathcal{R}}(p^3) \not\models \phi$$

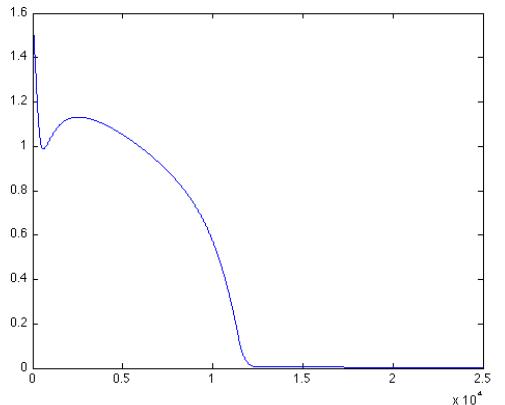
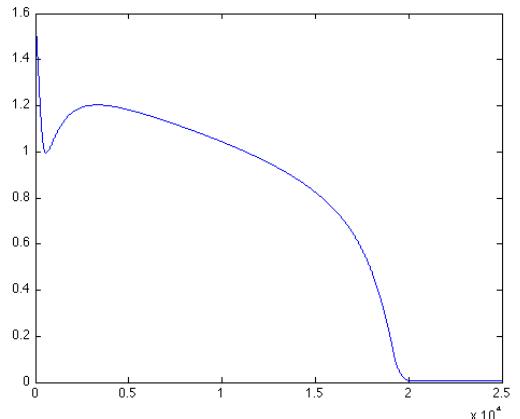
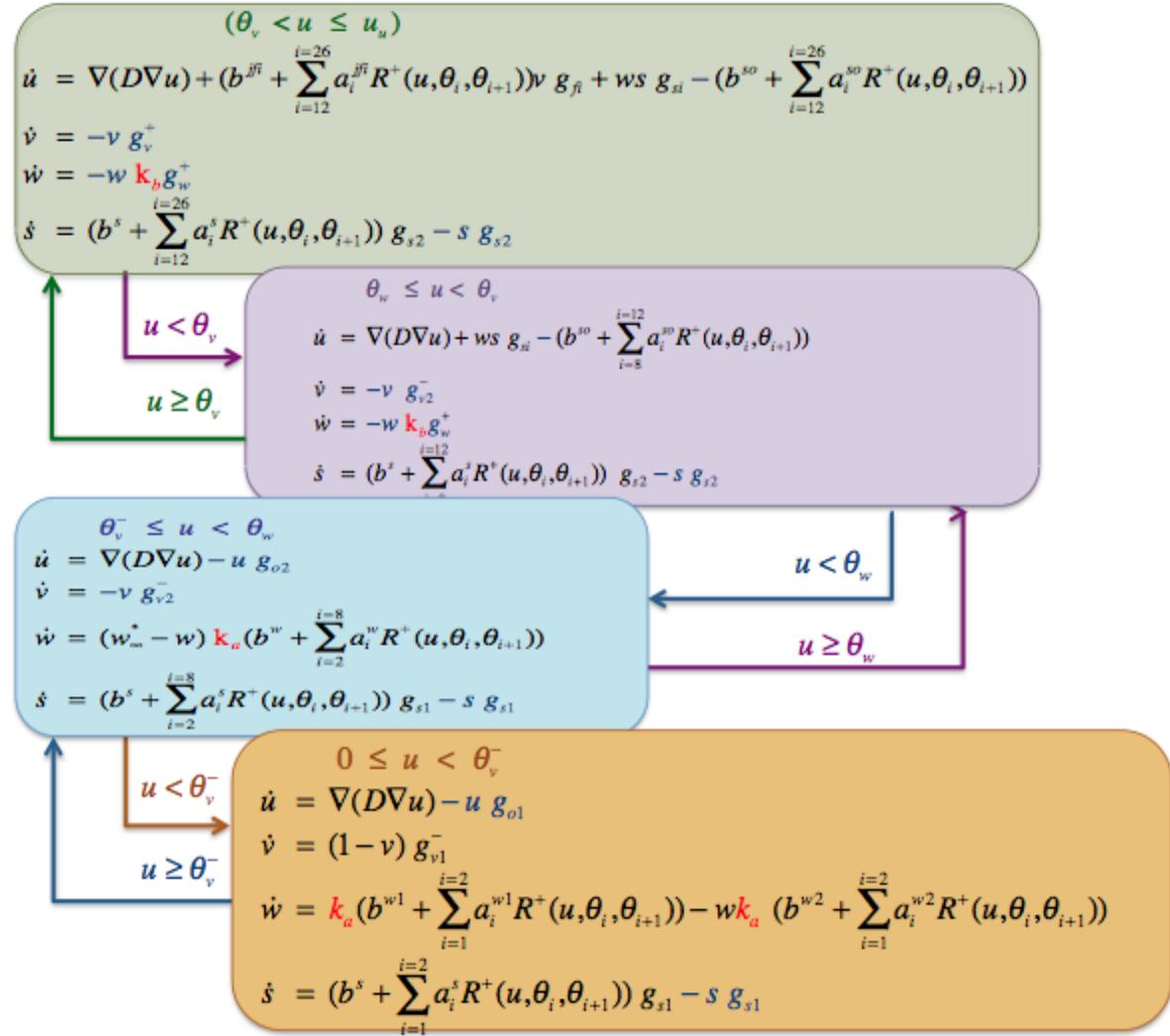
No conclusion

$$T_{\mathcal{R}}(p^9) \models \phi$$

Valid parameter set



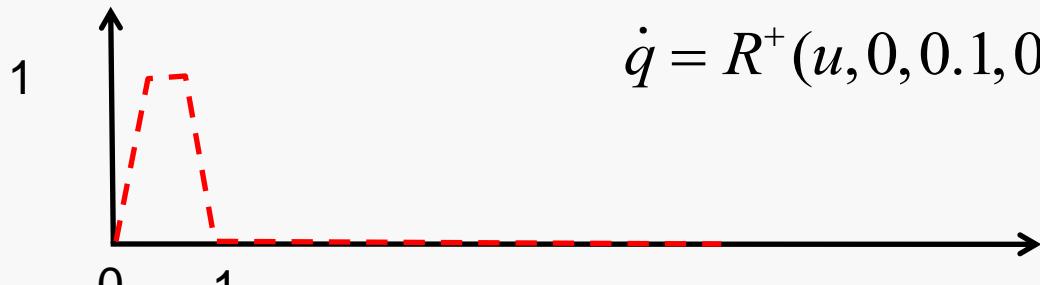
Parameter Identification for APD



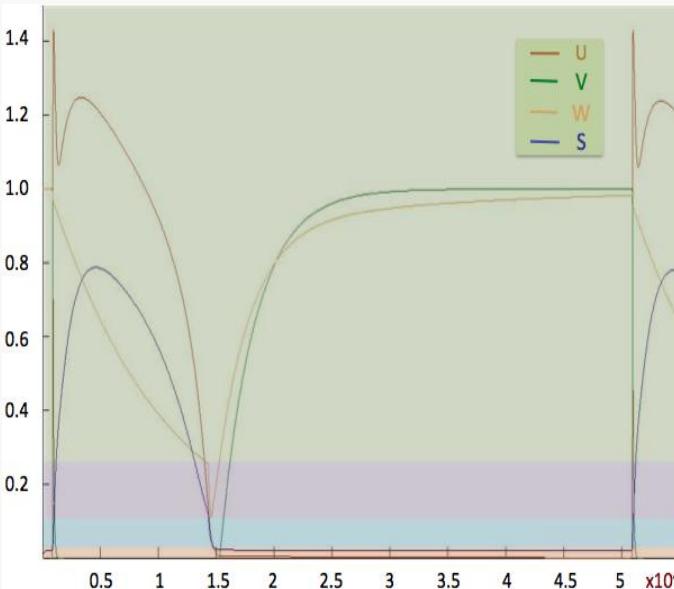


Encoding a Property on the APD

Introducing a new state variable for the stimulus:



Measuring the APD with another state variable: $\dot{z} = R^+(u, 0.3, 0.3, 0, 1)$



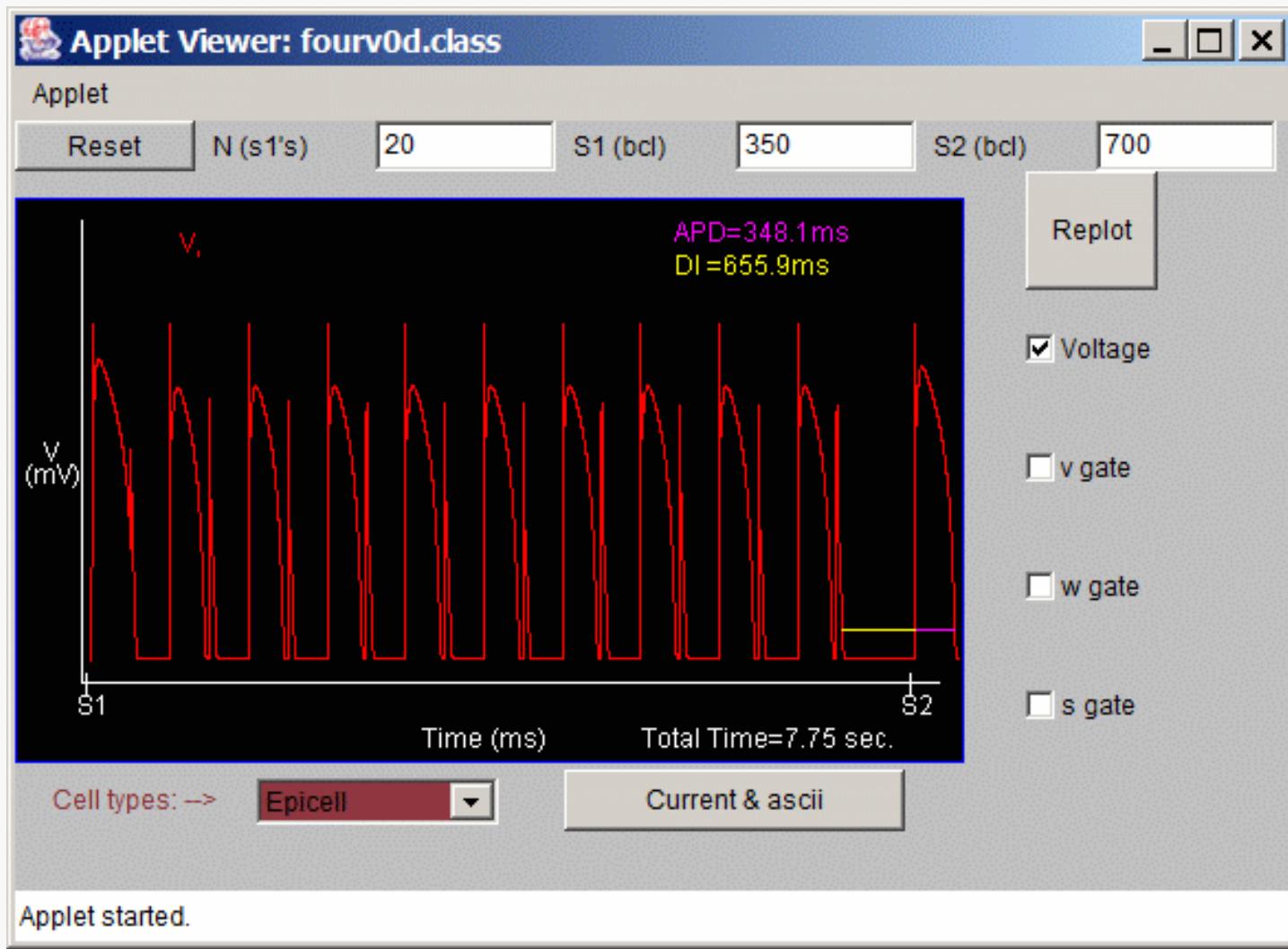
Property: starting with initial conditions $u=0, w=0.8, v=1.0, s=0.0$ the cell will fire an action potential with APD between 200 and 300 ms

LTL FORMULA:

$$\phi = (u = 0 \wedge w = 0.8 \wedge v = 1.0 \wedge s = 0.0) \rightarrow F(z > 200 \wedge z < 300)$$



Alternans



Future Works

- **Performing experiments with Rovergene**
- **Encoding more sophisticated properties**
- **Discriminate healthy/unhealthy tissues using model checking**

**Thanks for
the attention**



```
function [e,a,b,xb] = optimalLinearApproximation(x,y,S)
```

Input:

x,y: Curves given as an x-points vector and a vector of y-points vectors

S: Number >= 2 of desired segments

Output:

e: Errors matrix

a,b: Line-segment-coefficients matrix

xb: x-coordinate at breaking point matrix

Initialization

```
z1 = size(x); P = z1(2); Get number of points in each curve
```

```
z2 = size(y); C = z2(1); Get number of digitized curves
```

```
se = zeros(1,C); Initialize vector of errors, one error for each curve
```

Cost tables

```
cost = ones(P,S) * inf; cost(30,4) = min cost to pt 30 with 4-segm polyline
```

```
error = ones(P,P) * inf; error(i, n) = cached error of line segment (i,n)
```

```
cost(2,1) = 0; 1-segment-polyline cost of polyline (1,2) = 0
```

Predecessor table

```
father = ones(P,S) * inf; father(30,4) = pred of pt 30 on a 4-segm polyline
```

Computation of optimal segmentation

Initialize cost and father for 1-segment-polyline, from pt 1 to all other pts

```
for p = 2:P Traverse all other points
```

```
    for c = 1:C Traverse all curves
```

```
        se(c) = segmentError(x(1:p), y(c,1:p)); (1,p)-line-segment appr error
```

```
    end; for c
```

```
        cost(p,1) = max(se); Maximum error among all curves
```

```
        father(p,1) = 1; All 1-segment polylines have father point 1
```

```
    end; for p
```



```

Compute s-segm-polyline cost from point 1 to all other points
for s = 2:S           Number of segments in the polyline
  for p = 3:P          Next-point-number to consider
    minErr = cost(p-1,s-1); minIndex = p-1; Error of (p-1,p) = 0
    for i = s:p-2        Next-intermediate-point to consider
      if (error(i,p) == Inf)   Error of line segment (i,n) not cashed
        for c = 1:C          Next curve-number to consider
          se(c) = segmentError(x(i:p), y(k,i:p)); (i,p)-segment error
        end; for k
        error(i,p) = max(se); Maximum line segment error
      end; if
      currErr = cost(i,s-1) + error(i,p); s-segment-polyline error
      if (currErr < minErr)           Smaller error?
        minErr = currErr; minIndex = i; Update error and parent
      end; if
    end; for i
    cost(p,s) = minErr;       s-segment-polyline minimal cost
    father(p,s) = minIndex;  Last point's father on the polyline
  end; for p
end; for s
[e,a,b,xb] = ExtractAnswer;
end

```



function [e,a,b] = segmentError(x,y)

Input:

x,y: Digitized curve-segment as an x-vector and an y-vector

Output:

e: Error of the line segment between the first and last point

a,b: The coefficients defining this segment

Initialization:

z = size(x); P = s(2); Find out the number of points of x,y

Compute 1-segment linear-interpolation of (x,y) coefficients

a = (y(n) - y(1)) / (x(n) - x(1));

b = (y(1) * x(n) - y(n) * x(1)) / (x(n) - x(1));

Compute perpendicular-distance error for above line segment

e = 0; Initialize Error

for p = 1:P Compute error for the each point on the curve

e = e + (y(p) - a * x(p) - b)² / (a²+1); Accumulate least square

end;

end



```

function [e,a,b,xb] = ExtractAnswer
Output:
  e,a,b,xb: As in the output of optimalLinearApproximation
Initialization:
ib = zeros(S,S+1); xb = zeros(S,S+1); yb = zeros(C, S,S+1); Points matrices
a = zeros(C, S,S); b = zeros(C, S,S); er = zeros(C, S,S); Coefficients/error
Extract error and coefficient matrices
for s = S:-1:1    Traverse polyline segments in inverse order
  ib(s,s+1) = P;           Get last point number
  xb(s,s+1) = x(ib(s,s+1)); Get x-value for this point
  for c = 1:C    Traverse all curves
    yb(c,s,s+1) = y(c,ib(s,s+1)); Get y-value for this point
  end;
  for i = s:-1:1    Traverse predecessor points in inverse order
    ib(s,i) = father(ib(j,i+1),i);  Get predecessor point number
    xb(s,i) = x(ib(s,i));           Get x-value for this point
    for c = 1:C    Traverse all curves
      yb(c,s,i) = y(c,ib(s,i));    Get y-value for this point
      [er(c,s,i), a(c,s,i), b(c,s,i)] = Compute err, a and b for segm (x,y)
      segmentError( x(ib(s,i):ib(s,i+1)), y(c,ib(s,i):ib(s,i+1)) );
    end
  end;
end;
end;

```