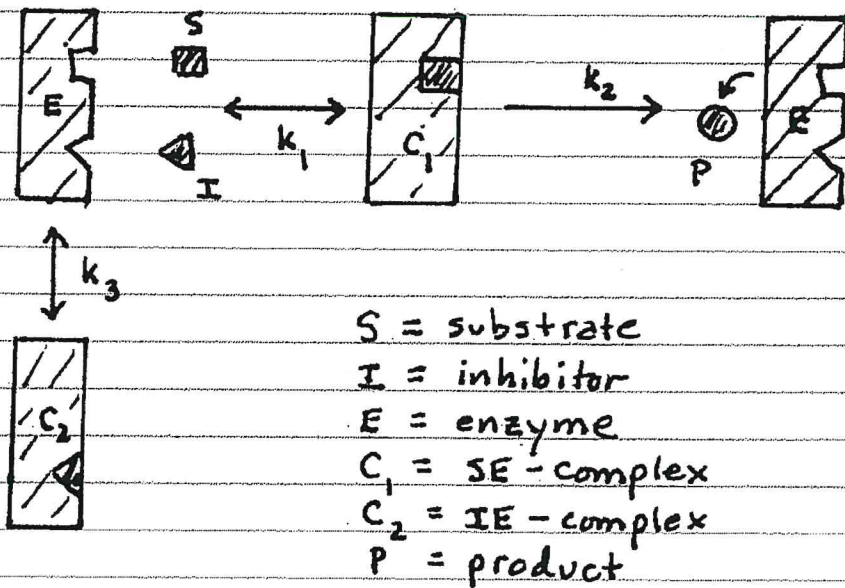


Competitive Inhibition



Enzyme has two binding sites but only SE complex can make product P so I acts to inhibit such production.

Law of mass action ($E = E_0/s_0 \ll 1$ assumed)

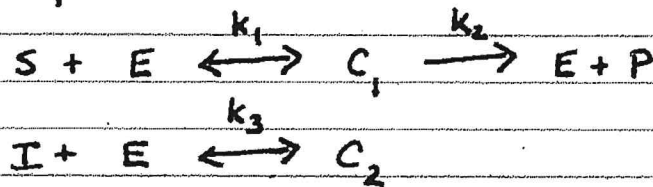
$$\left. \begin{aligned} (1) \quad S' &= -k_1 SE + k_{-1} C_1 \\ (2) \quad I' &= -k_3 IE + k_{-3} C_2 \end{aligned} \right\} \text{slow}$$

$$\left. \begin{aligned} (3) \quad C_1' &= k_1 SE - (k_{-1} + k_2) C_1 \\ (4) \quad C_2' &= k_3 IE - k_{-3} C_2 \end{aligned} \right\} \text{fast}$$

$$(5) \quad E' = -C_1' - C_2' \quad \text{conserved receptors}$$

$$(6) \quad P' = k_2 C_1 \quad \text{production rate}$$

Reaction Equations



With nondimensionalizing we know (3)-(4) will be fast. If E, C_k are scaled by E_0 and S by S_0 the dimensionless forms will be

$$E, C_k = O(1) \quad \epsilon = \frac{E_0}{S_0} \ll 1$$

Quasi Steady State

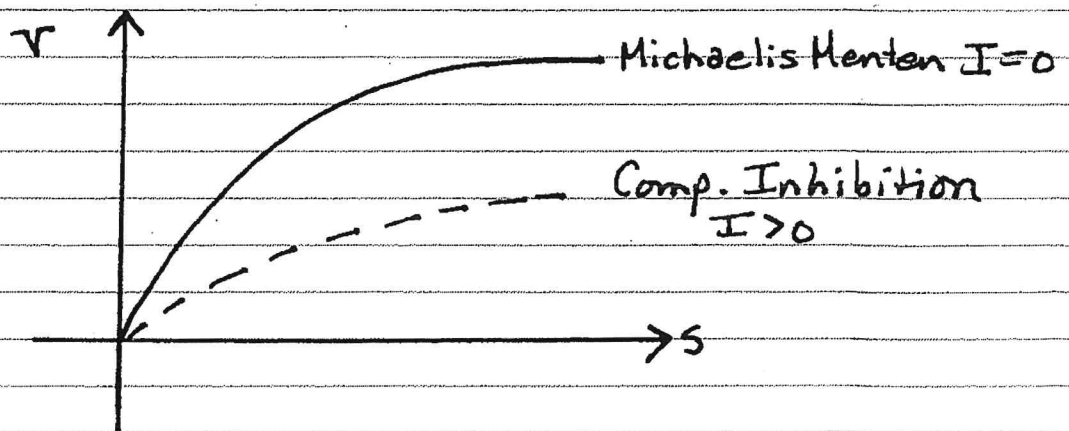
Set $C_1' = C_2' = 0$ and use conservation law (5) to solve for C_1 in terms of slow variables S, I to ultimately get production rate

$$\frac{dP}{dt} = v = \frac{\bar{v} S}{S + K_m (1 + I/K_i)}$$

↑ inhibition

where $\bar{v} = k_2 E_0$ maximal rate and

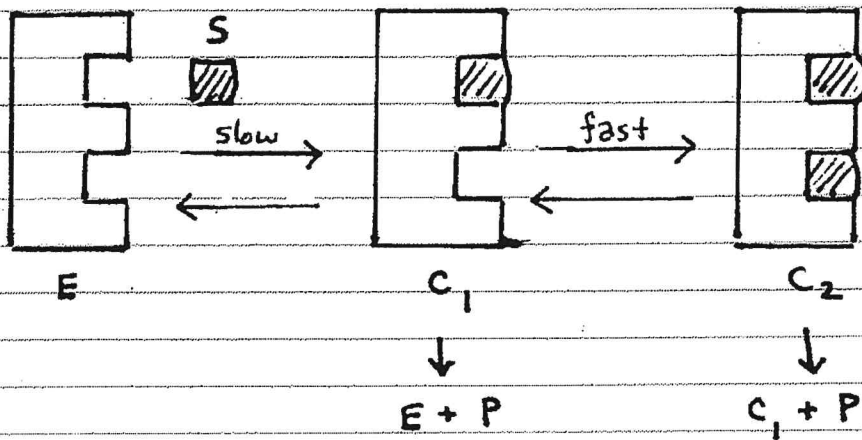
$$K_m = \frac{k_{-1} + k_2}{k_1} \quad K_i = \frac{k_{-3}}{k_3}$$



Cooperativity



Cartoon



Indicated rates k_1 and k_3 are, respectively, assumed to be small and large. System favors cooperative C_2 state.

QSS Approximation, conservation receptors, $\epsilon = \frac{E_0}{S_0} \ll 1$

$$\bar{V} = \frac{dP}{dt} = \frac{(k_2 K_2 + k_4 S') E_0 S'}{K_1 K_2 + K_2 S + S'^2}$$

where

$$K_1 = \frac{k_{-1} + k_2}{k_1} \quad K_2 = \frac{k_4 + k_{-3}}{k_3}$$

Large Positive Cooperativity

As indicated we assume (see cartoon) that k_1 is relatively small while k_3 is relatively large:

$$k_1 = \delta \ll 1 \quad k_3 = \frac{k}{\delta} \gg 1$$

for some constant k . Then V is a function of S

$$V = V(S) \quad \delta \ll 1$$

Since $V \sim V(0)$ we have an approximate production rate

$$(1) \quad V = \frac{\bar{V} S^2}{K + S^2} \quad K = K_1 K_2$$

where $\bar{V} = k_2 E_0$. $V(S)$ is a Hill function of Hill coefficient $n=2$.

n-site Positive Cooperativity

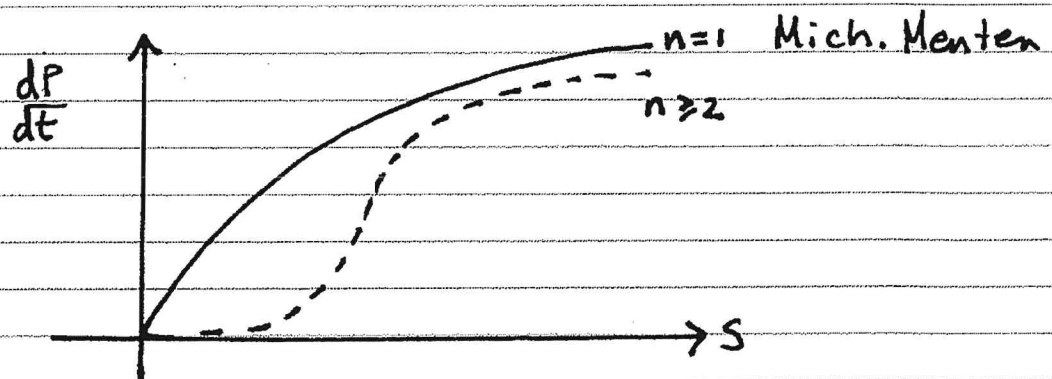
Assume formation of C_1 is slow ($k_1 \ll 1$)
but formation C_k , $k \geq 1$ fast.

Here E has n -sites and C_k = complex
of E with k -substrate molecules
attached.

Same analysis

$$\frac{dP}{dt} = v = \frac{\bar{V} S^n}{K + S^n} \quad \text{Hill eqn, coeff } n$$

$$\text{where } K = \prod_{j=1}^n K_j$$



Experimental Correlation

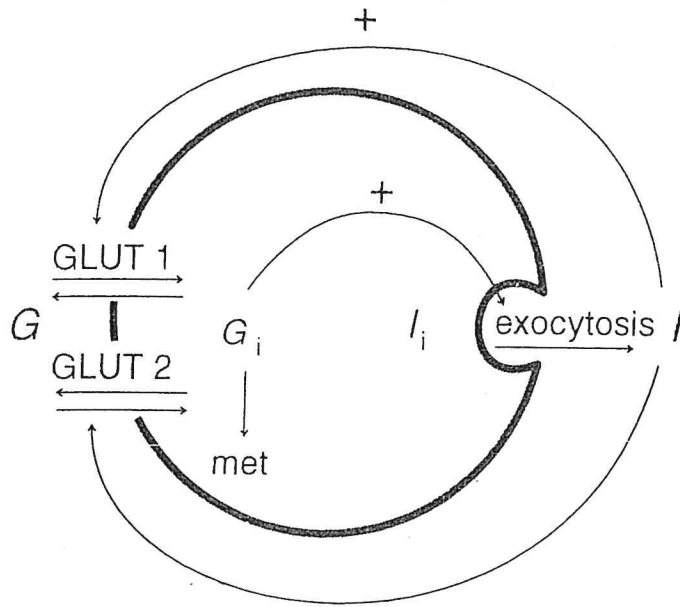
$$n \ln S = n \ln K + \ln \left(\frac{v}{v - \bar{v}} \right)$$

$$n \alpha_k = n \ln K + \beta_k$$

Data $\{(\alpha_k, \beta_k)\}$ and linear regression for n , K .

Glucose-dependent insulin secretion

Pancreatic β -cells secrete insulin I via exocytosis at a rate dependent on intra and extracellular glucose.



The following rates and assumptions apply

R_1 = rate of transport into cell through insulin activated GLUT 1 transporters

R_2 = rate of transport into cell through insulin inactivated GLUT 2 transporters

R_s = insulin production rate

R_m = intracellular glucose metabolism rate

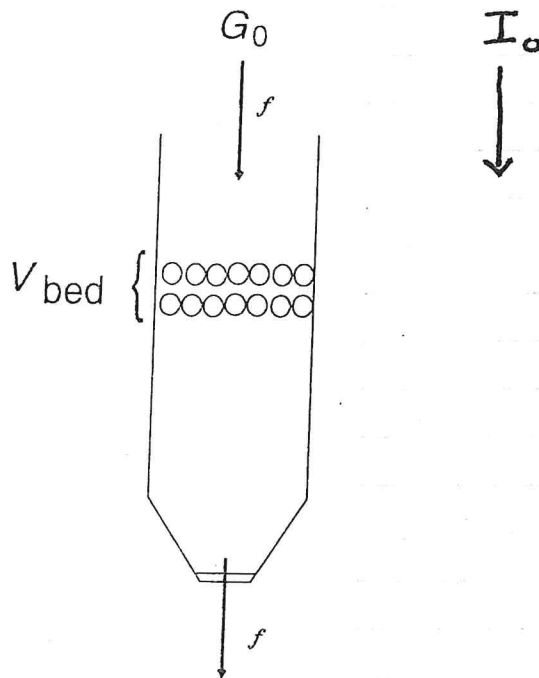
G = extracellular glucose

G_i = intracellular glucose

I = insulin

J = inhibition variable

Experimentally β -cells in a bed (V_{bed}) and glucose of concentration G_0 is continuously flushed through at flow rate f :



Controlled: inflow G_0, I_0

measured: outflow G, I

Nondimensionalized model (Fall et al, pg 87)

$$(1) \quad g' = -R_1 - R_2 - k_0 (g - g_0)$$

$$(2) \quad g_i' = R_1 + R_2 - R_m$$

$$(3) \quad i' = R_s - k_0 i$$

$$(4) \quad j' = j_\infty(i) - j \quad (\text{phenomenological})$$

where

$$R_1 = \frac{K_1 V_1 (g - g_i)}{(K_1 + g)(K_1 + g_i)} \cdot \frac{i^n \oplus}{K_i^n + i^n}$$

$$R_2 = \frac{K_2 V_2 (g_j^m \ominus - g_i)}{(K_2 + g)(K_2 + g_i)}$$

$$R_m = \frac{V_m g_i}{1 + g_i}$$

Mich-Menten QSS

$$R_s = \frac{V_s (R_m^4 + L^4)}{K_s^4 + R_m^4 + L^4}$$

(phenomenological)

The latter assumes insulin secretion rate that grows with glucose metabolism rate. Looks like 4-site cooperativity (were it not for L). Lastly

$$j_\infty(i) = \frac{1}{1+i}$$

Model Parameters

Table 4.1 Standard Dimensional Parameters

Fixed by experiment	V_m	0.24 mM/min
	K_m	9.8 mM
	V_s	0.034 mM/min
	K_s	0.13 mM/min
	V_{max2}	32.0 mM/min
	K_2	17.0 mM
	V_{max1}	120.0 mM/min
	K_1	1.4 mM
Experimentally variable	L	0.01 mM/min
	k_0	400.0/min
	l_0	0.0 mM
Adjustable in the model	G_0	8-22 mM
	K_l	1×10^{-6} mM
	K_i	4.0×10^{-5} mM
	τ	20.0 min

Table 4.2 Standard Dimensionless Parameters

Dimensionless parameter	Dimensional definition	Standard value
\hat{V}_m	$\tau V_m / K_m$	0.50
\hat{V}_s	$\tau V_s / K_l$	6.8×10^5
\hat{K}_s	$\tau K_s / K_m$	0.27
\hat{V}_{max2}	$\tau V_{max2} / K_m$	65.3
\hat{K}_2	K_2 / K_m	1.7
\hat{V}_{max1}	$\tau V_{max1} / K_m$	245.0
\hat{K}_1	K_1 / K_m	0.14
\hat{L}	$\tau L / K_m$	0.02
\hat{k}_0	τk_0	8×10^3
\hat{G}_0	G_0 / K_m	0.8-2.2
\hat{K}_i	K_i / K_l	40.0

large

large

Fast-slow dynamics

Since $k_0 \gg 1$ in

$$g' = -R_1 + R_2 - k_0(g - g_0)$$

$$i' = R_5 - k_0 i$$

yields (QSS) approximations:

$$0 = -R_1 + R_2 - k_0(g - g_0)$$

$$0 = R_5 - k_0 i$$

Using these one can eliminate (g, i) to get a slow subsystem of the form

$$g_i' = F_1(g_i, j)$$

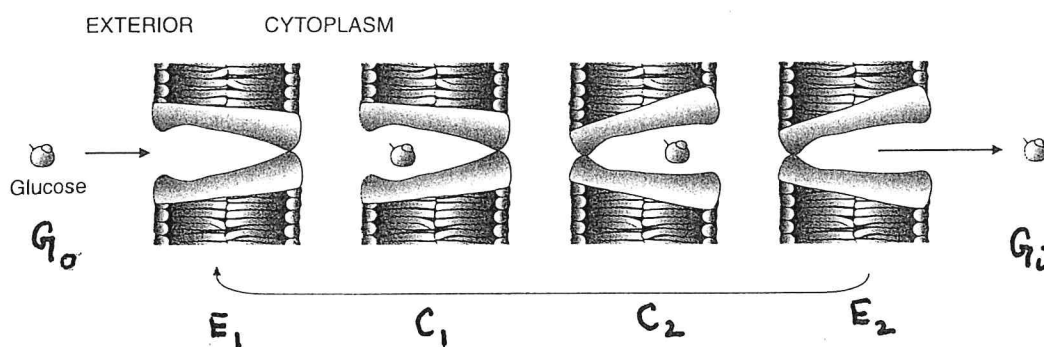
$$j' = F_2(g_i, j)$$

This system has oscillations!

Passive Transport

Transport of S from high to low concentrations through a membrane transporter such as extracellular glucose into cells. No energy input is required.

A



B

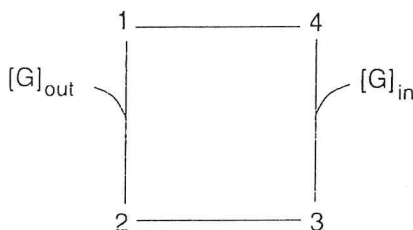


Figure 3.1 (A) Cartoon of four states of a GLUT transporter, showing the empty pore facing the exterior of the cell, glucose bound facing the exterior, glucose bound facing the interior, and the open pore facing the interior of the cell. Adapted from Leinhard et al. (1992). (B) Four-state kinetic diagram of a GLUT transporter based on the cartoon in (A).

- E_1 = one unbound conformational state
- C_1 = one bound conformational state
- C_2 = another bound conformational state
- E_2 = another unbound conformational state

(QSS)

$$V = \frac{dG_i}{dt} = \frac{\alpha_0 + \alpha_1 G_o + \alpha_2 G_i + \alpha_3 G_o G_i}{\beta_0 + \beta_1 G_o + \beta_2 G_i + \beta_3 G_o G_i}$$

SERCA Pumps

$$[Ca^{2+}]_o > [Ca^{2+}]_i$$

pump Ca^{2+}
up gradient

Rate of Ca^{2+} accumulation inside cell measured by rate of radioactive $^{45}Ca^{2+}$ uptake at ER (endoplasmic reticulum)
Conversion of ATP to ADP generates energy (loss of PO_4 phosphate group)

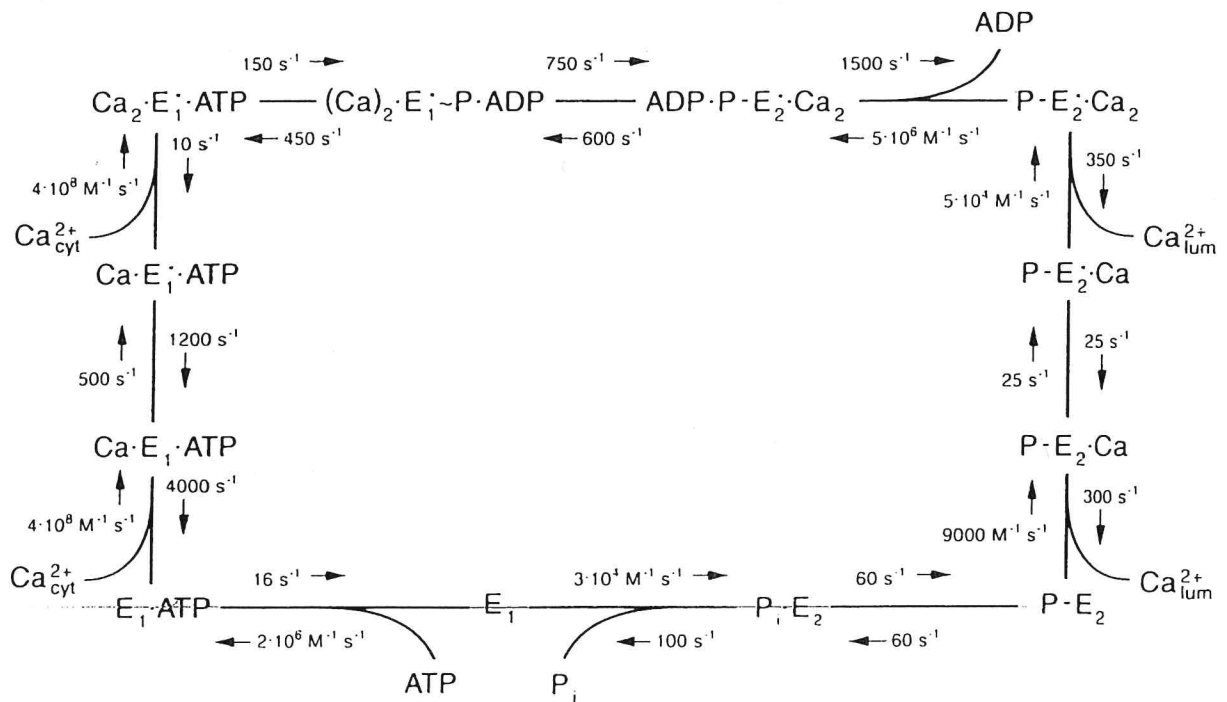


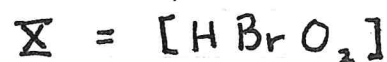
Figure 3.14 A twelve-state model of the SERCA pump. Note the two sequential Ca^{2+} binding steps on the left-hand side. Although the cycle is driven by the hydrolysis of ATP, all of the steps in the diagram contribute to the steady-state rate. Redrawn from Luger (1991).

Twelve State model well approximated by 2-site cooperativity model hence intracellular production rate (QSS)

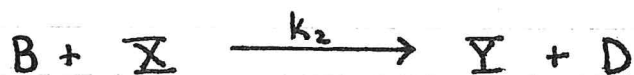
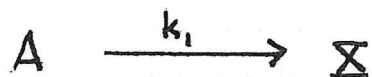
$$V = \frac{dCa_i}{dt} = \frac{\bar{V} [Ca_i]^2}{K^2 + [Ca_i]^2}$$

Chemical Oscillators - Brusselator (1968)

Mostly inorganic chemical species.



where the reactions are



Here D and E are other products.

It is assumed that the concentrations of A, B are artificially maintained at (nearly) constant levels. Resulting nondimensionalized eqns are called the Brusselator

Autocatalysis since X involved in its own production.

$$(1) \quad \dot{x} = a - (b+1)x + x^2 y$$

$$(2) \quad \dot{y} = bx - x^2 y$$

where (x, y) are dimensionless (\bar{X}, \bar{Y}) .

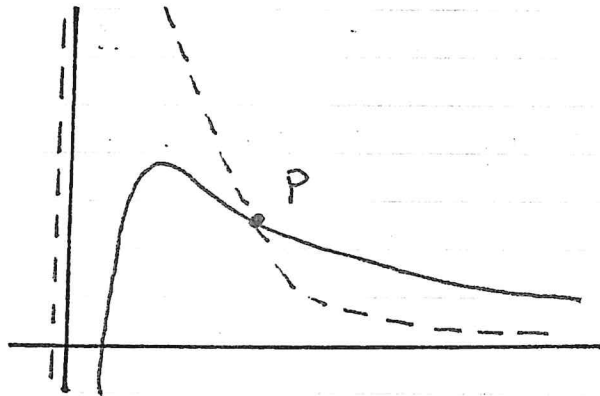
Nullclines, equilibria

$$\dot{x} = 0 \quad y = F(x) = \frac{(b+1)}{x} - \frac{a}{x^2}$$

$$\dot{y} = 0 \quad x = 0, \quad y = \frac{b}{x}$$

Sole equilibria

$$P = \left(a, \frac{b}{a}\right)$$



$$\dot{y} = 0$$

$$\dot{x} = 0$$

Jacobian at P

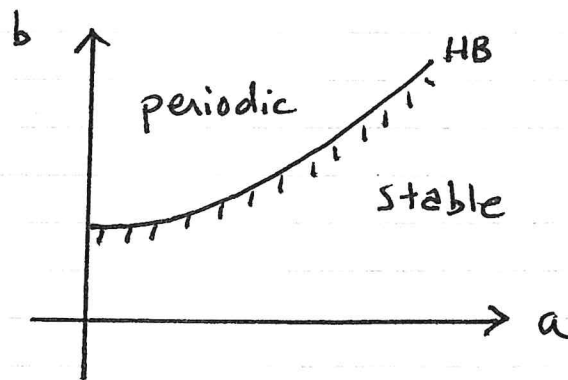
$$DF(P) = \begin{bmatrix} b-1 & a^2 \\ -b & -a^2 \end{bmatrix}$$

from which

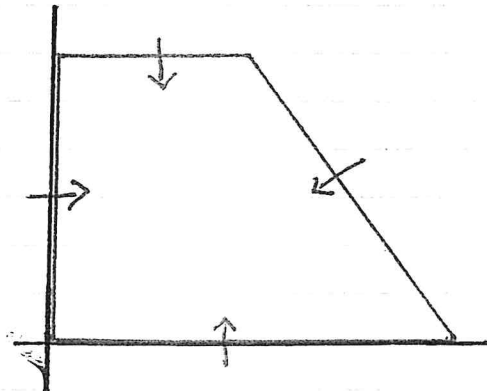
$$\det DF(P) = a^2 > 0$$

$$\text{Tr } DF(P) = b-1-a^2$$

Hopf bifurcations occur when $\text{Tr } DF(P)$



Hopf points alone only prove small amplitude periodic orbits



Trapping region
shape to prove
existence
given $\text{Tr } DF(P) > 0$