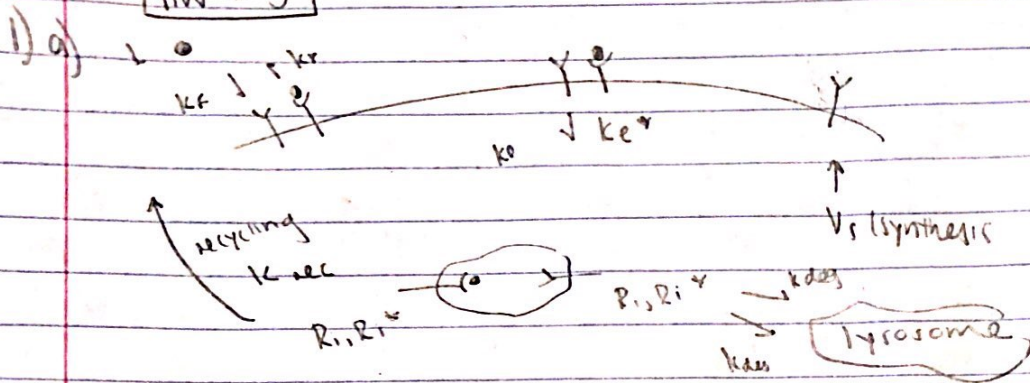


HW # 5

mass balances

free surface receptor: $\frac{dR_s}{dt} = -k_f L R_s + k_r R_s^* - k_i R_s + V_s + k_{rec} R_i^T$ (1)

bound surface receptor: $\frac{dR_s^*}{dt} = k_f L R_s - k_r R_s^* - k_i^* R_s^* + k_{rec} R_i^T$ (2)

total internal receptor: $\frac{dR_i^T}{dt} = k_i R_s + k_i^* R_s^* - k_{deg} R_i^T - k_{rec} R_i^T$ (3)

where $R_i^T = R_i^s + R_i^*$

internal active receptor: $\frac{dR_i^*}{dt} = k_i^* R_s^* - k_{deg} R_i^* - k_{rec} R_i^*$ (4)

at SS: $k_{rec} R_i + k_{rec} R_i^* = k_{rec} R_i^T$

$\frac{dR_s}{dt} = 0 = -k_f L R_s + k_r R_s^* - k_i R_s + V_s + k_{rec} R_i$

$\frac{dR_s^*}{dt} = 0 = k_f L R_s - k_r R_s^* - k_i^* R_s^* + k_{rec} R_i^*$

Following the same steps to derive the solution as done in class

@ SS $k_{deg} R_i^T = V_s$
from eqn(3)

$0 = k_i R_s + k_i^* R_s^* - V_s - k_{rec} R_i^T$

$0 = k_i R_s + k_i^* R_s^* - V_s - k_{rec} R_s - k_{rec} R_s^*$

Solve for R_s^*

$$k_e R_s - k_{rec} R_s = V_s + k_{rec} R_s^* - k_e^* R_s^*$$

$$R_s = \frac{V_s + k_{rec} R_s^* - k_e^* R_s^*}{k_e - k_{rec}}$$

plug into eqn (2)

$$0 = k_f L \left(\frac{V_s + k_{rec} R_s^* - k_e^* R_s^*}{k_e - k_{rec}} \right) - k_r R_s^* - k_e^* R_s^* + k_{rec} R_s^*$$

Solve for R_s^*

$$0 = \frac{k_f L V_s}{k_e - k_{rec}} + \frac{k_f L k_{rec} R_s^*}{k_e - k_{rec}} - \frac{k_f L k_e^* R_s^*}{k_e - k_{rec}} - k_r R_s^* - k_e^* R_s^* + k_{rec} R_s^*$$

$$\frac{k_f L V_s}{k_e - k_{rec}} = R_s^* \left[\frac{k_f L k_e^*}{k_e - k_{rec}} - \frac{k_f L k_{rec}}{k_e - k_{rec}} + \frac{(k_r + k_e^* - k_{rec})(k_e - k_{rec})}{k_e - k_{rec}} \right]$$

$$L R_s^* = \frac{k_f L V_s}{k_e - k_{rec} + (k_r + k_e^* - k_{rec})(k_e - k_{rec})}$$

$$k_f L k_e^* - k_f L k_{rec} + (k_e - k_{rec})(k_r + k_e^* - k_{rec})$$

divide everything by this term

$$R_s^* = \frac{k_f L V_s}{(k_e - k_{rec})(k_r + k_e^* - k_{rec}) + k_f L (k_e^* - k_{rec})}$$

Use $K_{ss} = \frac{k_f (k_e^* - k_{rec})}{(k_e - k_{rec})(k_r + k_e^* - k_{rec})}$

$$R_s^* = \frac{K_{ss} L}{1 + K_{ss} L} \left(\frac{V_s}{k_e^* - k_{rec}} \right)$$

Use eqn (4) to get R_s^*

$$0 = k_e^+ R_s^+ - k_{deg} R_i^+ - k_{rec} R_i^+$$

$$k_{deg} R_i^+ + k_{rec} R_i^+ = k_e^+ R_s^+$$

$$R_i^+ = \frac{k_e^+ R_s^+}{k_{deg} + k_{rec}}$$

$$R_{total}^+ = R_s^+ + R_i^+$$

$$= R_s^+ \left(1 + \frac{k_e^+}{k_{deg} + k_{rec}} \right)$$

$$= \frac{k_{ss} L}{1 + k_{ss} L} \left(\frac{V_s}{k_e^+ - k_{rec}} \right) \left(1 + \frac{k_e^+}{k_{deg} + k_{rec}} \right)$$

$$R_{total}^+ = \underbrace{\left(\frac{1}{k_e^+ - k_{rec}} + \frac{k_e^+}{(k_e^+ - k_{rec})(k_{deg} + k_{rec})} \right)}_{\text{distance to activation}} \underbrace{\left(\frac{k_{ss} L}{1 + k_{ss} L} \right)}_{\text{bound fraction (ranges from 0 to 1)}} \underbrace{V_s}_{\text{synthesis}}$$

max occurs when bound fraction = 1

$$R_{total}^{+, \max} = \left(\frac{1}{k_e^+ - k_{rec}} + \frac{k_e^+}{(k_e^+ - k_{rec})(k_{deg} + k_{rec})} \right) V_s$$

compare this to process w/o recycling

$$R_{total}^{+, \max} = \left(\frac{1}{k_e^+} + \frac{1}{k_{deg}} \right) V_s$$

looking at the first term, $\frac{1}{k_e^+}$ vs $\frac{1}{k_e^+ - k_{rec}}$

Recycling increases the max concentration of active receptors by keeping them in circulation which can increase overall binding

$$2) \quad \frac{dC_a}{dt} = -d_a C_a + \frac{r_{oa} + r_a C_a^2}{1 + C_a^2 + C_r^2}$$

$$\frac{dC_r}{dt} = -C_r + \frac{r_{or} + r_r C_a^2}{1 + C_a^2}$$

at SS: $\frac{dC_a}{dt} = \frac{dC_r}{dt} = 0$

$$(1) \quad d_a C_a = \frac{r_{oa} + r_a C_a^2}{1 + C_a^2 + C_r^2} \quad (2) \quad C_r = \frac{r_{or} + r_r C_a^2}{1 + C_a^2}$$

a) from this, we can draw the following conclusions

- ① R has no effect on R (no C_r on right side of eqn 2)
- ② A is an activator of R (indicated by increasing sigmoidal shape) ^{eqn 2}
- ③ R is an inhibitor of A (only appears in denominator of eqn 1)
- ④ A is an activator of A (sigmoidal shape of eqn 1 right side)

Basal rates:

$$C_A = C_R = 0$$

$$\frac{dC_a}{dt} = r_{oa}$$

$$\frac{dC_r}{dt} = r_{or}$$

d_a is a degradation constant for C_a

Maximal rates

$$\frac{dC_a}{dt} \rightarrow \text{max occurs when } C_r = 0$$

$$\frac{dC_a}{dt} = -d_a C_a + \frac{r_{oa} + r_a C_a^2}{1 + C_a^2}$$

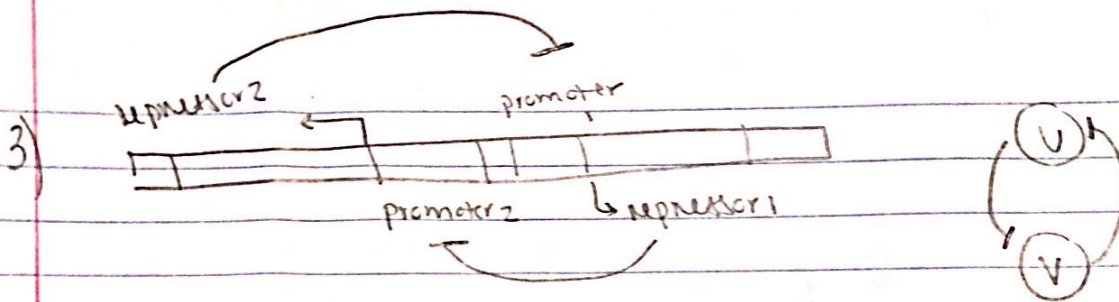
(b) used 'PS5_2_B.jl' and 'PhasePortrait V2.jl' to generate plot 1. The fixed point is where the two nullclines (red lines) intersect. Since the phase portrait shows that the vectors/trajectories do not all point towards this point, the fixed point is unstable.

(c) see plot 1

d) In the plot there is no clear relationship between C_r and C_n . This is due to the oscillatory nature of the system. There is a peak value of both C_n & C_r . The loop-like structure of the trajectory plotted for $C_{a0} = 1$ and $C_{r0} = 10$ exemplifies the oscillatory pattern that the concentrations follow over time.

e) see file "PS5N2e.m"

I attempted to use this code to solve the ODE system and plot. For some reason, the MATLAB solver was unable to solve this ODE system, though I tried it with other differential equations and the code worked.



$$\frac{dU}{dt} = \frac{\alpha}{1+V^n} - U = f(U, V)$$

$$\frac{dV}{dt} = \frac{\alpha}{1+U^n} - V = g(U, V)$$

a)

U, V = (i) conc of a repressor or gene expression

$\frac{dU}{dt}, \frac{dV}{dt}$ = (ii) effective rate of synthesis of repressor

n = (iii) cooperativity of repression

1 = (iv) degradation constant

α = effective rate of synthesis (according to Collins paper)

b + c) see "PS 5 Plots"

Linear stability analysis

d) (1) Find steady states: $f(U_s, V_s) = g(U_s, V_s) = 0$

(2) linearize system around steady state of interest for perturbations:

$$U = u - U_s$$

$$V = v - V_s$$

$$\frac{dU}{dt} = f(U_s, V_s) + \frac{df}{dU} \bigg|_{U_s, V_s} U + \frac{df}{dV} \bigg|_{U_s, V_s} V$$

$$\frac{dU}{dt} = -U - \frac{\alpha n V_s^{n-1}}{(V_s^n + 1)^2} V \bigg|_{U_s, V_s}$$

$$\frac{dU}{dt} = -U - \frac{\alpha n V_s^{n-1}}{(V_s^n + 1)^2} V$$

$$\frac{dV}{dt} = g(u_r, v_r) + \left. \frac{dg}{du} \right|_{u_r, v_r} U + \left. \frac{dg}{dv} \right|_{u_r, v_r} V$$

$$\frac{dV}{dt} = - \left. \frac{\alpha n u^{n-1}}{(u^n + 1)^2} \right|_{u_r, v_r} U - V$$

$$\frac{dV}{dt} = - \frac{\alpha n v_r^{n-1}}{(u_r^n + 1)^2} U - V$$

In matrix form

$$\dot{\vec{x}} = \vec{J} \vec{x}$$

where

$$\vec{x} = \begin{pmatrix} U \\ V \end{pmatrix},$$

$$\vec{J} = \begin{pmatrix} -1 & \dots & -\frac{\alpha n v_r^{n-1}}{(v_r^n + 1)^2} \\ -\frac{\alpha n u_r^{n-1}}{(u_r^n + 1)^2} & \dots & -1 \end{pmatrix}$$

* see next page for the rest of part d
jumping to (e)

$\alpha = 10$, $n = 1$ and 2

find numerical value of untr steady state ($u=v$)

$n=1$

$$\frac{10}{1+u} - u = 0$$

$$u = \frac{1}{2}(\sqrt{41} - 1) \approx 2.7016$$

(consistent w/ graph
from part b)

$n=2$

$$\frac{10}{1+u^2} - u = 0$$

$$u = 2 \quad (\text{consistent w/ part b})$$

Use MATLAB eig function to find λ 's
(see P55N3E3m)

for $n=1$

$$\lambda = -1.7298, -0.2702$$

both \ominus \therefore stable

2 d) Finding eigenvalues

$$\vec{J}\vec{x} = \lambda\vec{x} \Rightarrow (\vec{J} - \lambda\vec{I})\vec{x} = 0$$

Non trivial soln if

$$\det(\vec{J} - \lambda\vec{I}) = |\vec{J} - \lambda\vec{I}| = 0 \rightarrow \text{stability eqn}$$

$$\begin{vmatrix} -1 - \lambda & \frac{-\alpha n v_r^{n-1}}{(v_r^n + 1)^2} \\ \frac{-\alpha n u_r^{n-1}}{(u_r^n + 1)^2} & -1 - \lambda \end{vmatrix} = 0$$

$$\Rightarrow (-1 - \lambda)(-1 - \lambda) - \left[\frac{-\alpha n v_r^{n-1}}{(v_r^n + 1)^2} \right] \left[\frac{-\alpha n u_r^{n-1}}{(u_r^n + 1)^2} \right] = 0$$

$$\lambda_{\pm} = \frac{\text{tr}(\vec{J}) \pm \sqrt{\text{tr}(\vec{J})^2 - 4\det(\vec{J})}}{2}$$

where $\text{tr}(\vec{J}) = -2$

$$\det(\vec{J}) = 1 - \left(\frac{\alpha n v_r^{n-1}}{(v_r^n + 1)^2} \right) \left(\frac{\alpha n u_r^{n-1}}{(u_r^n + 1)^2} \right)$$

$$\lambda = \frac{-2 \pm \sqrt{-2 - 4 \left(1 - \left(\frac{\alpha n v_r^{n-1}}{(v_r^n + 1)^2} \right) \left(\frac{\alpha n u_r^{n-1}}{(u_r^n + 1)^2} \right) \right)}}{2}$$

2

for $n = 2$

$$\lambda = -2.65 \cdot 0.6$$

not both $\ominus \therefore$ unstable

change in cooperativity changes
the stability of the steady state of interest

$$f1) \quad \frac{dR_1^*}{dt} = k_f L R_1 - k_r R_1^* \quad \frac{dR_2^*}{dt} = k_f L R_2 - k_r R_2^*$$

$$\frac{dN_1^*}{dt} = k_f^{ND} N_1 D_2 - k_r^{ND} N_1^* \quad \frac{dN_2^*}{dt} = k_f^{ND} N_2 D_1 - k_r^{ND} N_2^*$$

$$\frac{dD_1}{dt} = k_D R_1^* - \gamma_D D_1$$

$$\frac{dD_2}{dt} = k_D R_2^* - \gamma_D D_2$$

$$\frac{dR_1}{dt} = \frac{B^n}{K^n + N_1^{*n}} - \gamma_R R_1$$

$$\frac{dR_2}{dt} = \frac{B^n}{K^n + N_2^{*n}} - \gamma_R R_2$$

since (3)-(5) assume fast

equilibrium

$$\frac{dR_1^*}{dt}, \frac{dR_2^*}{dt}, \frac{dN_1^*}{dt}, \frac{dN_2^*}{dt}, \frac{dD_1}{dt}, \frac{dD_2}{dt} = 0$$

$$0 = k_f^{ND} N_1 D_2 - k_r^{ND} N_1^*$$

$$N_1^* = \frac{k_f^{ND} N_1 D_2}{k_r^{ND}} \quad (N_1 = \text{const})$$

$$0 = k_D R_2^* - \gamma_D D_2$$

$$D_2 = \frac{k_D R_2^*}{\gamma_D}$$

$$0 = k_f L R_2 - k_r R_2^*$$

$$R_2^* = \frac{k_f L R_2}{k_r}$$

$$D_2 = \frac{k_D k_f L R_2}{\gamma_D k_r}$$

$$N_1^* = \frac{k_f^{ND} N_1 k_D k_f L R_2}{k_r^{ND}}$$

$$\frac{dR_1}{dt} = \frac{B^n}{K^n + \left(\frac{k_f^{ND} N_1 k_D k_f L R_2}{k_r^{ND}} \right)^n} - \gamma_R R_1$$

similar expression for $\frac{dR_2}{dt}$

$$\frac{dR_2}{dt} = \frac{B^n}{k^n + \left(\frac{k_f^{ND} N_2 k_D k_f R_1}{k_r^{ND}} \right)^n} - \gamma_R R_2$$

f2) non dimensional variables

$$u = \frac{R_1}{K}, \quad v = \frac{R_2}{K}, \quad \tau = \gamma_R t$$

$$\frac{dR_1}{dt} = \left(\frac{B^n}{k^n + \left(\frac{k_f^{ND} N_2 k_D k_f R_2}{k_r^{ND}} \right)^n} - \gamma_R R_1 \right)$$

$$\frac{1}{K} \frac{dR_1}{dt} = \left(\frac{B^n}{k^n + \left(\frac{k_f^{ND} N_2 k_D k_f R_2}{k_r^{ND}} \right)^n} - \gamma_R R_1 \right) \frac{1}{K}$$

$$\frac{dU}{dt} = \frac{B^n}{k^{n+1} + \left(\frac{k_f^{ND} N_2 k_D k_f L}{k_r^{ND}} \right)^n} V^n - \gamma_R U$$

$$\frac{dU}{d\tau} = \frac{B^n}{k^{n+1} + \left(\frac{k_f^{ND} N_2 k_D k_f L}{k_r^{ND}} \right)^n} V^n - \gamma_R U$$

B (concⁿ)
time

k conc

N_2 conc