

More IVP's:

Circadian rhythms.

"Circadian" means "occurring naturally on a 24-hour cycle."

One model (see Tyson et al. (1999), on our Canvas page) says: circadian rhythms are regulated by feedback of two proteins, PER, (short for "periodic") and TIM (short for "timeless"), on the "per" and "tim" mRNA that yield these proteins.

Variables:

$M = \text{per/tim mRNA}$

$P_1 = \text{PER/TIM monomers (basic protein building blocks)}$

$P_2 = \text{PER/TIM dimers (a dimer is built up from two monomers)}$

(A more detailed study might have six variables, by not grouping per and tim, or PER and TIM, together. But the above three are enough for us.)

The model says:

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$$\begin{aligned}
 \frac{dM}{dt} &= \frac{\overset{(i)}{a}}{1 + bP_2^2} - \overset{(ii)}{c}M \\
 \frac{dP_1}{dt} &= \overset{(iii)}{d}M - \frac{\overset{(iv)}{e}P_1}{f + P_1 + gP_2} - \overset{(v)}{h}P_1 - \overset{(vi)}{2k}P_1^2 + \overset{(vii)}{2l}P_2 \\
 \frac{dP_2}{dt} &= \frac{\overset{(viii)}{m}P_2}{f + P_1 + gP_2} - \overset{(ix)}{n}P_2 + \overset{(x)}{k}P_1^2 - \overset{(xi)}{l}P_2.
 \end{aligned}$$

(CR) {

(a, b, c, ..., l are positive parameters.)

Analysis of terms:

(i) P_2 inhibits production of M : (i) tells us that more P_2 means slower growth of M .

Note the P_2^2 here: in other situations, one might have P_2^h where h , the "Hill coefficient," reflects certain binding properties of the proteins.

(ii) M is degraded/used up at a rate proportional to the amount of M present.

(iii) P_1 is produced at a rate proportional to the amount of M present.

(iv) and (viii) represent phosphorylation: P_1 and P_2 combine with phosphates and are inactivated. Note that the decay implied by the numerators of (iv) and (viii) is tempered by the denominators.

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(v) and (ix) represent proteolysis: P_1 and P_2 decay (proteins have half-lives).

(vi) and (x) represent the reaction
monomer + monomer \rightarrow dimer.
[Note that (vi) cancels twice (x).]

(vii) and (xi) represent the reaction
dimer \rightarrow monomer + monomer.
[Note that (vii) cancels twice (xi).]

COOL FACTS:

(1) Solutions to (CR) are normally circadian, but

(2) A certain mutation per^L of per mRNA changes the "dimerization rate" k in (vi) and (x) above, resulting in M, P_1, P_2 having periods > 24 hours!

(3) Equations (CR) can further be reduced, under certain conditions, by combining P_1 and P_2 into a single protein variable P . See Sage worksheet Circadian.sws (on the Sage Page).