



Coupling Hair Follicle Cycles to Produce Traveling Waves

Megan Jeans ^{1,2} and Dr. G. Bard Ermentrout ³

¹Bioengineering and Bioinformatics Summer Institute, Department of Computational Biology, University of Pittsburgh, PA 15261
²Department of Bioengineering, Rice University, Houston, TX 77005
³Department of Mathematics, University of Pittsburgh, Pittsburgh, PA 15261

Examples of Biological Synchronization

- Pacemaker cells in the heart
- Discharging of brain cells during epileptic seizures
- Women's menstrual cycles

Hair growth in rodents – motivation for this study



Suzuki, *et al.*

The Hair Follicle Cycle

- Begins with catagen apoptosis
- Telogen rest, exogen usually occurs in this phase
- Anagen growth, longest phase



www.keratin.com/aa/aa008.shtml

Objectives

- Simulate the hair follicle cycle with two models.
 - Activator/ inhibitor
 - Substrate/ depletion
- Create networks of follicle oscillators via different modes of coupling
 - To observe the effects of certain variables
 - To produce synchronization.

The Activator/Inhibitor Model



The Substrate/Depletion Model

x - product

y - reactant

$$dx/dt = \frac{v(y-x)(\varepsilon^2 + x^2)}{1+x^2} - x$$
$$dy/dt = k - x$$

Fall, et al.

Hopf Bifurcation

- Fixed point loses stability as the eigenvalues cross the imaginary axis of the complex plane
- Stable limit cycle



Perturbations and the Interaction Function (H) $\frac{d\theta}{dt} = 2\pi / T_0 + \beta \sin(\omega t - \theta)$

 $\omega = 2\pi / T$

Phase difference between oscillator and stimulator

$$\phi = \omega t - \theta$$



Averaging

 Z(t) – response function • G(x) – type of coupling • Diffusive coupling x'-x Bath coupling f((x+0.01*x',y)-f(x,y))/0.01 • $H(\Phi)$ – interaction function; coupling as a function of phase difference between oscillators

$$H_1(\phi) = \frac{1}{T} \int_0^T Z(t) \cdot G_1(X_0(t+\phi), X_0(t)) dt$$

Diffusive coupling function via the activator



 $H(\phi) = 4 - 2\cos\theta - 2\cos 2\theta + 26\sin\theta + 5\sin 2\theta + \sin 3\theta$

Network Equation

20 oscillators in the network
Optional gradient (ε)
Coupling strength (a) – must be weak

$$\begin{aligned} x_{1} &= 1 + ah(x_{2} - x_{1}) \\ x_{2.19} &= 1 - \frac{(j-1)\varepsilon}{20} + a(h(x_{j-1} - x_{j}) + h(x_{j+1} - x_{j})) \\ x_{20} &= 1 - \frac{19\varepsilon}{20} + ah(x_{19} - x_{20}) \end{aligned}$$
 Neighbor influence

1-D Arrays

Kind of coupling

Via x or y variable
Via diffusive or bath coupling

Coupling constant (a)
Diffusion limit (number of neighbors)
Heterogeneity (ε)

$$x_{2.19} = 1 - \frac{(j-1)\varepsilon}{20} + a(h(x_{j-1} - x_j) + h(x_{j+1} - x_j))$$

Types of Coupling ($\epsilon = 0$, 2 nearest neighbors)

Activator

a = 1

a = 5

Inhibitor



Coupling Constant ($\epsilon = 0, 2$ nearest neighbors, diffusive coupling)



Diffusion Limit (a = 1, ε = 0, diffusive coupling via activator)

2 neighbors

4 neighbors

6 neighbors







Introducing Heterogeneity (a = 1, diffusive)



2-D Arrays

- Using Fourier approximation
- Simulates actual traveling waves in the mouse





Suzuki, et al.



Types of Coupling (a=1, eps=0, 2 nearest neighbor)



via reactant



Adding a gradient induces synchronization for bath coupling via product

eps = 0 eps = 1 eps = 5



2-D Arrays

Diffusive coupling via

Product



Reactant



Bath coupling did not produce any patterns.

Results Summary – Types of Coupling

 Only 4 networks synchronized for both 1 and 2 D arrays.

	AI Model	1-D	2-D	SD Model	1-D	2-D
		Array	Array		Array	Array
Diffusive	Activator	×	*	Product	*	*
	Inhibitor	*		Reactant	*	*
Bath	Activator	×	*	Product	*	
	Inhibitor			Reactant		

Results Summary - Variables

• 1-D Arrays

- The only coupling mode which never synchronized was bath coupling via the reactant.
- Changing the gradient can induce traveling waves, reverse wave direction and produce synchrony.
- The coupling constant and the diffusion limit are proportional to synchronization speed.
- 4 nearest neighbor coupling can initially produce unusual patterns.
- 2-D Arrays

Conclusions

- The hair follicle cycle could be represented by either model.
- Most likely networks are those which synchronize for 1 and 2 D networks.
 - Diffusive coupling via activator, product, and reactant
 - Bath coupling via activator

Who cares??



Acknowledgments

- Dr. G. Bard Ermentrout, advisor
- Judy Wieber and Rajan Munshi, coordinators
- BBSI participants
- BBSI Department of Computational Biology, Pittsburgh
 NIH/NSF

References

- Edelstein- Keshet, Leah. <u>Mathematical Models in</u> <u>Biology</u>, 1st ed. Boston: McGraw Hill, 1988.
- Ermentrout, G.B. and J. Rinzel. "Beyond a pacemaker's entrainment limit: phase walk-through." Am J Physiol Regulatory Integrative Comp Physiol. 246 (1984): 102-106.
- Fall, Christopher P., et. al., eds. <u>Computational Cell</u> <u>Biology</u>. New York: Springer-Verlag, 2002.
- Maini, Philip K. "How the mouse got its stripes." PNAS 100:17(2003): 9656-9657.
- Paus, Ralf and Kerstin Foitzik. "In search of the 'hair cycle clock': a guided tour." *Differentiation*. 72(2004):489-511.
- Suzuki, Noboru, et al. "Traveling strips on the skin of a mutant mouse." PNAS. 100:17 (2003): 9680-9685.