

# On the channel capacity of channel rhodopsin (and other biological signal transduction pathways)

Peter Thomas  
Case Western Reserve University  
Joint work with Andrew Eckford, York University

Don't forget to press the record button.



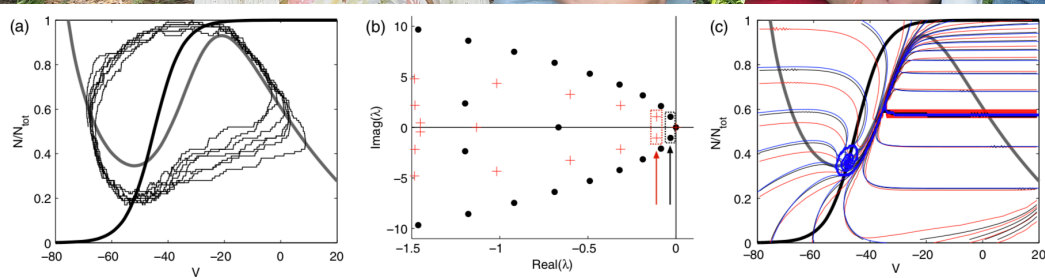


Computational Biomathematics Laboratory  
at Case Western Reserve University

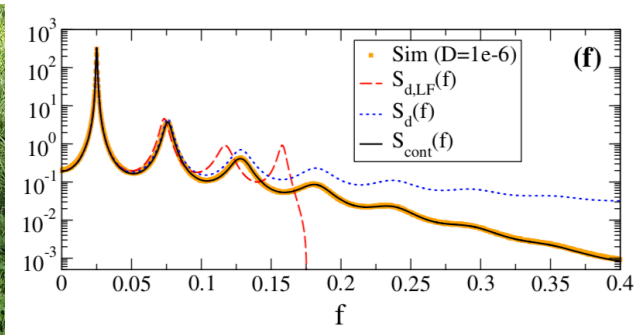
(Cleveland, Ohio)



# Computational Biomathematics Laboratory at Case Western Reserve University

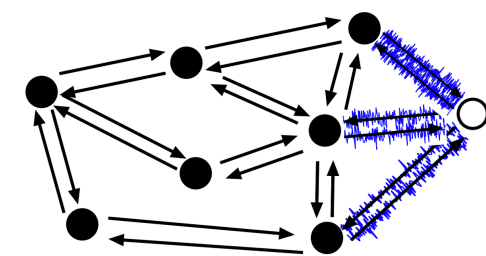
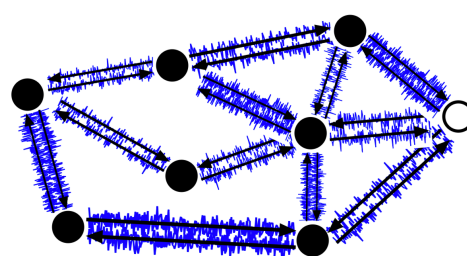
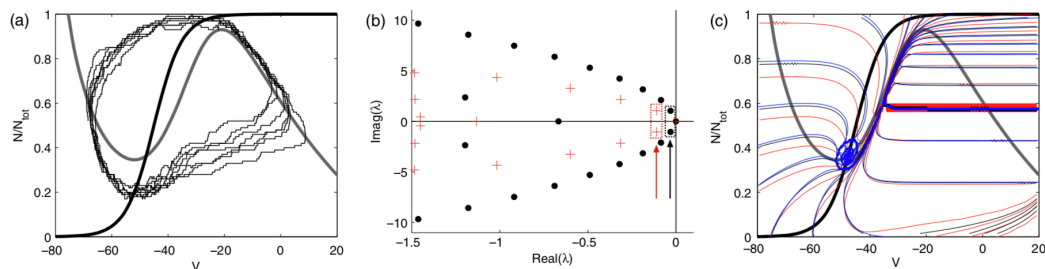


stochastic dynamics in neural oscillators



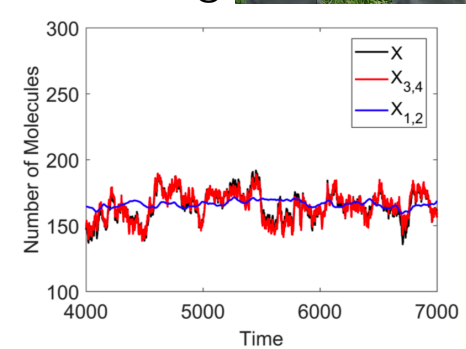
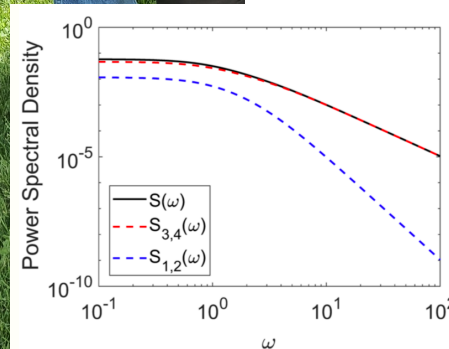
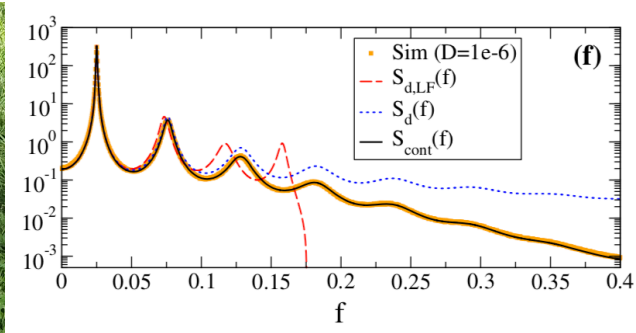


# Computational Biomathematics Laboratory at Case Western Reserve University

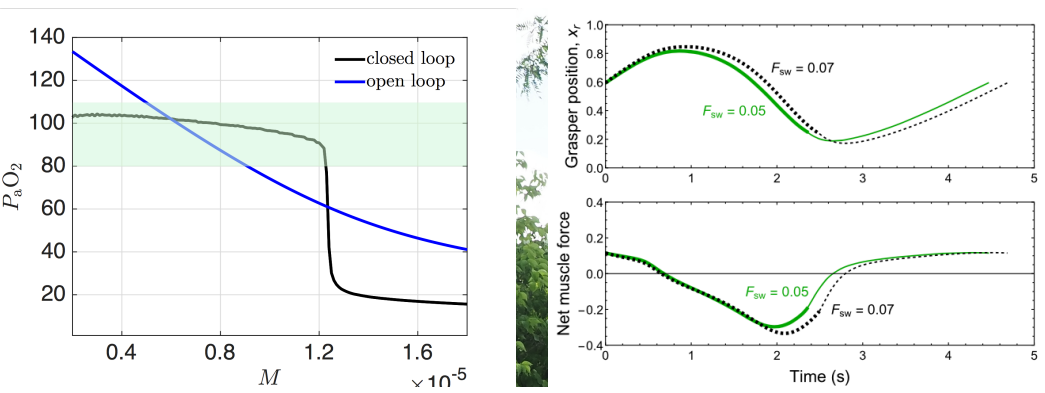


stochastic dynamics in neural oscillators

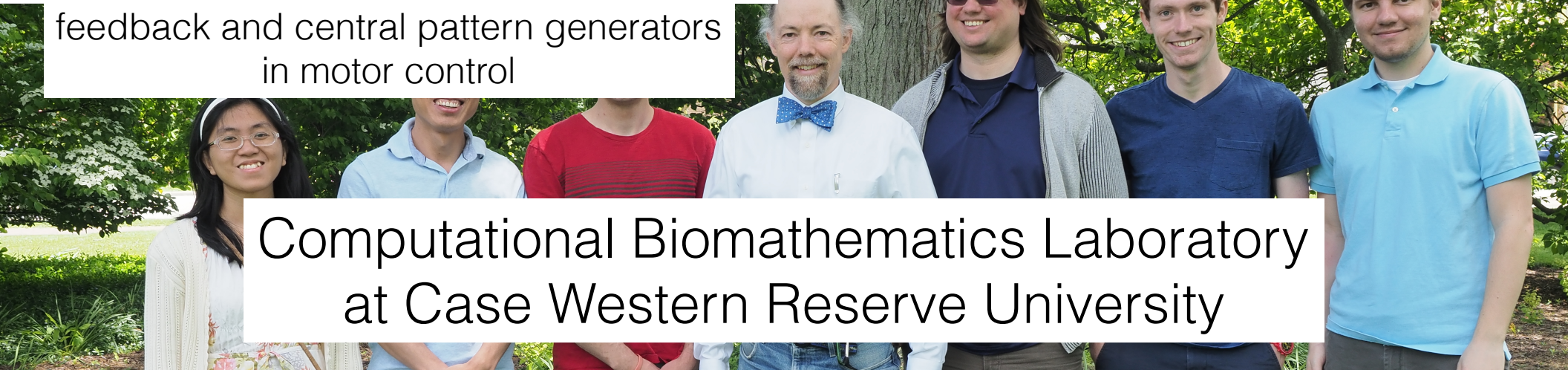
stochastic shielding



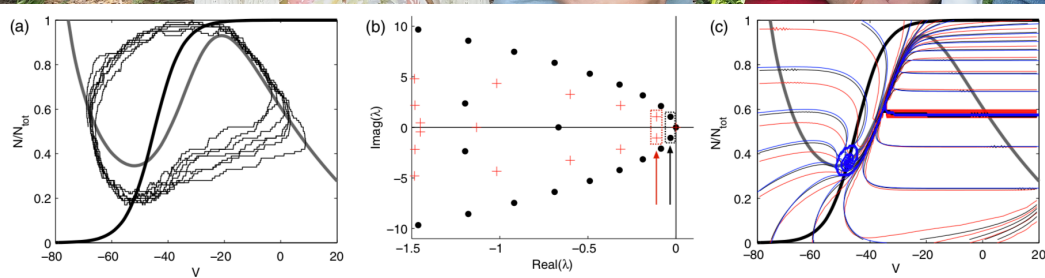




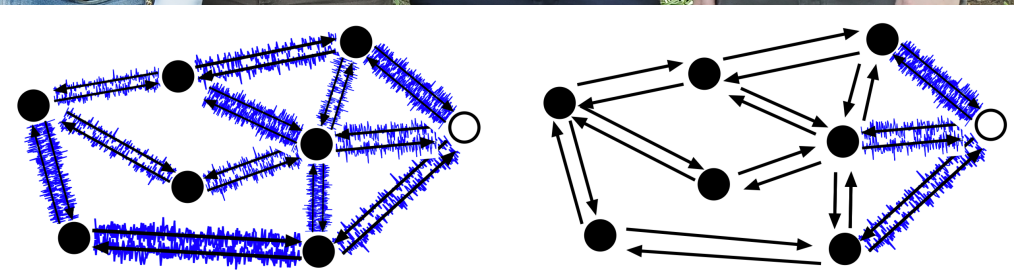
feedback and central pattern generators in motor control



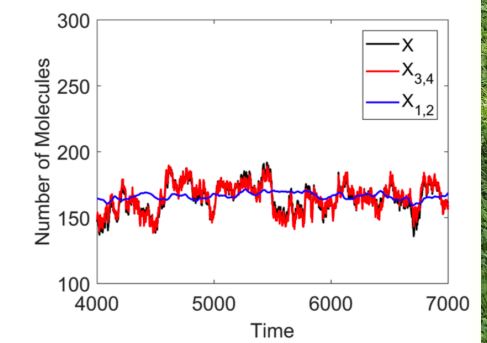
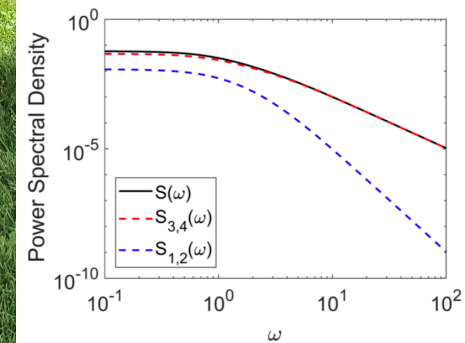
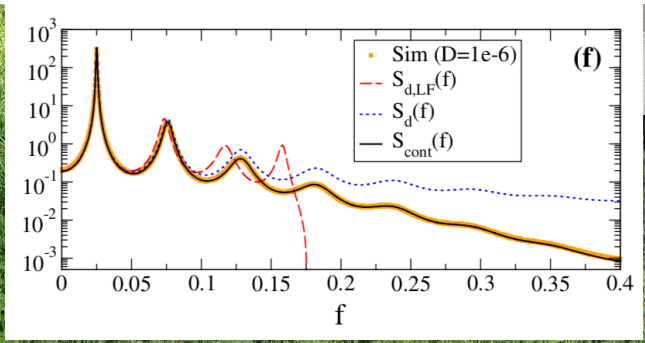
# Computational Biomathematics Laboratory at Case Western Reserve University



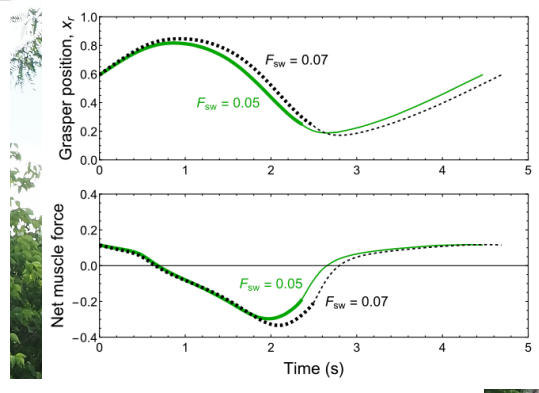
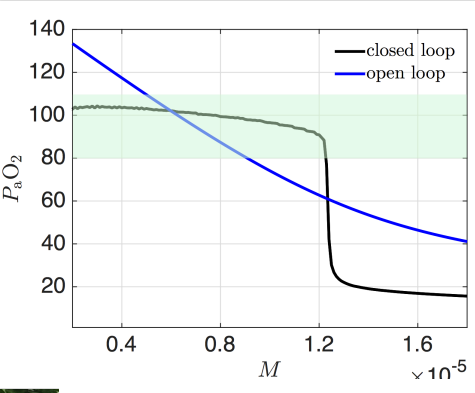
stochastic dynamics in neural oscillators



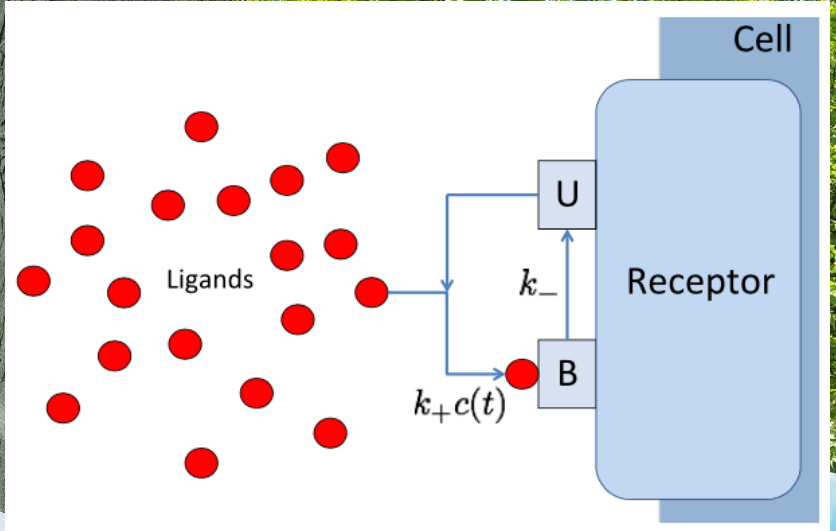
stochastic shielding





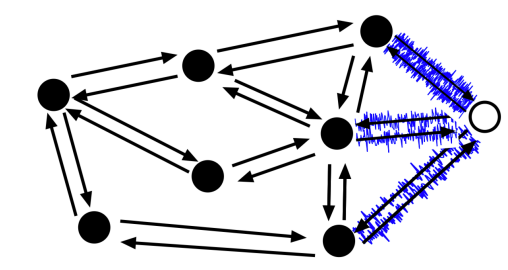
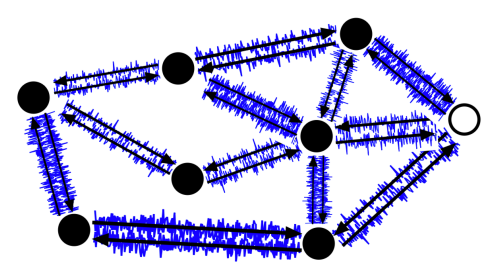
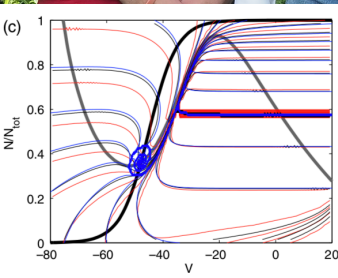
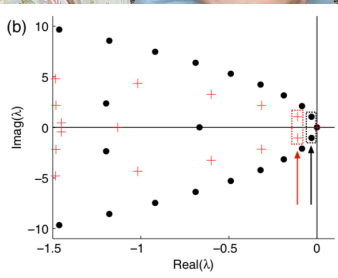
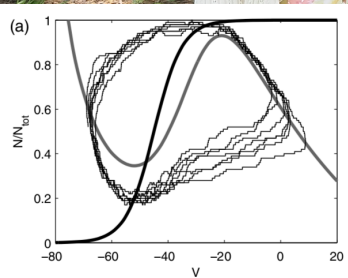


feedback and central pattern generators in motor control



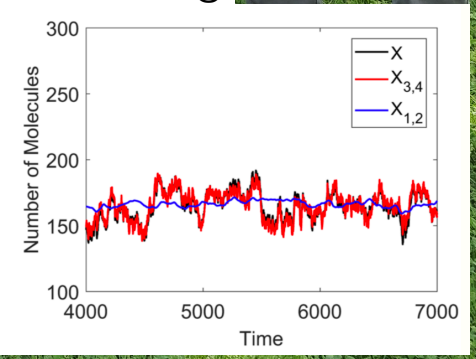
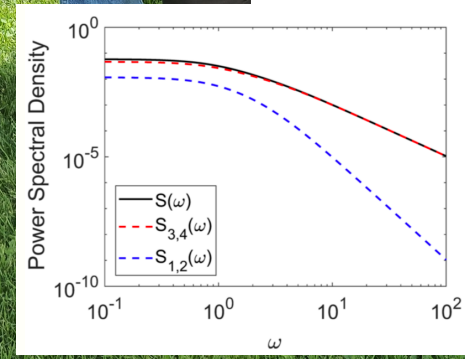
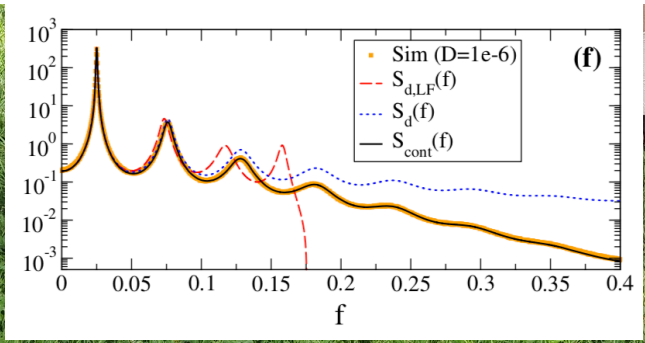
information theory in signal transduction systems

# Computational Biomathematics Laboratory at Case Western Reserve University



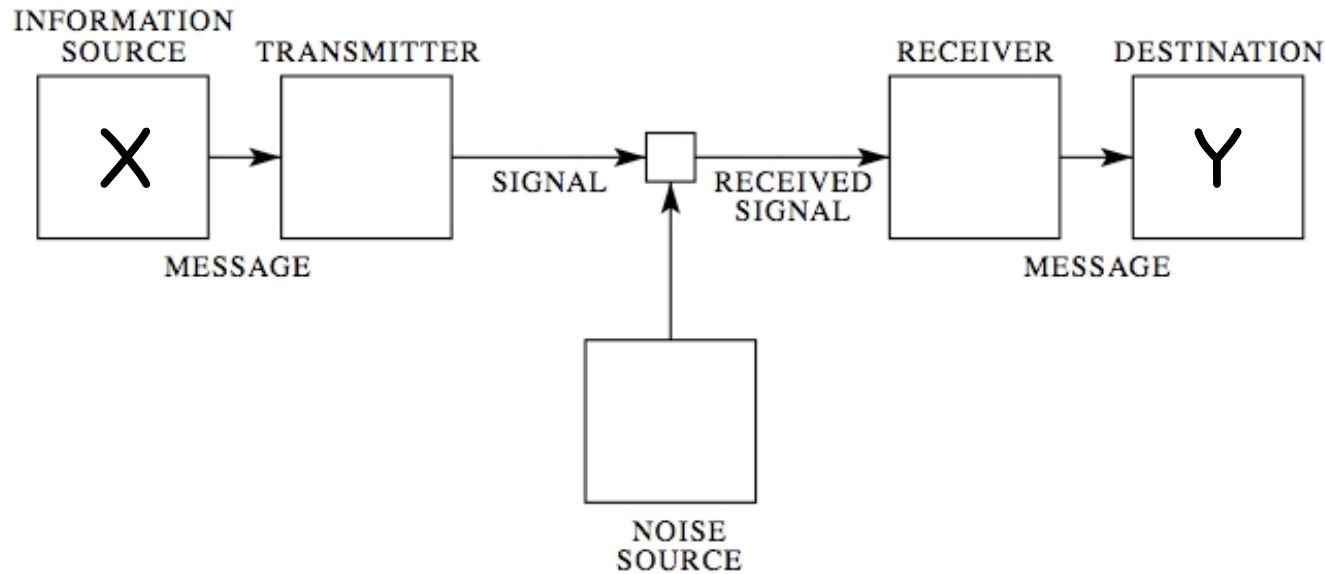
stochastic dynamics in neural oscillators

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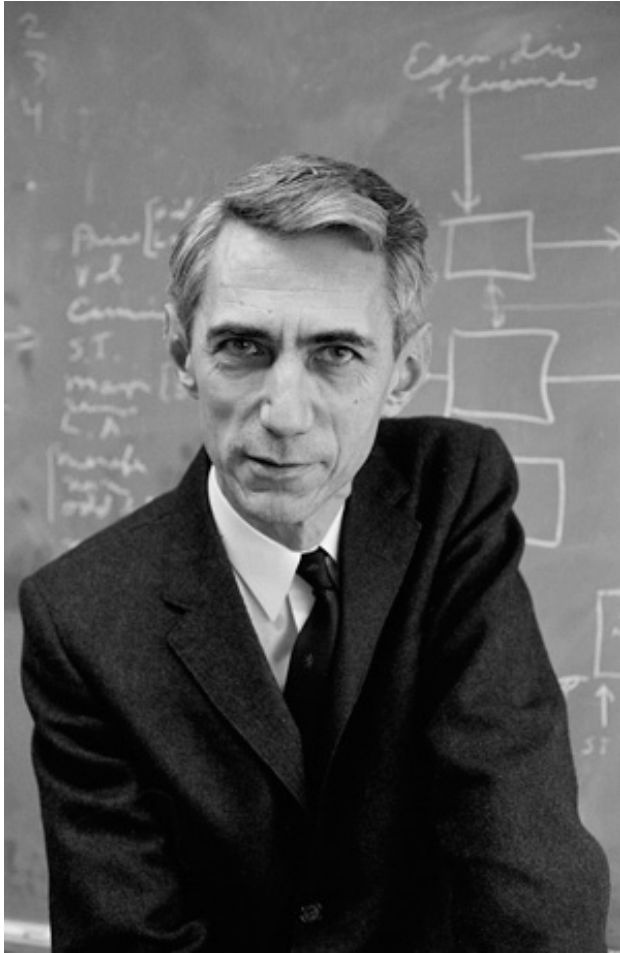
The fundamental question of information theory:  
*How much information can a communications system  
communicate?*



Shannon (1948) A Mathematical Theory of Communication

Answer: Channel Capacity =  $\text{Max}(\text{Mutual Information } I(X:Y))$ ;  
Maximize over input ensembles  $P(X)$



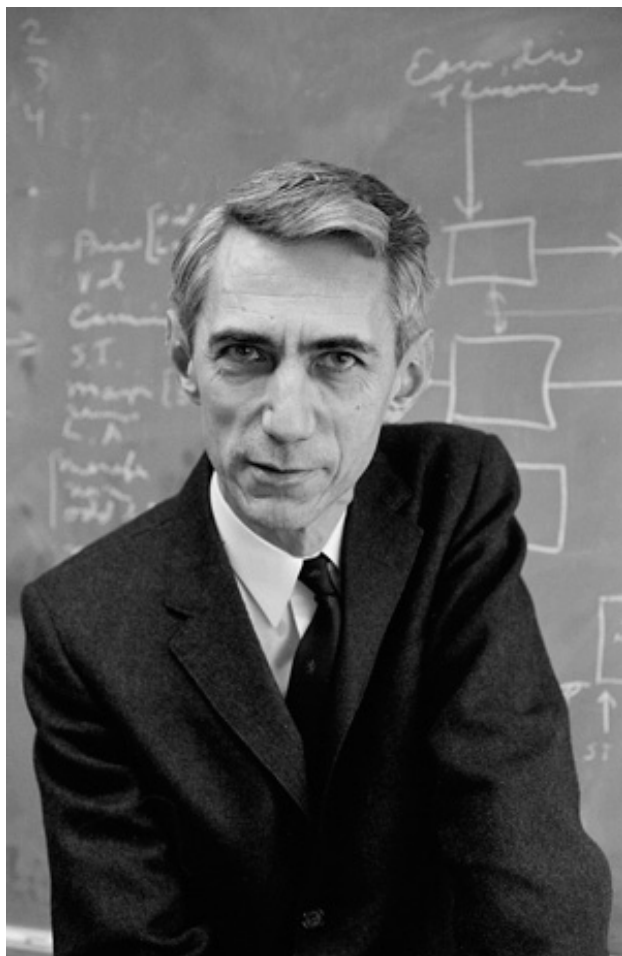
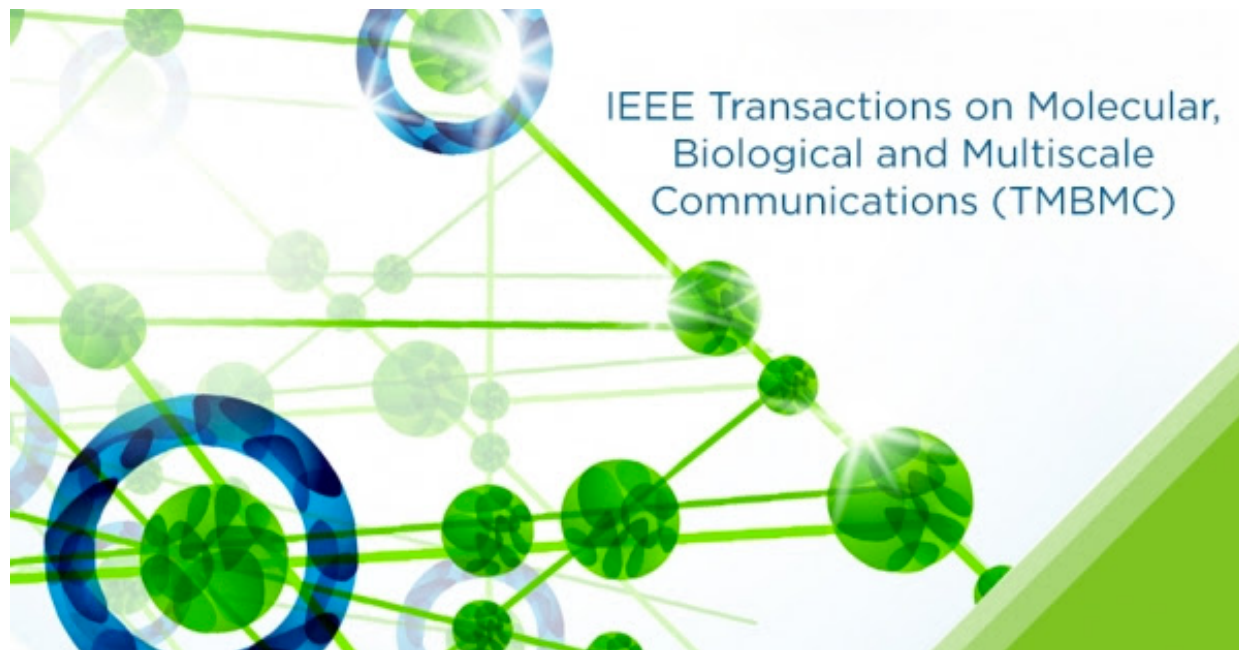


# Claude Shannon

## 1916-2001

Alfred Eisenstaedt/The LIFE Picture Collection/Getty Images

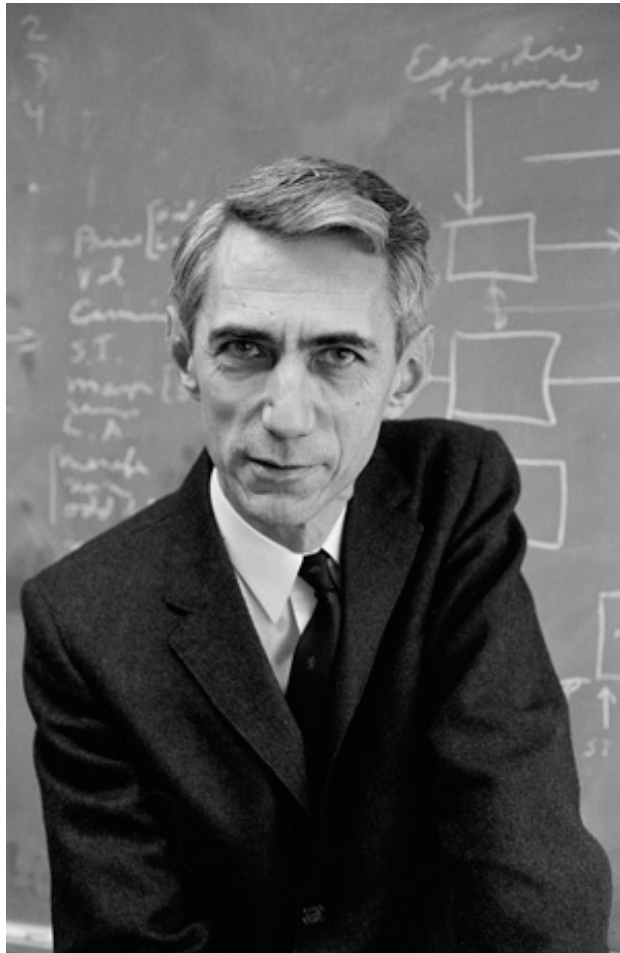




Claude Shannon  
1916-2001

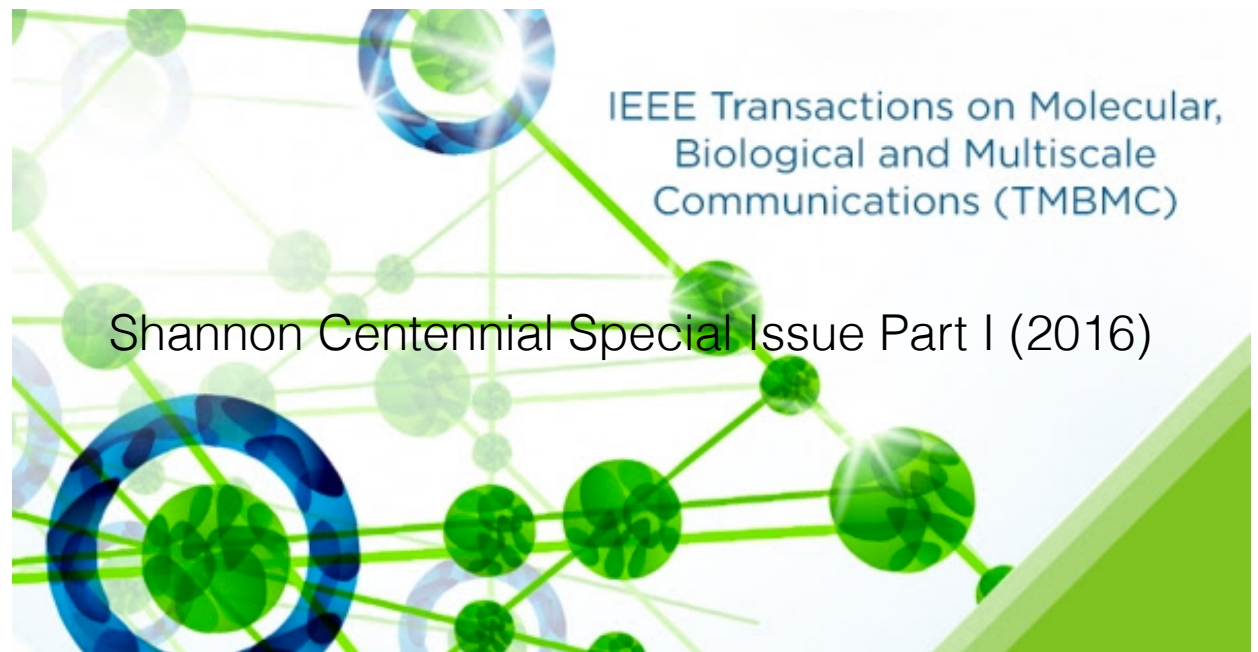
Alfred Eisenstaedt/The LIFE Picture Collection/Getty Images





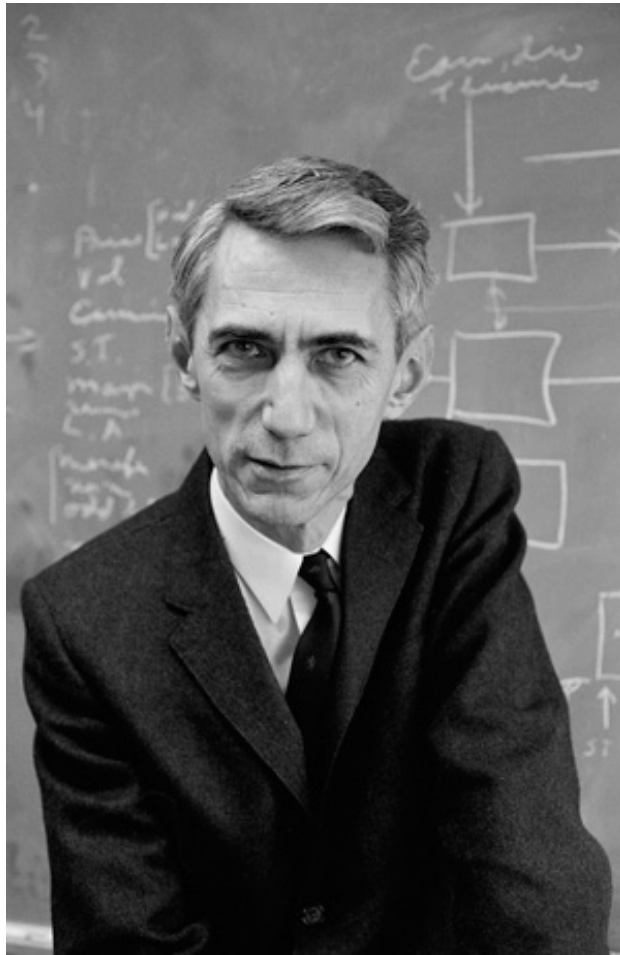
Claude Shannon  
1916-2001

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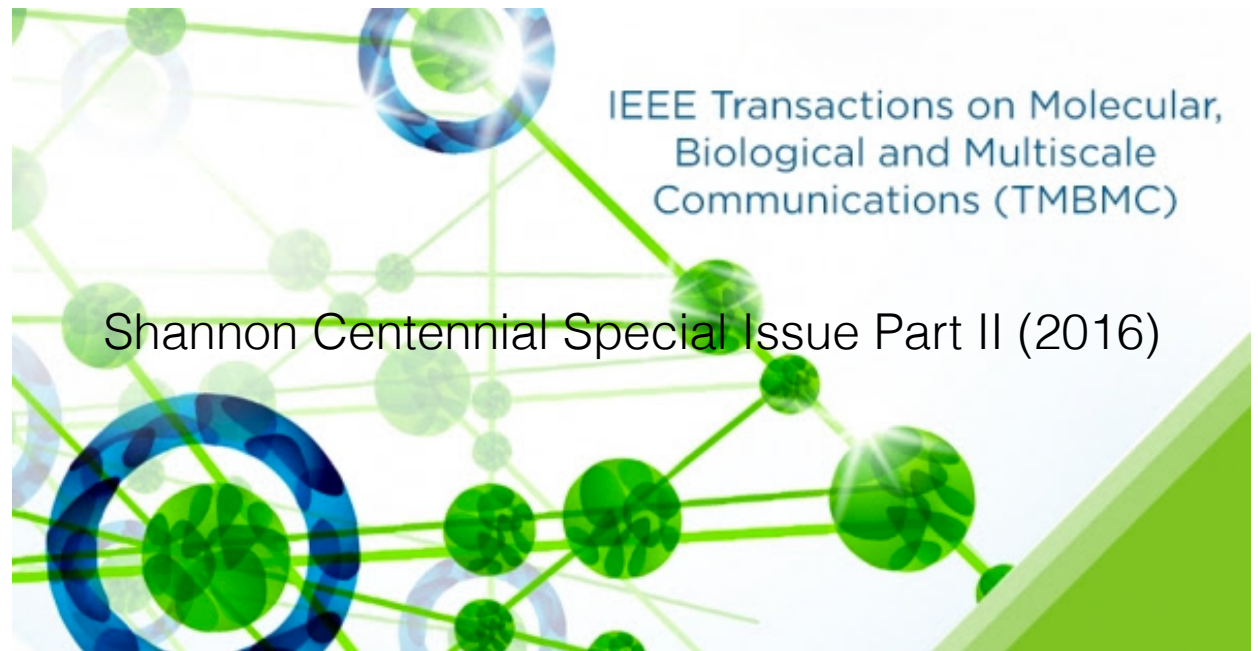
- Mechanisms of information filtering in neural systems (Linder)
- Noise Filtering and Prediction in Biological Signaling Networks (Hathcock, Sheehy, Weisenberger, Ilker, and [Hinczewski](#))
- The Use of Rate Distortion Theory to Evaluate Biological Signaling Pathways (Iglesias)
- Nonlinear Stochastic Dynamics of Complex Systems, III: Nonequilibrium Thermodynamics of Self-Replication Kinetics (Saakian and Qian)
- Inferring Biological Networks by Sparse Identification of Nonlinear Dynamics (Mangan, Brunton, Proctor and Kutz)
- Info-Clustering: A Mathematical Theory for Data Clustering (Chan, Al-Bashabsheh, Kaced, Zhou and Liu)
- Fundamental Bounds for Sequence Reconstruction from Nanopore Sequencers (Magner, Duda, Szpankowski and Grama)
- Inference of Causal Information Flow in Collective Animal Behavior (Lord, Sun, Ouellette and Bolt)





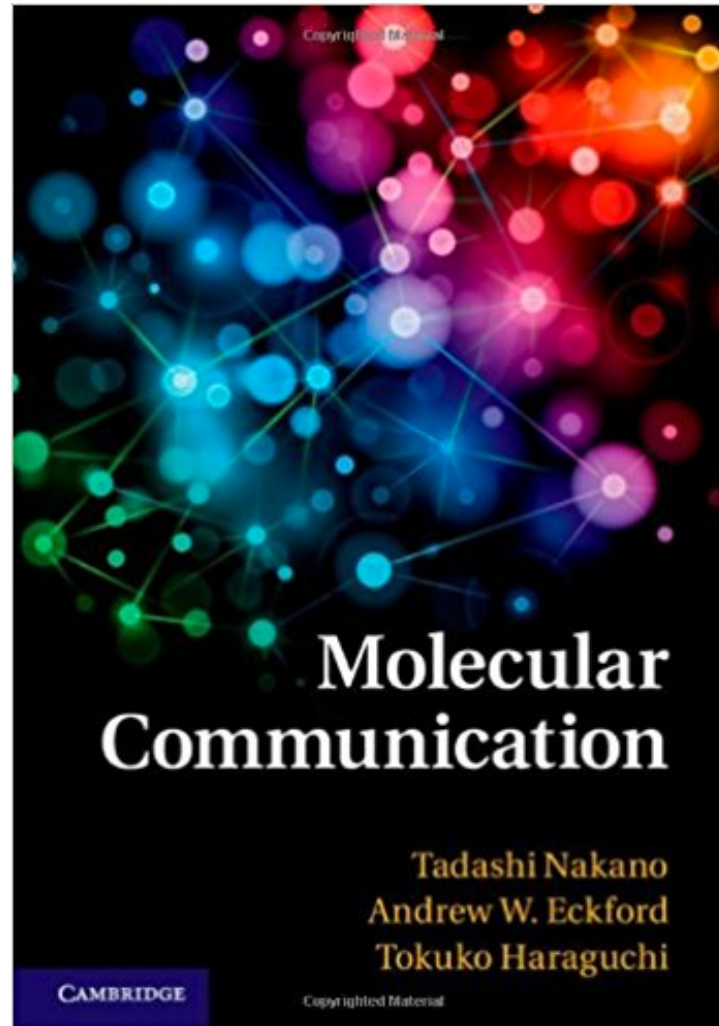
Claude Shannon  
1916-2001

Alfred Eisenstaedt/The LIFE Picture Collection/Getty Images



- Information Theory of Molecular Communication: Directions and Challenges (Gohari, Mirmohseni, and Nasiri-Kenari)
- On Palimpsests in Neural Memory: An Information Theory Viewpoint ([Varshney](#), Kusuma, and Goyal)
- Neural Computation from First Principles: Using the Maximum Entropy Method to Obtain an Optimal Bits-Per-Joule Neuron (Berger, Levy and Sungkar)
- Mutual Information and Parameter Estimation in the Generalized Inverse Gaussian Diffusion Model of Cortical Neurons (Sungkar, Berger, and Levy)
- Identifying Multisensory Dendritic Stimulus Processors (Lazar and Zhou)
- Fundamental Limits of Genome Assembly under an Adversarial Erasure Model (Shomorony, Courtade, and Tse)
- Inscribed Matter Communication: Part I ([Rose](#) and Mian)
- Inscribed Matter Communication: Part II ([Rose](#) and Mian)
- Process Information and Evolution (Chastain and Smith)





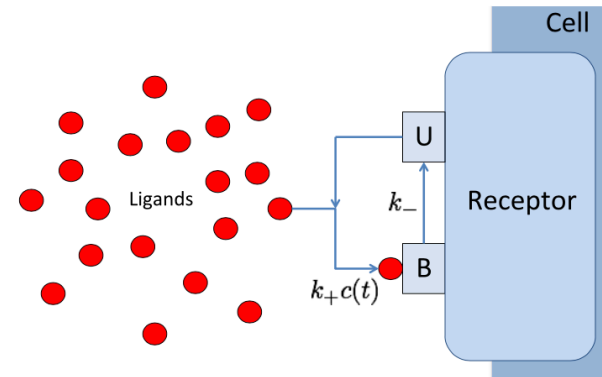
(2013)



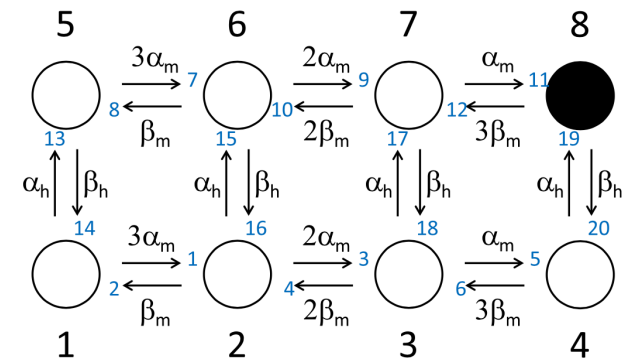
# Signal Transduction:

## Transforming Extracellular Signals into Intracellular Responses

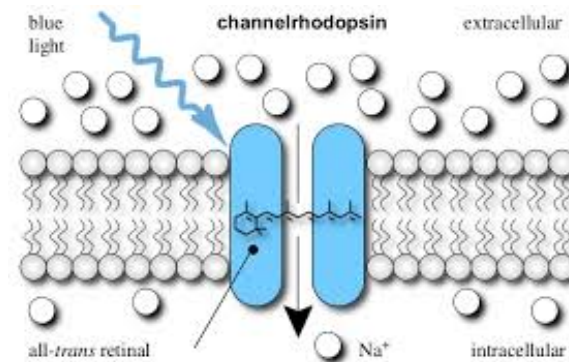
- \* Ligand-receptor systems
  - \* cyclic AMP receptor
  - \* acetylcholine receptor
  - \* calcium signaling (calmodulin)



- \* Voltage-gated ion channels
  - \* Hodgkin-Huxley sodium channel
  - \* gap junction mediated sync.



- \* Light-gated ion channels
  - \* channelrhodopsin
  - \* light-driven cAMP synthesis

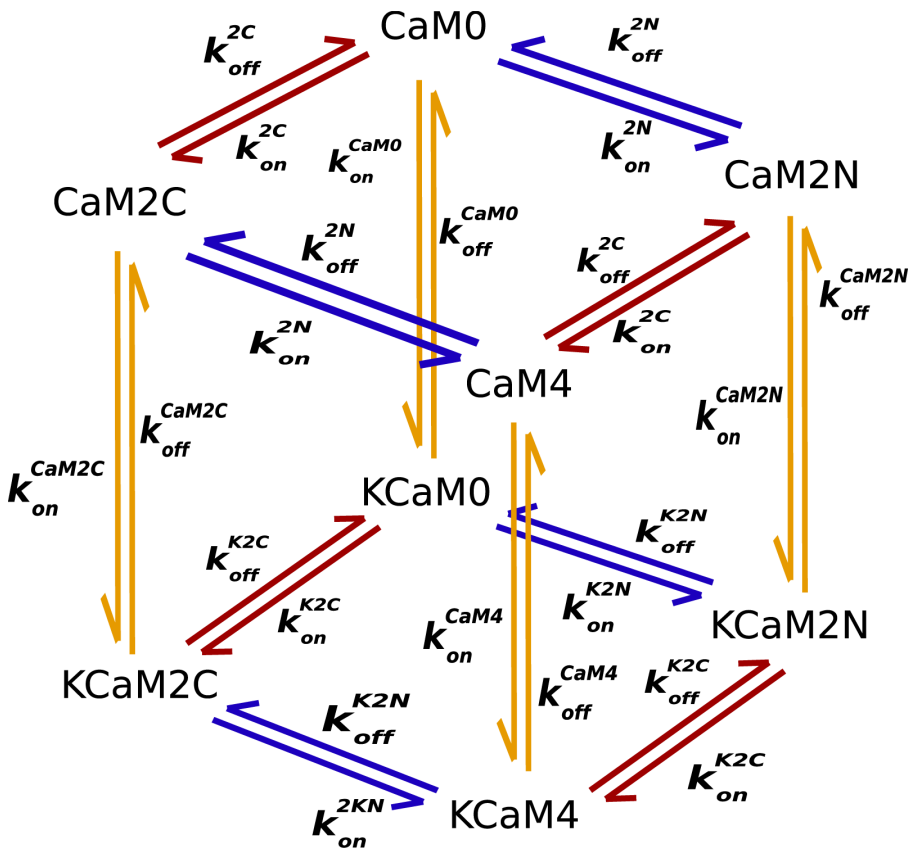




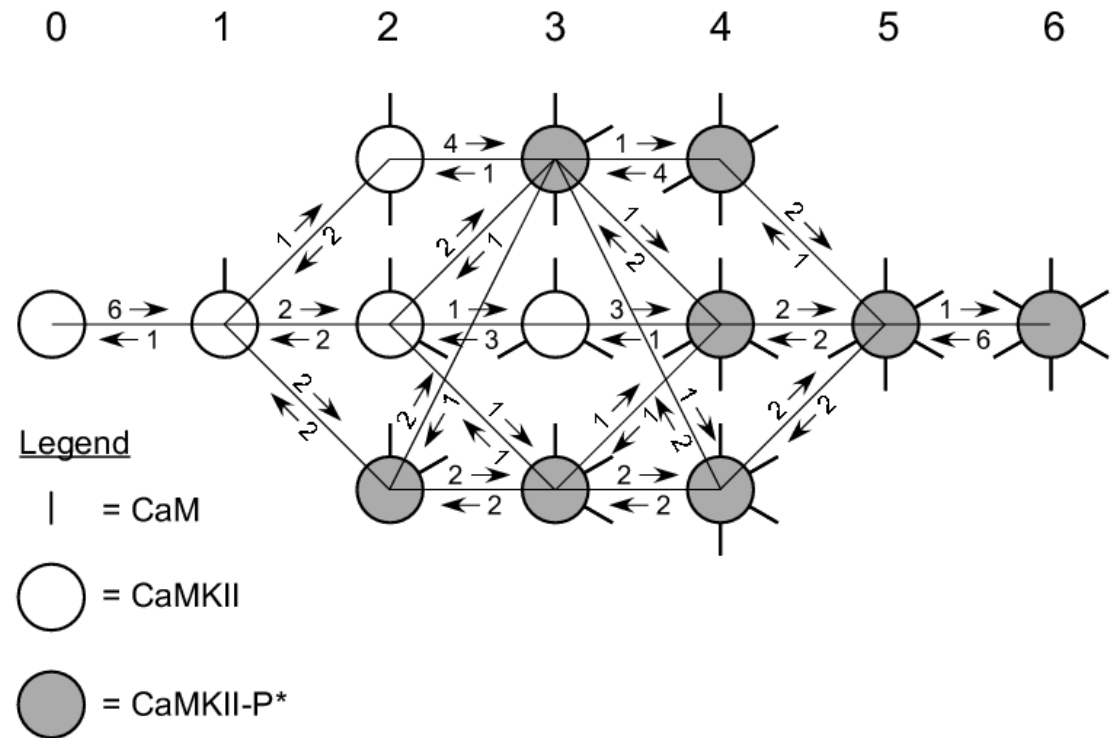
# Information Capacity of a Signal Transduction Channel

Transitions between receptor states  $\{1, \dots, j, \dots, S\}$  driven by signal concentration  $X(t)$ .

$$\frac{dp_k}{dt} = \sum_{j=1}^n p_j(t) q_{jk}(X(t))$$



Calcium-Calmodulin Binding Graph

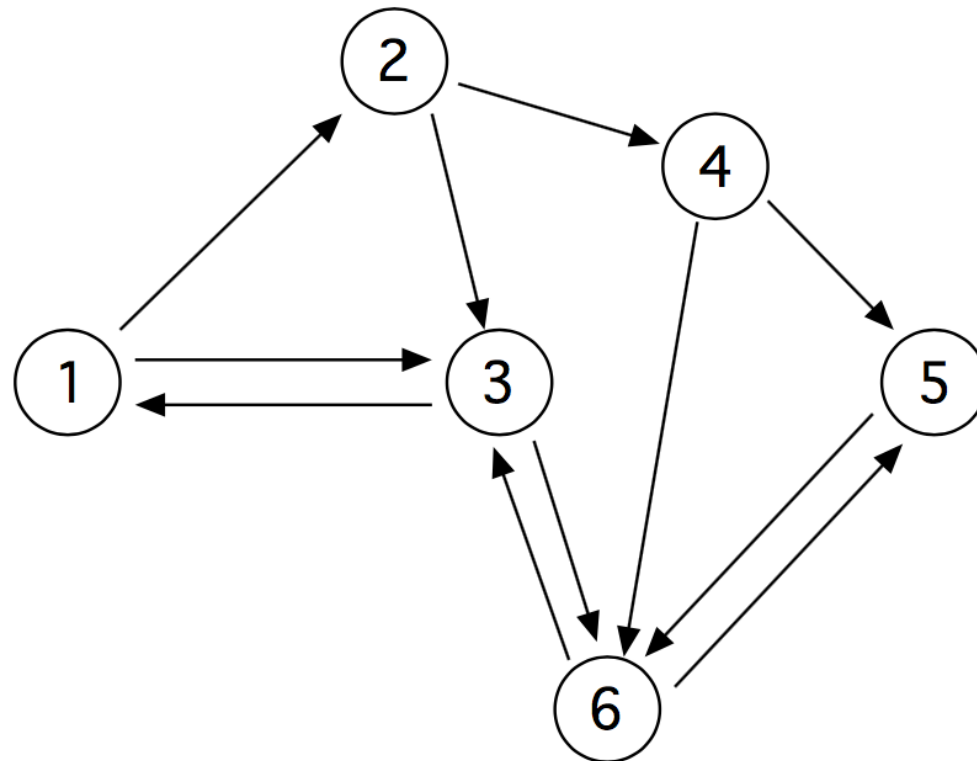


Calmodulin/CaMKII state graph



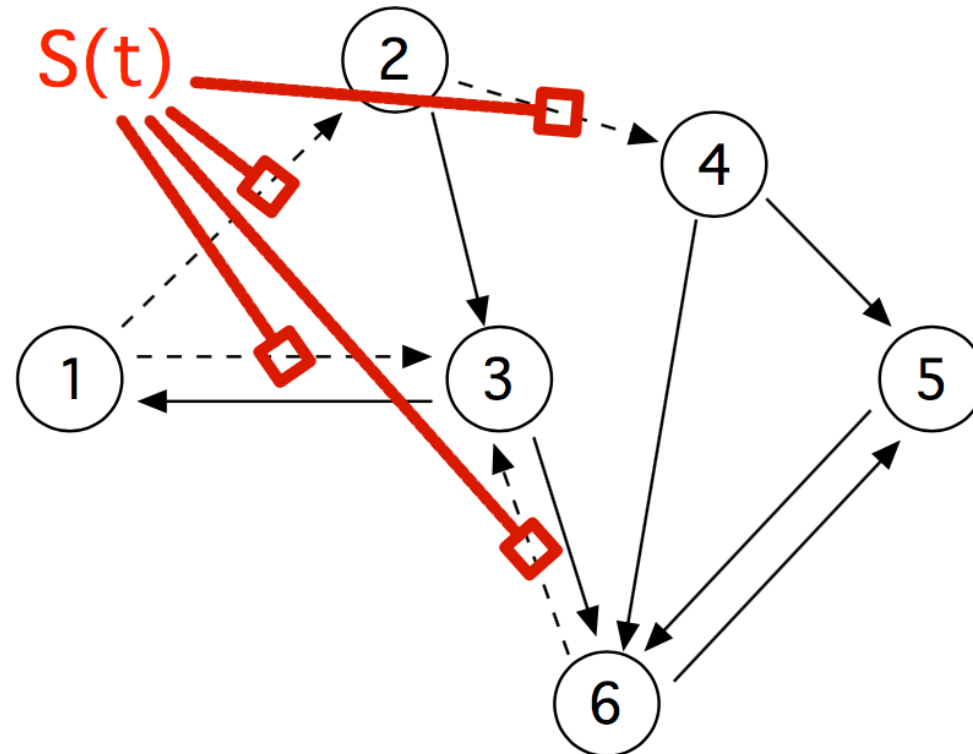
# Graphical representation of signal transduction

- ▶ We represent an individual receptor's state as a node in a directed graph  $\mathcal{G} = (\mathcal{V}, \mathcal{E}) = (\{\text{vertices}\}, \{\text{edges}\})$ .
- ▶ Edge  $i \rightarrow j$  represents a transition from state  $i$  to state  $j$ .
- ▶ If the per capita transition rate  $\alpha_{ij}$  depends on the input signal  $S(t)$ , the state  $i$  is *sensitive*.



# Graphical representation of signal transduction

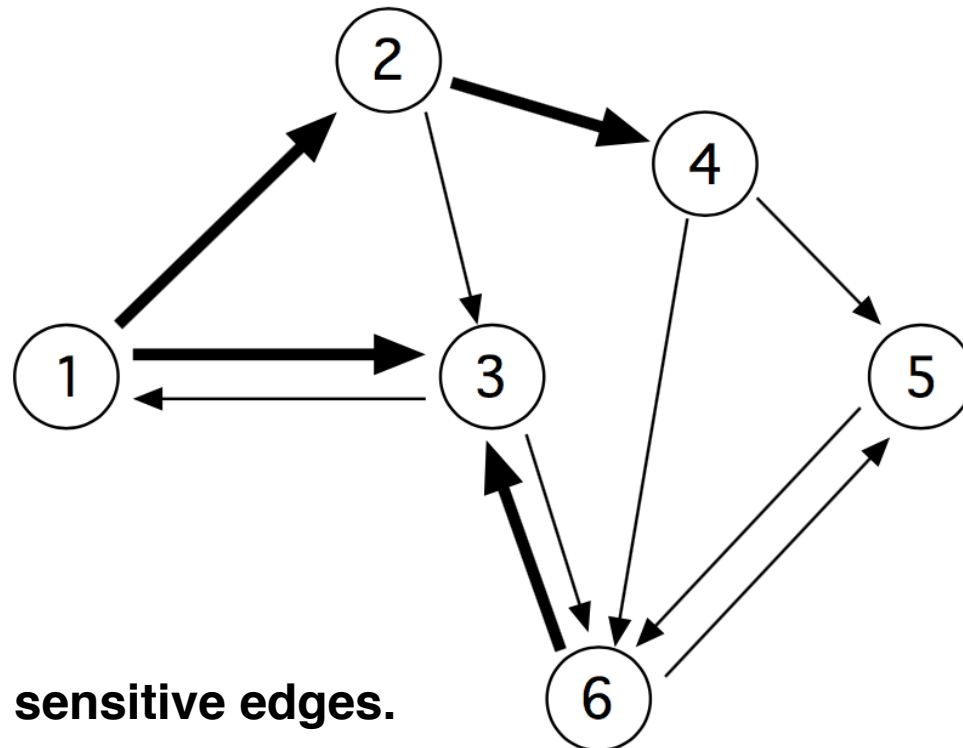
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# Graphical representation of signal transduction

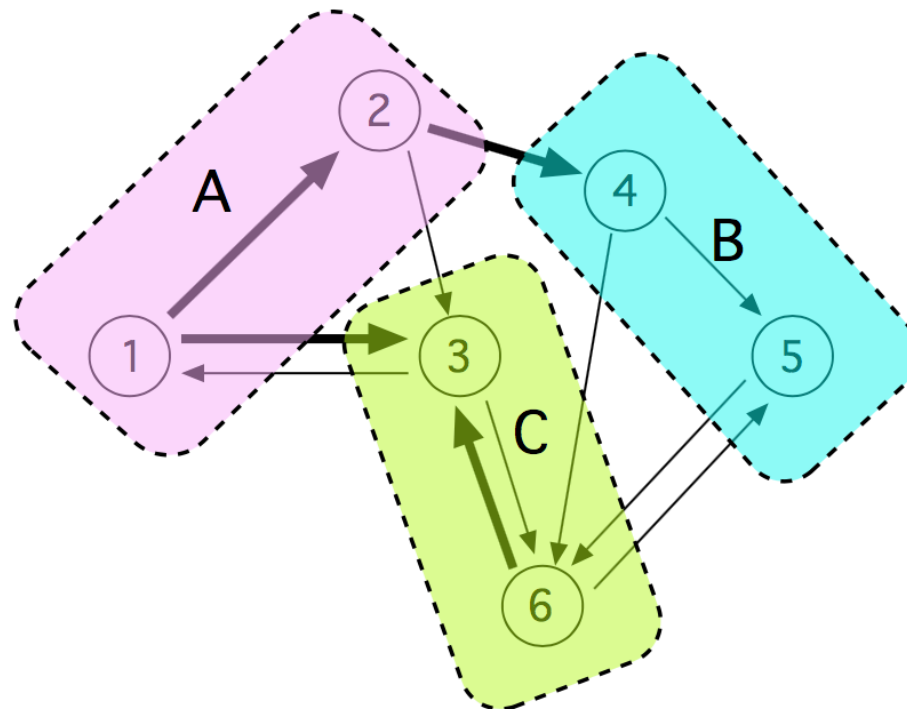
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**Bold arrows denote sensitive edges.**

# Graphical representation of signal transduction

- ▶ The observable state of the receptor may be more coarse grained than the underlying state graph, e.g. observables  $A = \text{Ind}(\{1, 2\})$ ,  $B = \text{Ind}(\{4, 5\})$ ,  $C = \text{Ind}(\{3, 6\})$ .
- ▶ If the transition  $i \rightarrow j$  changes the coarse-grained state, then edge  $i \rightarrow j$  is *observable*.
- ▶ We distinguish the mutual information and capacity for the *fully observed* versus the *partially observed* receptor.



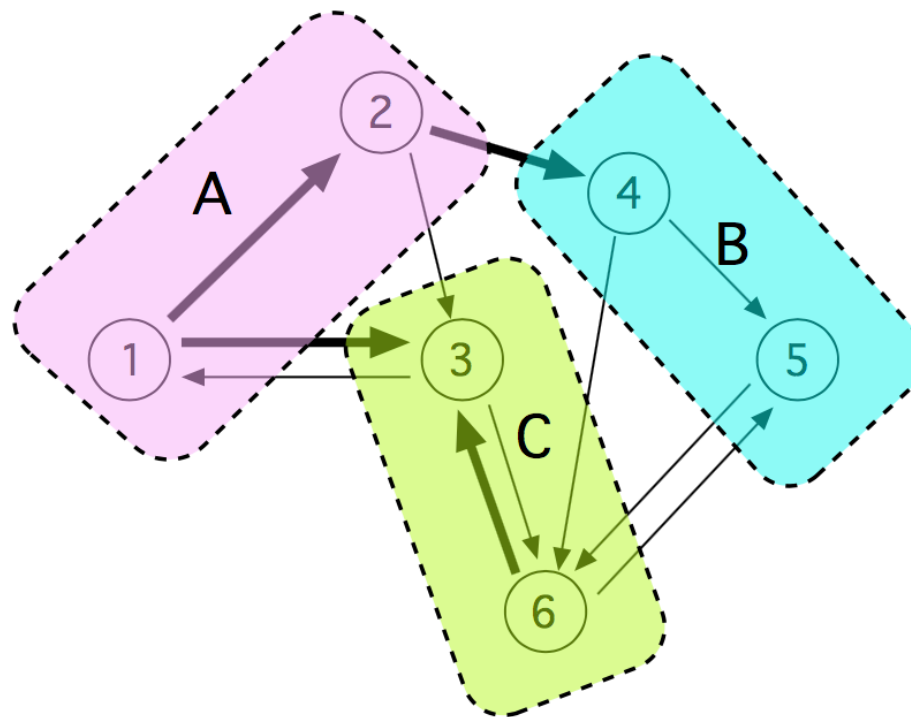


# Continuous-time channel model

- ▶ Input:  $X(t) : [0, \infty) \rightarrow [x_{\min}, x_{\max}]$  with  $0 \leq x_{\min} \leq x_{\max}$ .
- ▶ Channel State:  $Y(t) \in \{1, \dots, K\}$ .  $p_i(t) = \Pr(Y(t) = i)$ .

$$\frac{dp_j}{dt} = \sum_{i=1}^K p_i \alpha_{ij}(X(t)), \text{ with } \alpha_{jj} = - \sum_{k \neq j} \alpha_{jk}.$$

- ▶ Observable Output:  $Z(t) = C \cdot Y(t)$  for an  $M \times K$  matrix  $C$ .

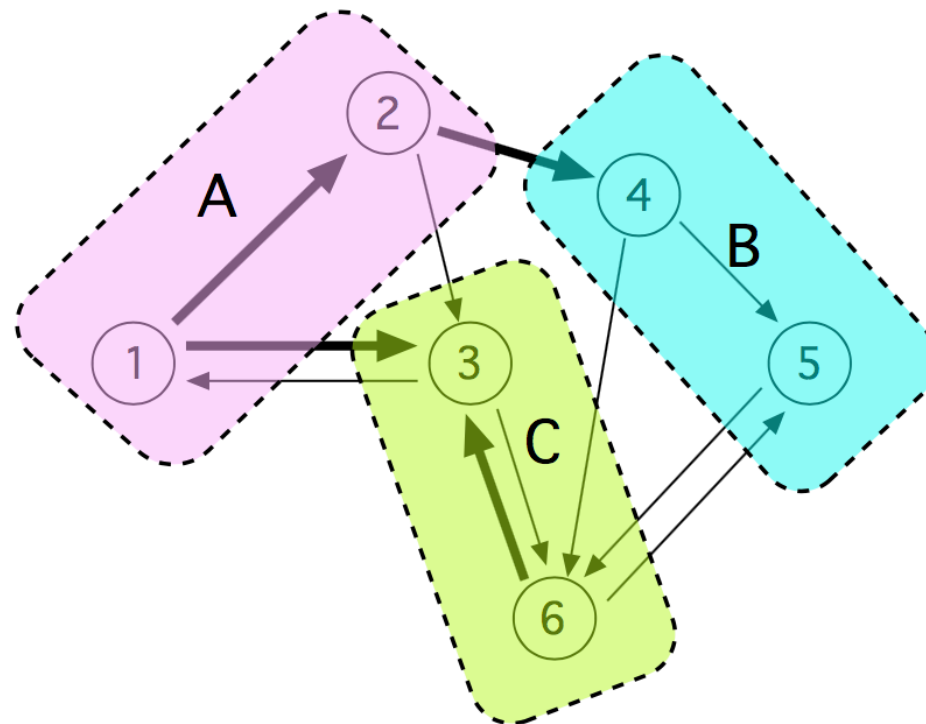


# Discrete-time channel model ( $0 < \Delta t \ll 1$ )

- ▶ Input:  $X(t) : \{0, \Delta t, 2\Delta t, \dots\} \rightarrow [x_{\min}, x_{\max}]$ .
- ▶ Channel State:  $Y(t) \in \{1, \dots, K\}$ .  $p_i(t) = \Pr(Y(t) = i)$ .

$$p_j(t + \Delta t) = p_j(t)(1 - \alpha_{jj}\Delta t) + \sum_{i \neq j} p_i(t)\alpha_{ij}(X(t)).$$

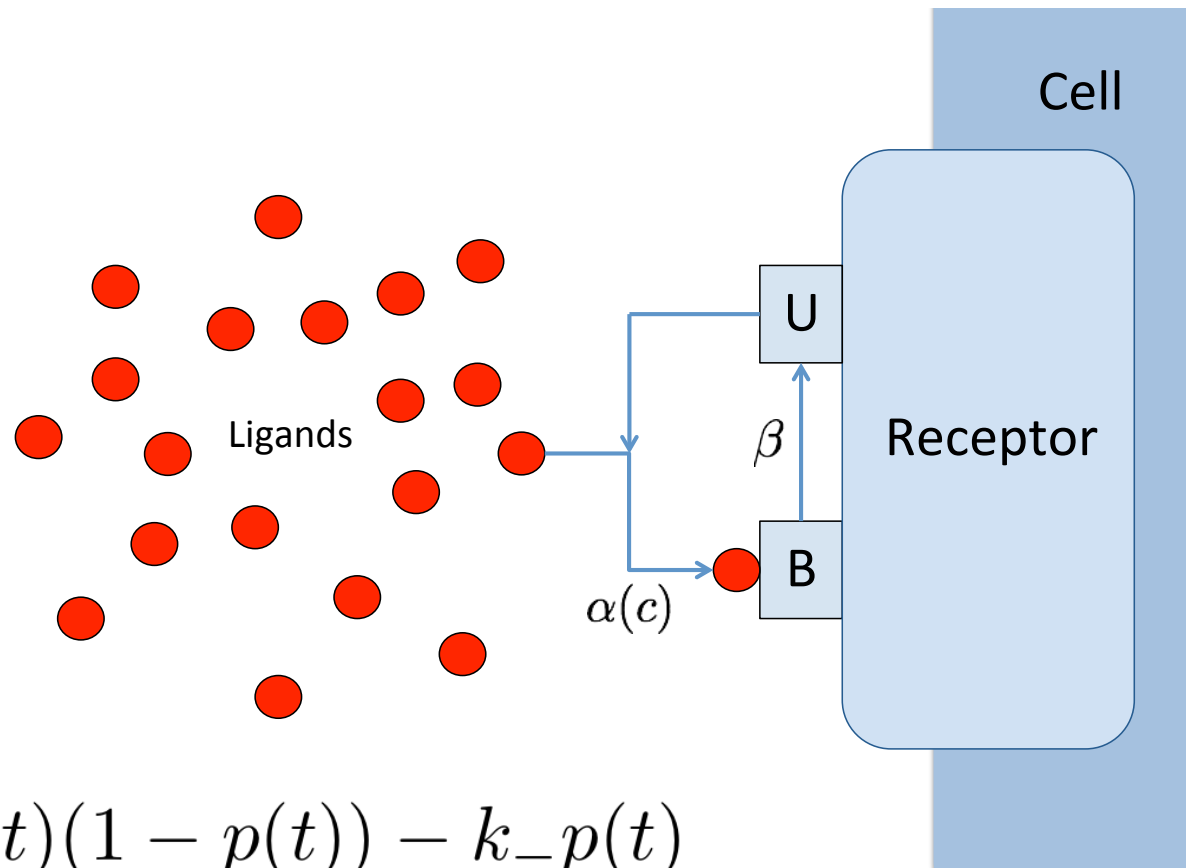
- ▶ Observable Output:  $Z(t) = C \cdot Y(t)$  for an  $M \times K$  matrix  $C$ .





# Example 1: the BIND channel (a single binary receptor)

Receptor has only two states (Bound/Unbound).



$$\frac{dp}{dt} = k_+ c(t)(1 - p(t)) - k_- p(t)$$

$p$ : probability receptor is bound.

$k_+$ : binding rate constant.

$k_-$ : unbinding rate constant.

$$\alpha_L = k_+ c_{\min} \Delta t$$

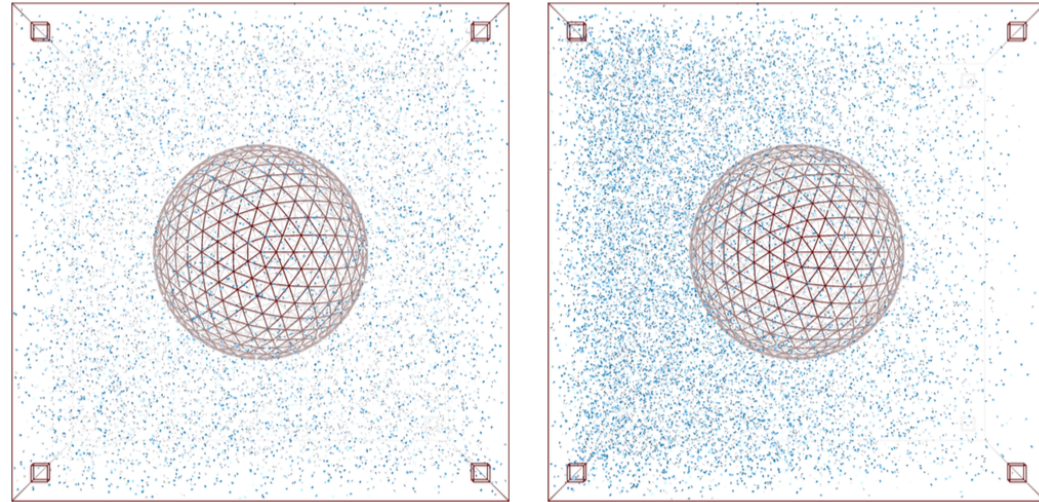
$$\alpha_H = k_+ c_{\max} \Delta t$$

$$\beta = k_- \Delta t$$

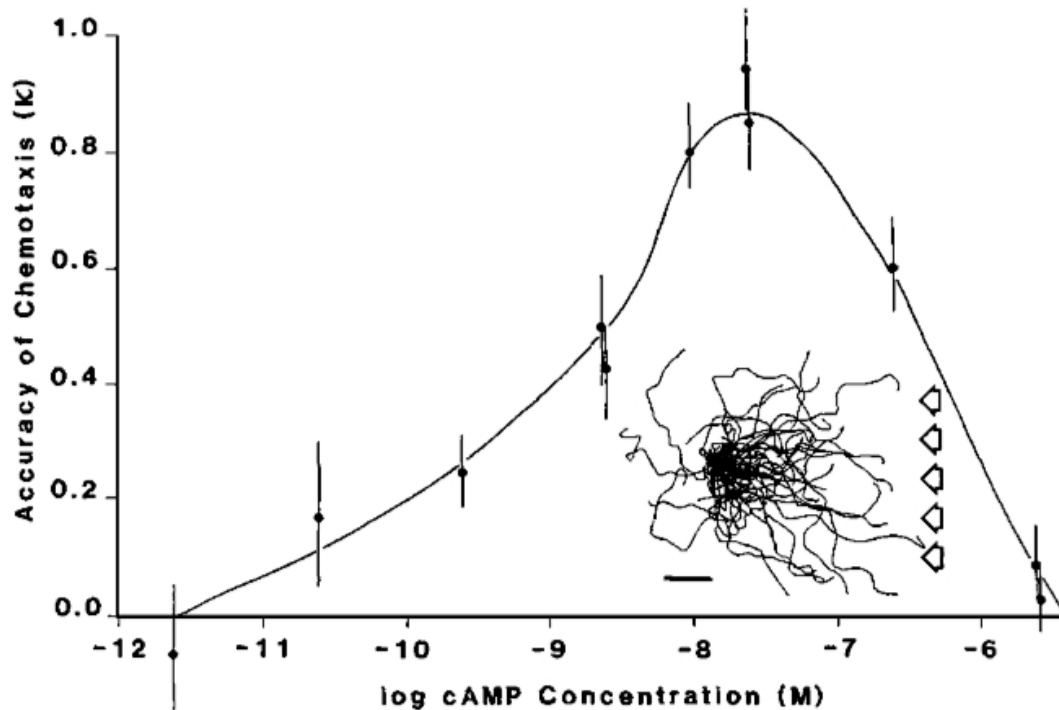
# Information Theory and Directed Cell Migration (Chemotaxis)



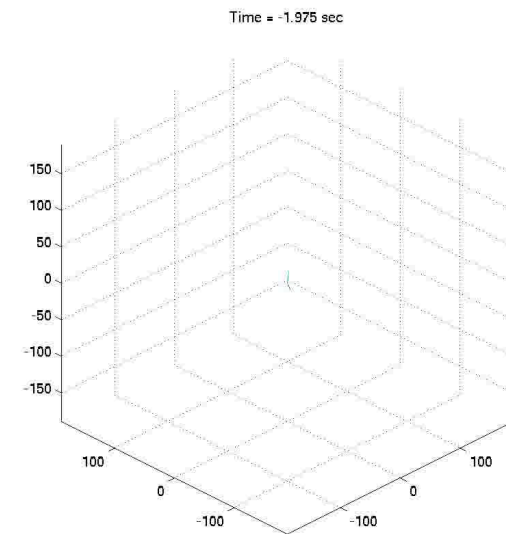
*Dictyostelium discoideum* amoeba detecting cAMP gradients. G. Gerisch.



Gradient sensing *via* cAMP receptors  
Kimmel, Salter & Thomas 2006 NIPS



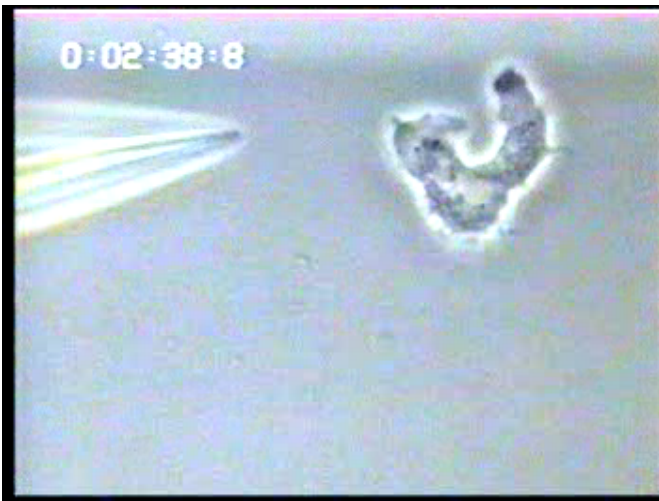
Cell tracks provide individual unit responses.  
Fischer et al., J. Cell Biology, 1989



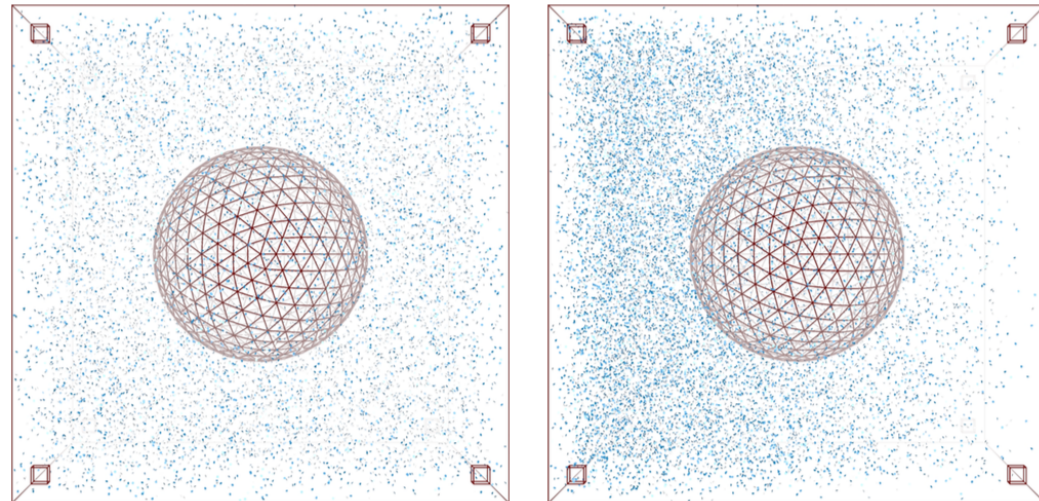
Ensemble of directional estimates:  
shallow gradient



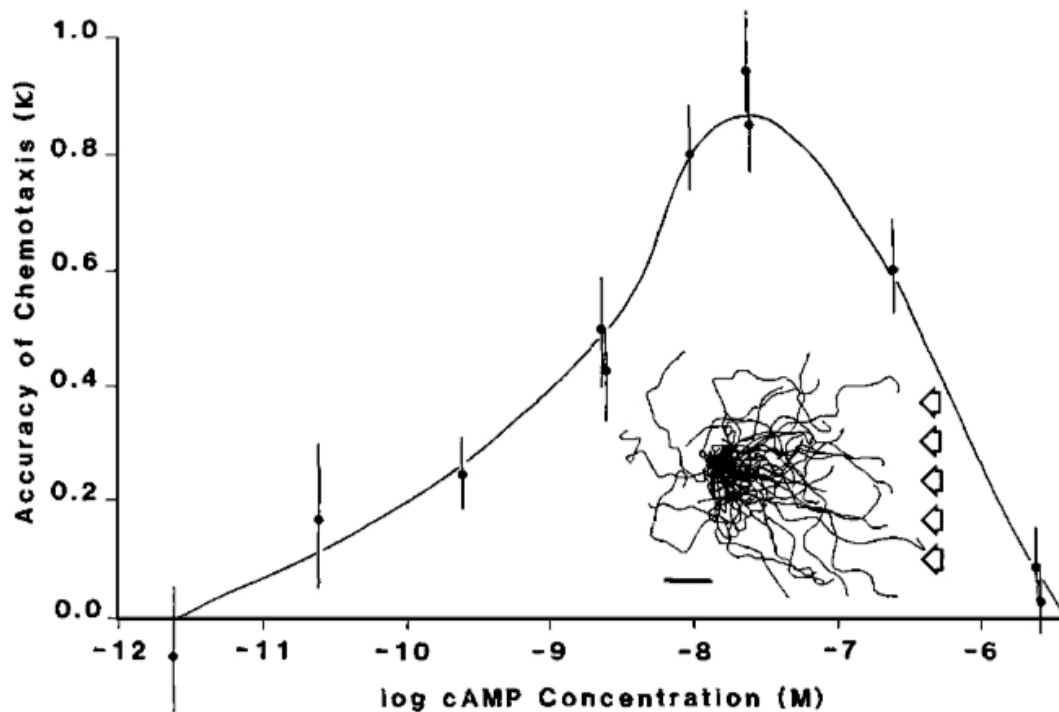
# Information Theory and Directed Cell Migration (Chemotaxis)



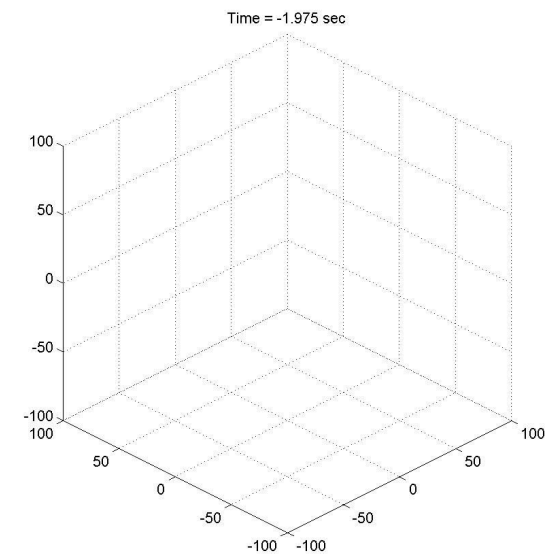
*Dictyostelium discoideum* amoeba detecting cAMP gradients.  
G. Gerisch.



Gradient sensing *via* cAMP receptors  
Kimmel, Salter & Thomas 2006 NIPS

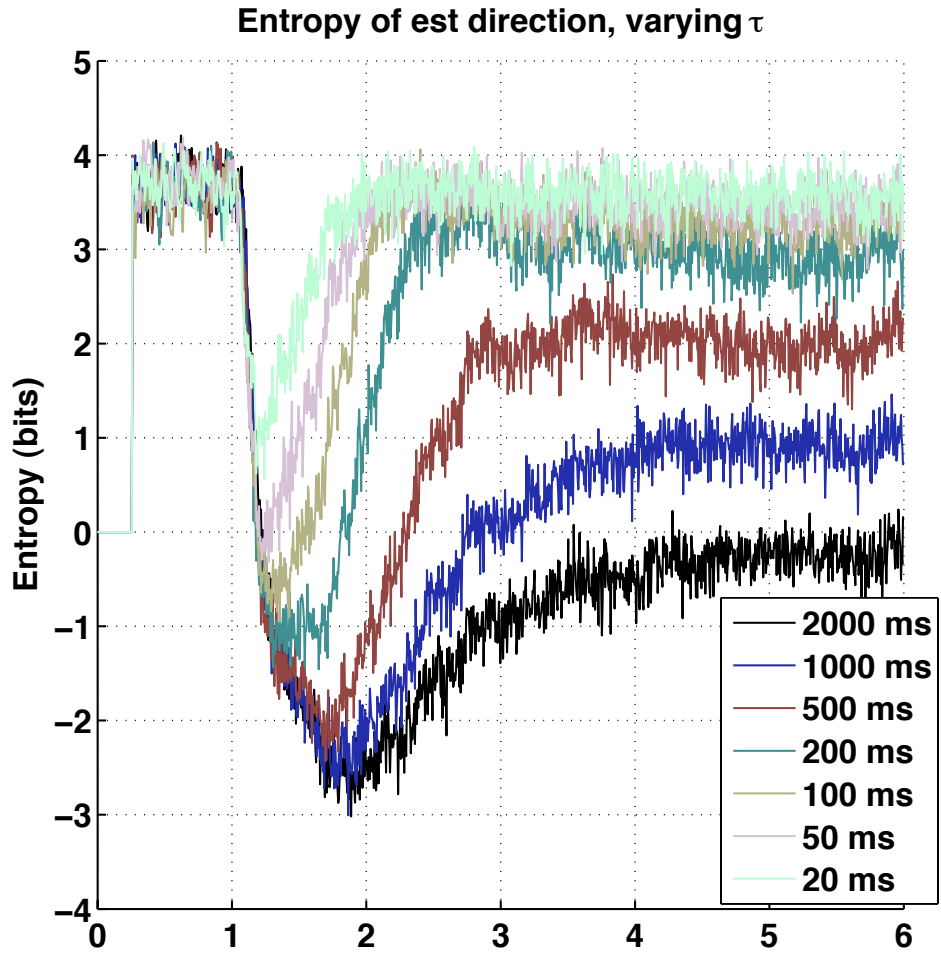


Cell tracks provide individual unit responses.  
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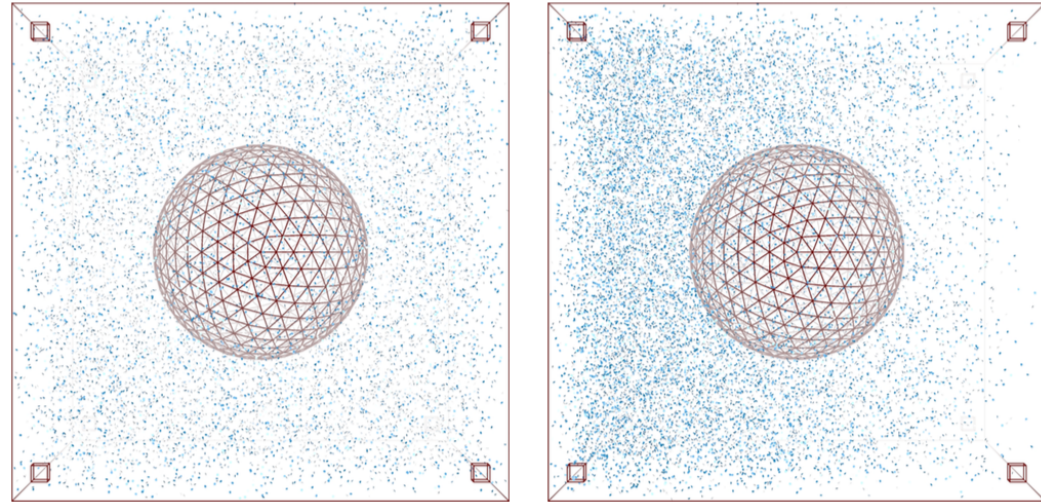
Ensemble of directional estimates:  
steep gradient

# Information Theory and Directed Cell Migration (Chemotaxis)

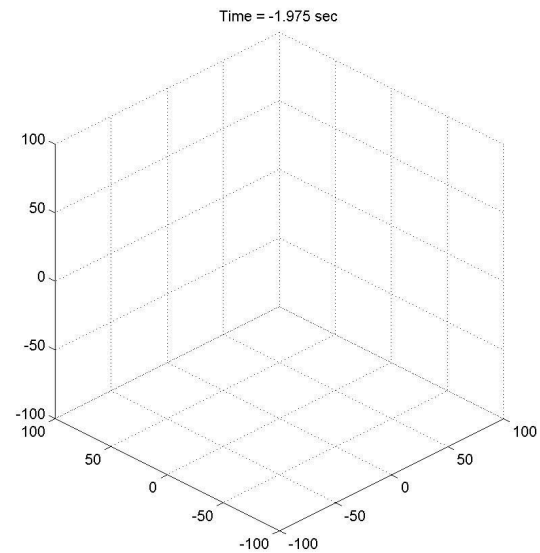


Mutual information peaks then recedes for saturating gradient signal (200 nM mean concentration;  $K_{eq}=25\text{nM}$ ).

Tau = filtering time scale for estimate.



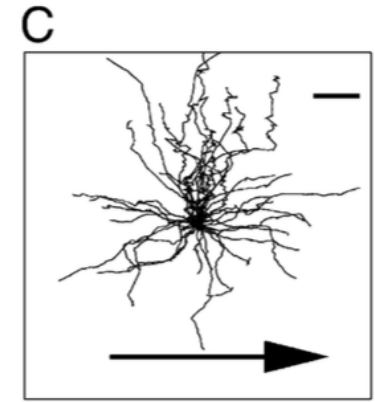
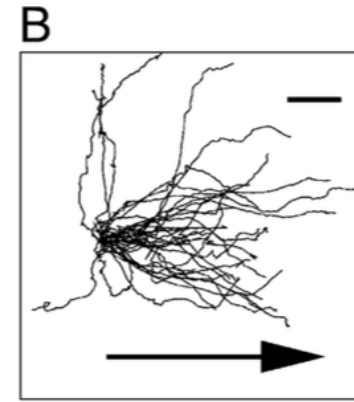
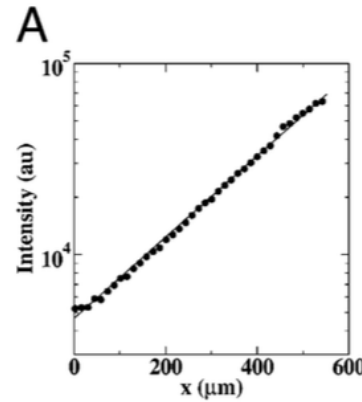
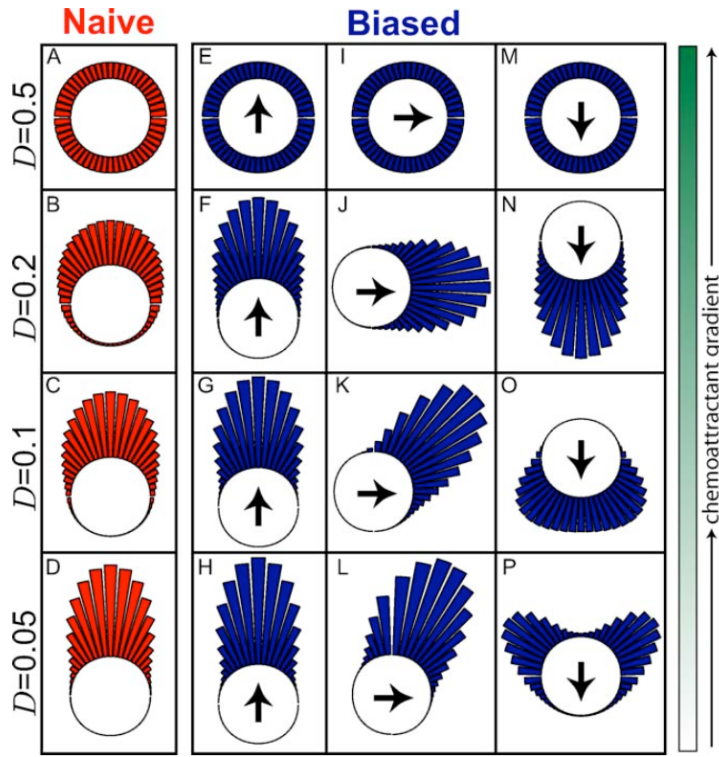
Gradient sensing *via* cAMP receptors  
Kimmel, Salter & Thomas 2006 NIPS



Ensemble of directional estimates:  
steep gradient



# Information Theory and Directed Cell Migration (Chemotaxis)

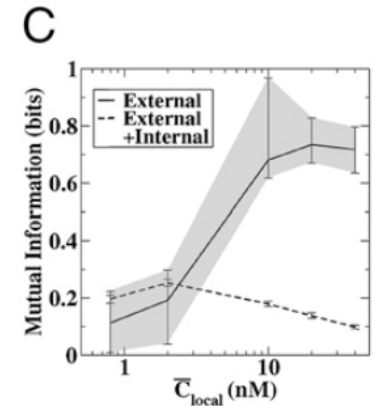
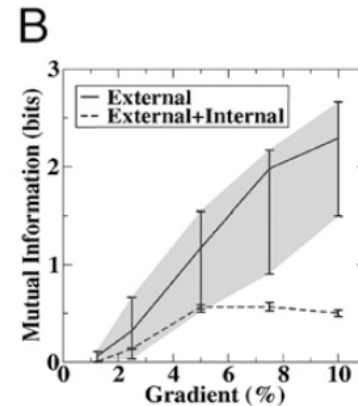
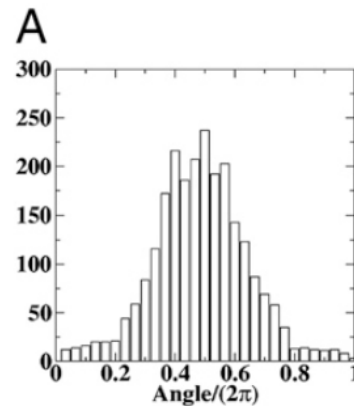


Cell tracks in a microfluidic device (exponential gradient)

(Fuller et al 2010 PNAS)

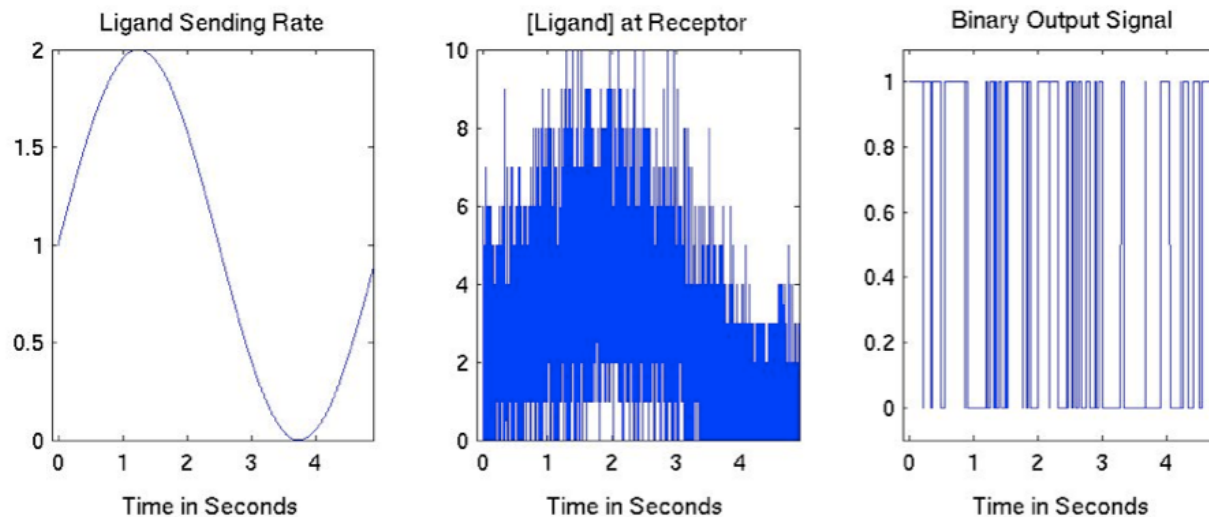
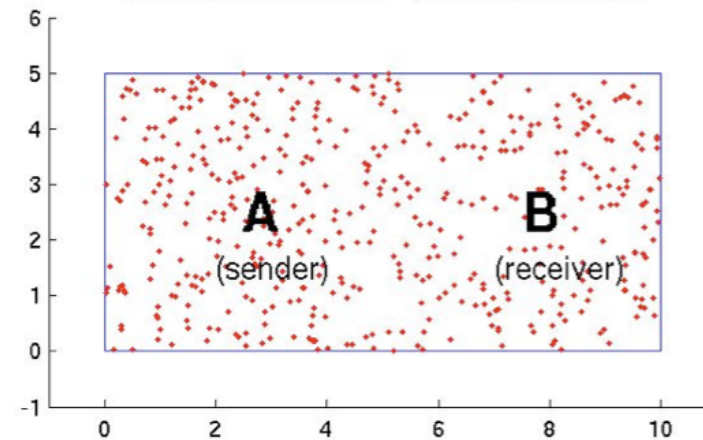
Mutual Information of chemotactic response;  
resolution of internal versus external noise sources.

Optimal Gradient Sensing Response  
Andrews & Iglesias 2007 PLoS CB



# Information Capacity of a Signal Transduction Channel

1. Secretion of signaling molecule
2. Diffusion from sender to receiver
3. Ligand binding to receptor protein

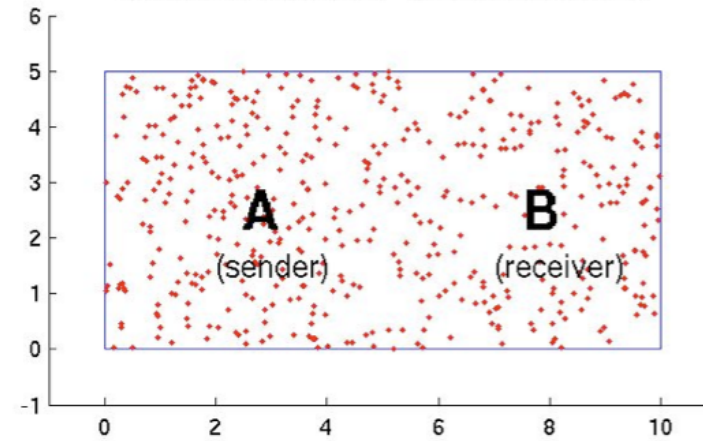


Thomas et al 2003 NIPS

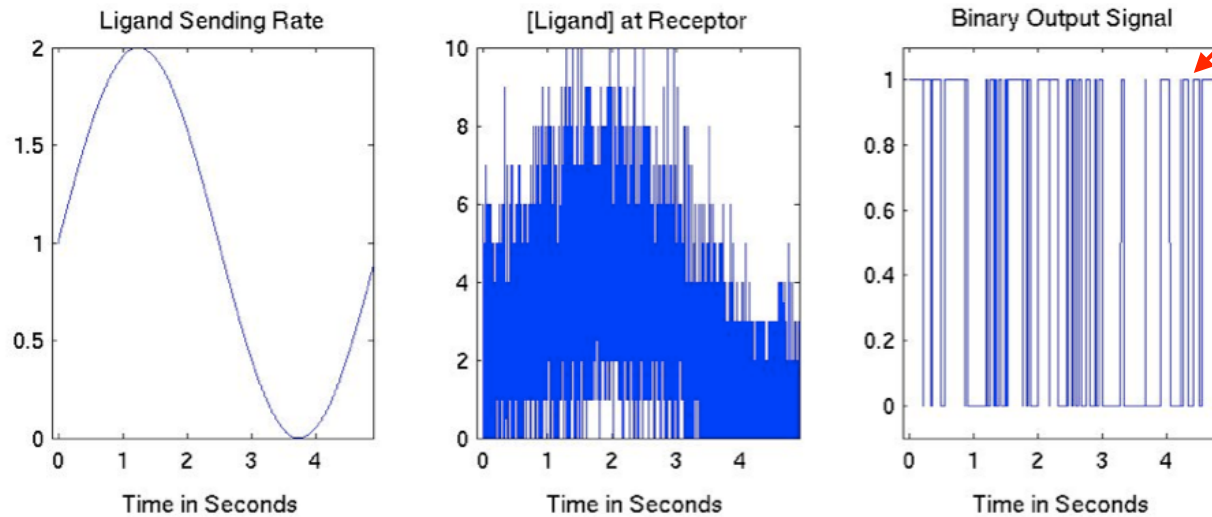


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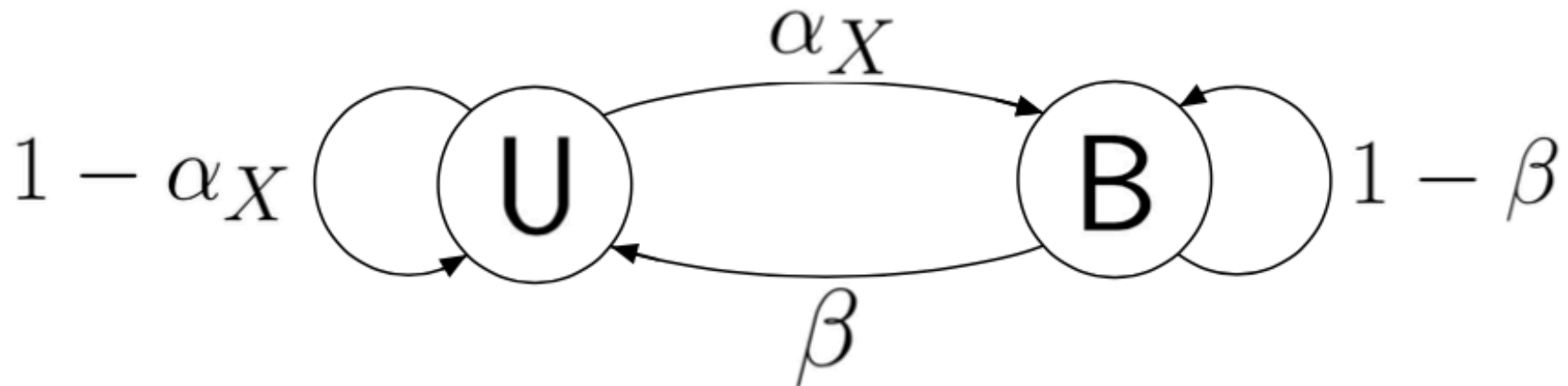
Ligand-receptor binding



Thomas et al 2003 NIPS

## Example 1: the BIND channel (a single binary receptor)

Receptor has only two states (Bound/Unbound).



$$\frac{dp}{dt} = k_+ c(t)(1 - p(t)) - k_- p(t)$$

$p$ : probability receptor is bound.

$k_+$ : binding rate constant.

$k_-$ : unbinding rate constant.

$$\alpha_L = k_+ c_{\min} \Delta t$$

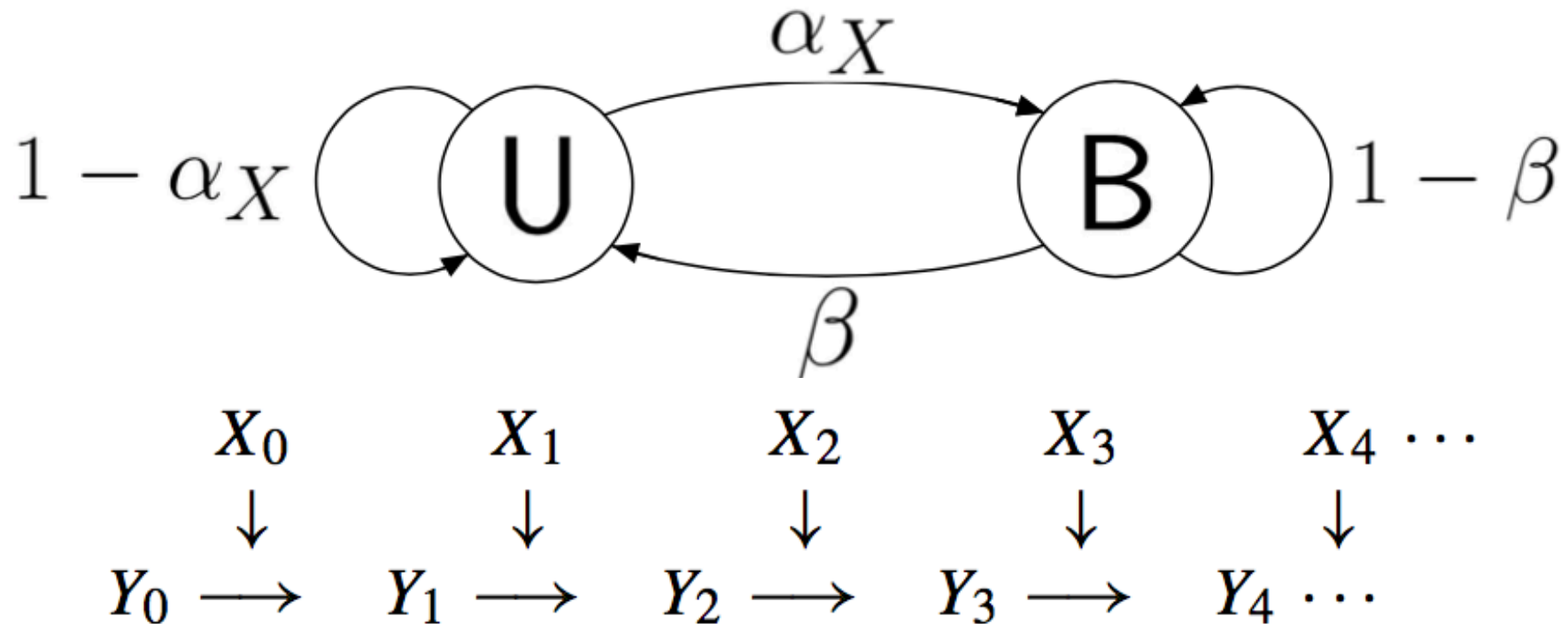
$$\alpha_H = k_+ c_{\max} \Delta t$$

$$\beta = k_- \Delta t$$



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## Example 1: the BIND channel (a single binary receptor)

Applying a general theorem due to Chen and Berger to the two-state discrete time Markov channel, we can show that

- 1 Capacity  $C$  of the discrete channel model is

$$C = \max_{p_H} \frac{\mathcal{H}(p_L \alpha_L + p_H \alpha_H) - p_L \mathcal{H}(\alpha_L) - p_H \mathcal{H}(\alpha_H)}{1 + (p_L \alpha_L + p_H \alpha_H)/\beta},$$

where  $p_L = 1 - p_H$  and  $\mathcal{H}(p) = p \log \frac{1}{p} + (1 - p) \log \frac{1}{1-p}$ .

- 2 The capacity cannot be increased by feedback.
- 3 The capacity can be realized by an IID input source.

Eckford & Thomas, 2013 International Symposium on Information Theory (ISIT)

Thomas & Eckford, 2016 IEEE Transactions on Information Theory



# Example 1: the BIND channel (a single binary receptor)

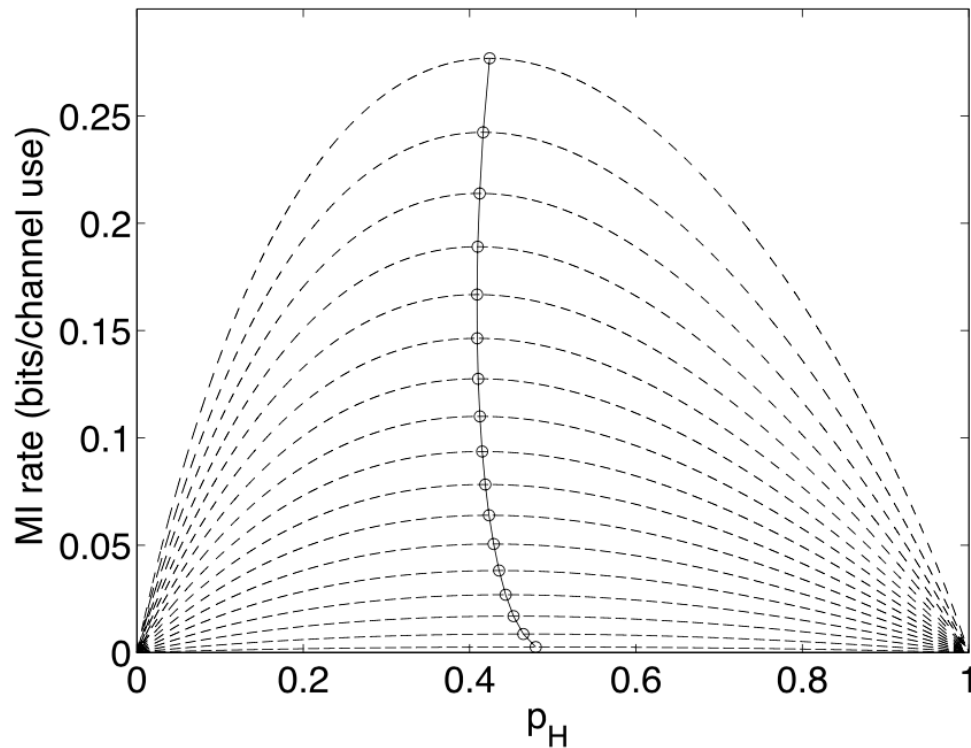


Fig. 3. Information maximizing values of  $p_H$ , with  $\alpha_L = 0.1$  and  $\beta = 0.9$ . Each dashed curve corresponds to a particular value of  $\alpha_H$ : from the bottom,  $\alpha_H = 0.15$ ; each higher curve increases  $\alpha_H$  by 0.05, up to  $\alpha_H = 0.95$  in the topmost curve. The maxima are circled and connected with a solid line.

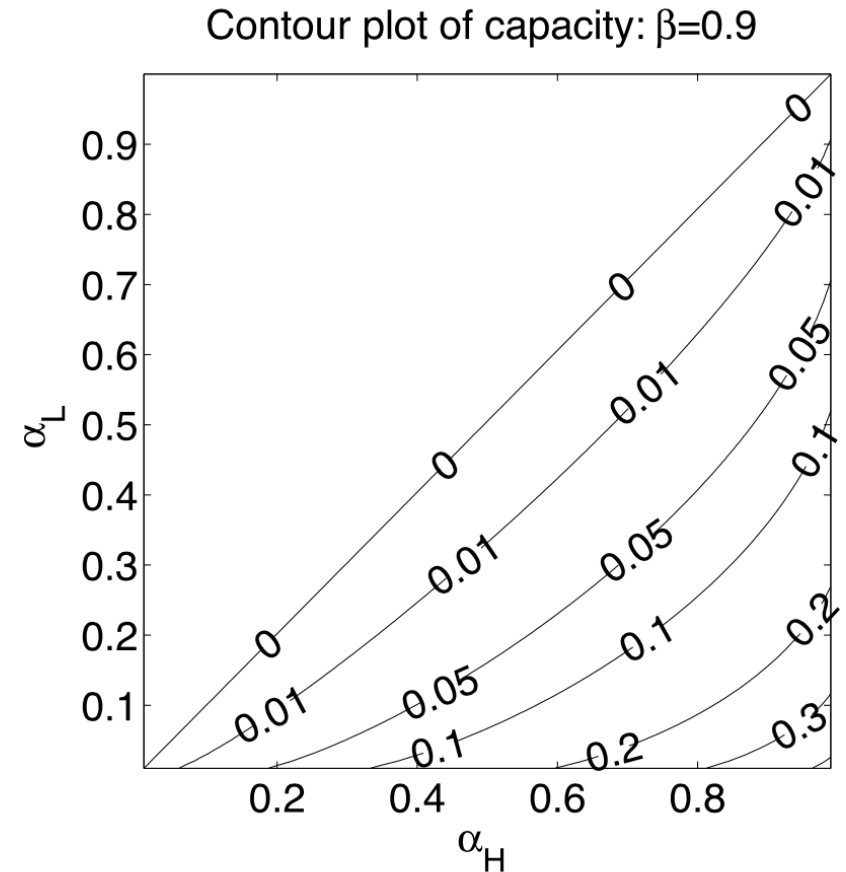


Fig. 4. Contour plot of capacity with respect to  $\alpha_L$  and  $\alpha_H$ , fixing  $\beta = 0.9$ . Note that  $\alpha_L > \alpha_H$  in the upper left triangle, so capacity here is undefined.

Thomas & Eckford 2016 IEEE Transactions on Information Theory

## Example 1: the BIND channel (a single binary receptor)

The mutual information decomposes:

$$I(X : Y) = \langle \mathbf{1}_{Y=U} \rangle (\mathcal{H}(\langle \alpha \rangle) - \langle \mathcal{H}(\alpha) \rangle)$$

As  $\Delta t \rightarrow 0$  we obtain a continuous time mutual information rate

$$\mathcal{I}(x) = - \left( \frac{\beta}{\beta + \bar{\alpha}} \right) (\bar{\alpha} \log(\bar{\alpha}) - (x\alpha_H \log \alpha_H + (1-x)\alpha_L \log \alpha_L))$$

of the same form.

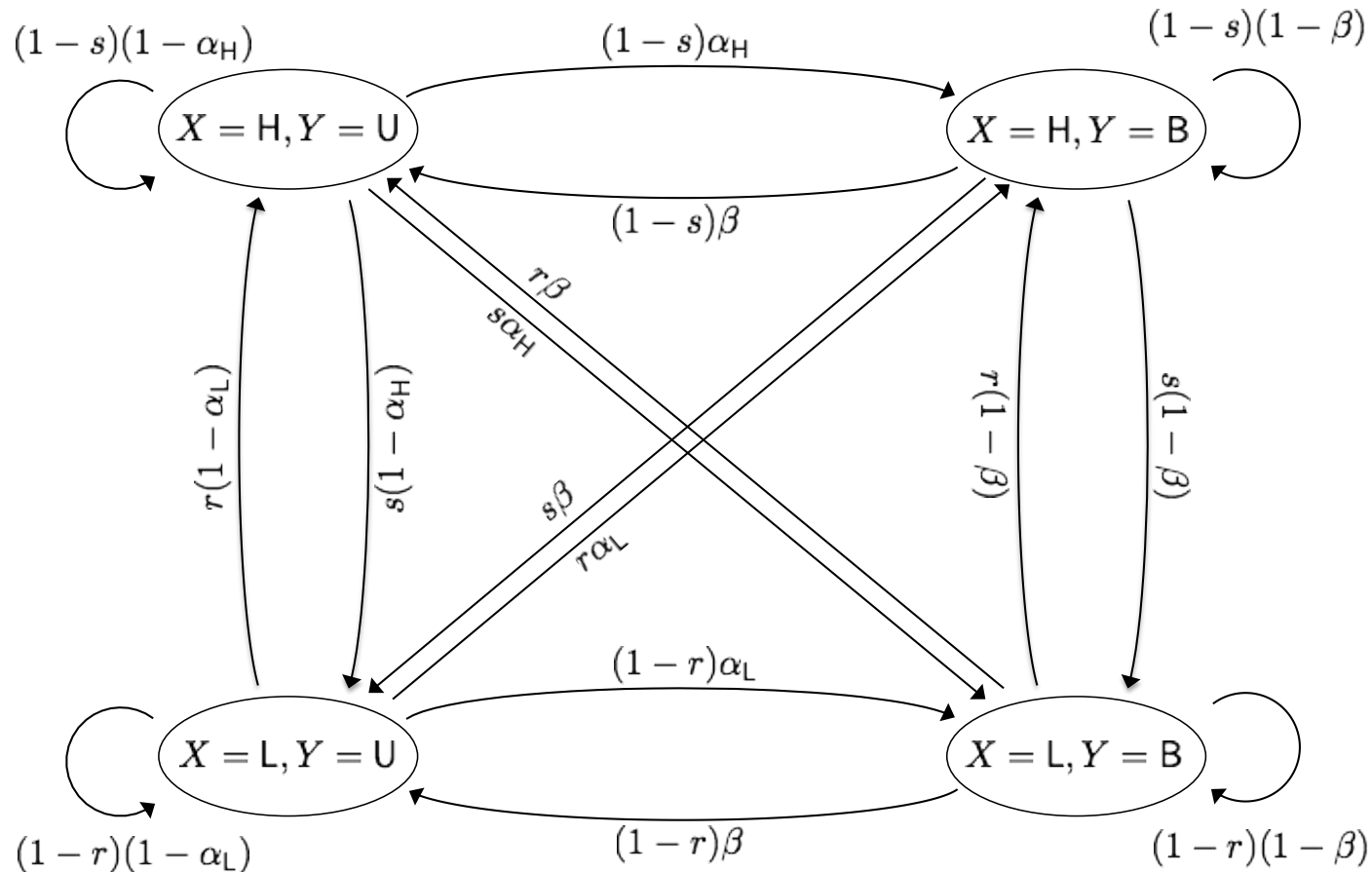
Here  $x$  is fraction of time input concentration signal is “high”. In the limit of rapid unbinding ( $\beta \rightarrow \infty$ ) we recover Kabanov’s capacity for the Poisson channel:

$$C_{\text{Kab}}(m, \lambda) = \frac{1}{e} (\lambda + m) \left( 1 + \frac{m}{\lambda} \right)^{\lambda/m} - \lambda \left( 1 + \frac{\lambda}{m} \right) \log \left( 1 + \frac{m}{\lambda} \right)$$

where  $\lambda = \alpha_L$  and  $m = \alpha_H - \alpha_L$ .

# Example 1: the BIND channel (a single binary receptor)

## Special Case: Markov Inputs



Transition probabilities

$X : L \rightarrow H$  with prob.  $r$ .

$X : H \rightarrow L$  with prob.  $s$ .

$Y : U \rightarrow B$  with prob.  $\alpha_{L/H}(X)$ .

$Y : B \rightarrow U$  with prob.  $\beta$ .



# Example 1: the BIND channel (a single binary receptor)

Joint process is Markov on four states:  $X \in \{L, H\}, Y \in \{U, B\}$ .

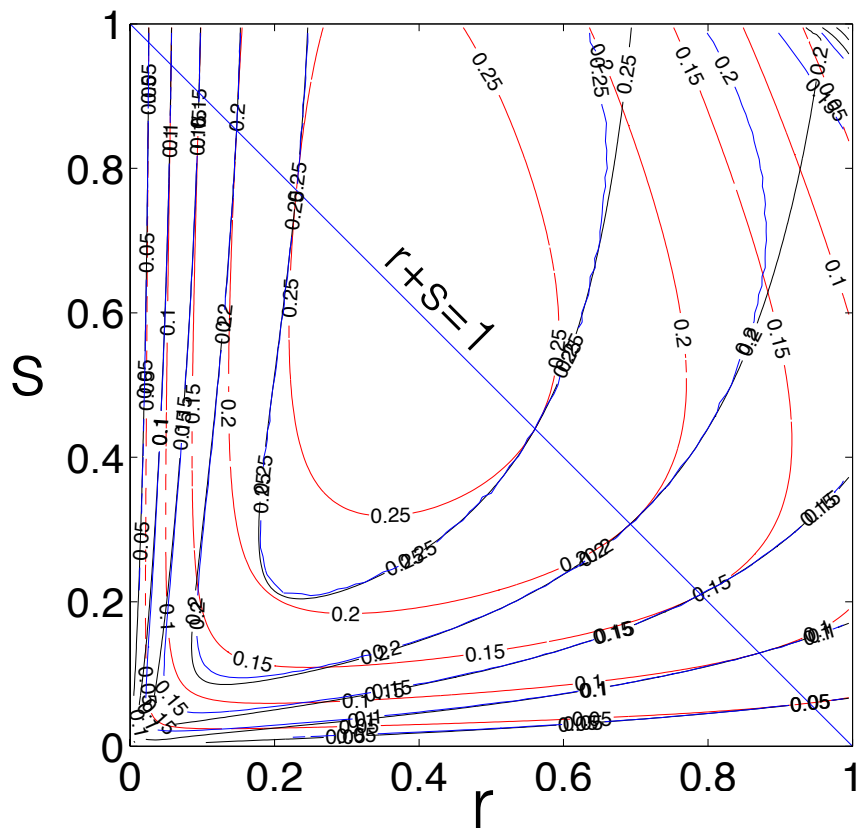
$$\mathcal{I}(X : Y) = \mathcal{H}(X, Y) - \mathcal{H}(X) - \mathcal{H}(Y)$$

Entropy rates  $\mathcal{H}(X, Y)$  and  $\mathcal{H}(X)$  are known in closed form.

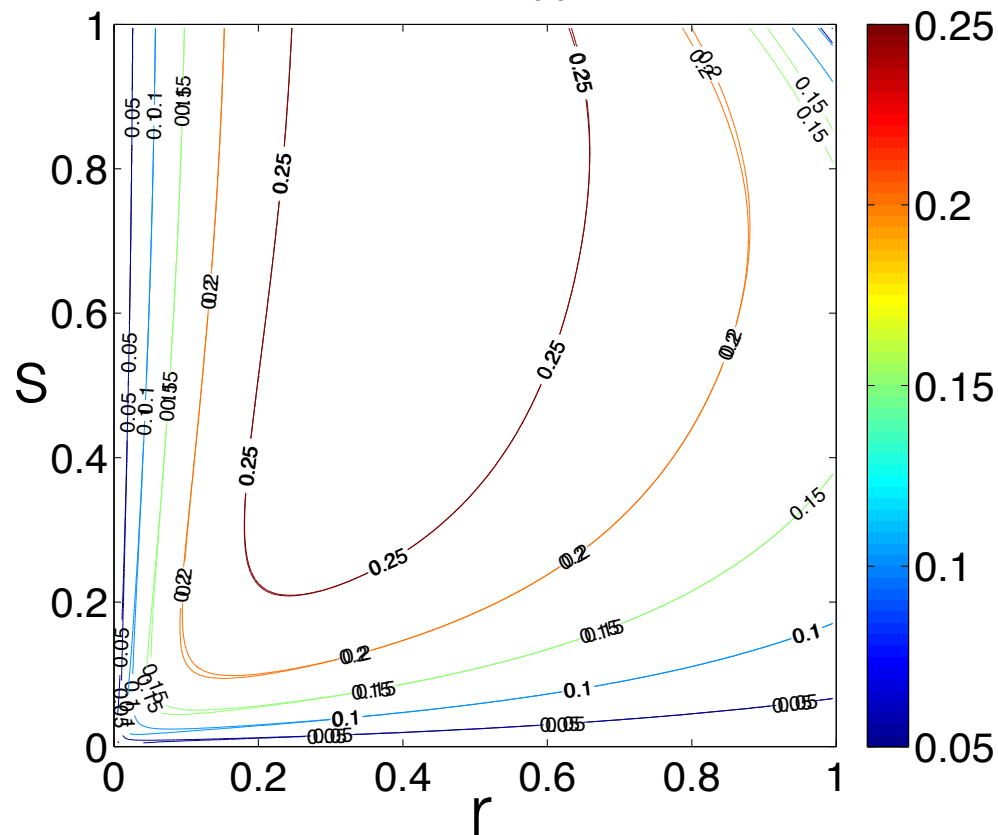
Approximate  $\mathcal{H}(Y) = \lim_{n \rightarrow \infty} H(Y_n | Y_{n-1}, \dots, Y_0)$  with

$$H(Y_n | Y_{n-1}, \dots, Y_0, X_0) \leq \mathcal{H}(Y) \leq H(Y_n | Y_{n-1}, \dots, Y_0)$$

Lower and Upper Bounds of MI (Depth=2)

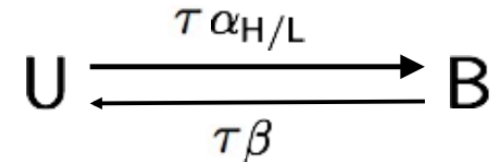


Lower Bound 5 and Upper Bound 5

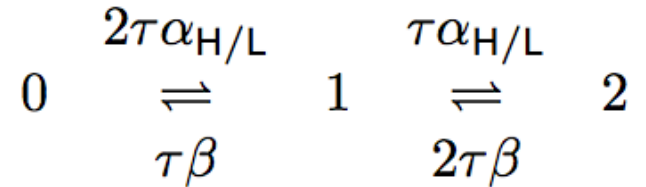


# Multiple Independent Receptors

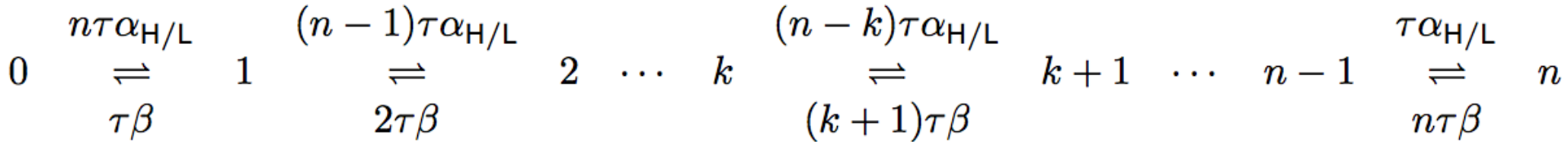
\* One BIND receptor.



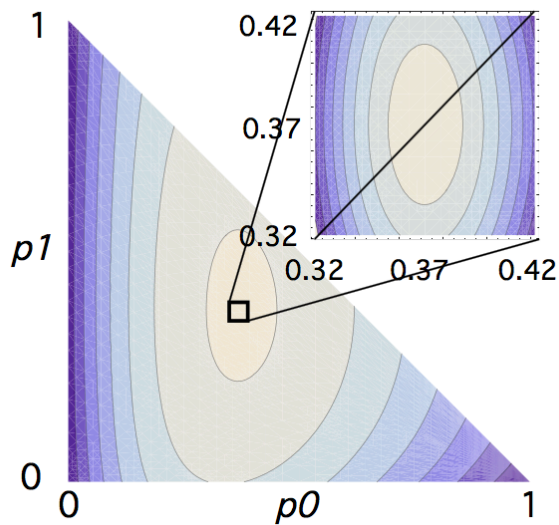
\* Two independent BIND receptors



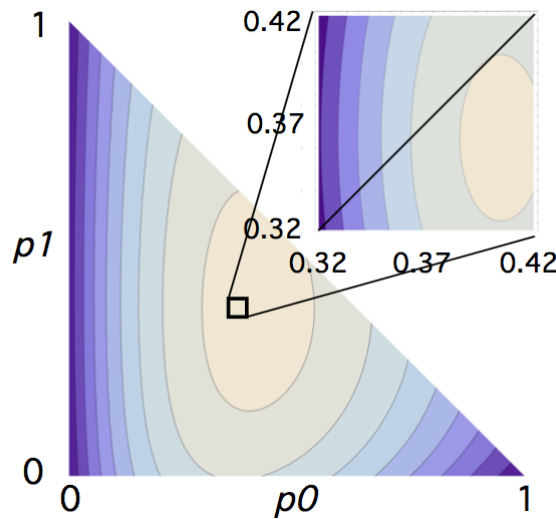
\* n independent BIND receptors



two independent receptors



two non-independent receptors

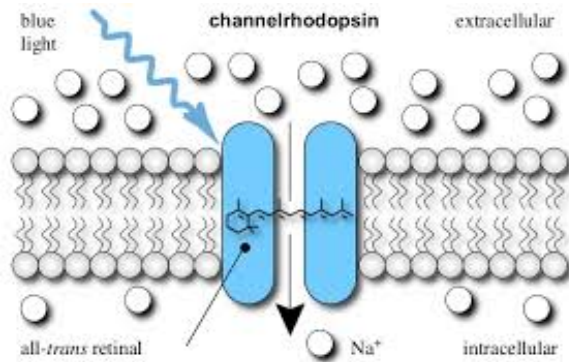


Capacity for n independent receptors is n times the single-receptor capacity.

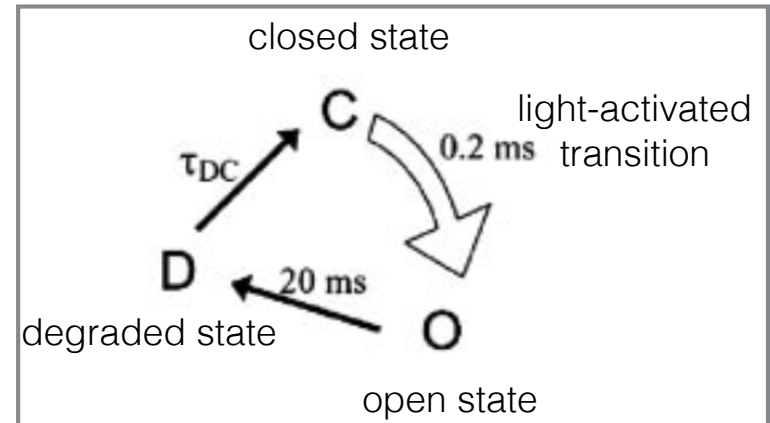
Thomas and Eckford 2016 ISIT

# Example 2: Channelrhodopsin

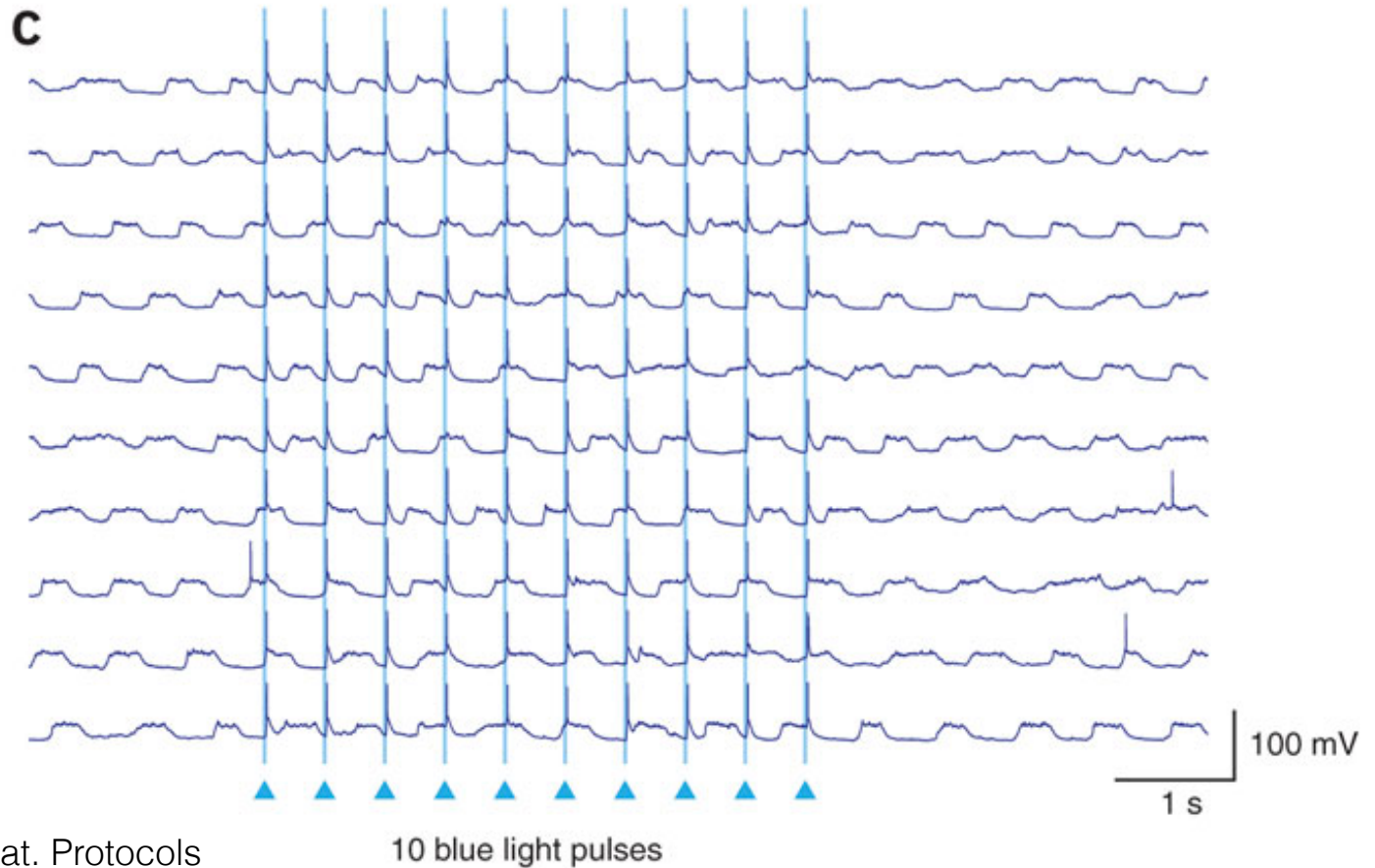
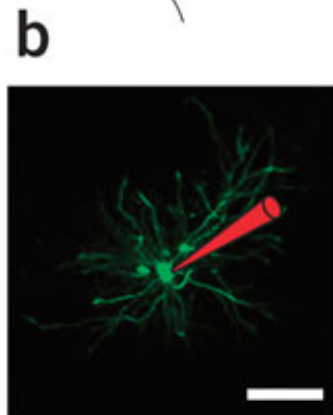
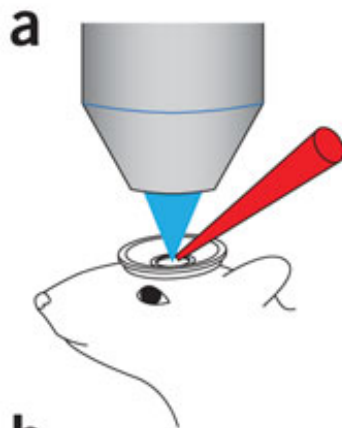
(Eckford & Thomas, under review)



openoptogenetics.org



Nagel et al 2003 PNAS

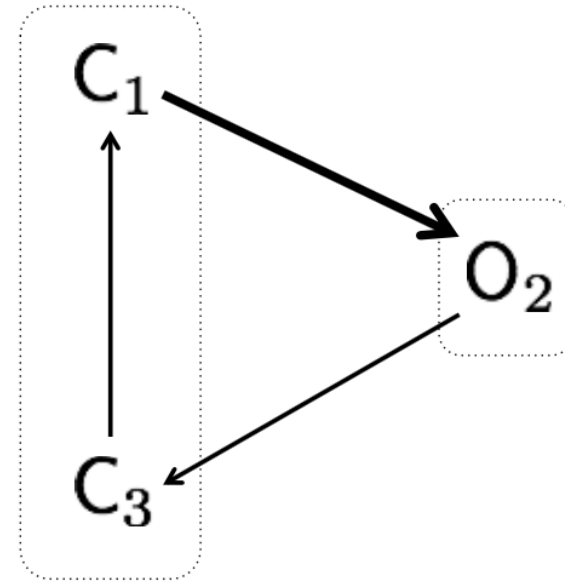
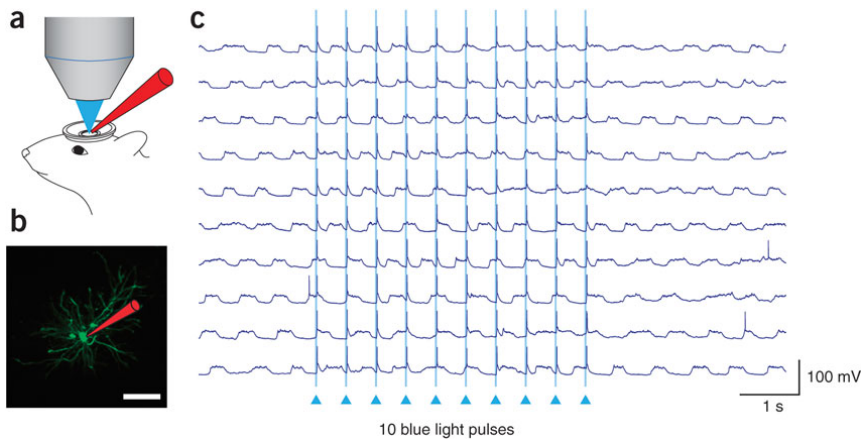


Judkewitz et al 2009 Nat. Protocols



# Example 2: Channelrhodopsin

(Eckford & Thomas, under review)



$$Q = \begin{bmatrix} R_1 & q_{12}x(t) & 0 \\ 0 & R_2 & q_{23} \\ q_{31} & 0 & R_3 \end{bmatrix}$$

Parameter	from [2]	Units
$q_{12}x(t)$	$(5 \times 10^3)x(t)$	$s^{-1}$
$q_{23}$	50	$s^{-1}$
$q_{31}$	17	$s^{-1}$

Input: Light intensity

Channel States: 2 closed, 1 open.

One sensitive state; one observable transition.

# Channelrhodopsin under IID inputs

(Eckford & Thomas, under review)

- ▶ If the input sequence is IID, the channel state  $\{Y(k\Delta t)\}_{k \geq 0}$  forms a Markov chain<sup>1</sup> with stationary distribution  $\pi_y$ .
- ▶ The mutual information between input  $X$  & channel state  $Y$  is

$$\mathcal{I}(X; Y) = \sum_{(y_{i-1}, y_i) \in \mathcal{S}} \pi_{y_{i-1}} \left( \sum_{x_i \in \mathcal{X}} p(x_i) \phi(p(y_i | x_i, y_{i-1})) - \phi \left( \sum_{x_i \in \mathcal{X}} p(x_i) p(y_i | x_i, y_{i-1}) \right) \right).$$

Here  $\mathcal{S}$  is the set of sensitive transitions and  $\phi(p) = p \log p$ .

- ▶ For channelrhodopsin we find

$$\begin{aligned} \mathcal{I}(X; Y) &= \pi_{C_1} \left( \mathcal{H}(p_L \Delta t q_{12}^{x_L} + p_H \Delta t q_{12}^{x_H}) - p_L \mathcal{H}(\Delta t q_{12}^{x_L}) - p_H \mathcal{H}(\Delta t q_{12}^{x_H}) \right) \\ &= \frac{\mathcal{H}(p_L \Delta t q_{12}^{x_L} + p_H \Delta t q_{12}^{x_H}) - p_L \mathcal{H}(\Delta t q_{12}^{x_L}) - p_H \mathcal{H}(\Delta t q_{12}^{x_H})}{1 + E[x](q_{12}/q_{23} + q_{12}/q_{31})}, \end{aligned}$$

where  $E[x] = p_L x_L + p_H x_H$  is the average input intensity.

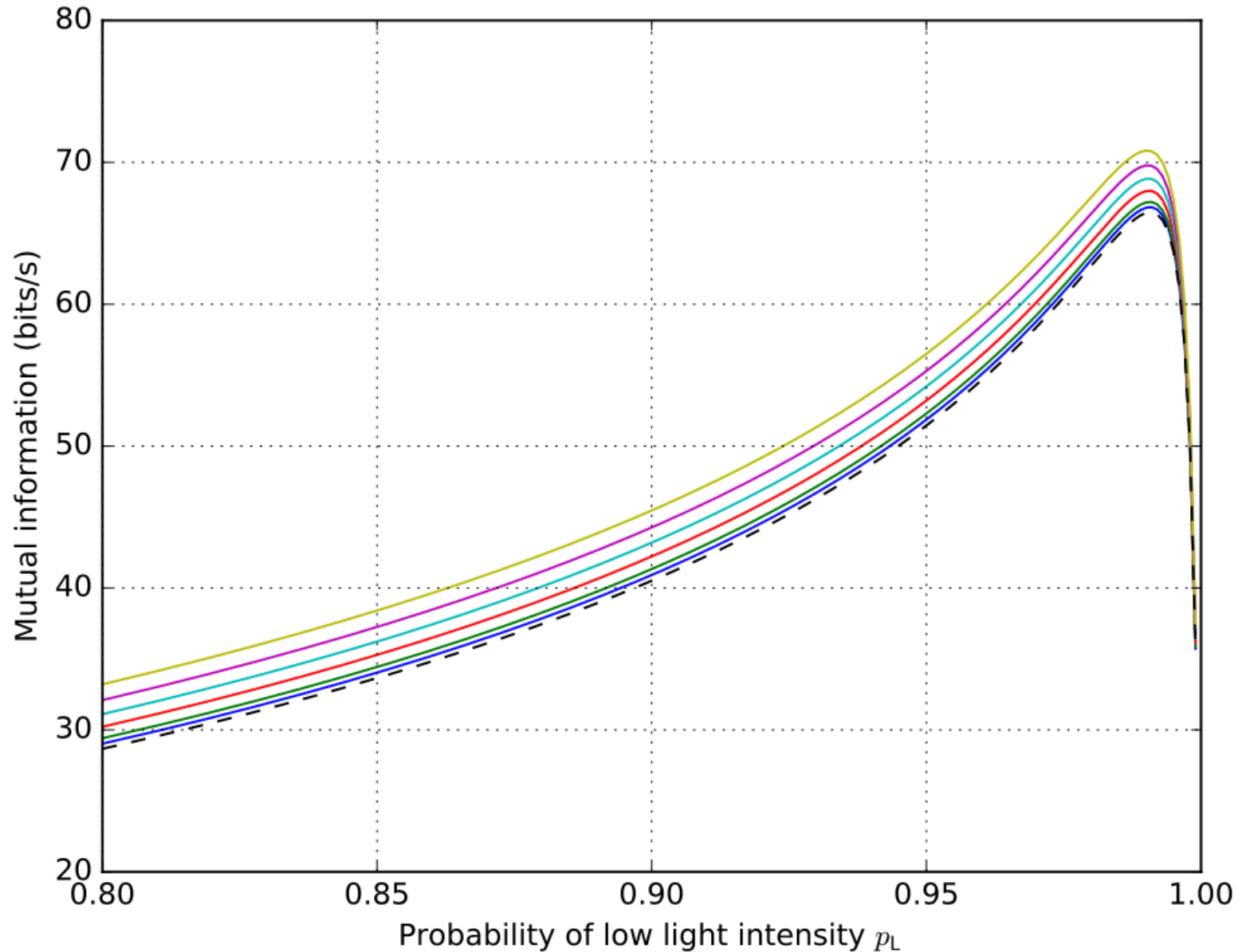
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<sup>1</sup>  $Y$  is time-homogeneous, irreducible, aperiodic, and positive recurrent.

# Channelrhodopsin under IID inputs

*(Eckford & Thomas, under review)*

As  $\Delta t \rightarrow 0$  the mutual information rate  $I(X; Y)$  converges.





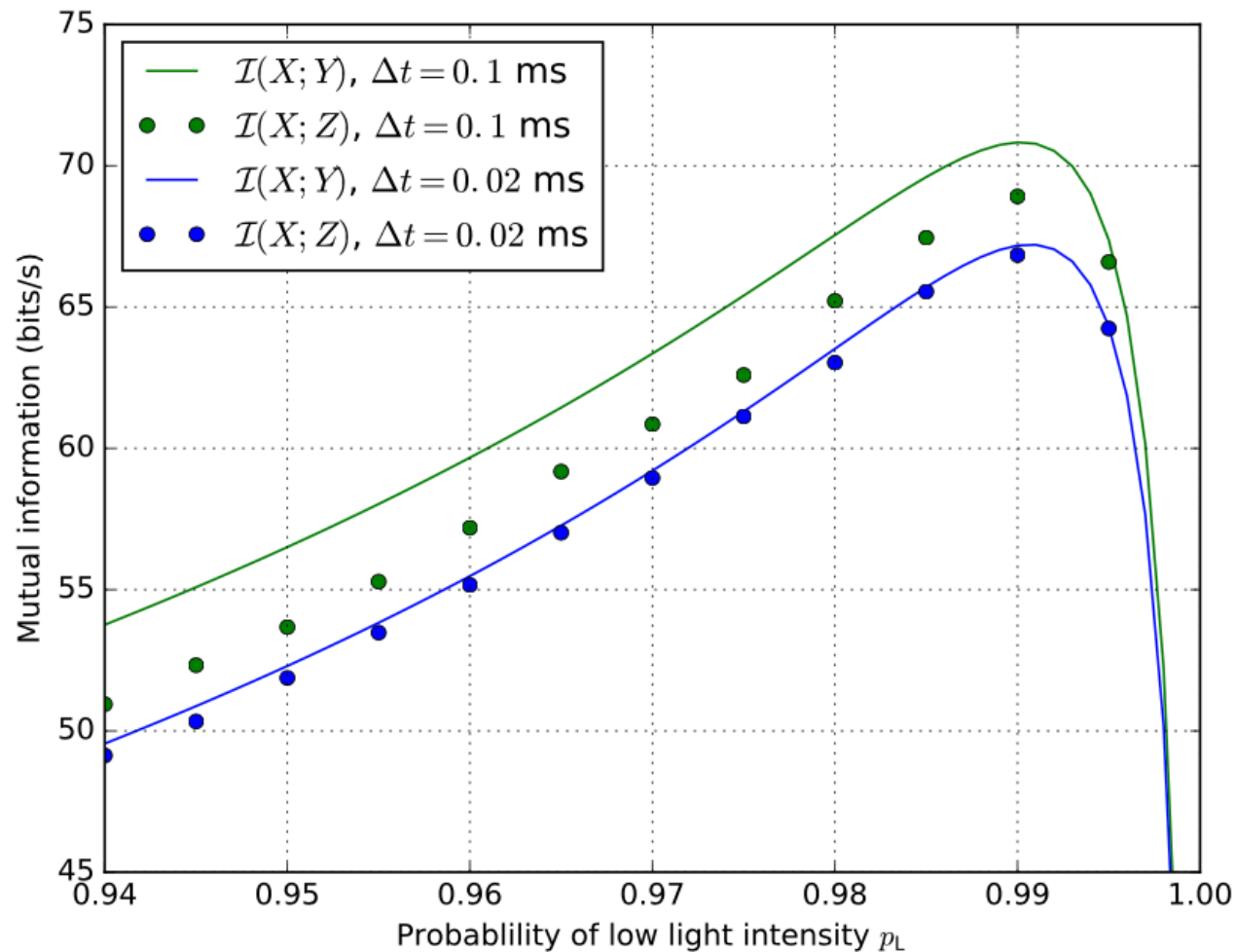
# Channelrhodopsin under IID inputs

(Eckford & Thomas, under review)

For the *partially observed* channel,  $I(X; Z) \leq I(X; Y)$ .

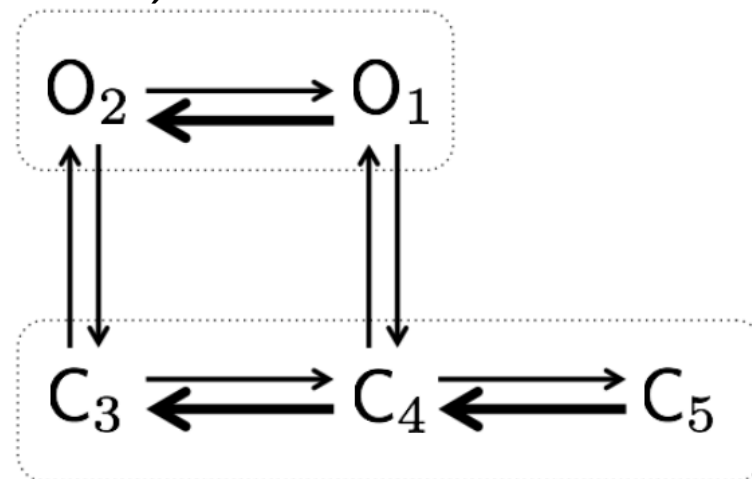
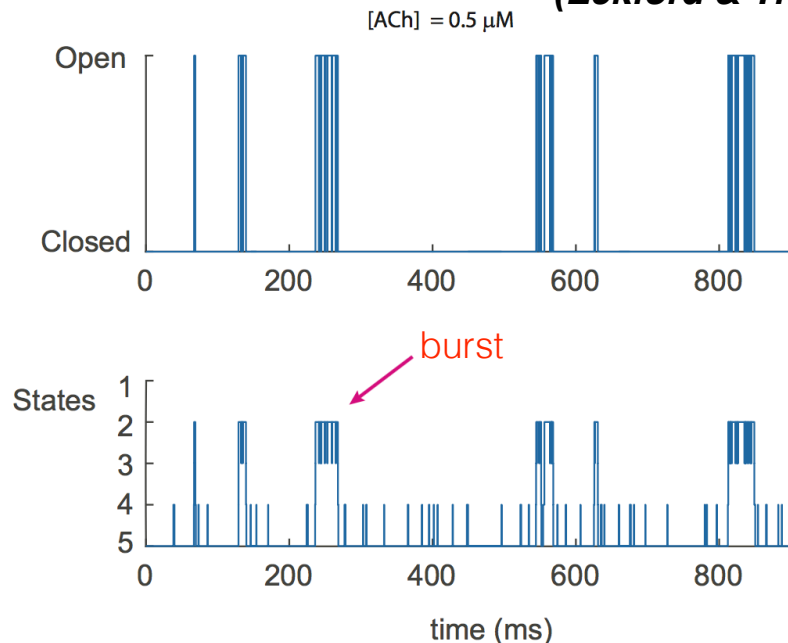
For channelrhodopsin, all sensitive transitions are observable.

Monte Carlo estimates suggest that as  $\Delta t \rightarrow 0$ , we have  $I(X; Z) \rightarrow I(X; Y)$ .



# Example 3: Acetylcholine receptor

(Eckford & Thomas, under review)



$$Q = \begin{bmatrix} R_1 & q_{12}x(t) & 0 & q_{14} & 0 \\ q_{21} & R_2 & q_{23} & 0 & 0 \\ 0 & q_{32} & R_3 & q_{34} & 0 \\ q_{41} & 0 & q_{43}x(t) & R_4 & q_{45} \\ 0 & 0 & 0 & q_{54}x(t) & R_5 \end{bmatrix}$$

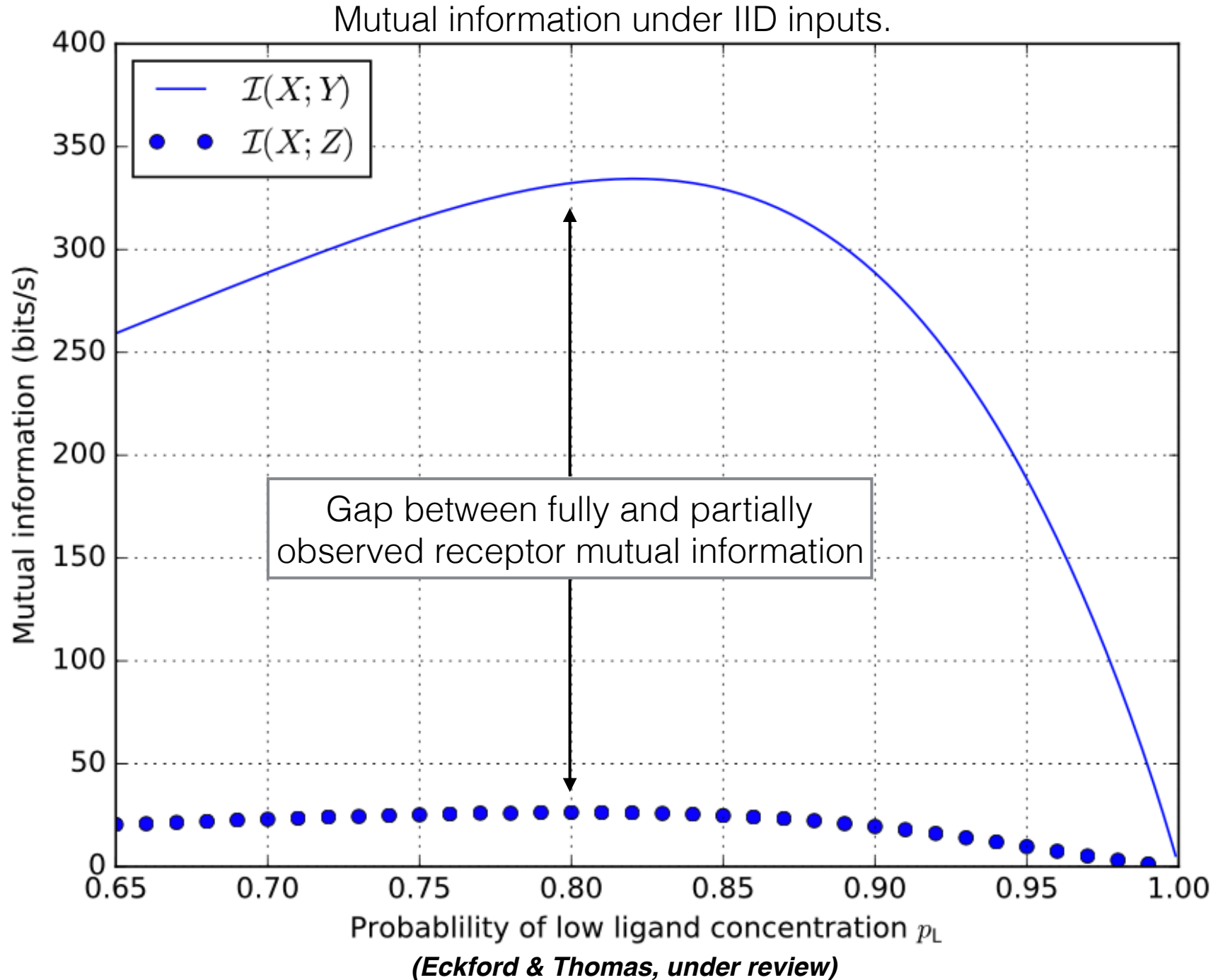
Input: Acetylcholine [ACh] concentration

Channel States: 3 closed, 2 open

Three sensitive states; 4 observable transitions.

Parameter	Name in [3]	Value/range	Units
$q_{12}x(t)$	$k_{+2}x$	$(5 \times 10^8)x(t)$	$s^{-1}$
$q_{14}$	$\alpha_1$	$3 \times 10^3$	$s^{-1}$
$q_{21}$	$2k_{-2}^*$	0.66	$s^{-1}$
$q_{23}$	$\alpha_2$	$5 \times 10^2$	$s^{-1}$
$q_{32}$	$\beta_2$	$1.5 \times 10^4$	$s^{-1}$
$q_{34}$	$2k_{-2}$	$4 \times 10^3$	$s^{-1}$
$q_{41}$	$\beta_1$	15	$s^{-1}$
$q_{43}x(t)$	$k_{+2}x$	$(5 \times 10^8)x(t)$	$s^{-1}$
$q_{45}$	$k_{-1}$	$2 \times 10^3$	$s^{-1}$
$q_{54}x(t)$	$2k_{+1}x$	$(1 \times 10^8)x(t)$	$s^{-1}$

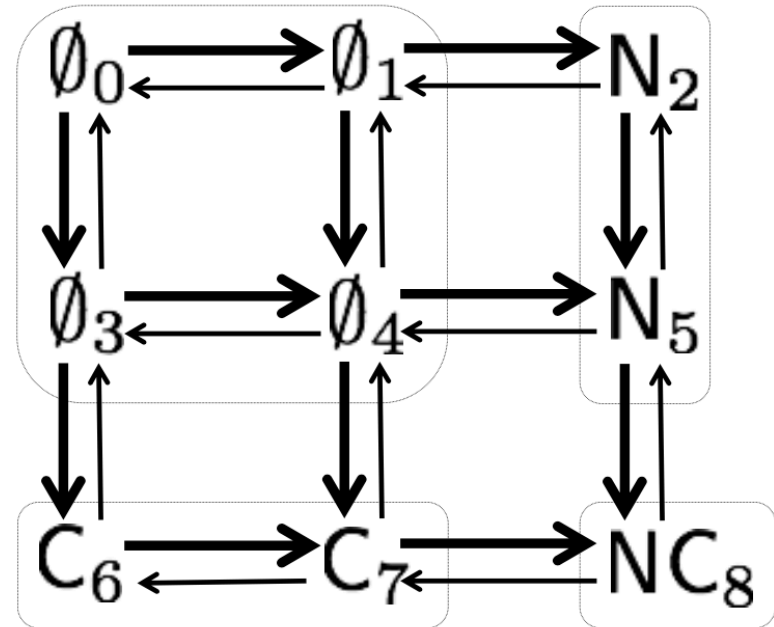
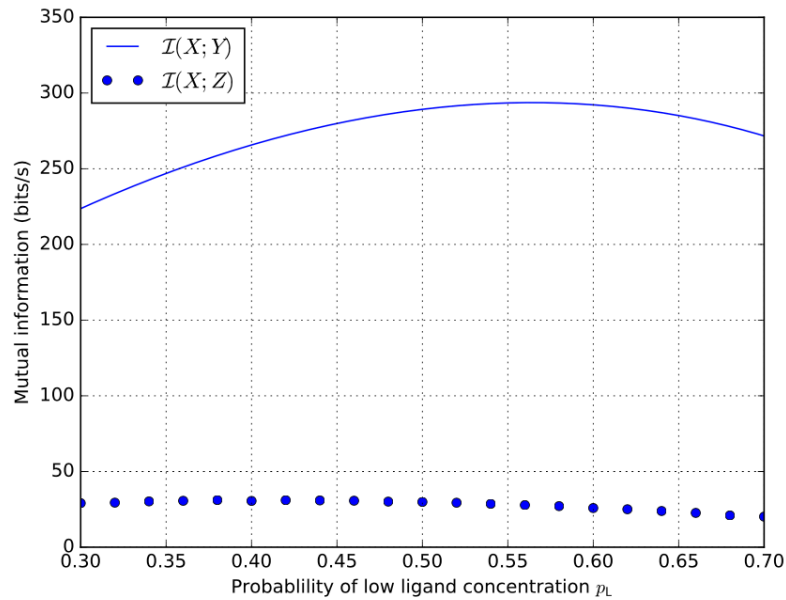
# Example 3: Acetylcholine receptor





# Example 4: Calmodulin (a Calcium binding protein)

(Eckford & Thomas, under review)



$$\begin{bmatrix}
 R_0 & q_{01}x(t) & 0 & q_{03}x(t) & 0 & 0 & 0 & 0 & 0 \\
 q_{10} & R_1 & q_{12}x(t) & 0 & q_{14}x(t) & 0 & 0 & 0 & 0 \\
 0 & q_{21} & R_2 & 0 & 0 & q_{25}x(t) & 0 & 0 & 0 \\
 q_{30} & 0 & 0 & R_3 & q_{34}x(t) & 0 & q_{36}x(t) & 0 & 0 \\
 0 & q_{41} & 0 & q_{43} & R_4 & q_{45}x(t) & 0 & q_{47}x(t) & 0 \\
 0 & 0 & q_{52} & 0 & q_{54} & R_5 & 0 & 0 & q_{58}x(t) \\
 0 & 0 & 0 & q_{63} & 0 & 0 & R_6 & q_{67}x(t) & 0 \\
 0 & 0 & 0 & 0 & q_{74} & 0 & q_{76} & R_7 & q_{78}x(t) \\
 0 & 0 & 0 & 0 & 0 & q_{85} & 0 & q_{87} & R_8
 \end{bmatrix}$$

Parameter	Name in [4]	Value/range	Units
$q_{01}x(t), q_{34}x(t), q_{67}x(t)$	$k_{\text{on}(T),N}$	$(7.7 \times 10^8)x(t)$	$\text{s}^{-1}$
$q_{10}, q_{43}, q_{76}$	$k_{\text{off}(T),N}$	$1.6 \times 10^5$	$\text{s}^{-1}$
$q_{12}x(t), q_{45}x(t), q_{78}x(t)$	$k_{\text{on}(R),N}$	$(3.2 \times 10^{10})x(t)$	$\text{s}^{-1}$
$q_{21}, q_{54}, q_{87}$	$k_{\text{off}(R),N}$	$2.2 \times 10^4$	$\text{s}^{-1}$
$q_{03}x(t), q_{14}x(t), q_{25}x(t)$	$k_{\text{on}(T),C}$	$(8.4 \times 10^7)x(t)$	$\text{s}^{-1}$
$q_{30}, q_{41}, q_{52}$	$k_{\text{off}(T),C}$	$2.6 \times 10^3$	$\text{s}^{-1}$
$q_{36}x(t), q_{47}x(t), q_{58}x(t)$	$k_{\text{on}(R),C}$	$(2.5 \times 10^7)x(t)$	$\text{s}^{-1}$
$q_{63}, q_{74}, q_{85}$	$k_{\text{off}(R),C}$	6.5	$\text{s}^{-1}$

Input: Calcium [Ca<sup>2+</sup>] concentration

Channel States: 9 states of 4 types

8 sensitive states

6 of 12 sensitive transitions are also observable.

## Conclusions

- ▶ Bio to Engineering: Intensity-driven signal transduction systems provide a broad class of biologically motivated communications channel models.
- ▶ Engineering to Bio: Fully and partially observed channel models under IID inputs are amenable to capacity analysis.
- ▶ The mutual information gap between fully and partially observed channels depends on the observability of those edges which are sensitive to the input.

## Ongoing work

- ▶ Capacity of general  $N$ -state intensity-driven receptor.
- ▶ Gap between partially & fully observed receptors: theory?
- ▶ Net capacity of ligand secretion, diffusion, binding channel.
- ▶ Energetics: metabolic burden; information cost vs fitness.

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