Why are chemotaxis receptors clustered but other receptors aren't?

With thanks to: Monica Skoge (UCSD) and Yigal Meir (Ben Gurion) Konstantin Doubrovinski (Princeton), Anirvan Sengupta (Rutgers), Robert Endres (Imperial College), Juan Keymer (Delft), Clinton Hansen (Harvard) Victor Sourjik, Olga Oleksiuk (Heidelberg)

Support from NIH and HFSP

Outline

- Introduction to chemotaxis in *E. coli*The "engineering" challenge
 The chemotaxis network and receptors
- FRET reveals that receptors cooperate

 Cooperativity adapts
- Other receptors don't cluster or cooperate
- Cooperativity and noise
- So why do chemotaxis receptors cooperate?
- Conclusions

E. coli chemotaxis: runs and tumbles



(Thanks to Howard Berg.)

Principles of Chemoreceptor "Engineering"

- High gain via receptor cooperativity
- Broad range via adaptation (integral feedback)



The chemotaxis network (best studied network in biology)



http://www.rowland.harvard.edu/l abs/bacteria/projects_fret.html

Chemoreceptor clustering

Receptors are clustered globally, and locally form trimers of dimers, arranged in a honeycomb lattice.

al.

(2009)



In vivo FRET studies of receptor activity



Real-time measurement of rate of phosphorylation of CheY.

Sourjik and Berg (2002)

Receptors cooperate in teams

PCA of FRET data (Tar-only strains)

Ŧ

Δ

盇

 10^{-1}

 $\Delta \Delta$

Â

team size





 10°



 10^{-2}

MeAsp [mM]

 \triangle QQQQ

QEQQ

QEQE

OEEE

 \times CheRB⁺, 0 mM

CheRB⁺, 0.1 mM

 10^{-3}

Receptor Activity

0.8

0.6

0.4

 $\rightarrow 0.2$

0

 10^{-4}

Why don't other receptors cooperate?

Dictyostelium cells have ~uniform cAR1-GFP distribution (cAMP receptor)



Traynor et al. (2007)

V. harveyi quorumsensing doseresponse curves have no gain



Could cooperativity increase noise?

Model for cooperative receptors with switching noise



Ising model of receptor clusters:

With Glauber's "heat bath" dynamics:







1D chain with n = 10 receptors

- Signal increases with coupling J (cooperativity).
- Then signal decreases as response slows.

Noise



- Noise increases with coupling J.
- Longer *T*_{avg} reduces noise.

Signal-to-Noise Ratio (SNR)



SNR is best for independent receptors!

Scaling relations for SNR



SNR is always best for independent receptors!

- Slow ligand dynamics?
- Extrinsic noise?
- We're asking the wrong question?

- Slow ligan dynamics?
- Extrinsic noise?
- We're asking the wrong question?

Slow ligan dynamics?

• Extrinsionoise?

We're asking the wrong question?

We should be asking how to optimize chemotactic velocity, not SNR.

Effect of signal and noise on chemotaxis



- Noise reduces < v>_{drift}, but only gradually
- Signal increases <*v*>_{drift} *linearly*

Maximizing chemotactic velocity



"Run and tumble" strategy implies noise threshold, which sets optimal cooperativity.

Open questions

- When is cooperativity advantageous?
- In *E. coli* chemotaxis, what controls cooperativity and its adaptation, and why does cooperativity adapt?
- Why ~5,000 chemoreceptors?



Conclusions

- *E. coli* chemotaxis receptors cooperate
 - increase of gain
 - cooperativity adapts
- Cooperativity is bad for SNR
- Chemotactic velocity optimized by increasing signal, up to noise threshold
- Do other receptors optimize SNR?

Keymer *et al.*, PNAS (2006) Endres and Wingreen, PNAS (2006) Skoge, Endres, and Wingreen, Biophys J (2006) Hansen, Endres, and Wingreen, PLoS CB (2007) Endres *et al.*, MSB (2008)

Wang *et* al., PRL (2008) Greenfield *et al.*, PLoS Biology (2009)

Slow ligand dynamics



• SNR still best for independent receptors



 Correlation time is multiplied by ligand binding time, due to domain wall pinning

Correlation time





$$T_c \sim \exp(2J)$$

Static, extinsic receptor noise

E.g. noise from receptor methylation:





- Cooperativity amplifies static noise
- SNR still best for independent receptors

Correlation time in 2D



 $T_c \sim \exp(?)$

FRET data: two regimes of activity

Sourjik and Berg (2002)



Regime I:

- Activity low at zero attractant
- K_i small and ≈ constant

Regime II:

- Activity high (saturated) at zero attractant
- *K_i* large and increases with methylation

Two regimes of receptor activity consistent with 2-state receptor model.

Two regimes of a 2-state receptor



Data collapse

Receptor activity depends only on difference between on and off state free energies.



And data collapses and yields K_D s: $K_D^{off} = 25 \ \mu M, \ K_D^{on} \approx 0.5 \ mM$

Recovery time – confirms K_D values



Berg and Tedesco (1975)



$$t \sim \Delta f \sim \log \left(\frac{1 + C / K_D^{off}}{1 + C / K_D^{on}} \right)$$

 $\rightarrow K_D^{off} = 27 \,\mu\text{M}, \quad K_D^{on} \approx 0.9 \,\text{mM}$

Slow ligan dynamics?

• Extrinsionoise?

We're asking the wrong question?

We should be asking how to optimize chemotactic velocity, not SNR.