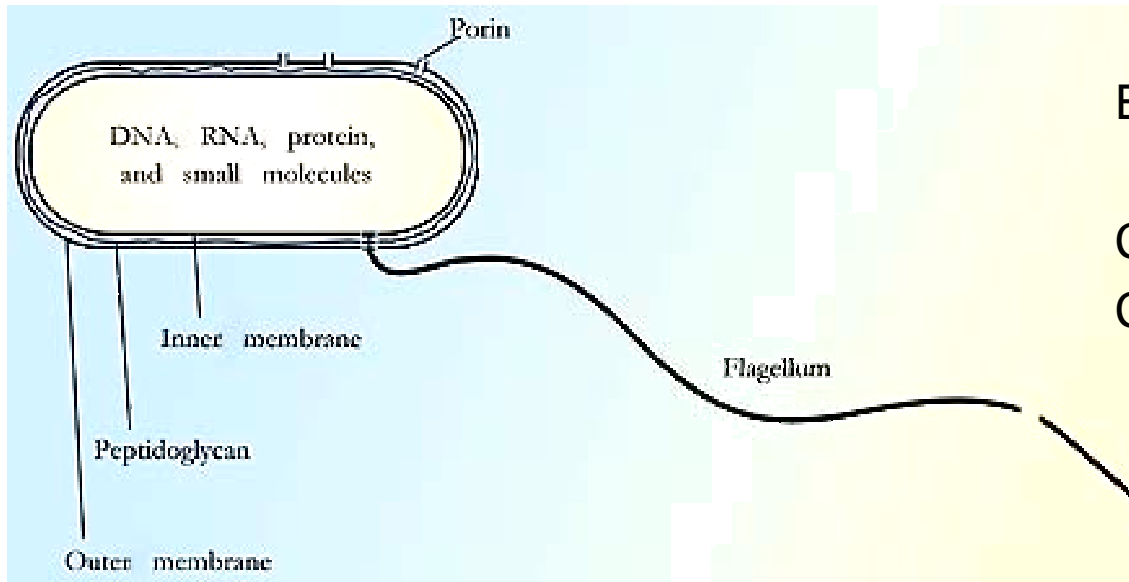


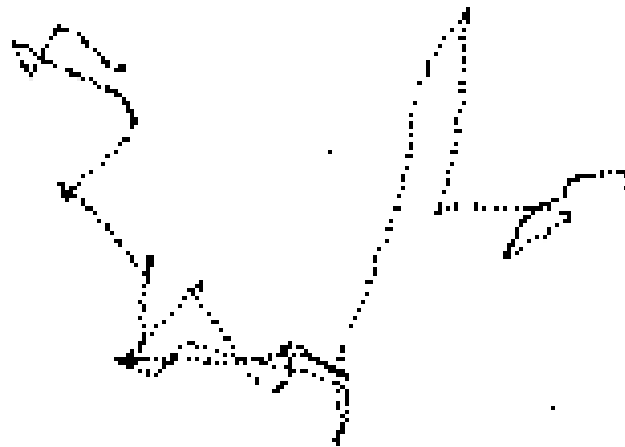
Bacterial chemotaxis and the question of high gain in signal transduction

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E. coli lives in the gut and takes up nutrients through its pores

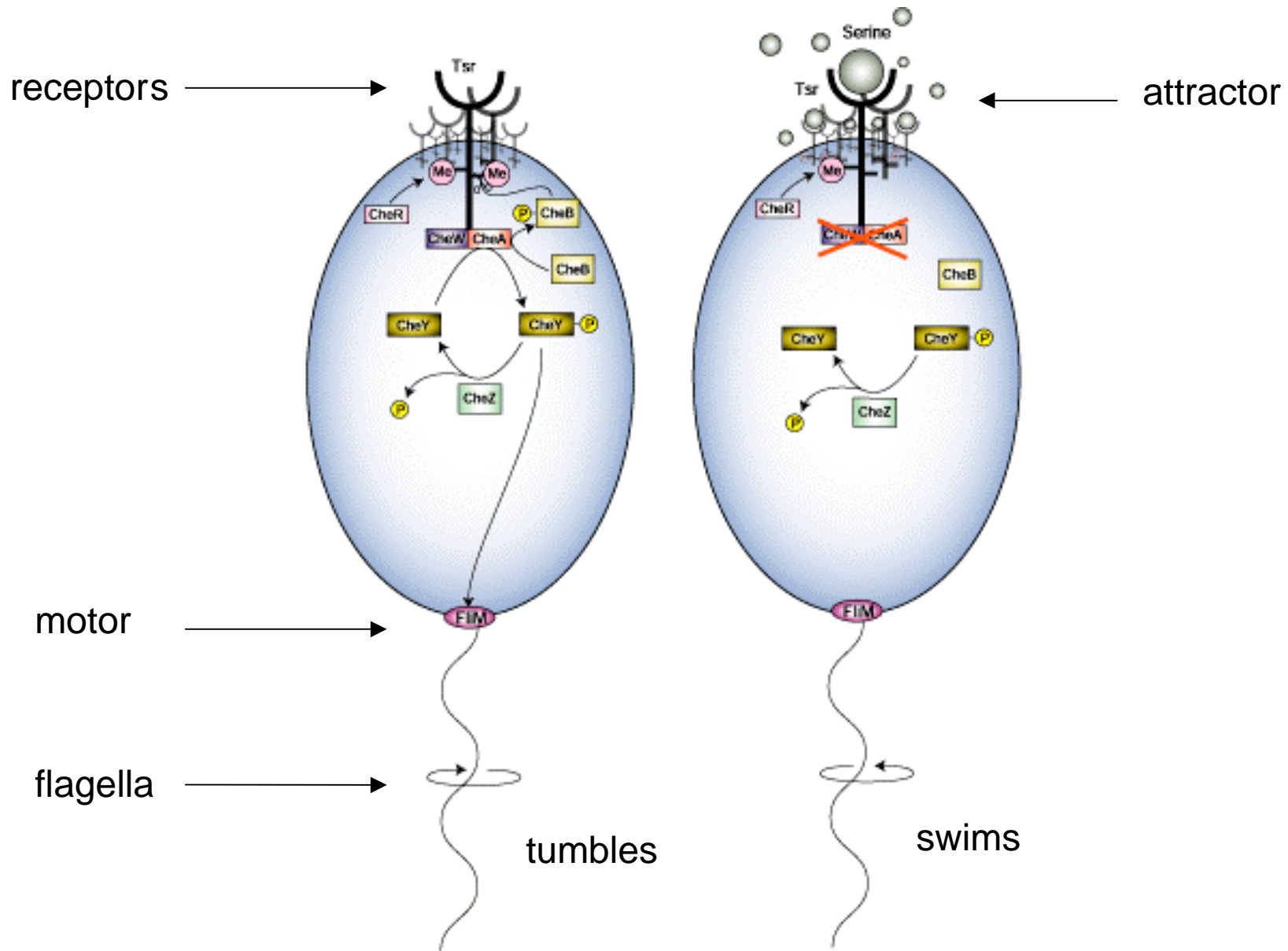


E. coli moves by rotating its flagella
CCW- flagella rotate together
CW- flagella independent

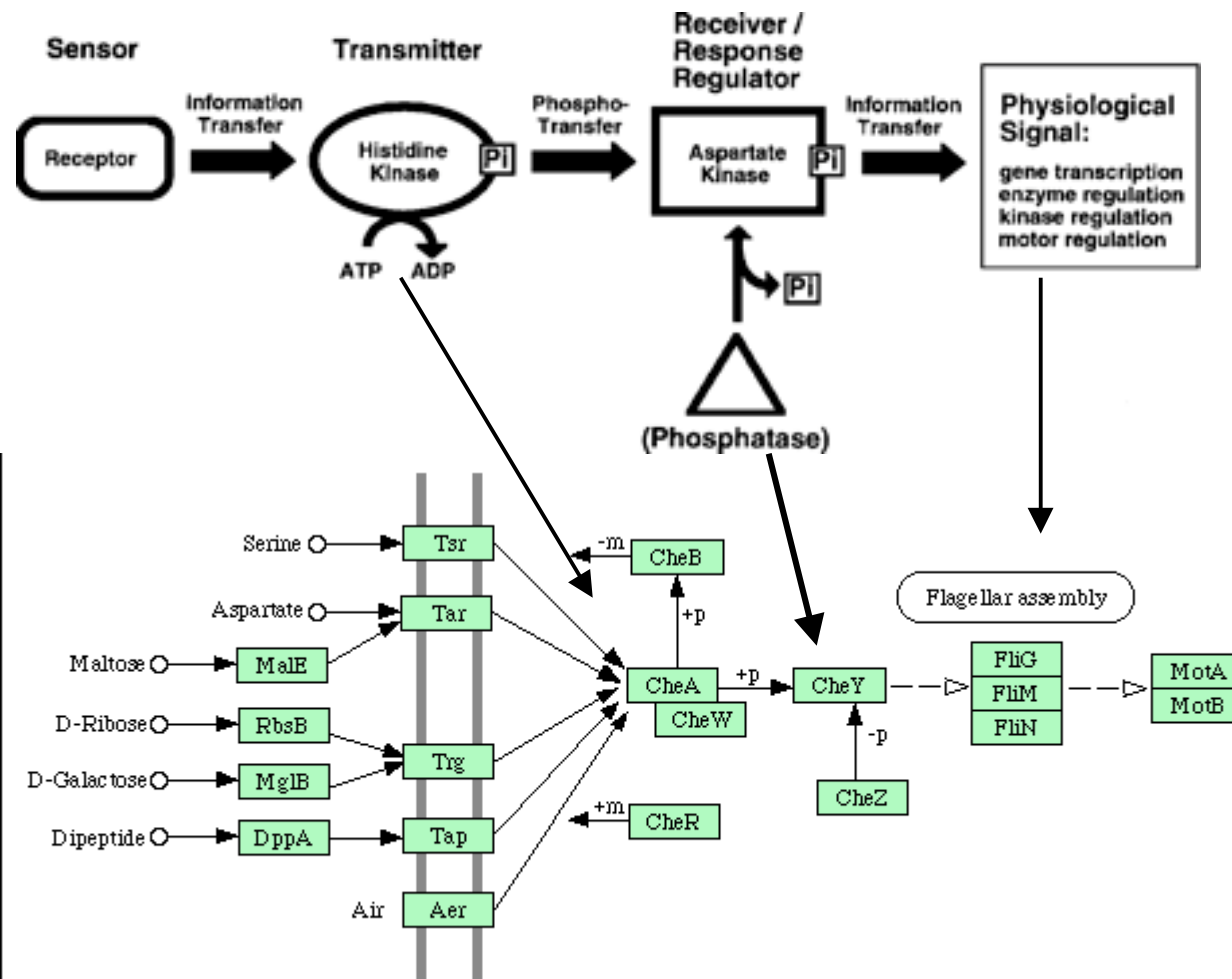


In homogenous environments the motion is a random combination of runs and tumbles

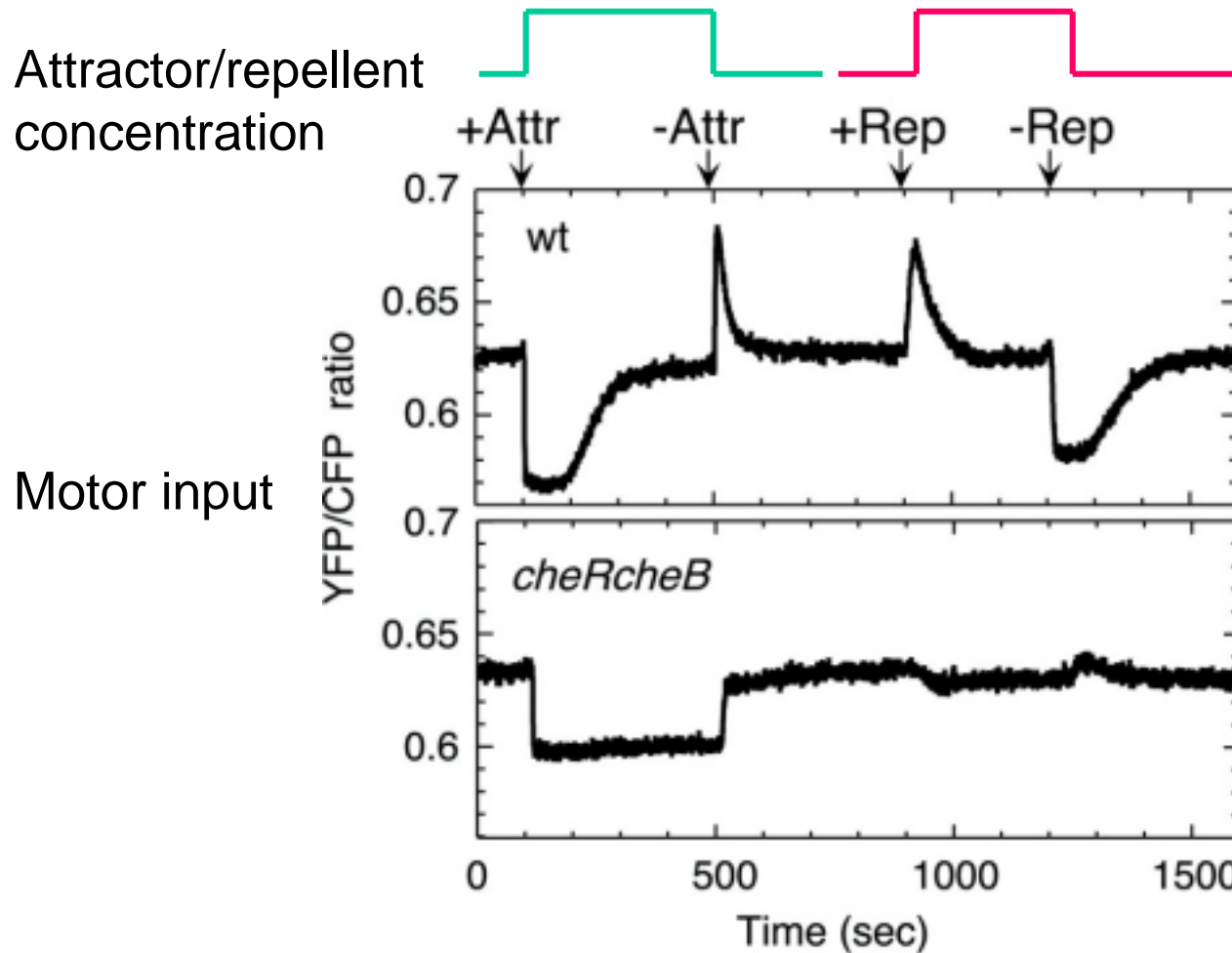
Bacteria change the direction of their motion in response to chemical signals



The signal transduction network is an example of a two-component pathway

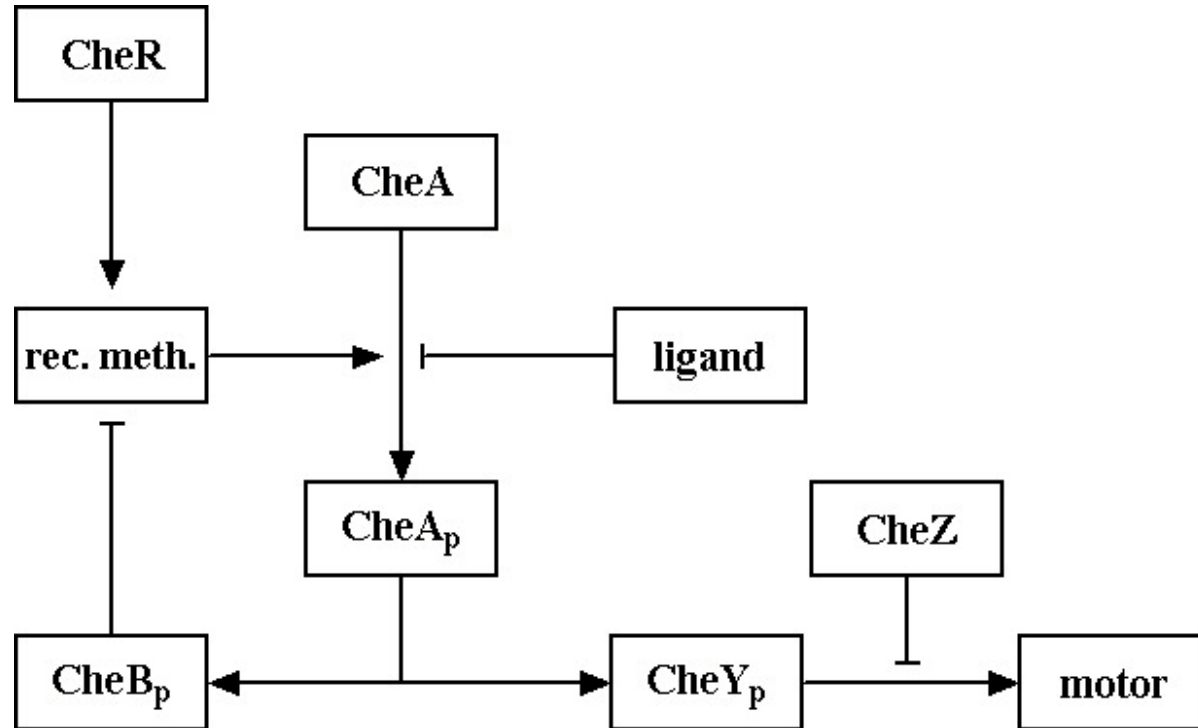


Bacteria respond to concentration changes but adapt to a constant stimulus



Adaptation is based on changes in the methylation levels.

Negative feedback as the source of adaptation



The steady state concentrations are determined by the equilibrium between activation and inhibition.

Modeling the signal transduction network

Idea: The cell contains a large set of receptors and other proteins

Input: ligand concentration

Variables: concentrations of

- the different (ligand bound, methylated) states of the receptor complex
- CheR, CheB, CheB_p, CheY, CheZ

Output: the concentration of CheY_p

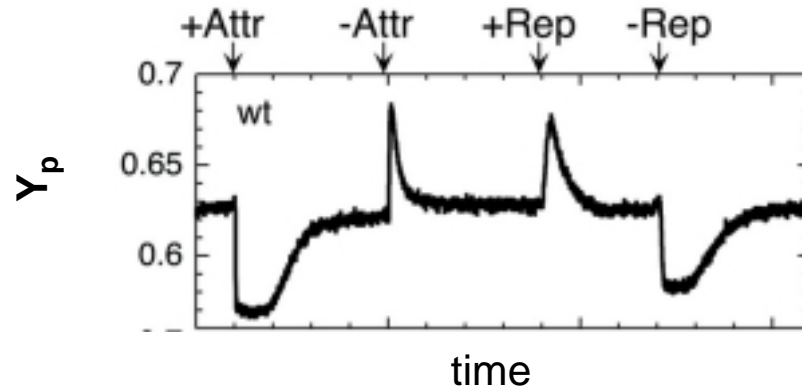
Method: differential equations describing the reaction kinetics.

Three types of reaction: ligand binding, phosphorylation, methylation.

N. Barkai and S. Leibler, Nature 387, 913 (1997)

P. A. Spiro, J. S. Parkinson, H. G. Othmer, PNAS 94, 7263 (1997)

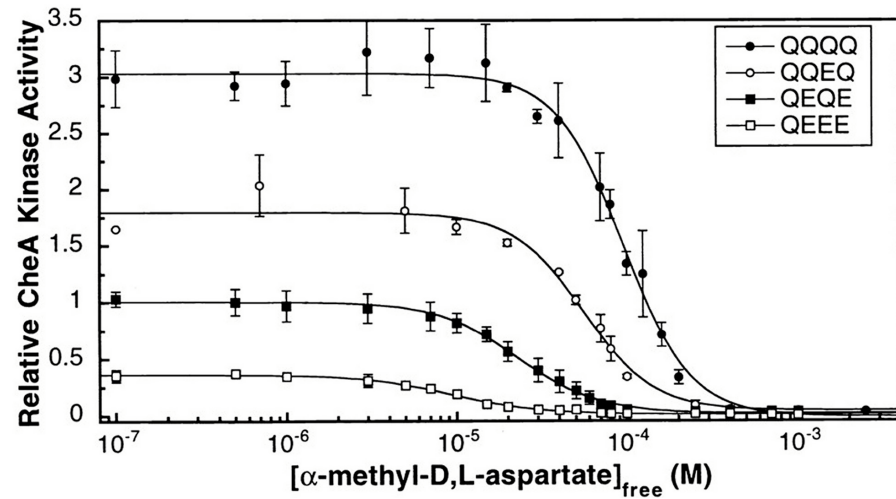
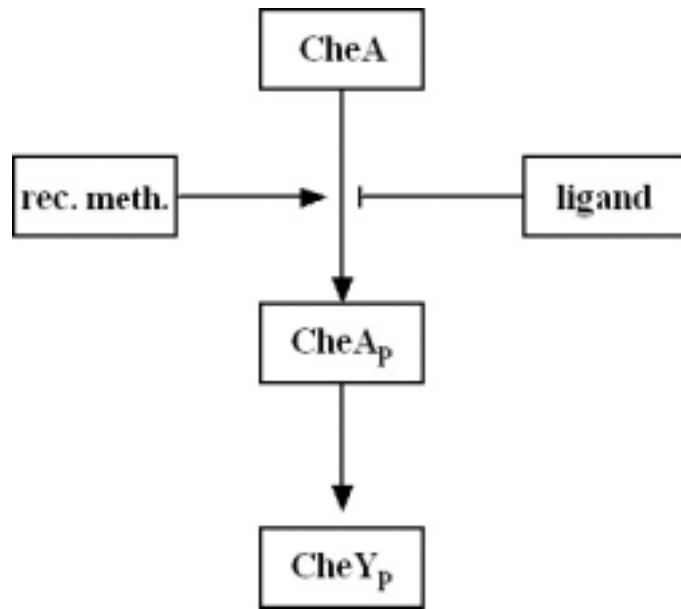
The immediate response of the system to ligand addition



- The amplitude of the initial peaks depends on the added ligand, and also on the methylation state of the receptors.
- Idea: keep the receptors in a fixed methylation state, and study the variation of the initial response with added ligand.

J. A. Bornhorst and J. J. Falke, *Journ. Gen. Physiol.* 118, 693 (2001)

Scaling with ligand concentration



- The data can be fitted with Hill functions

$$\frac{Y_p(L)}{Y_p(0)} = 1 - \frac{L^H}{K_A^H + L^H}, \quad H \in [1,2)$$

- The initial amplitudes and K_A values depend on the methylation level.

Simplest assumption

Assume that in ligand-bound receptor complexes CheA cannot autophosphorylate.

Then $Y_p(L)$ is proportional with R

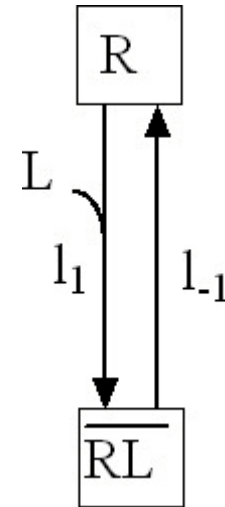
Mass-action kinetics:
$$\frac{dR}{dt} = -l_1 L R + l_{-1} \overline{RL}$$

$$R_t = R + \overline{RL}$$

Result:
$$\frac{R}{R_t} = 1 - \frac{L}{K_D + L}, \quad K_D = \frac{l_{-1}}{l_1}$$

Experiments:
$$\frac{Y_p(L)}{Y_p(0)} = 1 - \frac{L^H}{K_A^H + L^H}, \quad H \in [1, 2)$$

methylation-dependent $H, K_A, \quad K_A \neq \frac{l_{-1}}{l_1}$



Gain in signal transduction

Input: ligand concentration

Transduction: CheA autophosphorylation

Output: CheY_p

Intuitive assumption: output depends linearly on the input

$$\text{gain: } g = \frac{\text{relative change in output}}{\text{relative change in input}} = \frac{\frac{dO}{O}}{\frac{dI}{I}}$$

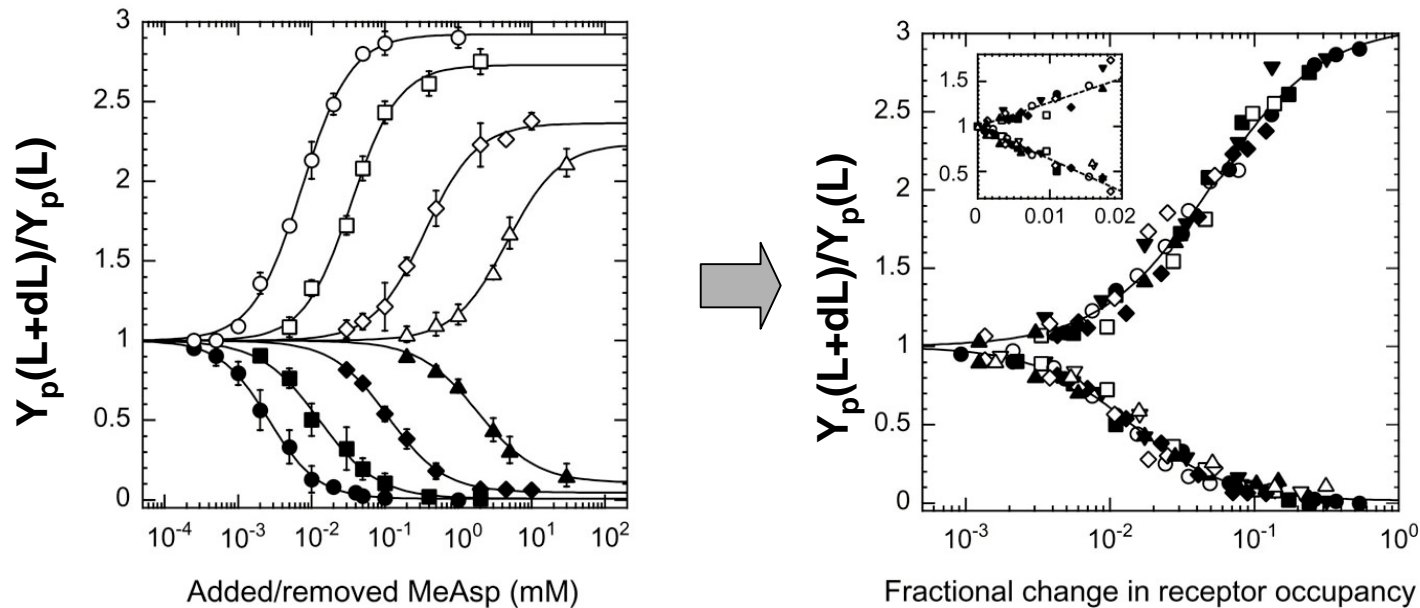
If the output is proportional with the input, $g = \text{constant}$.

A high gain indicates an amplification in signal transduction.

Implicit assumption: there is a constant gain between

$$\text{ligand occupancy, } o = \frac{L}{K_D + L} \quad \text{and CheY}_p$$

Response to additional ligand scales with fractional change in receptor occupancy



The initial slope of the curve is ~ 35 , indicating a surprisingly high amplification for the signal transduction network.

V. Sourjik and H. Berg, PNAS 99, 123 (2002)

General explanation of the high amplification

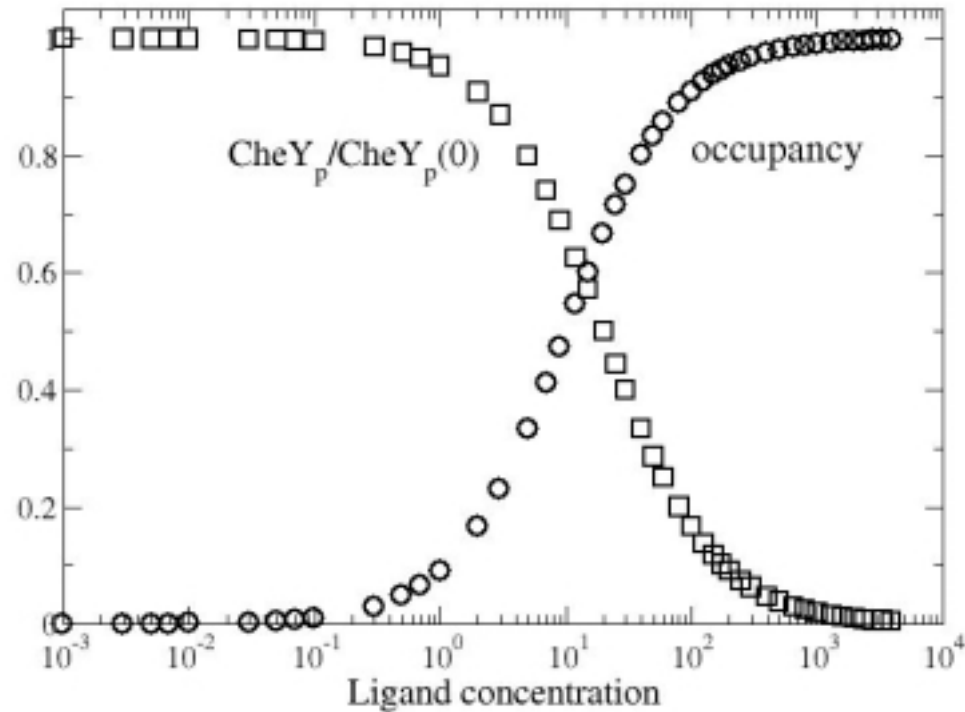
Assume that the functional form of the dependence of $CheY_p$ on the ligand concentration is $\frac{Y_p(L)}{Y_p(0)} = 1 - \frac{L^H}{K_A^H + L^H}$

The receptor occupancy is defined as $o(L) = \frac{L}{K_D + L}$

Then the signal amplification is $\frac{d \ln Y_p}{d \ln o} = -\frac{H}{K_D} \frac{L^H (K_D + L)}{K_A^H + L^H}$

The amplification is a continuously increasing function of the ligand concentration, so large amplifications are possible at high ligand levels.

CheY_p decreases with increasing occupancy



In the high occupancy region the relative change in occupancy is small, but since CheY_p is small, its relative change is significant.