

Short Course: Computation of Olfaction

Lecture 5

Lecture 5: Heteroclinic vs. HH the pheromone sub-system

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University of Sussex

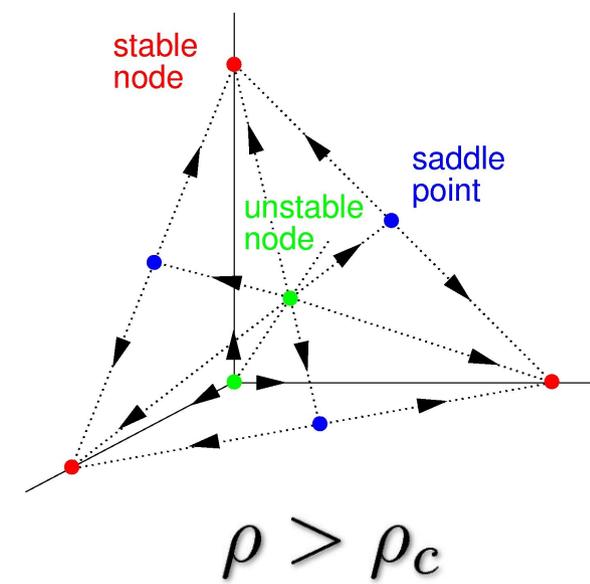
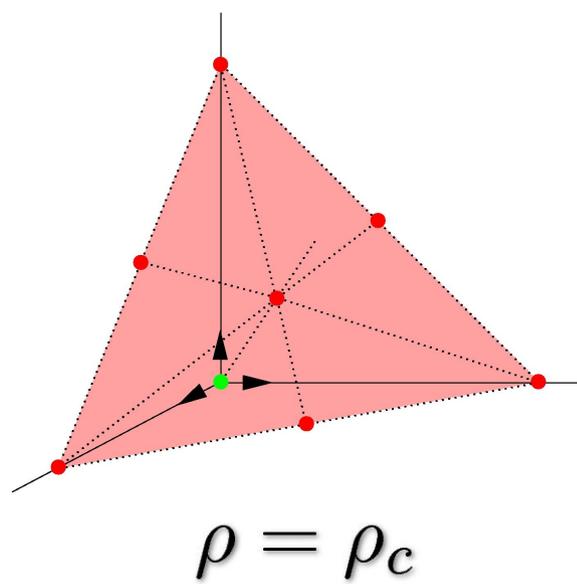
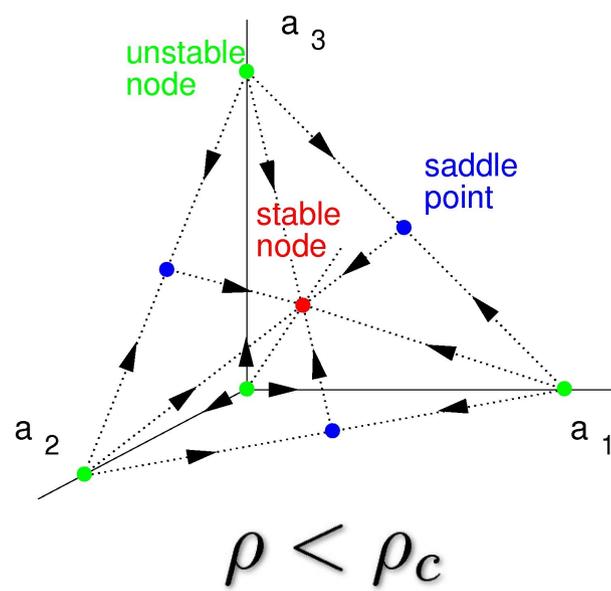
Yesterday ...

- Heteroclinic orbits exist in the generalized Lotka-Volterra model with asymmetric synaptic connections
- Under certain conditions the heteroclinic can be a global attractor
- Now ... does something like this exist for more realistic Hodgkin-Huxley neurons?

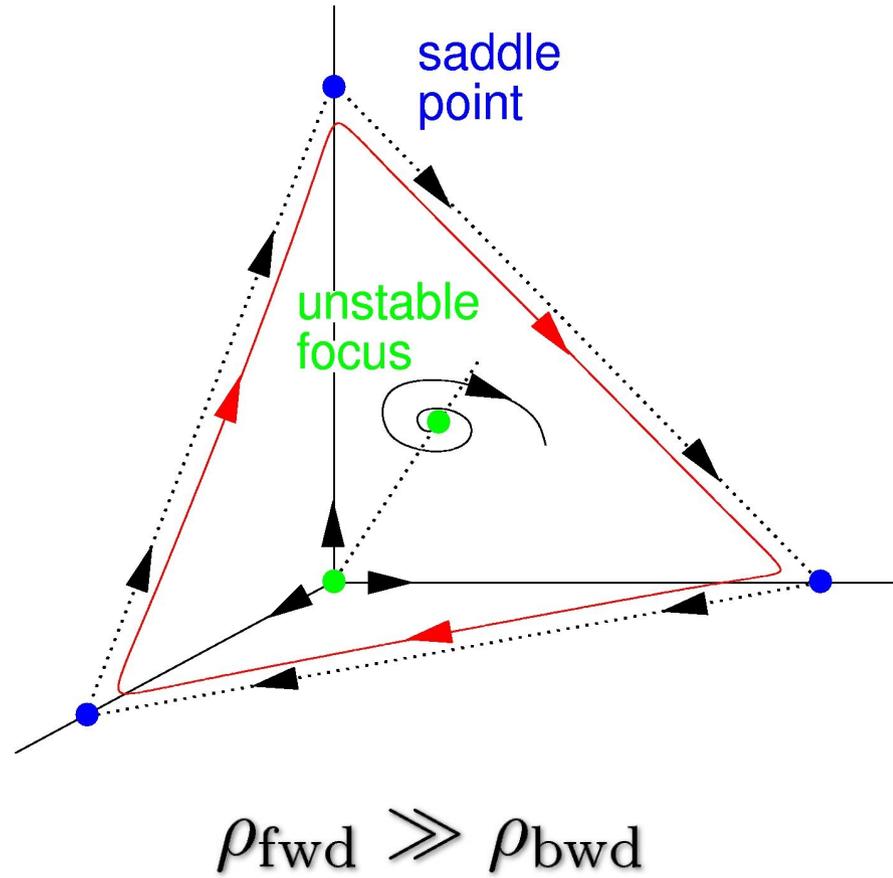
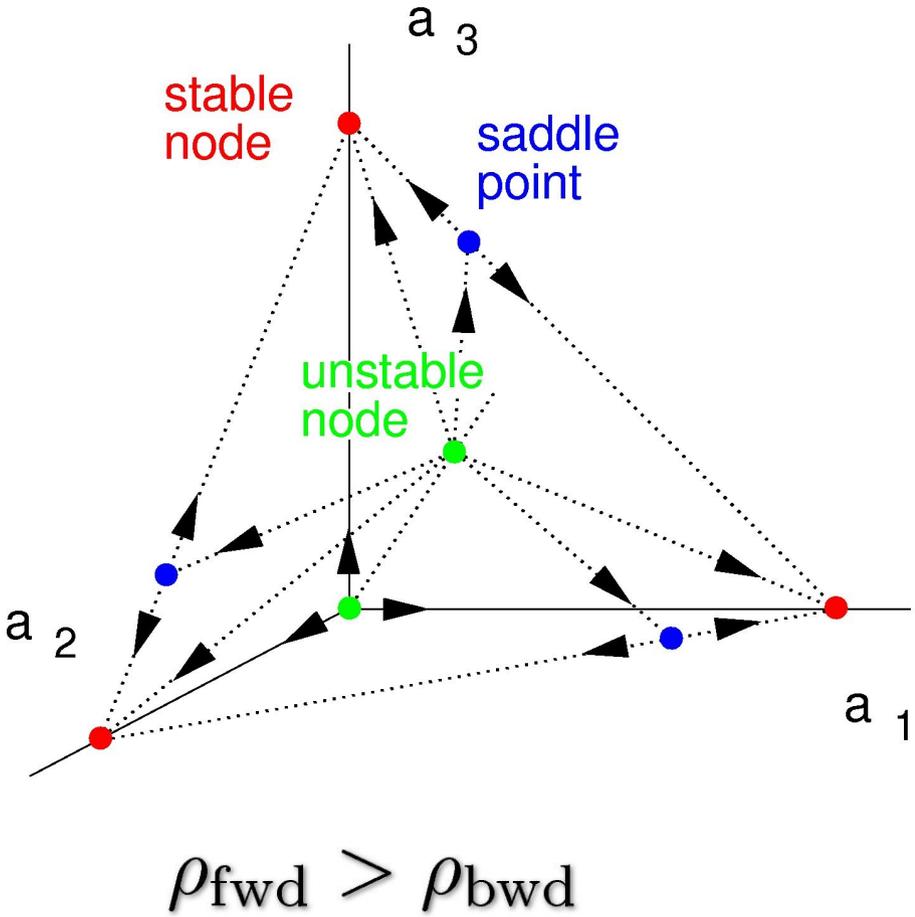
Reminder: Lotka-Volterra Model

$$\frac{a_i(t)}{dt} = a_i \left[1 - \left(a_i + \sum_{i \neq j}^N \rho_{ij} a_j \right) \right]$$

Symmetric connections: $\rho_{ij} = \rho$



Reminder: Asymmetric connections



Heteroclinic for HH neurons?

- Problem: For HH neurons, saddle points are “saddle limit cycles” and analytically and numerically intractable
- Approach: Find an “equivalent rate model” that has heteroclinic structure:

(The following is published in:

Nowotny and Rabinovich, *Dynamical Origin of Independent Spiking and Bursting Activity in Neural Microcircuits*, Phys Rev Lett **98**, 128106 (2007))

Individual neurons

$$C \frac{dV(t)}{dt} = -I_{\text{Na}} - I_{\text{K}} - I_{\text{leak}} + I_{\text{syn}}$$

$$I_{\text{Na}}(t) = g_{\text{Na}} m(t)^3 h(t) (V(t) - E_{\text{Na}})$$

$$I_{\text{K}}(t) = g_{\text{K}} n(t)^4 (V(t) - E_{\text{K}})$$

$$I_{\text{leak}}(t) = g_{\text{leak}} (V(t) - E_{\text{leak}})$$

Traub and Miles, Neural Networks of the Hippocampus,
Cambridge University Press, 1991

Individual neurons: Activation and Inactivation

$$\frac{dy(t)}{dt} = \alpha_y(V(t))(1 - y(t)) - \beta_y(V(t))y(t)$$

$$\alpha_m = 0.32(-52 - V)/(\exp((-52 - V)/4) - 1)$$

$$\beta_m = 0.28(25 + V)/(\exp((25 + V)/5) - 1)$$

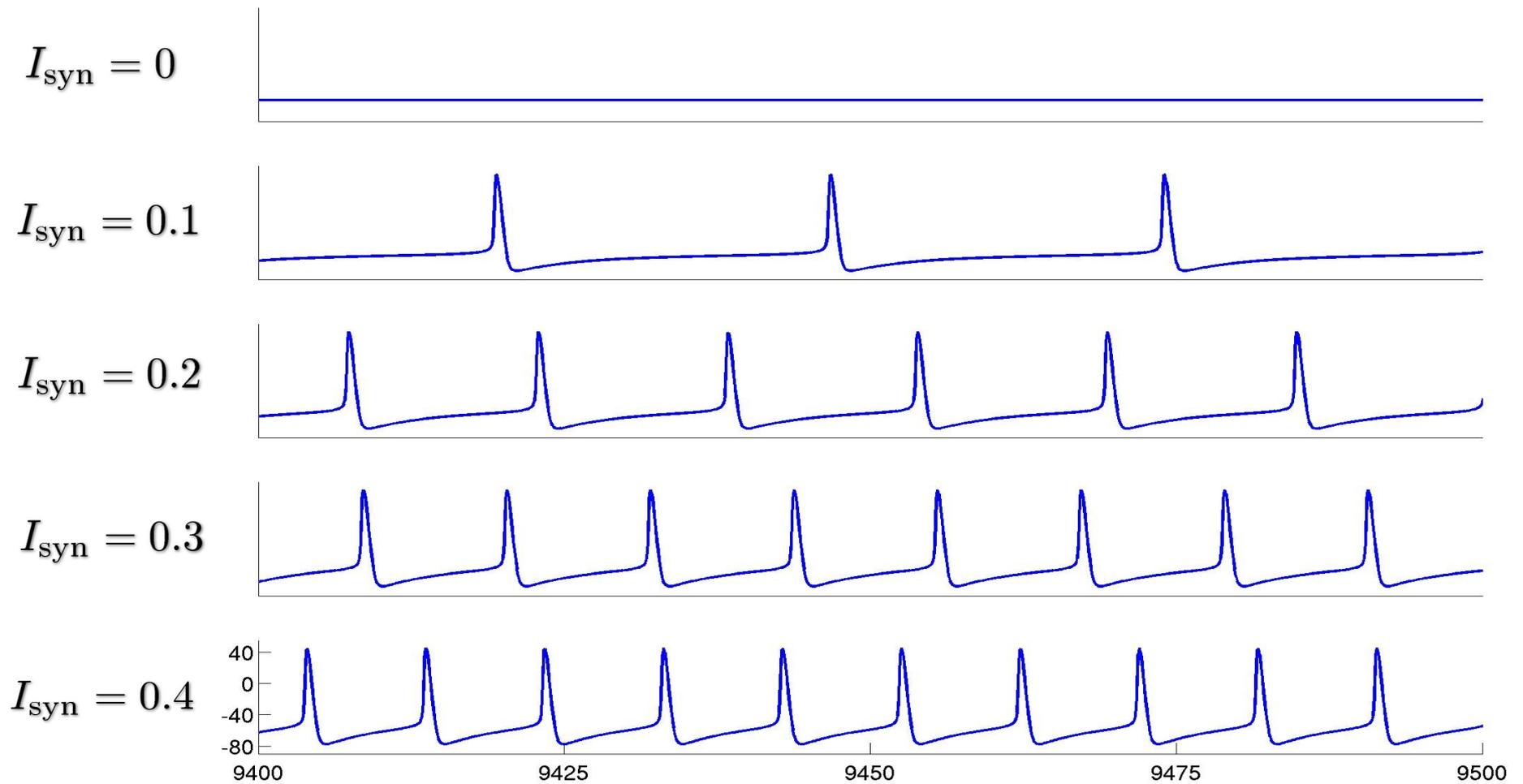
$$\alpha_h = 0.128 \exp((-48 - V)/18)$$

$$\beta_h = 4/(\exp((-25 - V)/5) + 1).$$

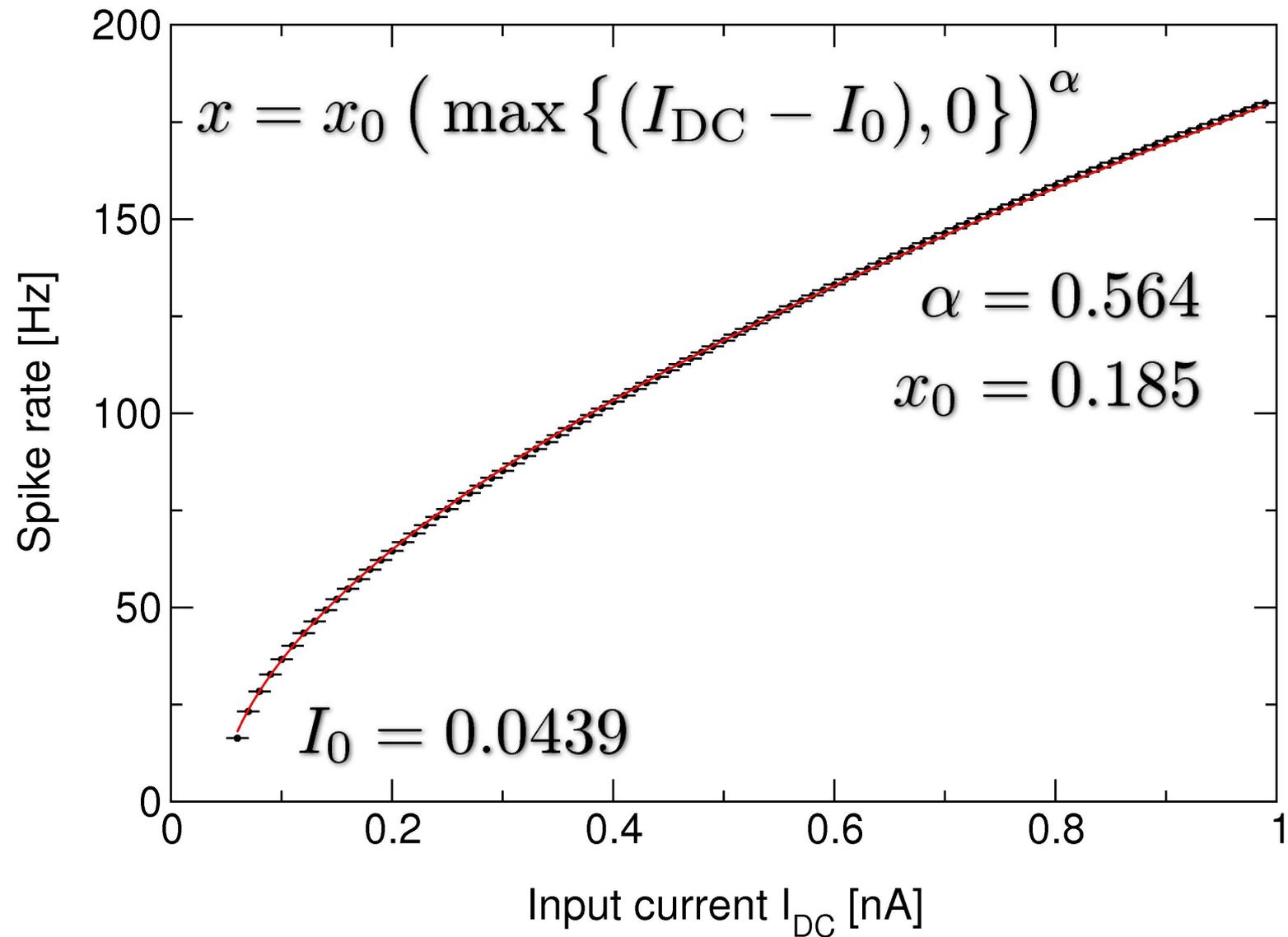
$$\alpha_n = 0.032(-50 - V)/(\exp((-50 - V)/5) - 1)$$

$$\beta_n = 0.5 \exp((-55 - V)/40)$$

Example waveforms



Spike rate



Synapses

$$I_{\text{syn},ji} = g_{ji} S_i (V_j - V_{\text{rev}})$$

$$\tau \frac{dS_i}{dt} = (R_i - \kappa S_i) \frac{S_{\text{max}} - S_i}{S_{\text{max}}} \quad \tau = 50 \text{ ms}$$

$$\tau \frac{dR_i}{dt} = \Theta(V_i - V_{\text{th}}) - R_i,$$

Modified from: W. Rall, J Neurophysiol
30:1138 (1967).

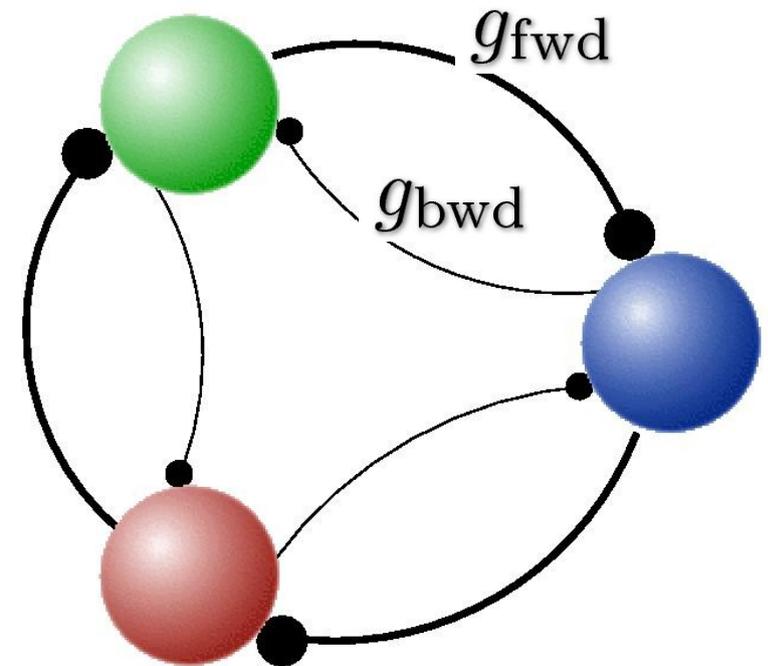
Summary

Neurons: $V_i(t), m_i(t), h_i(t), n_i(t) \quad i = 1 \dots 3$

Synapses: $R_i(t), S_i(t) \quad i = 1 \dots 3$

Total: 18 dimensional system

Main parameters: $g_{\text{fwd}} \quad g_{\text{bwd}}$



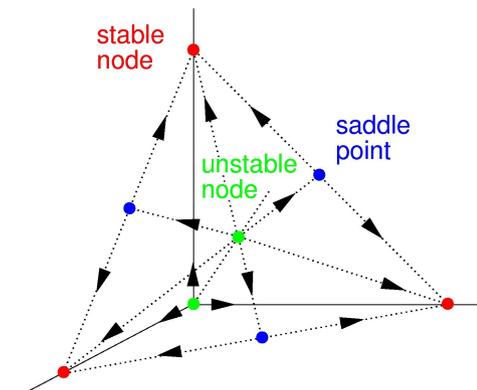
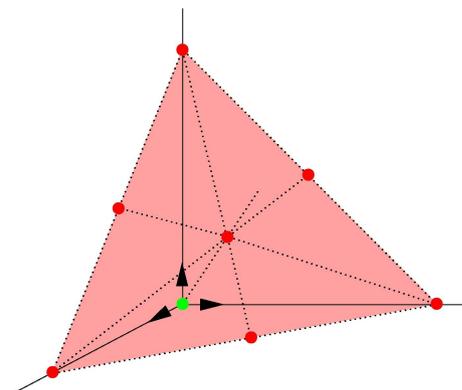
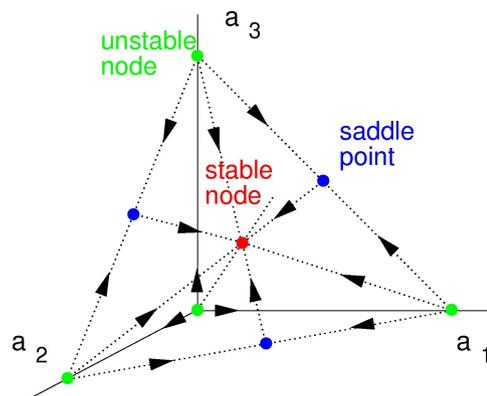
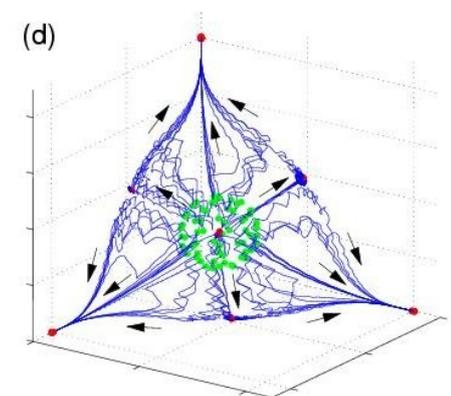
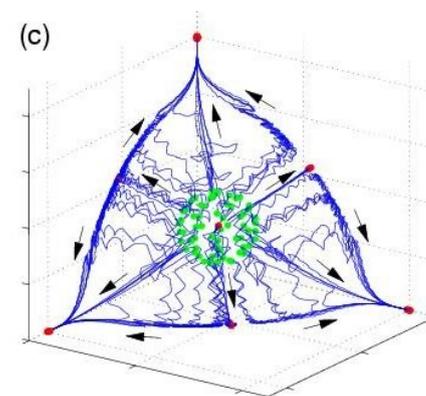
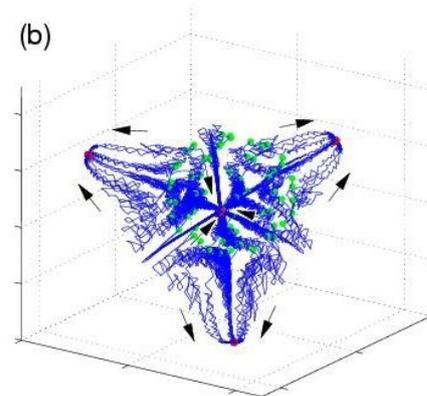
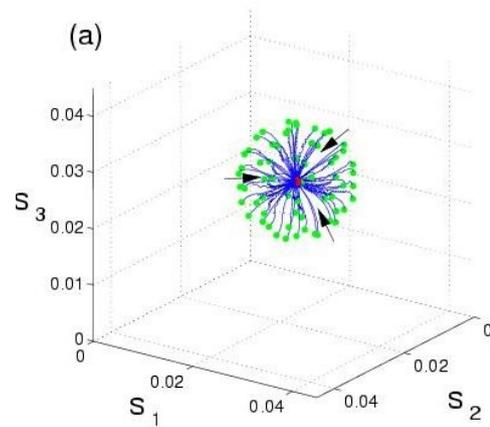
Symmetric inhibition

$g = 10 \text{ nS}$

$g = 30 \text{ nS}$

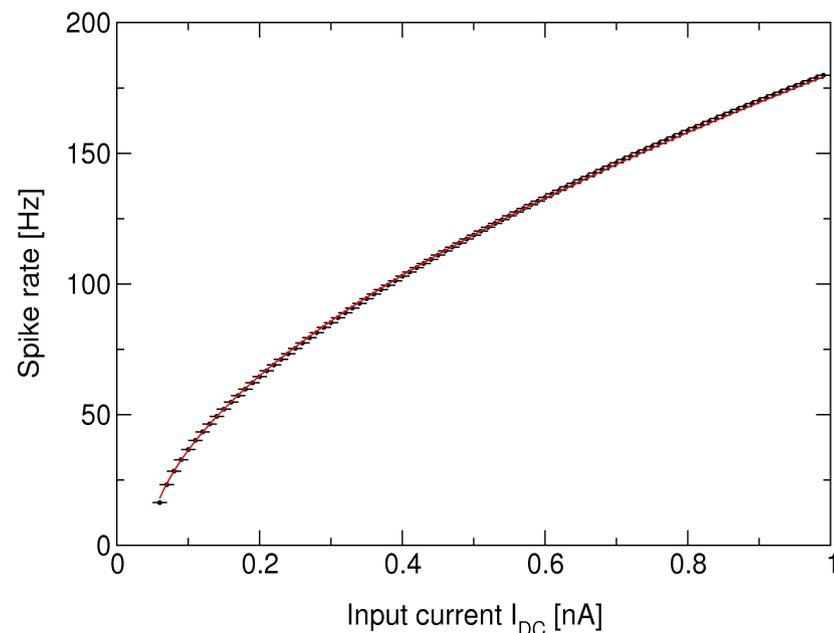
$g = 50 \text{ nS}$

$g = 60 \text{ nS}$



Lotka-Volterra model

Reduction to a rate model



$$x = x_0 \left(\max \{ (I_{DC} - I_0), 0 \} \right)^\alpha$$

$$\text{Ansatz: } \tau \frac{dr_i}{dt} \approx a(x_i) - r_i$$

Then require that for tonic spiking the spiking model and rate model match, leading to

$$a(x_i) = \frac{1 - \exp(-\tau_{\text{spike}}/\tau)}{1 - \exp(-1/(x_i\tau))} \approx c x_i \tau$$

Equivalent rate model

Pre-synaptic
activation

Post-synaptic
activation

$$\tau \frac{ds_i}{dt} = (r_i - \kappa s_i) \frac{S_{\max} - s_i}{S_{\max}}$$
$$\tau \frac{dr_i}{dt} = \tilde{x}_0 \left(\tilde{I} - \sum_j \tilde{g}_{ij} s_j \right)_+^\alpha \tau - r_i$$

The variables in this “rate model” are the synapse activation variables!

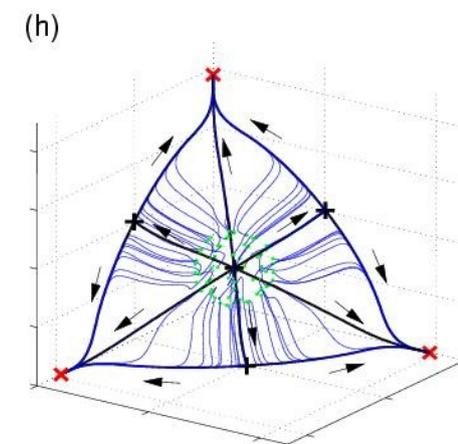
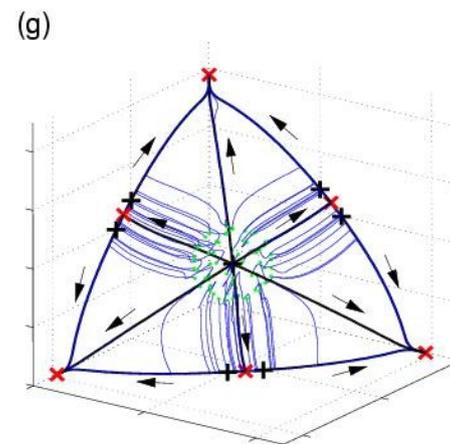
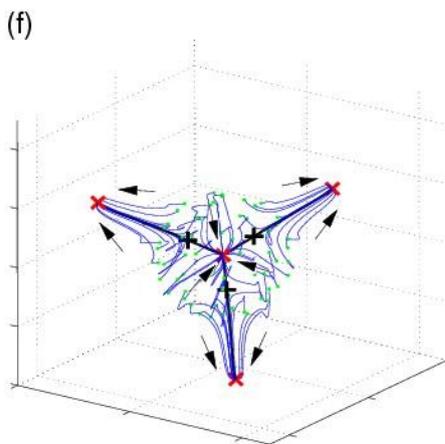
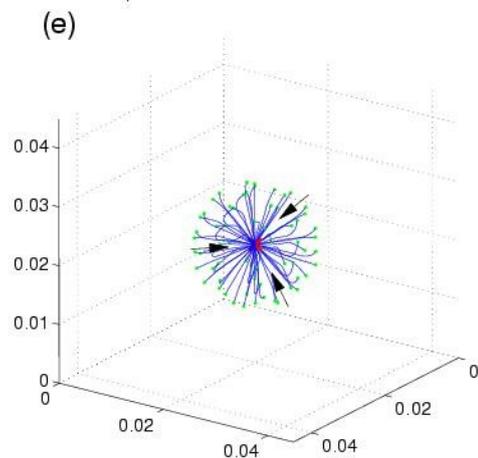
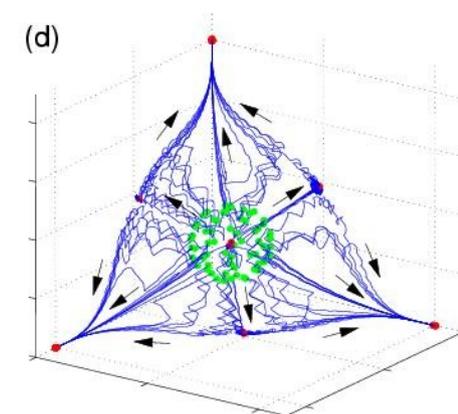
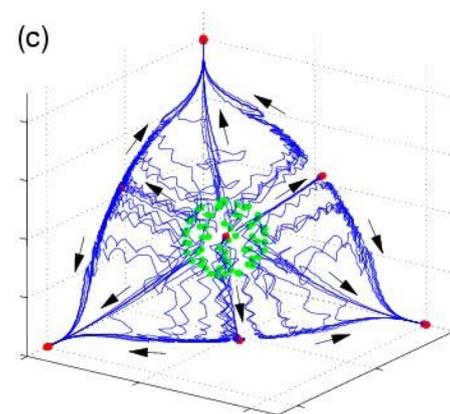
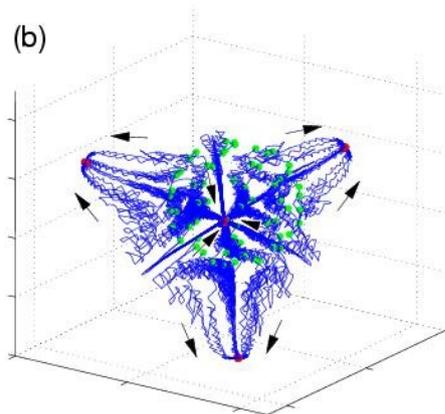
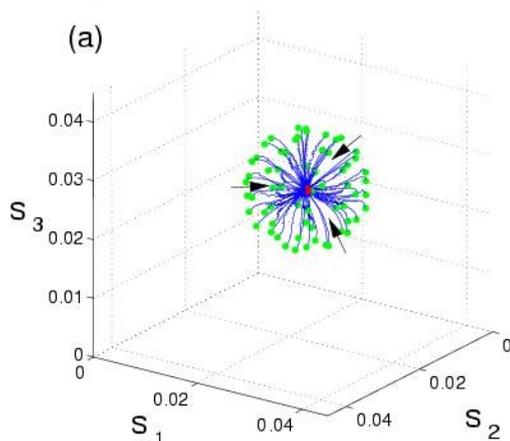
Comparison

$g = 10 \text{ nS}$

$g = 30 \text{ nS}$

$g = 50 \text{ nS}$

$g = 60 \text{ nS}$



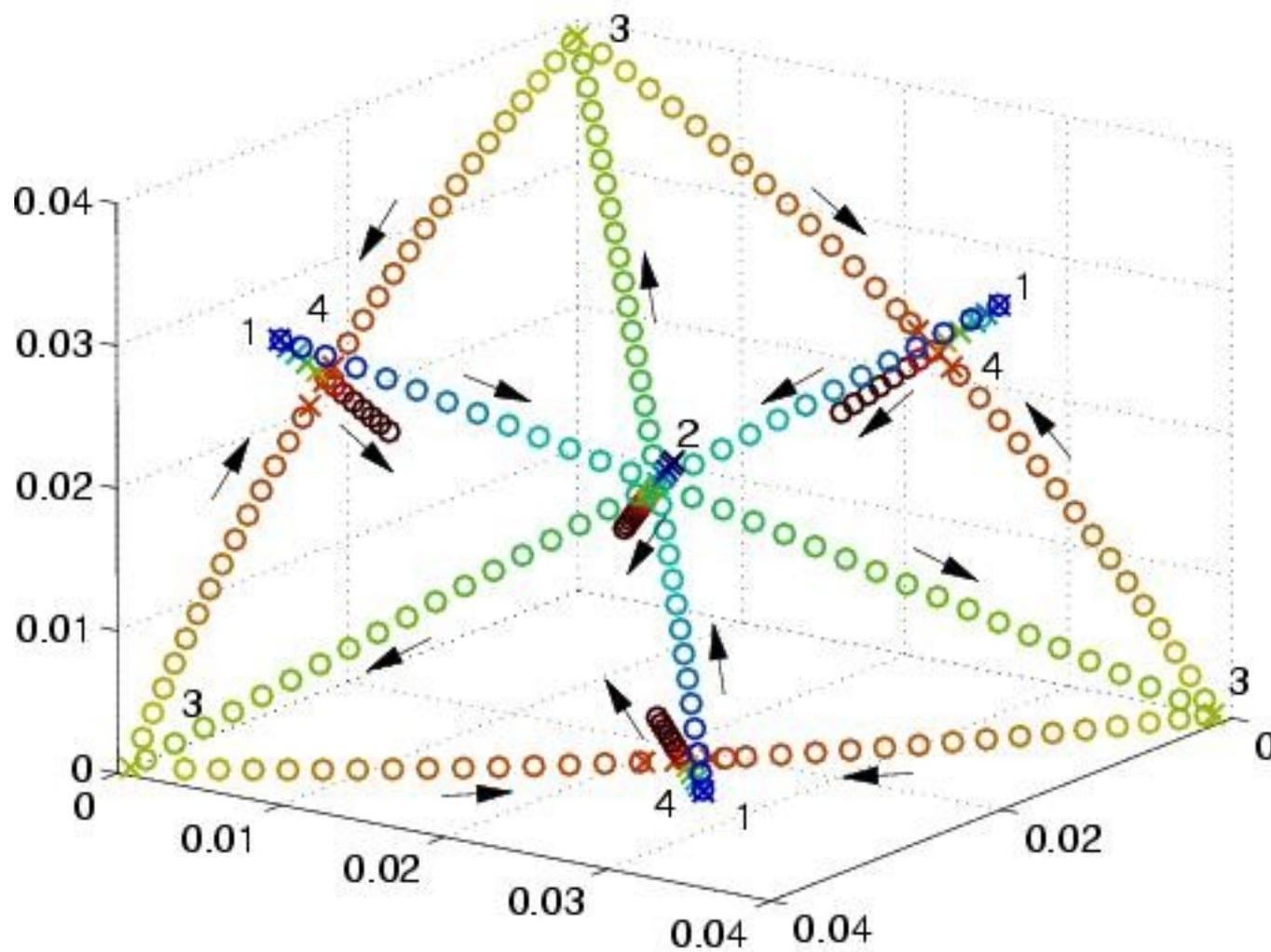
$g = 30 \text{ nS}$

$g = 40 \text{ nS}$

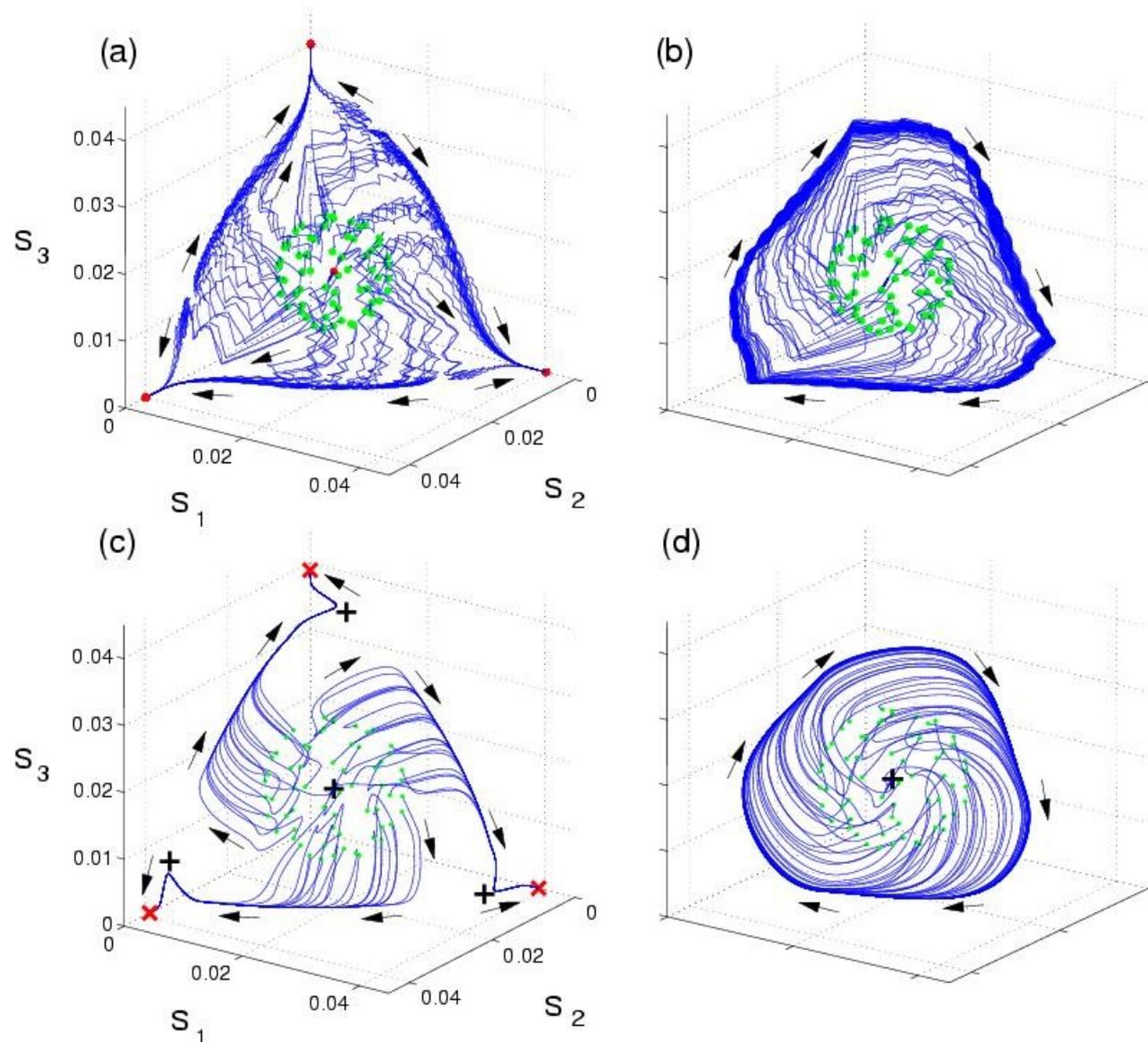
$g = 51.4 \text{ nS}$

$g = 60 \text{ nS}$

Numerical bifurcation analysis

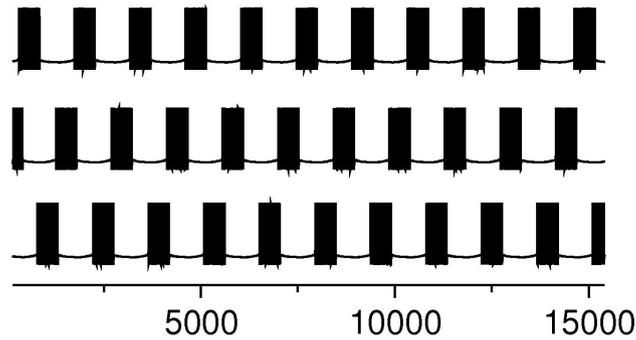


Asymmetric connections

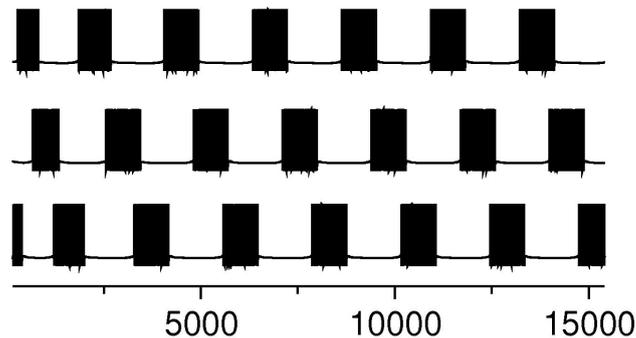


Stable heteroclinic orbit (?)

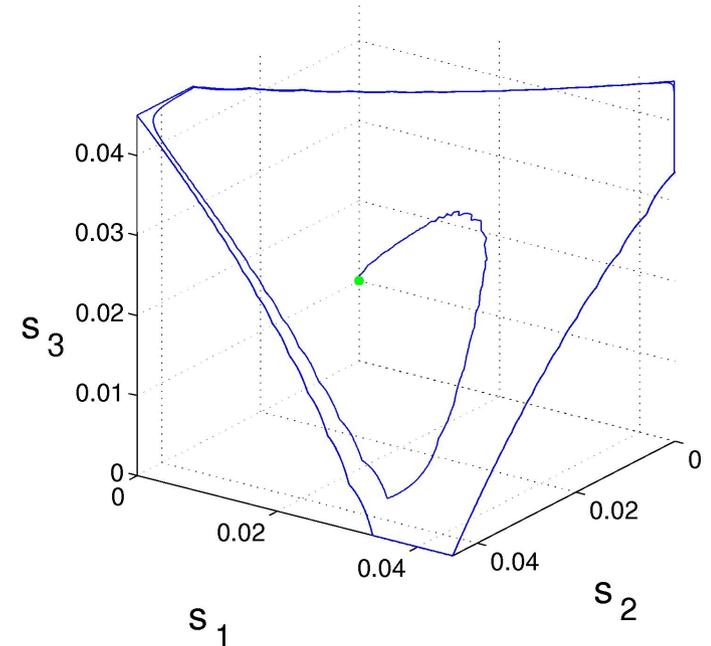
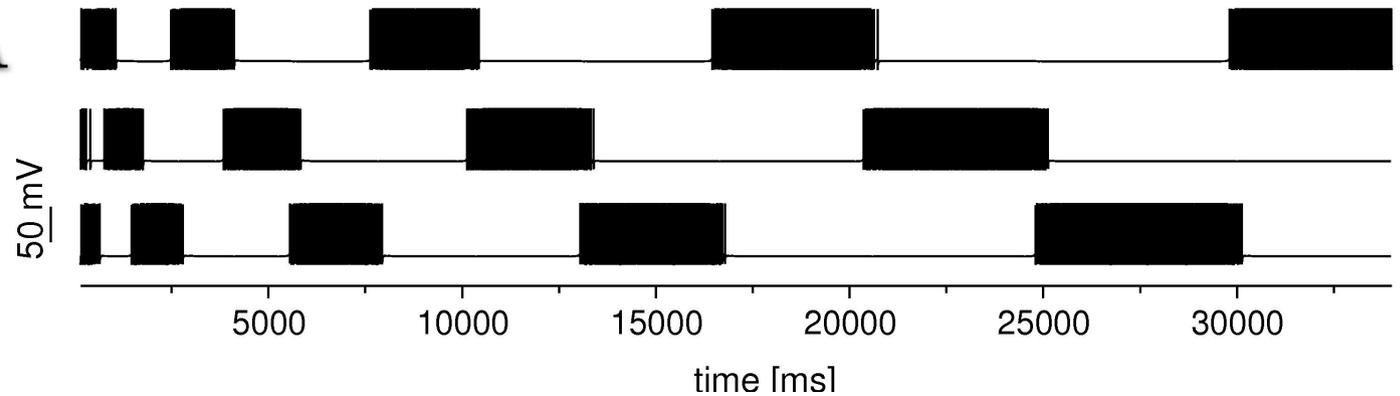
$I_{DC} = 0.08 \text{ nA}$



$I_{DC} = 0.16 \text{ nA}$



$I_{DC} = 0.22 \text{ nA}$



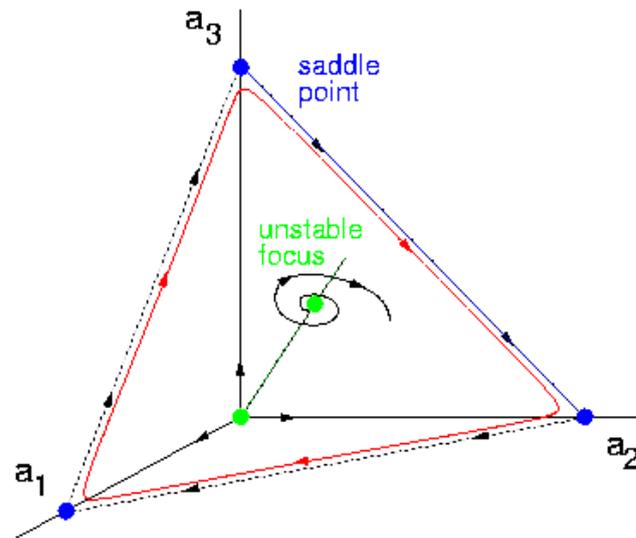
Conclusions: Heteroclinic in HH model

- The reduction from the spiking model (18D) to the rate model (6D) seems successful
- The bifurcation analysis in 6D shows the expected structures, similar to LVm
- The spiking model can be pushed into a regime where its trajectory seems to approach a heteroclinic.
- However, *the last point is contingent on the modification of the synapse model*

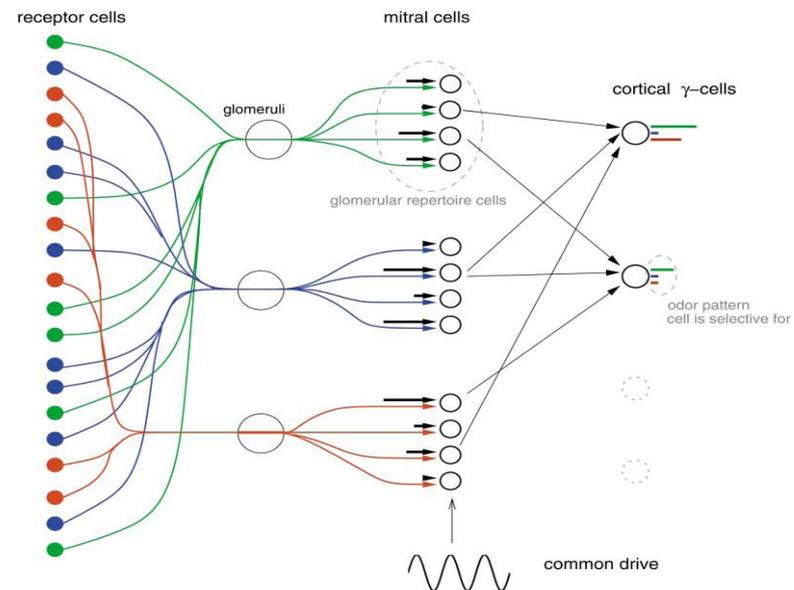
Pheromone Subsystem

Last time ... general olfactory system

Winnerless competition



Hopfield's model of olfaction



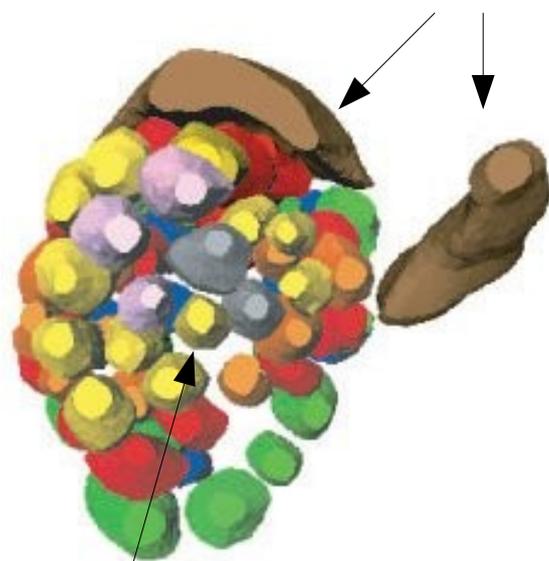
- This was all about the general olfactory system
- Today: Pheromone sub-system

Role of pheromones

- Pheromones are substances that animals secrete to communicate with each other
- Examples
 - Pheromone trails laid by ants
 - Pheromones in urine to mark territory
 - Sexual pheromones to communicate mating status
 - Sexual pheromones to attract mates
 - ...

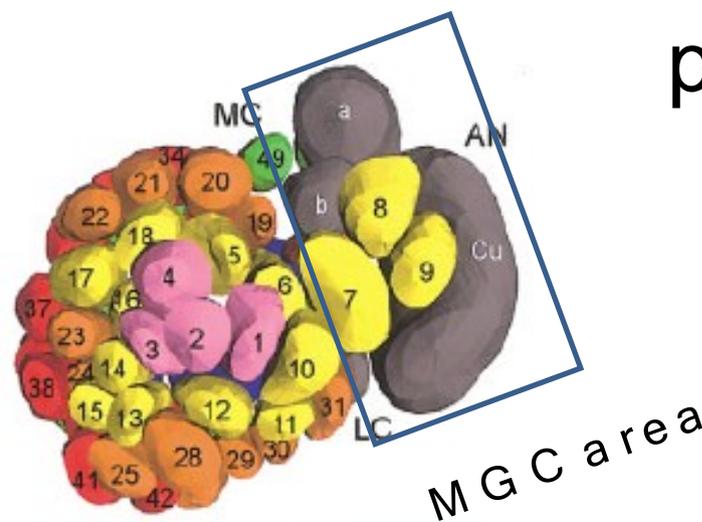
Pheromone sub-system: Anatomy

Pheromone sub-system:
Macro-glomerular Complex (MGC)



General antennal
lobe

In insects there are typically 2-3 large glomeruli exclusively dedicated for pheromone processing



MGC area

Pheromone sub-system in moths

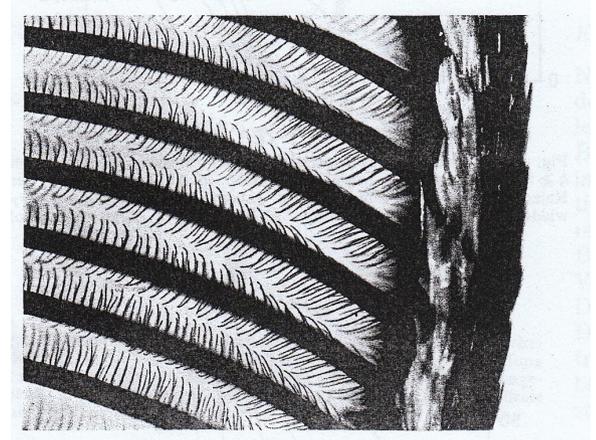
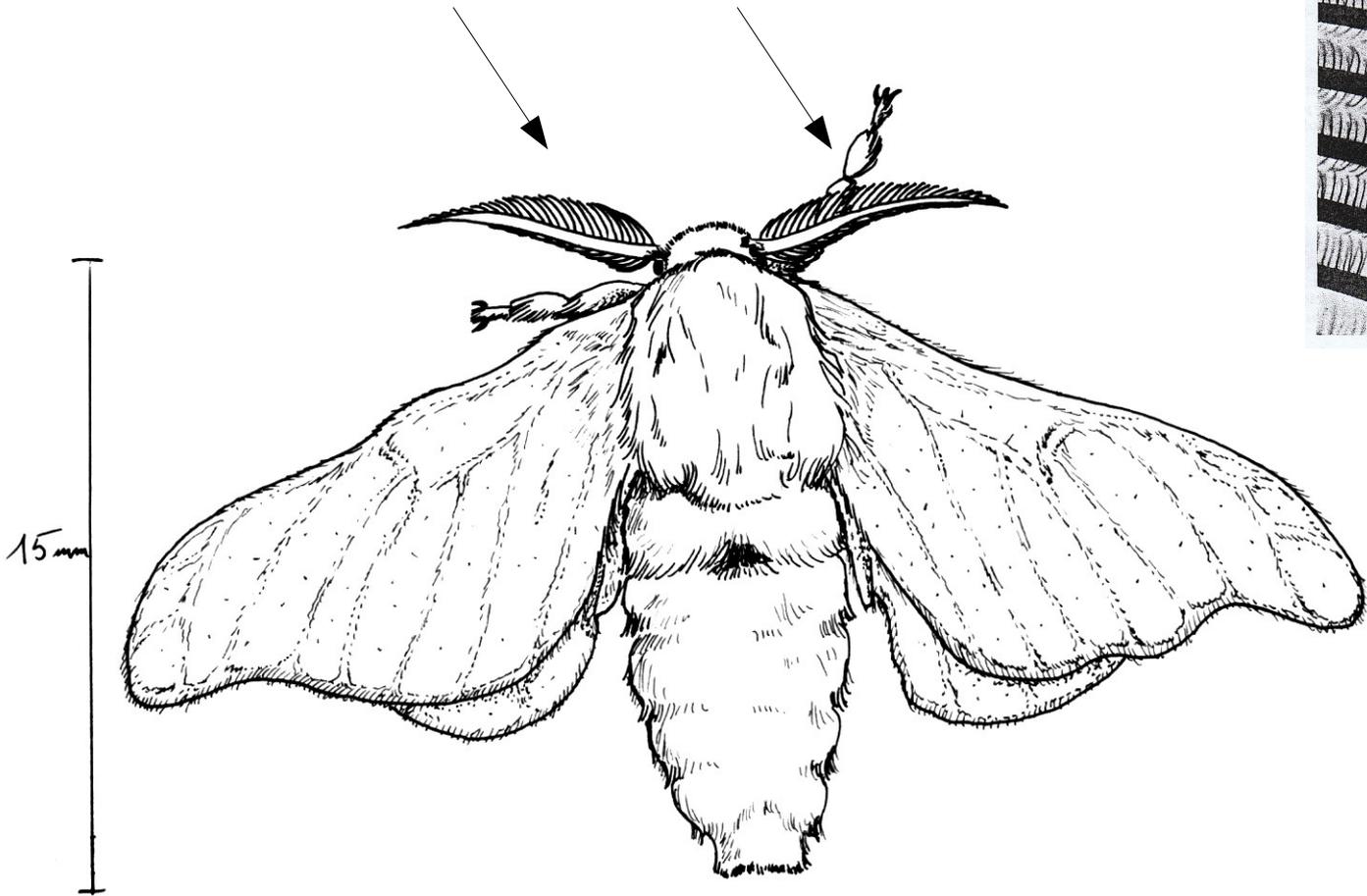
- Here I will concentrate on the pheromone system of moths
- Pheromone is secreted by females to attract males
- Males can smell the pheromone at distances of up to 2-3 miles
- Pheromone blend consists of several chemicals (components), in a typical ratio
- Related (but distinct) species may use the same chemicals but in a different ratio (!)

Pheromone subsystem is different

- General olfactory system
 - ORN broadly tuned
 - One ORN type – one glomerulus
 - Need to recognize pure odors, mixtures, concentrations
- Pheromone subsystem
 - ORN very narrowly tuned to 1 chemical
 - One ORN type – one glomerulus
 - Need to recognize one specific mixture of pheromone components

Bombyx Mori (Silk Moth)

Antennae

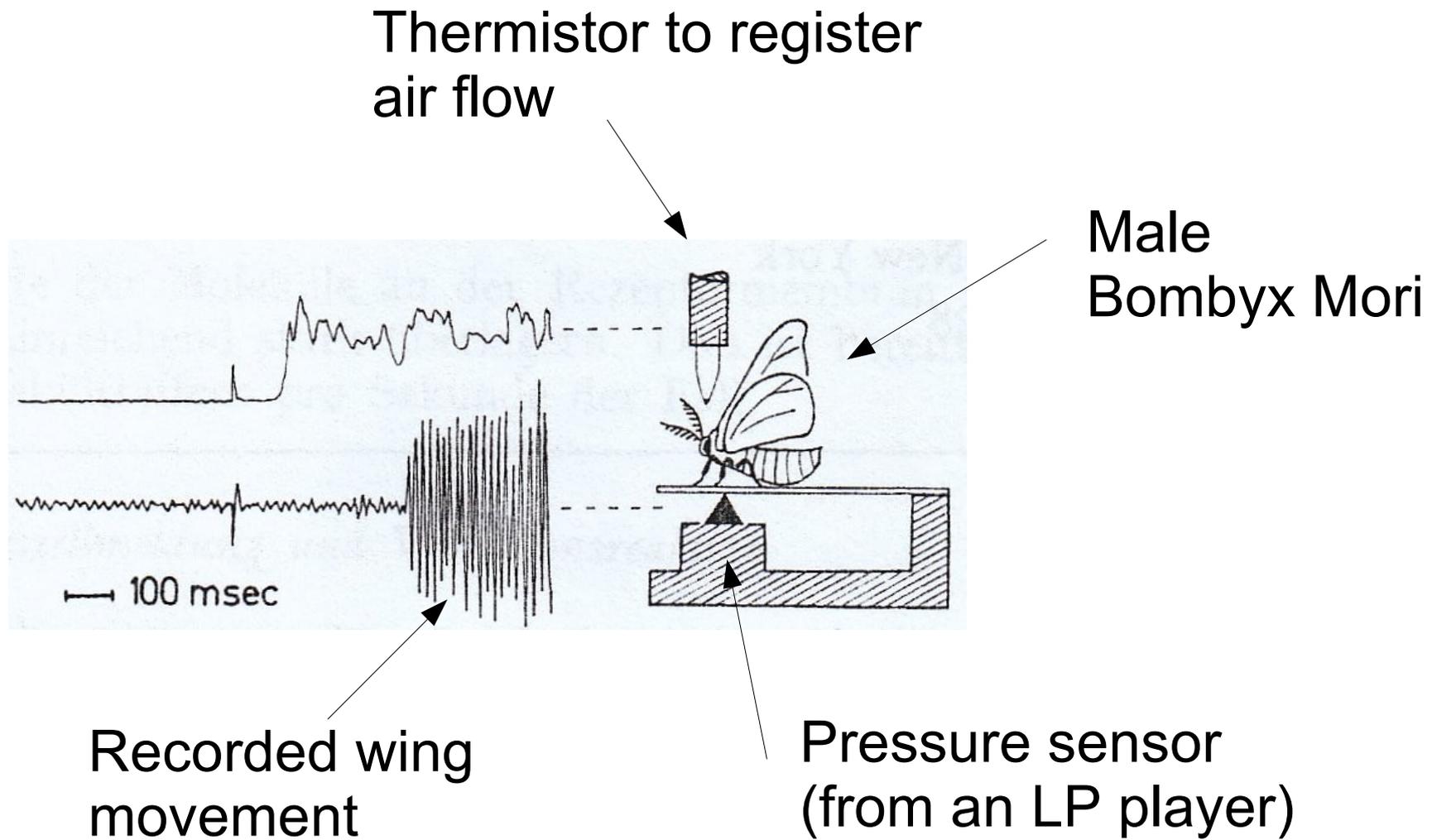


Classic results on Bombyx Mori

In the following we will have a look at classic results from

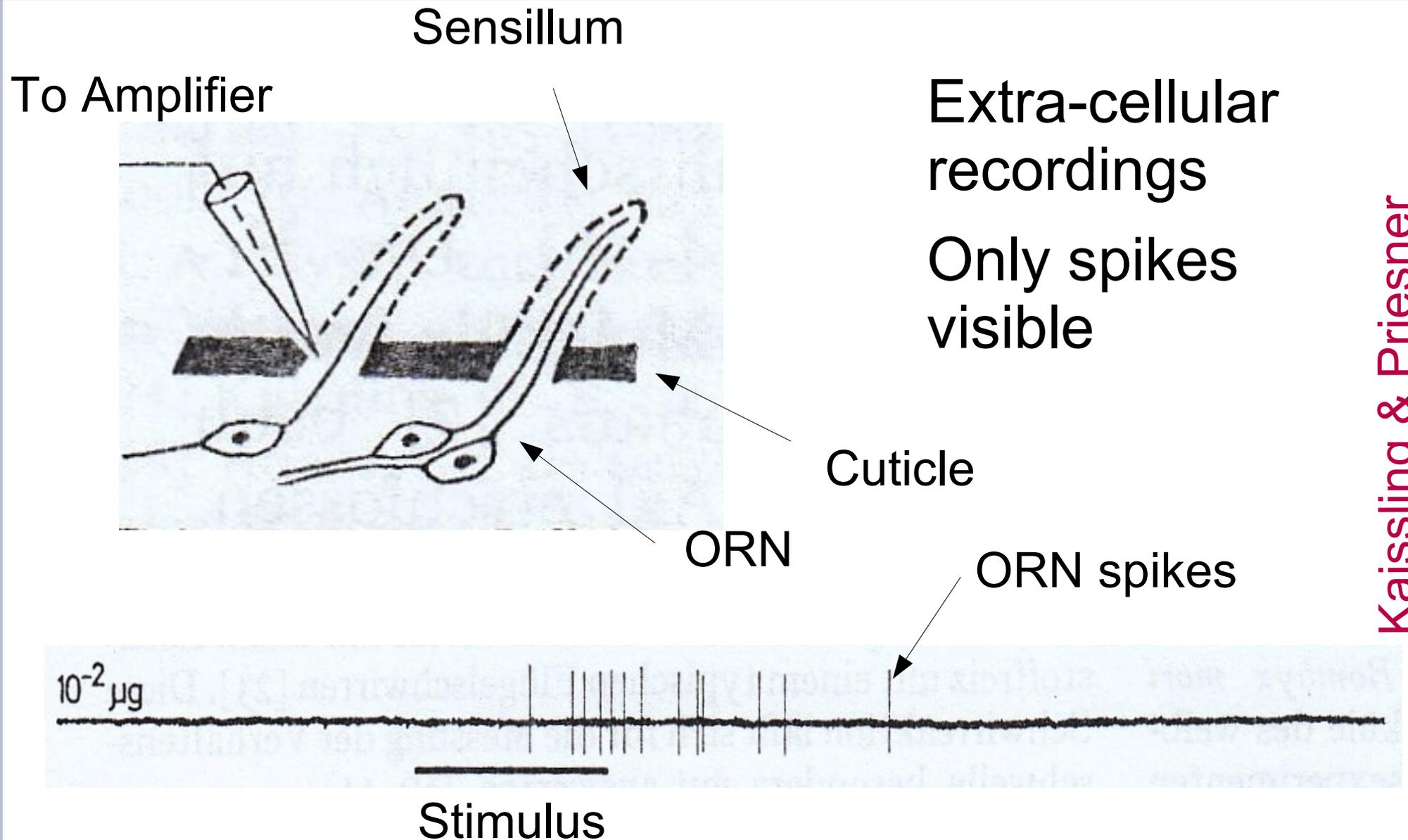
Kaissling K-E and Priesner E, “Die Riechschwelle des Seidenspinners”, Die Naturwissenschaften **57**(1): 23-28, 1970.

Type of experiments: Behavioral



Kaissling & Priesner

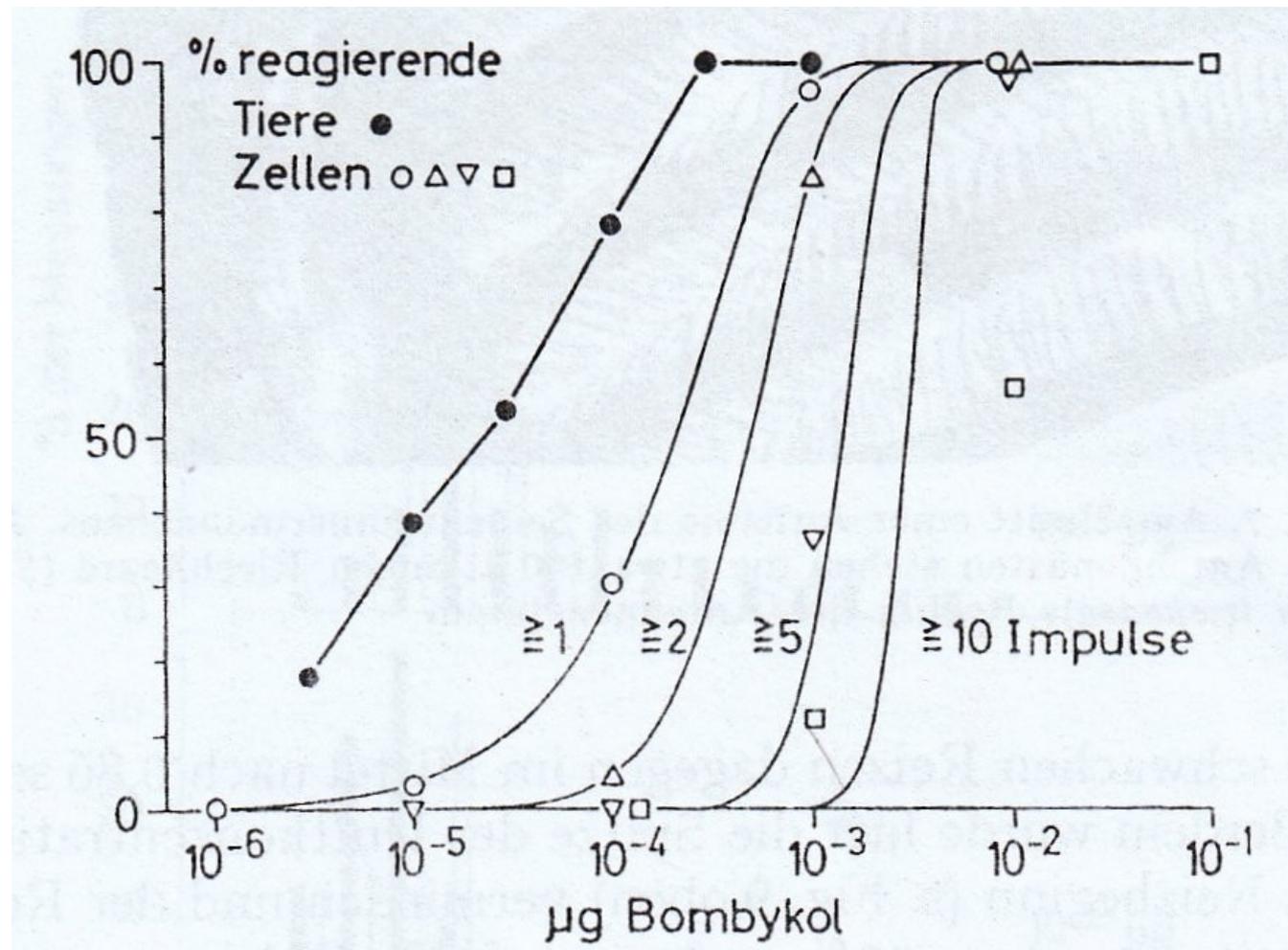
Type of experiments: Electrophysiology



Extra-cellular recordings
Only spikes visible

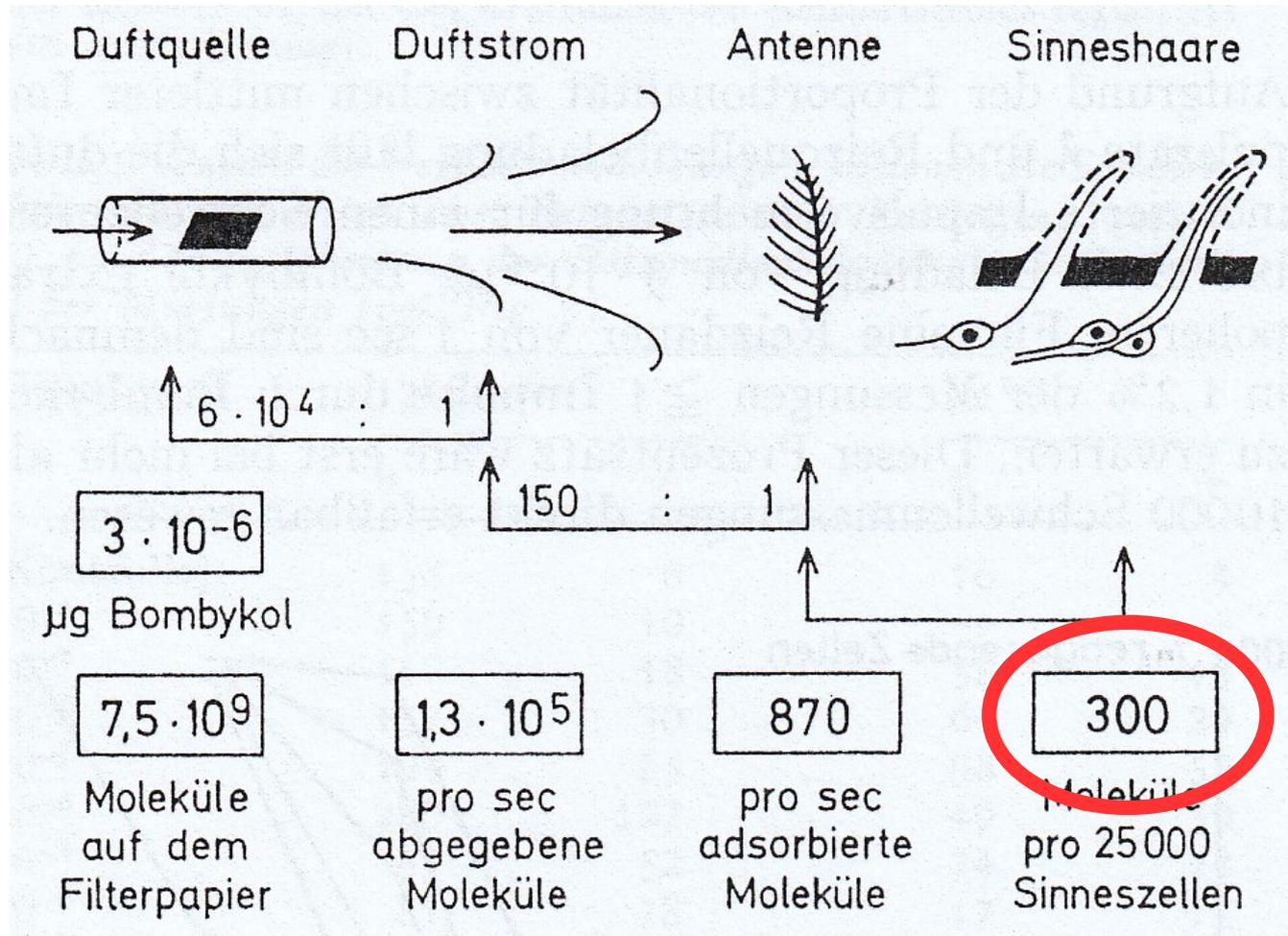
Kaissling & Priesner

Sensitivity (behavior and ORN response)



Kaissling & Priesner

Analysis with radio-actively labeled pheromone

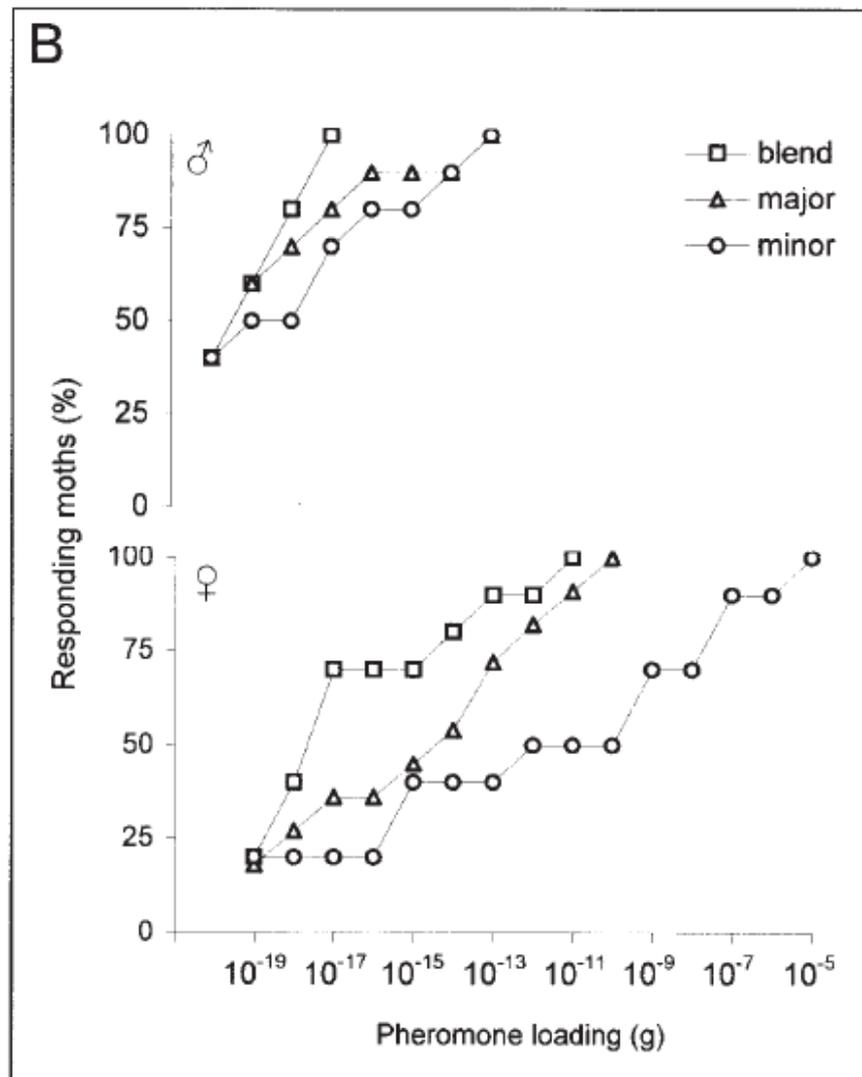


More recent study

Angioy AM et al. “Extreme Sensitivity in an Olfactory System”, *Chem Senses* **28**:279-284 (2003)

- Similar experiments but measuring the heartbeat of the moths (rather than the actual behavior)
- Experiments on *Spodoptera Littoralis* (cotton leafworm)

Results



Male moths seem to respond from $< 10^{-18}$ g (10^{-9} ng);
6 molecules on antenna (!)

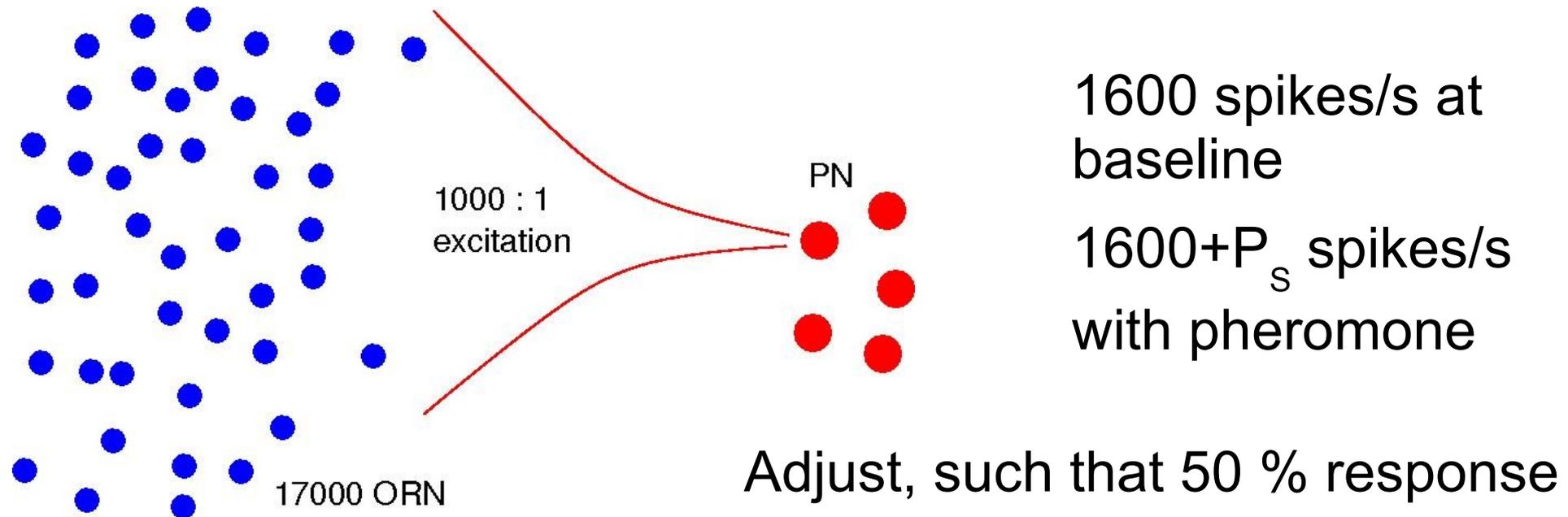
(Kaissling: $3 \cdot 10^{-3}$ ng;
300 molecules on antenna)

Angioy et al, 2003

Sensitivity analysis

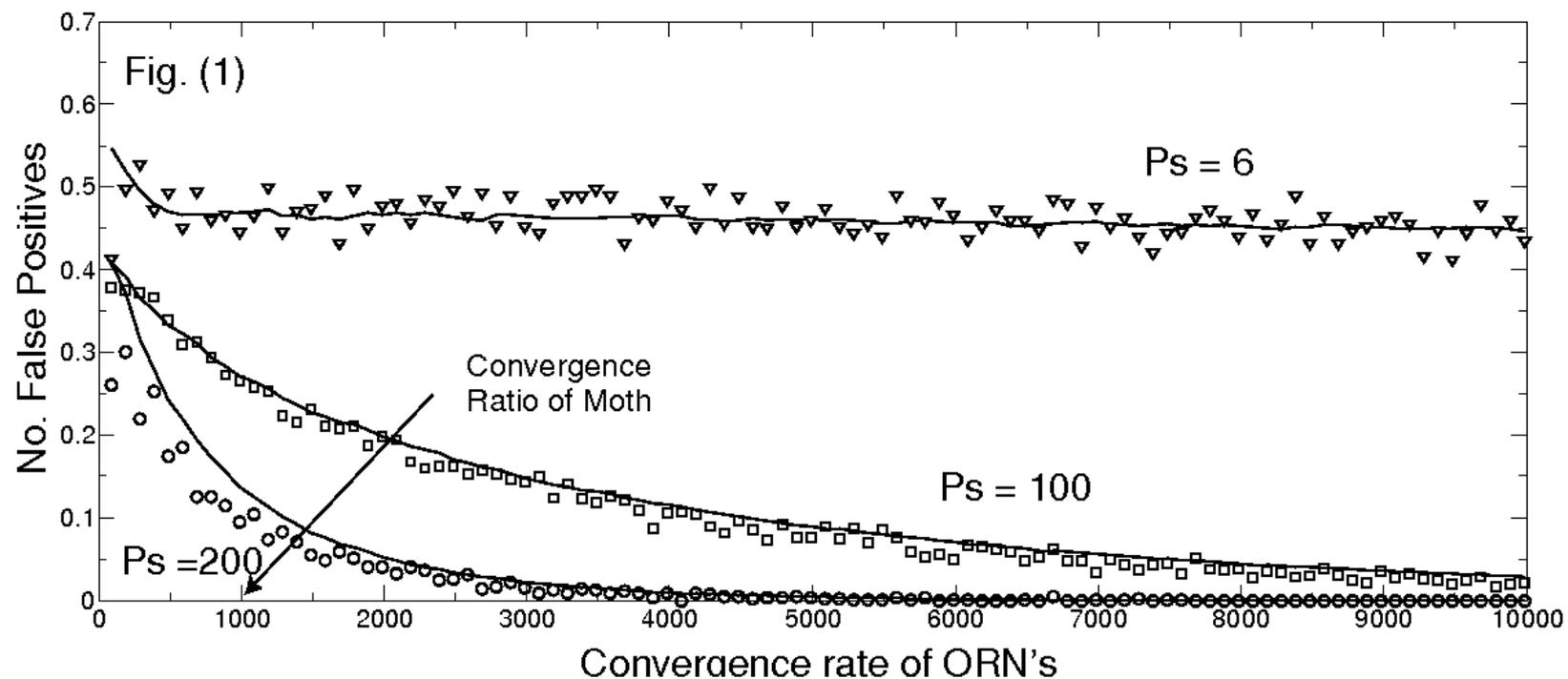
- Is the convergence of ~20000 ORN to less than 100 PN sufficient to explain response threshold at 300 molecules?

Simple statistical model:



Result on convergence sufficiency

Calculating the number of false positives:



Dr. Chris Buckley (Sussex)

<http://www.informatics.sussex.ac.uk/research/projects/PheroSys>

Conclusion so far

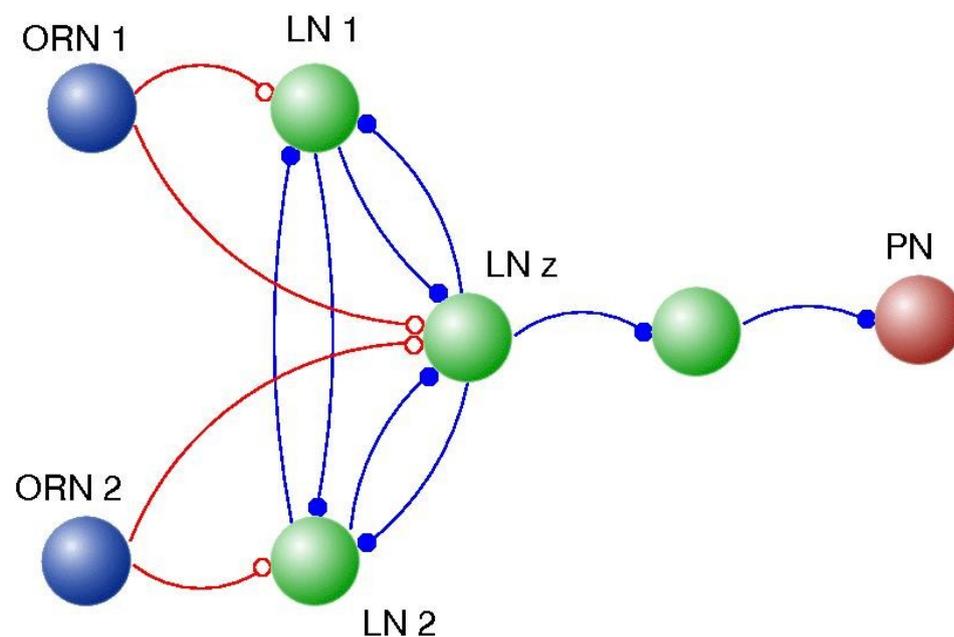
- s convergence enough to explain 200 molecule threshold? - maybe
- Is convergence enough to explain 6 molecule threshold? - no!
- ... to be continued

<http://www.informatics.sussex.ac.uk/research/projects/PheroSys/>

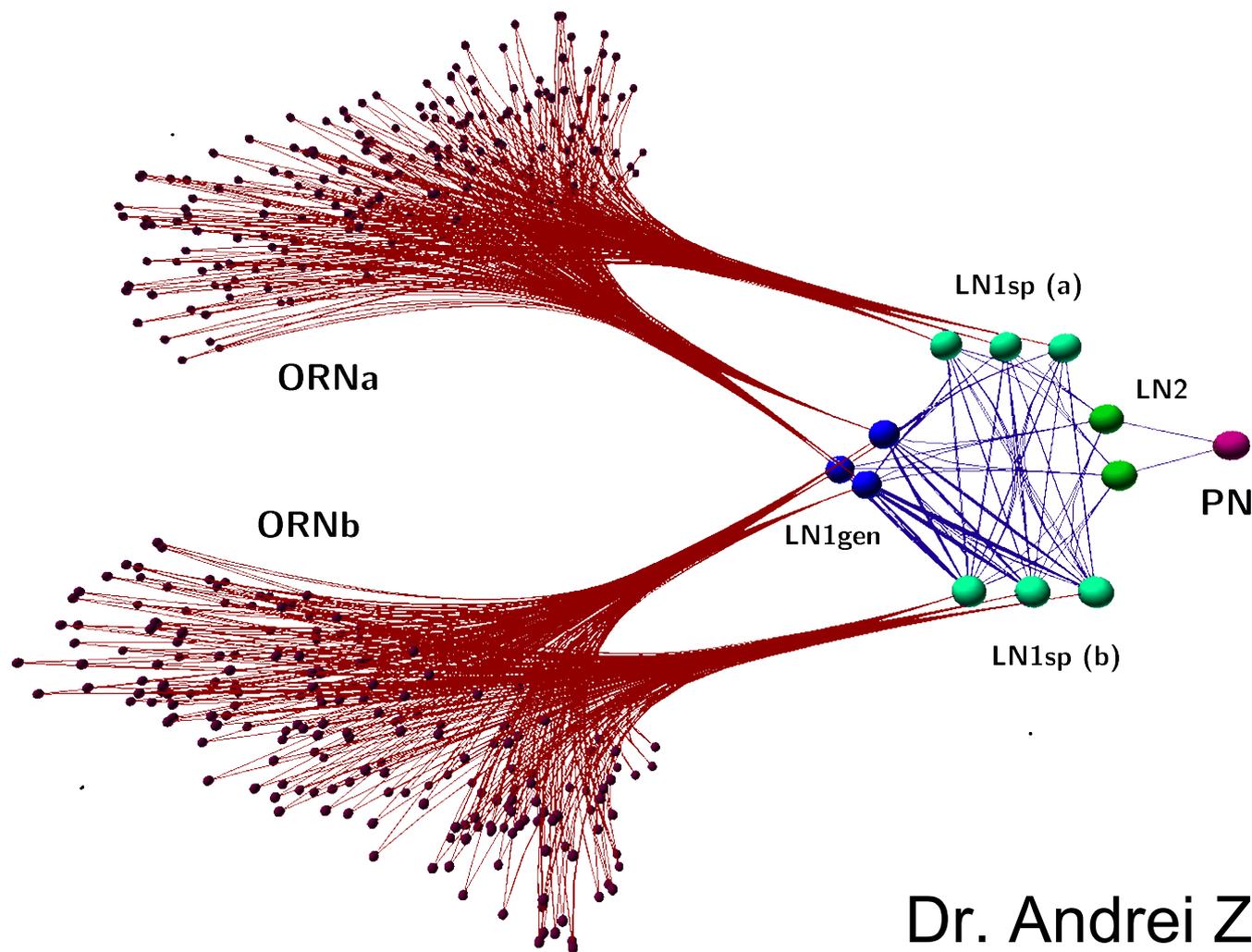
Ratio coding

- It is essential that moths can recognize the correct ratio over a large range of different concentrations
- In principle, this can be solved by winner-take-all competition, e.g.

See, e.g. Kwok YC, Encoding of Odor Blends in the Moth Antennal Lobe, PhD Thesis, University of Leicester, 2007



HH based model



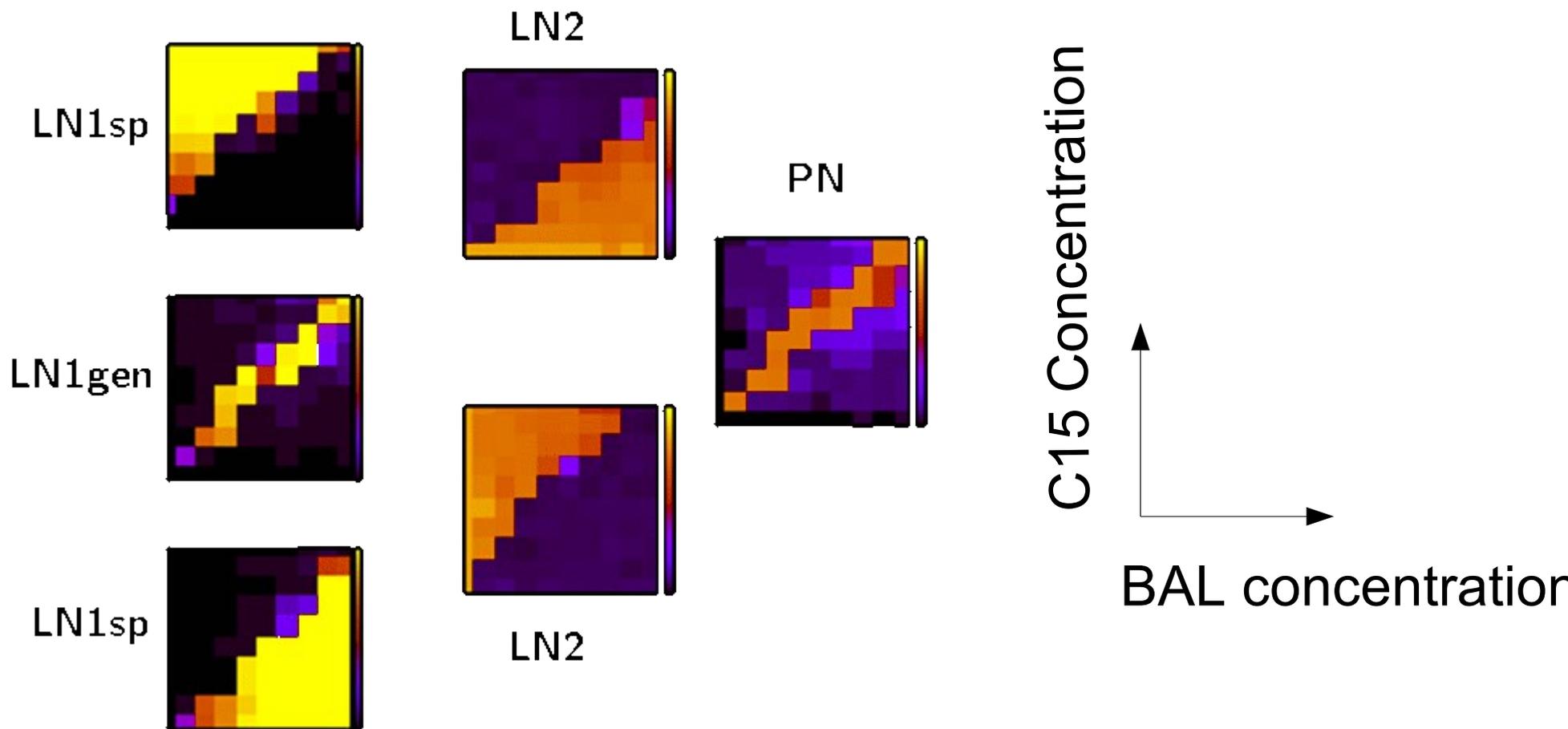
Dr. Andrei Zavada

<http://www.informatics.sussex.ac.uk/research/projects/PheroSys/>

Preliminary Results

b

$0.0118000 - 0.0092000$



Dr. Andrei Zavada

Ongoing work

- Optimize model with automated parameter estimation
- Analyze emerging synchronization phenomena
- Different ratios
- Generalization to multiple ratios

<http://www.informatics.sussex.ac.uk/research/projects/PheroSys/>

Discussion

- Existence of human sexual pheromones still debated
- Note: If they exist, it is still unclear what role they may play: It is unlikely that it is to find females like moths do