

Workshop: *Dynamical Complex Systems*

INTECIN -- Facultad de Ingeniería (U.B.A.)

December 17th, 2009 -- 3rd floor, EGRIT's room

Solving the inverse problem: deducing a kinetic model from electrophysiology data

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Direct problem

- We have an ion channel kinetic model (transitions probabilities between states)
- We know the experimental conditions

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We deduce the expected electrophysiology data

Inverse problem

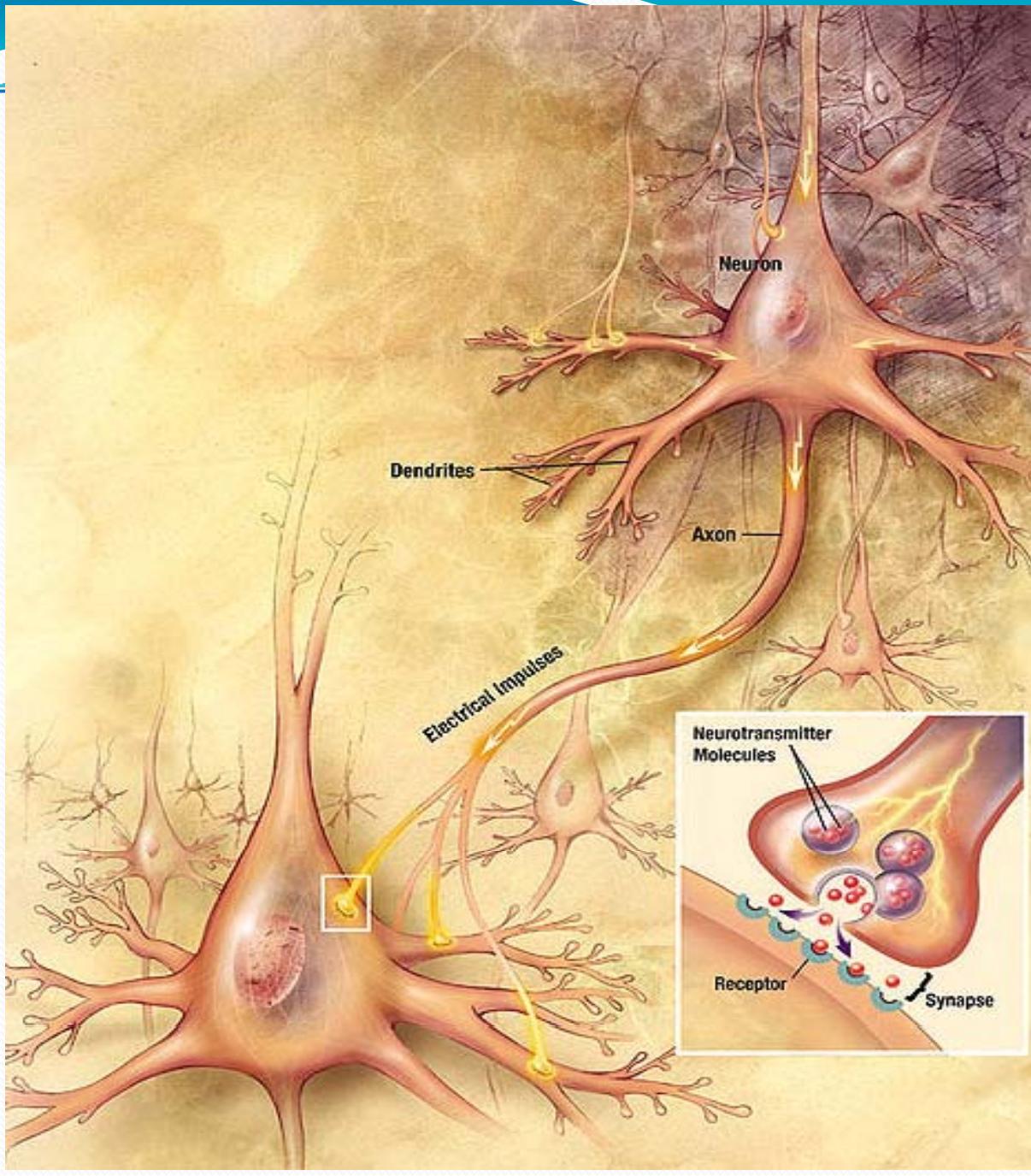
- We have the electrophysiology data
- We have the experimental conditions

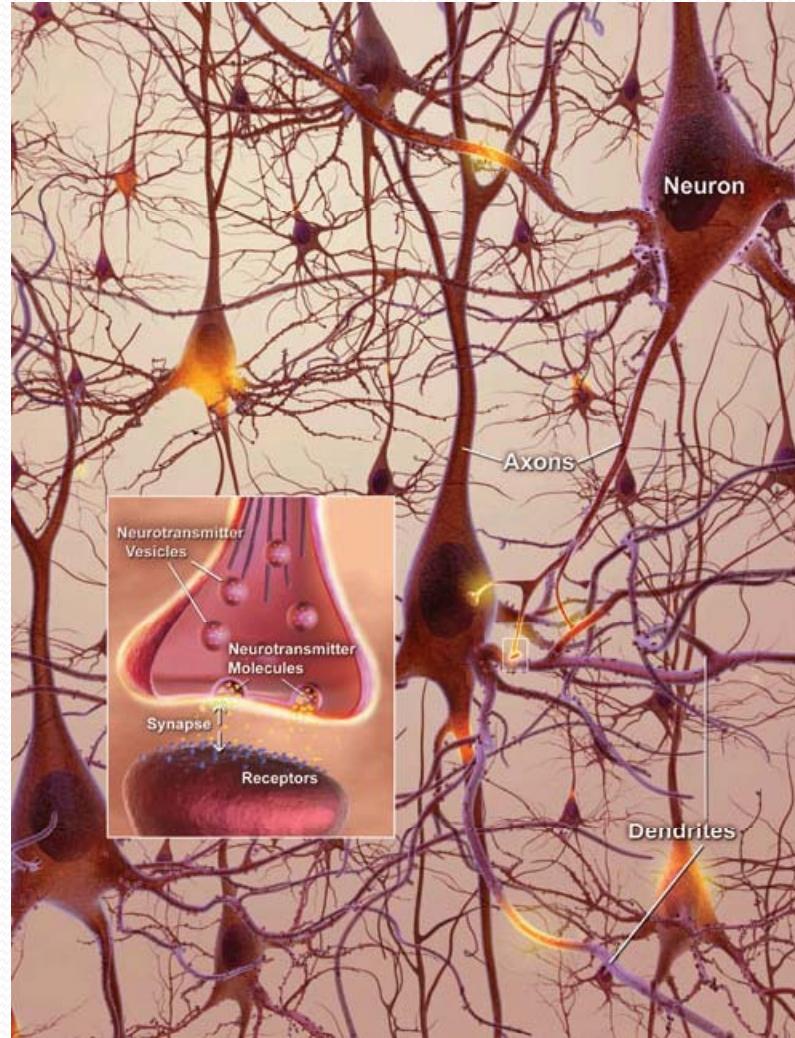
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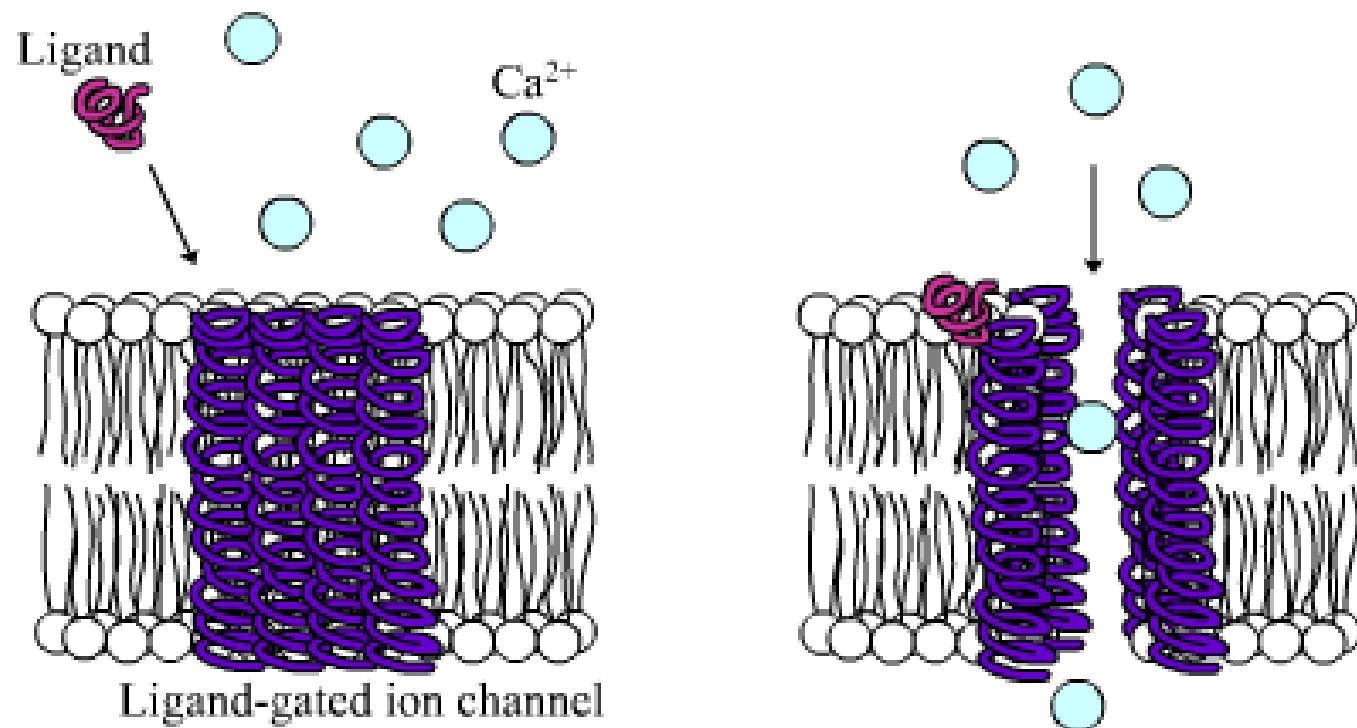
We deduce the kinetic model



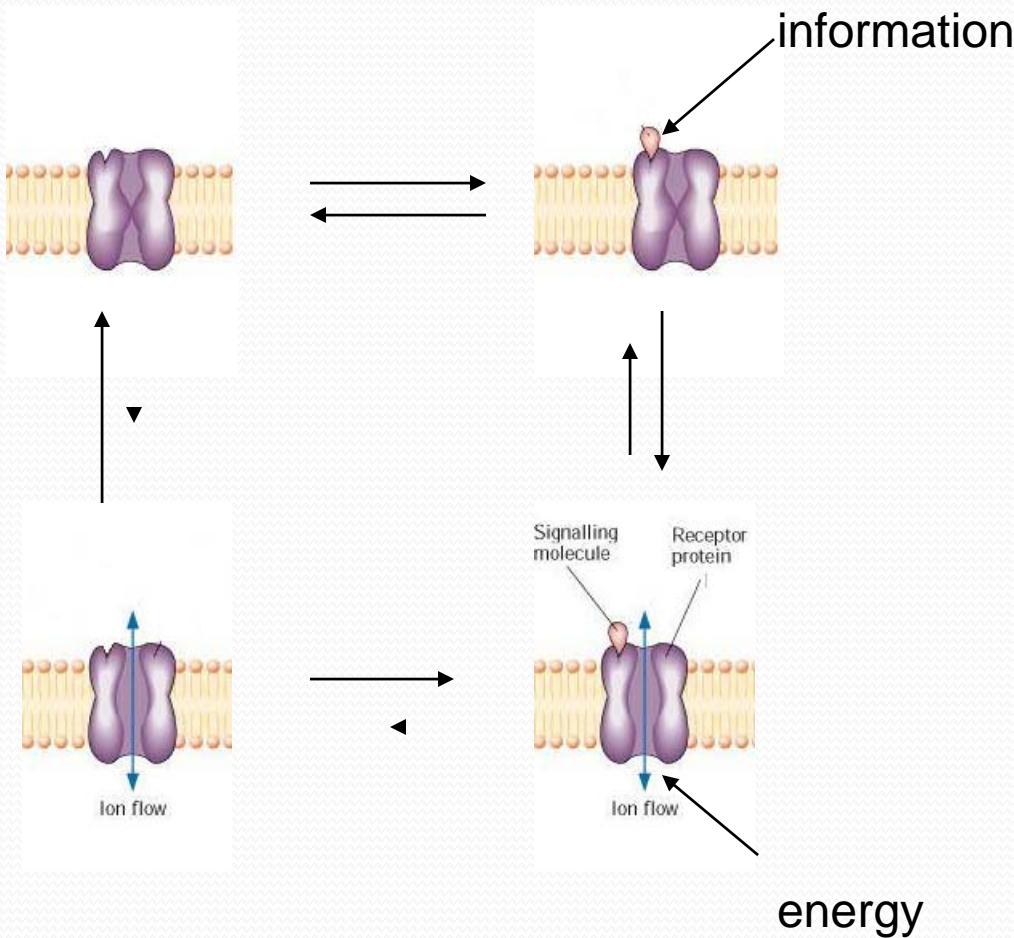
Some background information



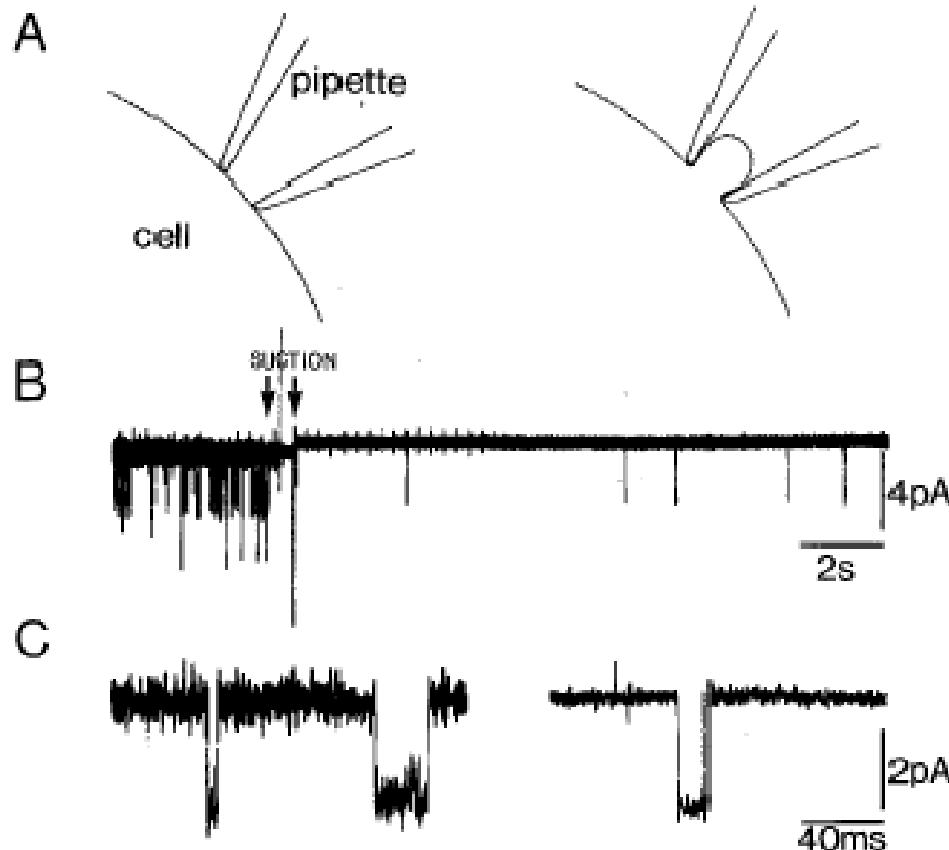




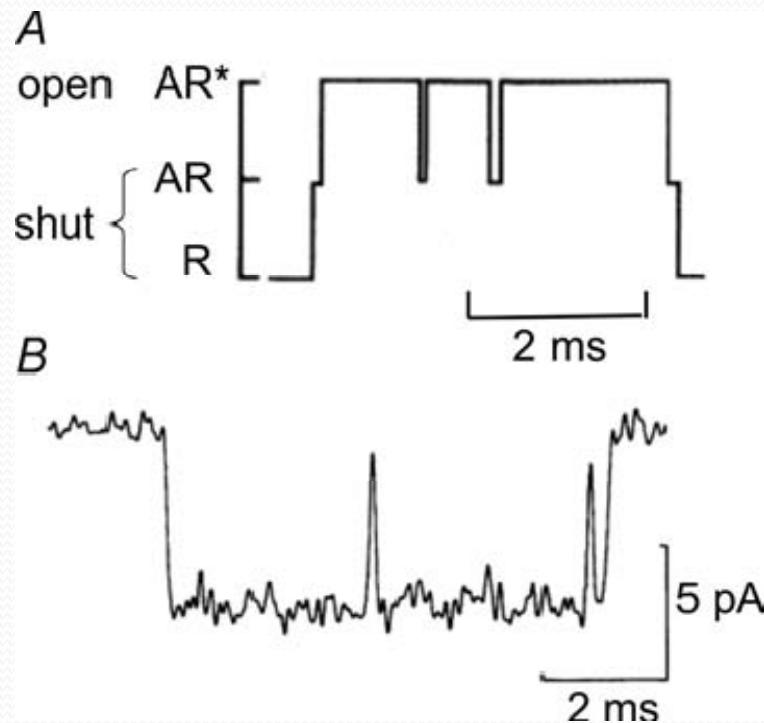
Allosteric coupling



Patch clamp technique

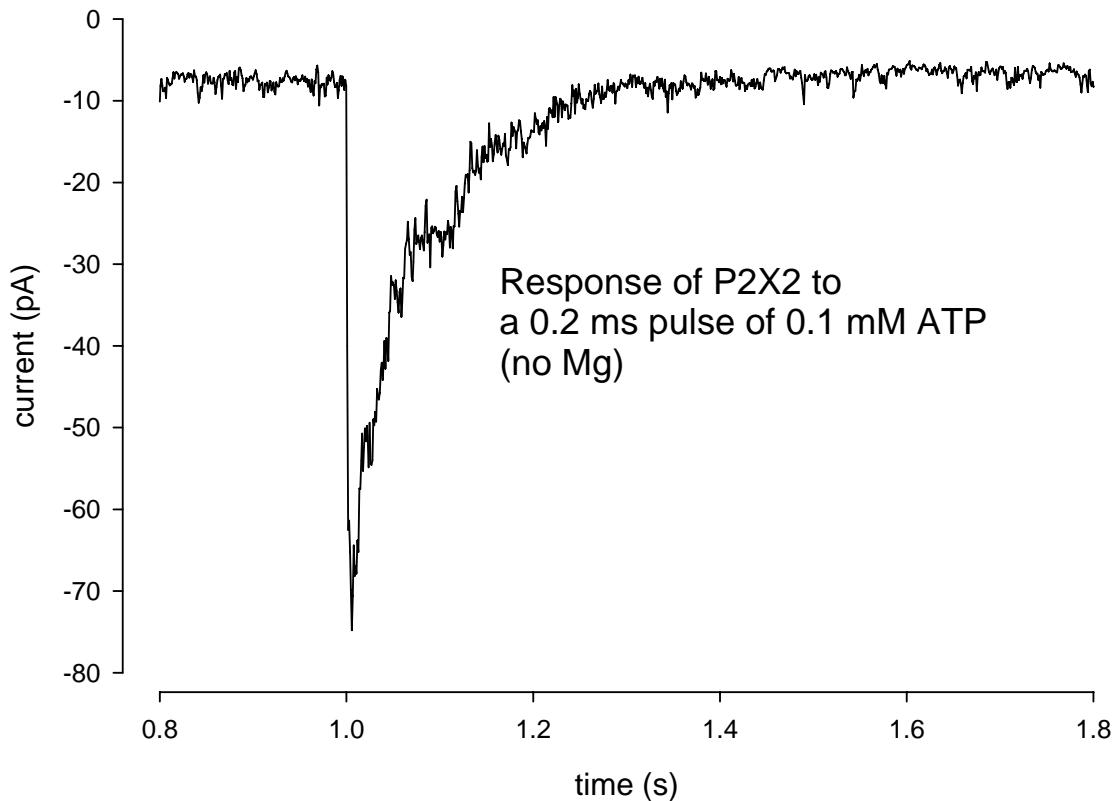


Ligand gated channels



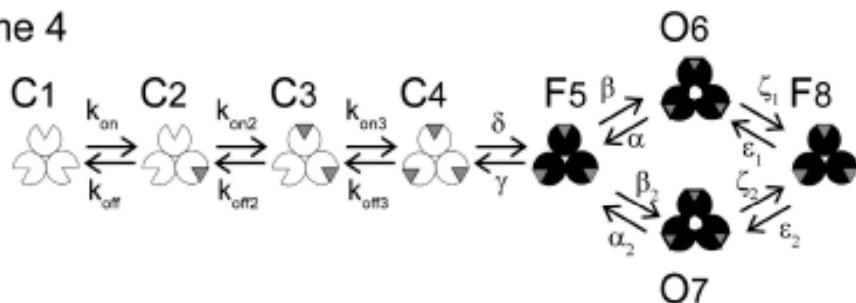
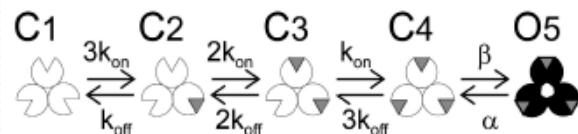
- Closed without agonist
- On agonist presence, ion channels open and close at random

Macroscopic currents

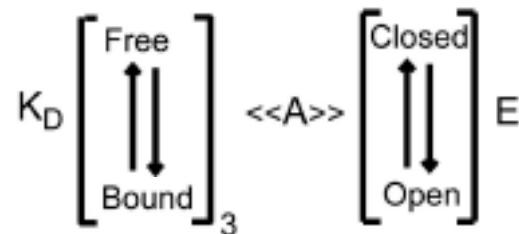


Kinetic models

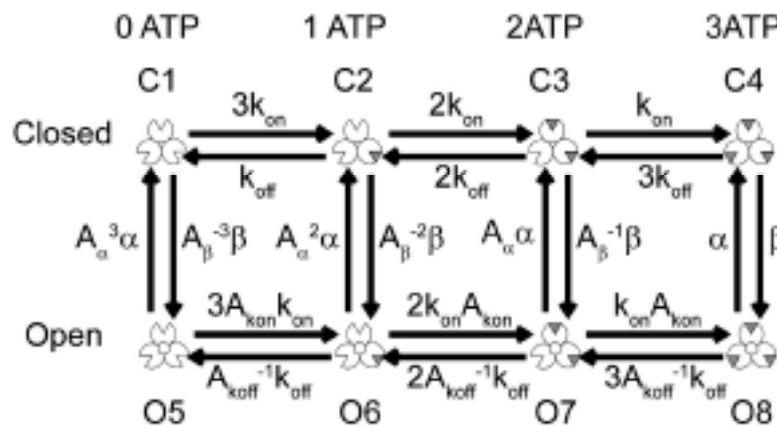
Scheme 4



Scheme 5, synthetic view



Scheme 5, expanded view



Direct problem

- Markov process master equation

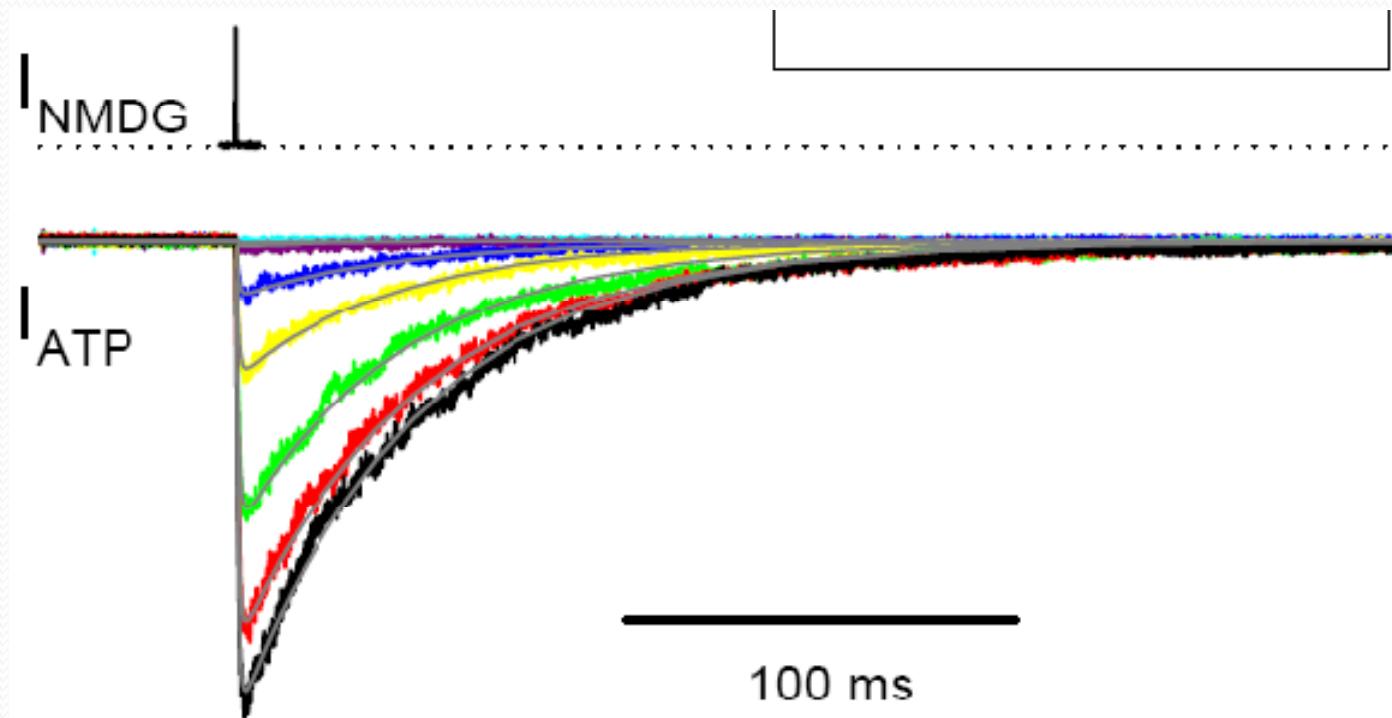
$$\dot{p}(t) = p(t) \cdot \mathbf{Q}(t)$$

$$p(t + \Delta t) = p(t) \cdot \exp(\mathbf{Q} \cdot \Delta t).$$

$$\mathbf{Q}(t) = (\mathbf{Q}_0 + \mathbf{Q}_a \cdot ATP(t))$$

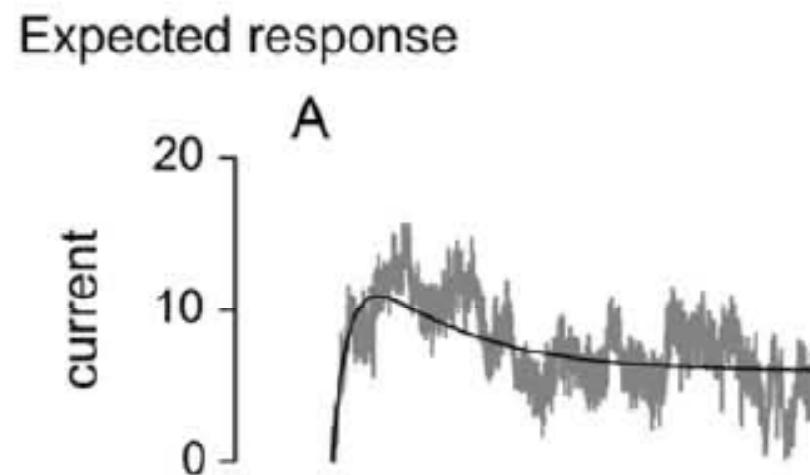
$$y(t) = p(t) \cdot g$$

Inverse problem

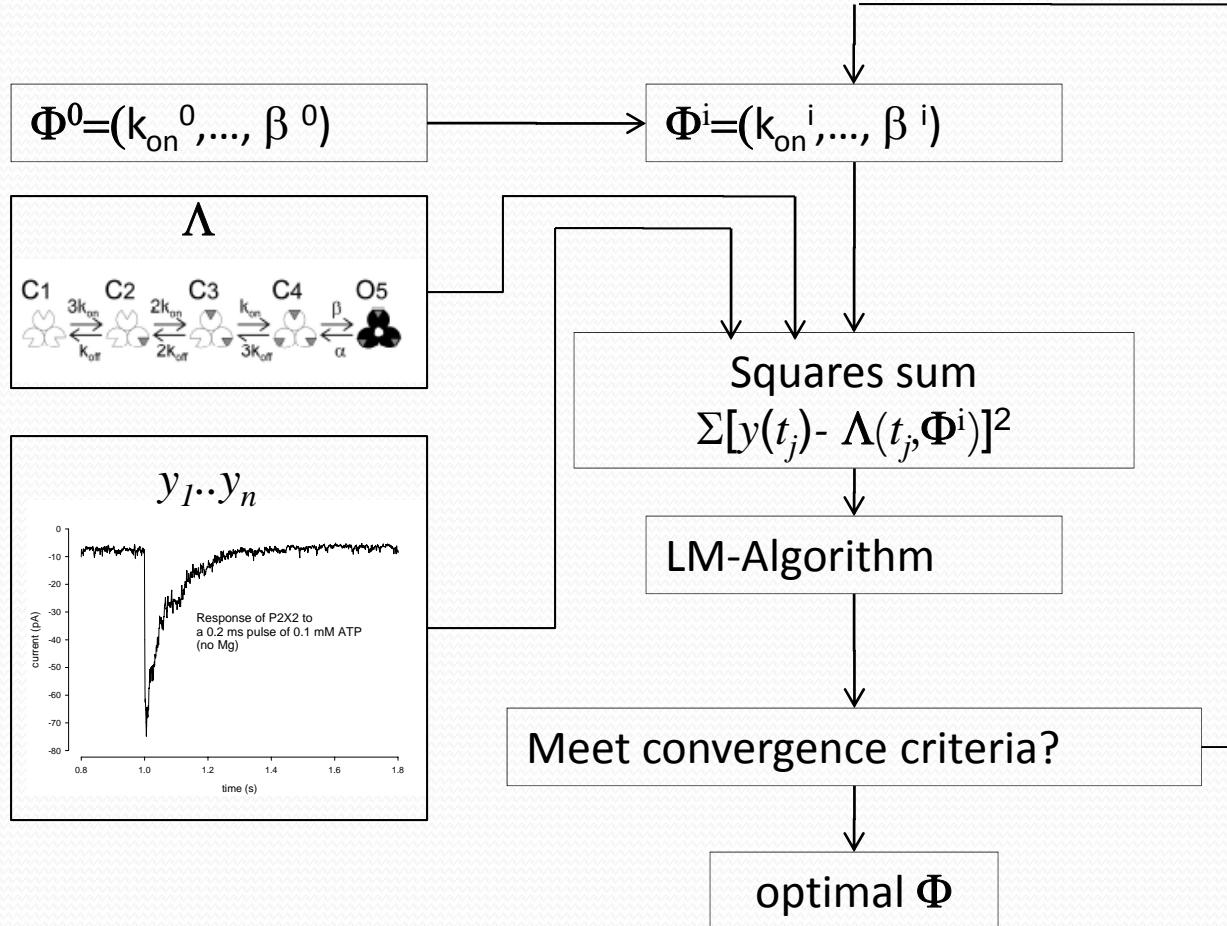


Least squares approach

- Given a kinetic model, make an initial guess of the kinetic constants and minimize the squared difference between expected and measured.

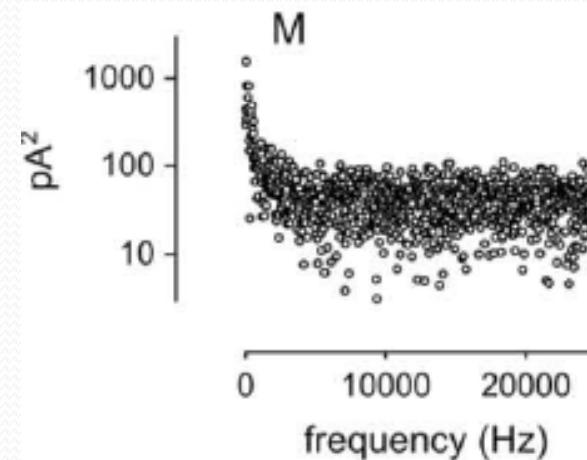


Least squares estimation



Problems with this approach

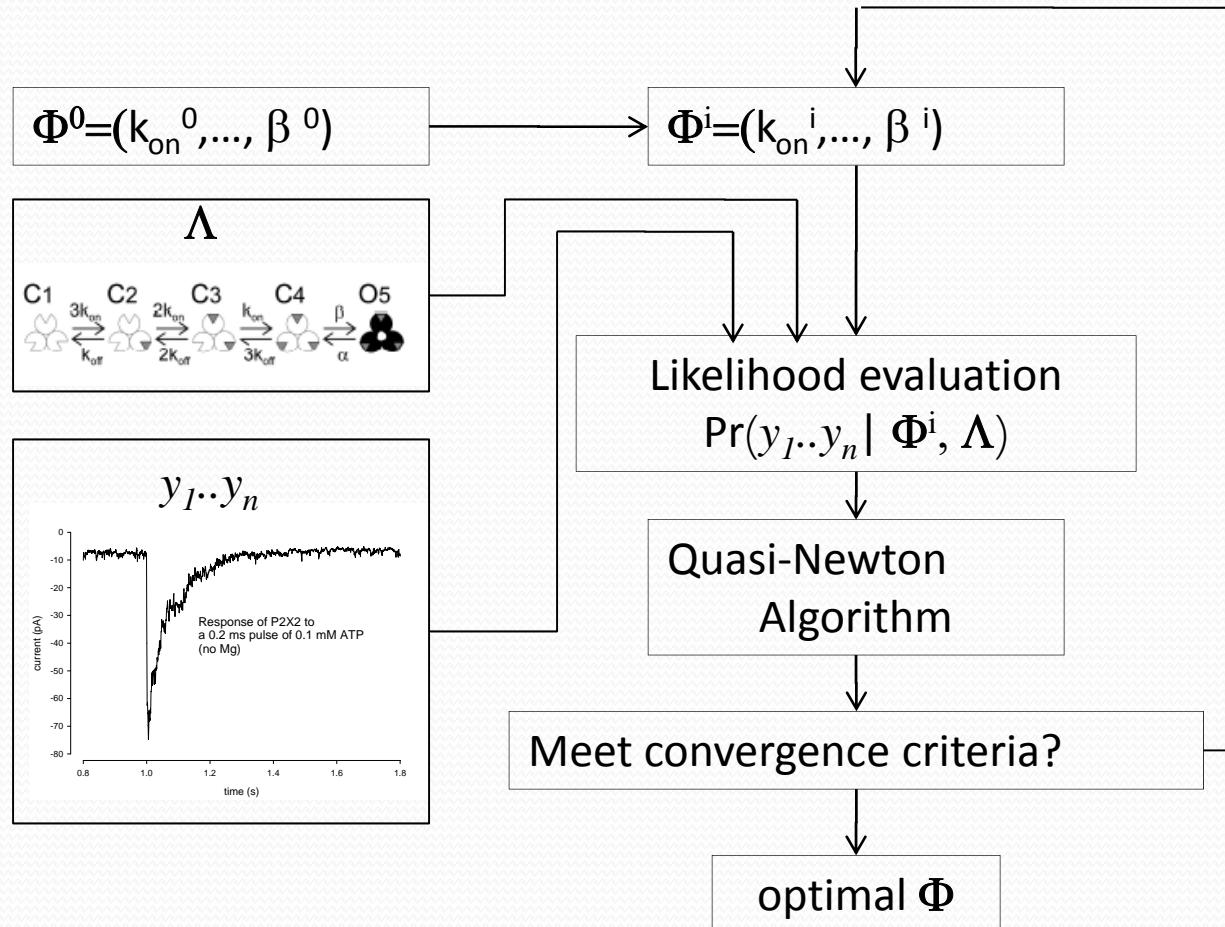
- Time correlation on the residuals-> NO VALID STATISTICAL ANALYSIS



A possible solution

- Maximize the likelihood.

Maximum likelihood estimation

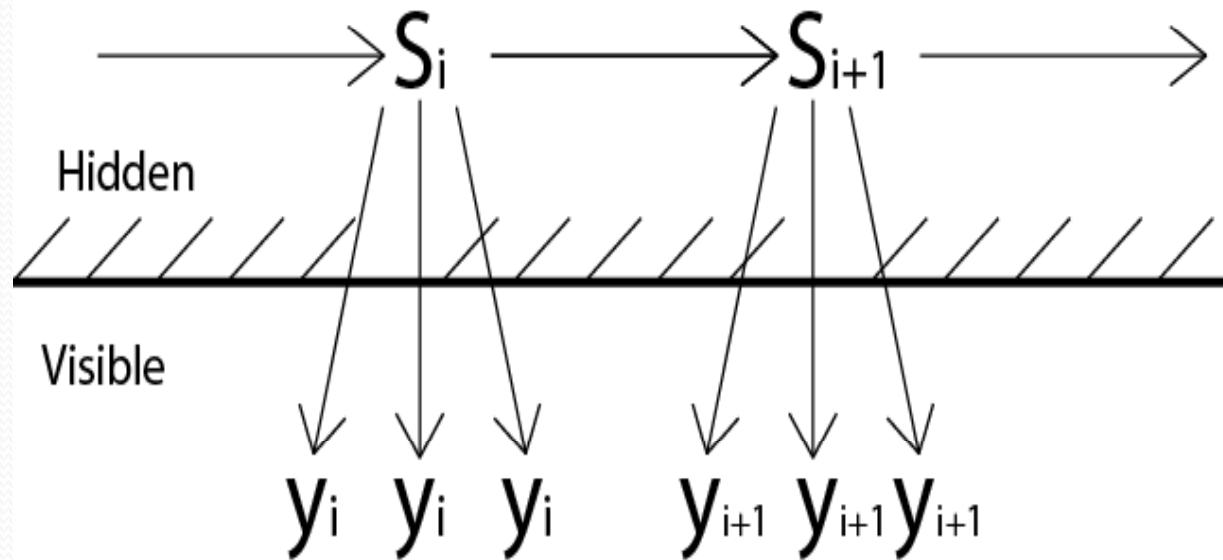


Using the incremental likelihood

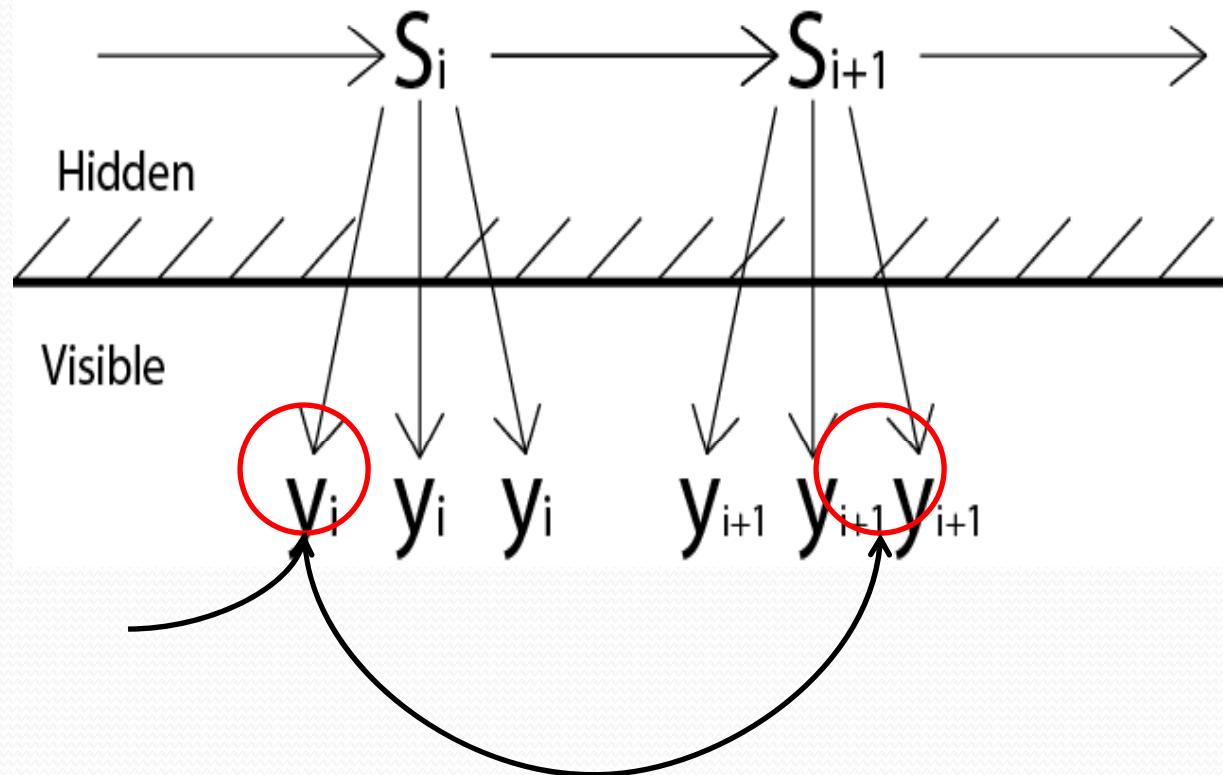
$$\ell_T = \ell(\Theta | y_1 \dots y_T) = \prod_{t=1}^T \ell'_t,$$
$$\ell'_t = Pr(y_t | y_1 \dots y_{t-1}, \Theta).$$

- It decomposes the likelihood of the joint measurement on a product of the partial likelihoods of each measurement, given the previous ones.

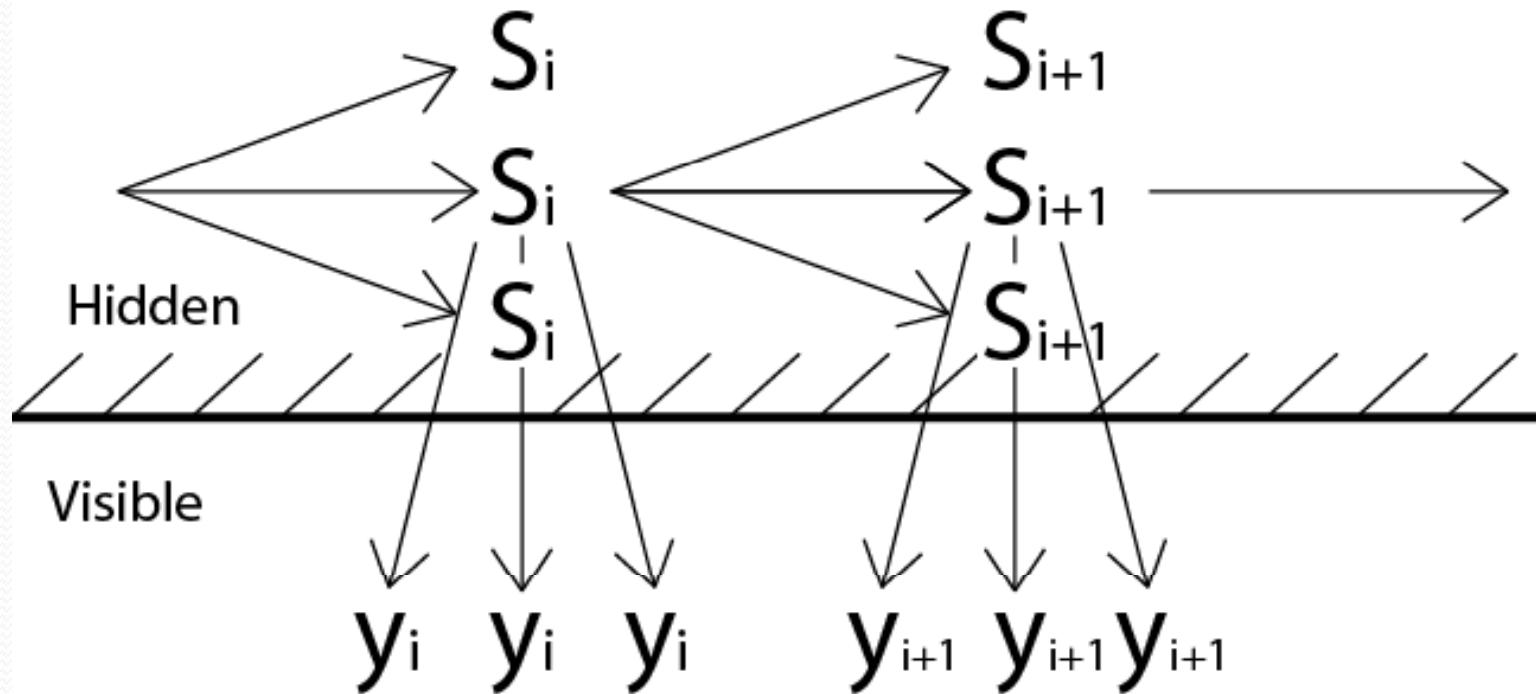
Deterministic models



Deterministic models



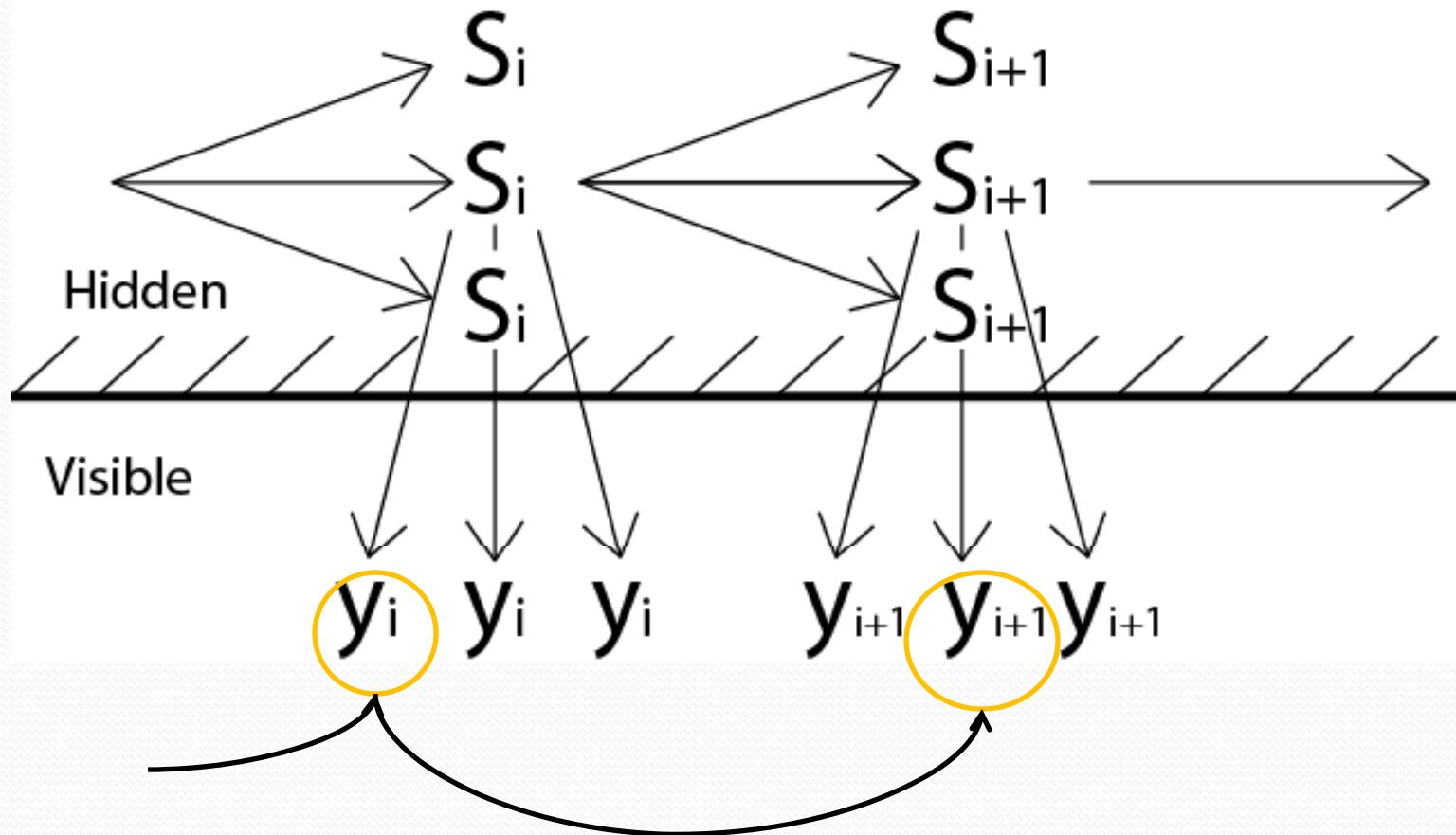
Stochastic models



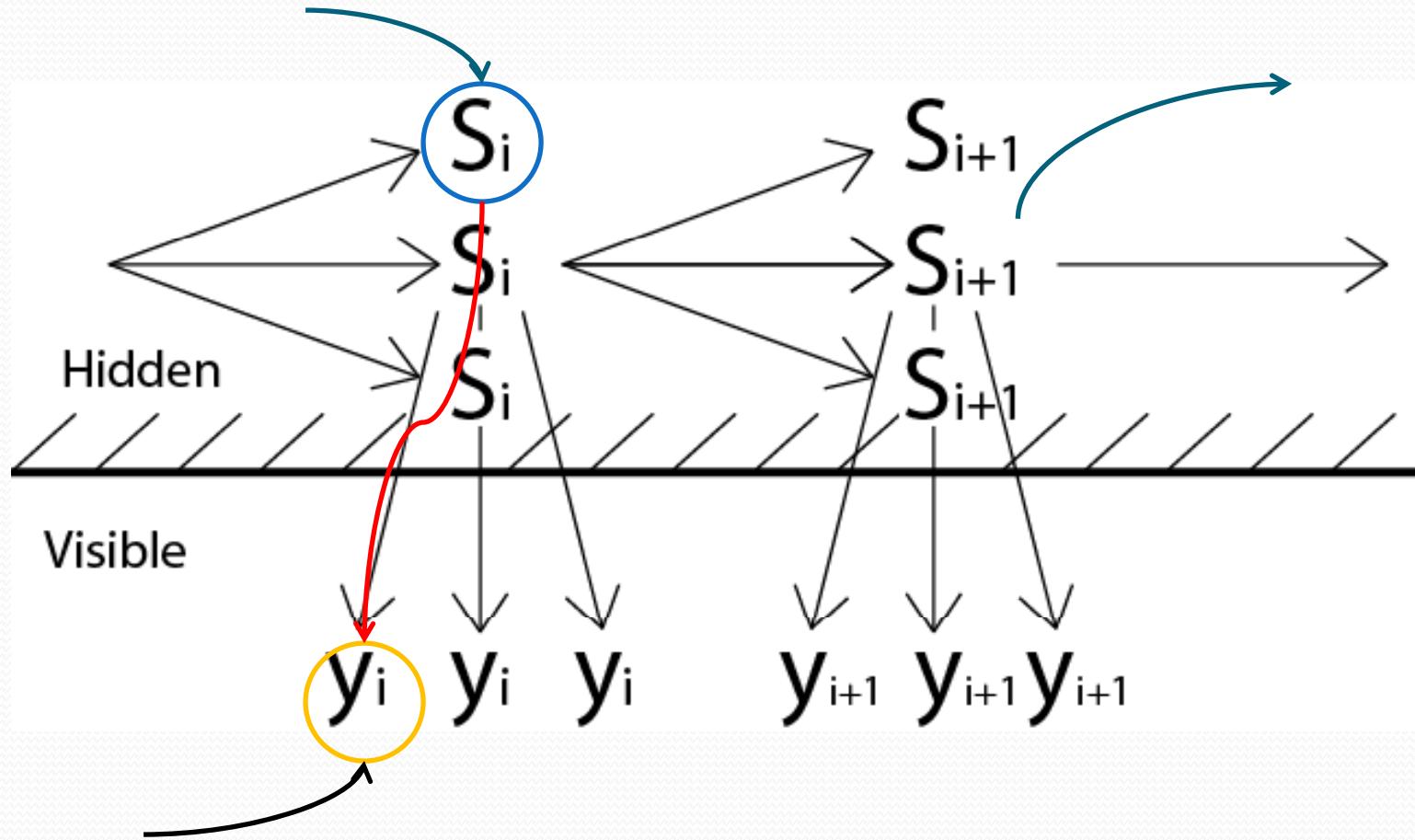
Classical Hidden Markov Modeling

- **Markov model**: rules of change of the hidden system
- **Observational model**: the relationship between the hidden and the observed
- The sequence of **hidden states**
- The sequence of **observations**

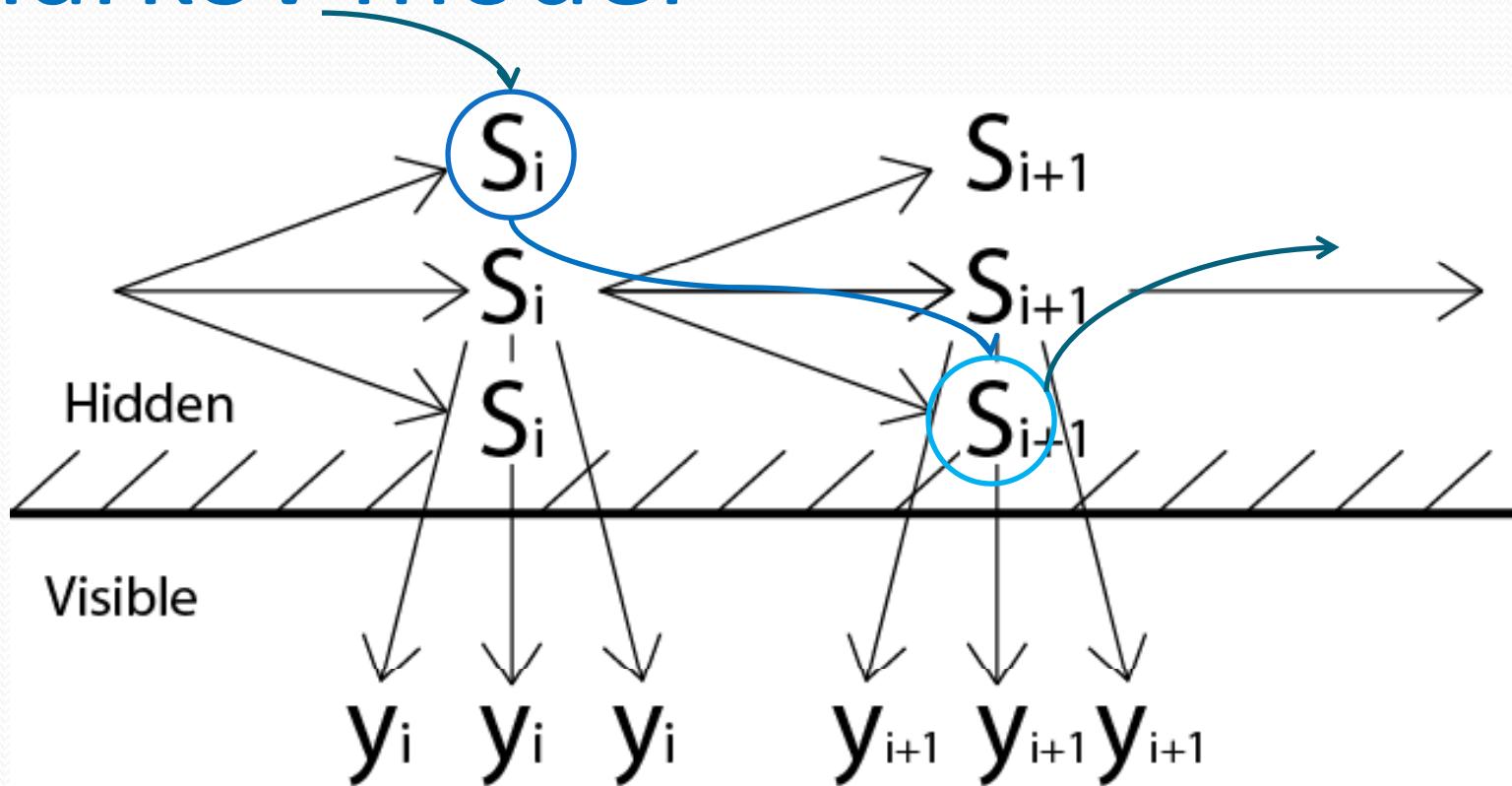
Observations



Observational model



Markov model



Bayesian statistics

- Probability is the measure of the plausibility of something given a series of hypothesis and data.
- Bayes theorem

$$\Pr(H_i/D, X) = \Pr(H_i/X) \frac{\Pr(D/H_i, X)}{\Pr(D/X)}$$

Hi-> Series of
alternative
Hypothesis
D-> new data
X-> old data

Bayes theorem to HMM

- Incremental Likelihood from Observational model

$$\Pr(y_n/y_{1..n-1}, \Theta) = \sum_j \Pr(y_n/s_j(t_n), y_{1..n-1}, \Theta) \Pr(s_i(t_n)/y_{1..n-1}, \Theta)$$

- Posterior

$$\Pr(s_i(t_n)/y_{1..n}, \Theta) = \Pr(s_i(t_n)/y_{1..n}, \Theta) \frac{\Pr(y_n/s_i(t_n), y_{1..n-1}, \Theta)}{\Pr(y_n/y_{1..n-1}, \Theta)}$$

Markov chain rule

- Markov model

$$\Pr(s_i(t_{n+1})/s_j(t_n)) = a_{i,j}$$

$$a_{i,j} = \exp(\mathbf{Q} \cdot (t_{n+1} - t_n))$$

- Next prior

$$\Pr(s_j(t_{n+1})/y_{1...n}, \Theta) = \sum_j a_{i,j} \Pr(s_i(t_n)/y_{1...n}, \Theta)$$

Approaches to many channels

Definition of the “meta state”

Microscopic approach

- It lists all the possible combination of states:
 $M = \binom{N+k-1}{N}$
N,0,0,0,...,0
N-1,1,0,...,0
N-1,0,1,...,0
0,0,0,...N
- The knowledge we have about the state of the system is a M-vector

Microscopic recursive

- It uses the same equations of single channel applied to the M-state.

Macroscopic approach

$$Pr(\mathbf{r}_t) = N\left(\mathbf{r} - \boldsymbol{\mu}_t, \frac{1}{N}\boldsymbol{\Sigma}_t\right) = N^{k/2}(2\pi)^{-k/2}|\boldsymbol{\Sigma}_t|^{-1/2} \\ \exp\left(-\frac{N}{2}(\mathbf{r} - \boldsymbol{\mu}_t) \cdot \boldsymbol{\Sigma}_t^{-1} \cdot (\mathbf{r} - \boldsymbol{\mu}_t)^T\right),$$

- The knowledge about the state of the channels (i.e., \mathbf{r}) is provided by a continuous multinormal distribution.
- Two parameters define the distribution: the mean $\boldsymbol{\mu}$ and covariance $\boldsymbol{\Sigma}$

Incremental Likelihood

$$\ell'_t = \int_s Pr(y_t | s_t = \mathbf{r}, \varepsilon^2) \cdot p_t^*(\mathbf{r}) \cdot d\mathbf{r}.$$

$$\ell'_t = \int_s N(y_t - N\mathbf{r}_t \cdot \boldsymbol{\gamma}, \varepsilon^2) N\left(\mathbf{r}_t - \boldsymbol{\mu}_t^*, \frac{1}{N} \boldsymbol{\Sigma}_t^*\right) d\mathbf{r}_t,$$

$$\ell'_t = N(y_t - y_t^*, \sigma_t^2), \quad y_t^* = N\boldsymbol{\mu}_t^* \cdot \boldsymbol{\gamma}, \\ \sigma_t^2 = \varepsilon^2 + N\boldsymbol{\gamma}^T \cdot \boldsymbol{\Sigma}_t^* \cdot \boldsymbol{\gamma}.$$

- We integrate the product of the measurement probability by the prior

Posterior

$$p_t(\mathbf{r}) = \frac{N(y_t - N\mathbf{r}_t \cdot \boldsymbol{\gamma}, \varepsilon^2)}{\ell'_t} N\left(\mathbf{r}_t - \boldsymbol{\mu}_t^*, \frac{1}{N} \boldsymbol{\Sigma}_t^*\right).$$

$$p_t(\mathbf{r}) = N\left(\mathbf{r}_t - \boldsymbol{\mu}_t, \frac{1}{N} \boldsymbol{\Sigma}_t\right),$$

$$\boldsymbol{\mu}_t = \boldsymbol{\mu}_t^* + \frac{(y_t - y_t^*)}{\sigma_t^2} \cdot \boldsymbol{\gamma}^T \cdot \boldsymbol{\Sigma}_t^*,$$
$$\boldsymbol{\Sigma}_t = \boldsymbol{\Sigma}_t^* - \frac{N}{\sigma_t^2} \boldsymbol{\Sigma}_t^* \cdot \boldsymbol{\gamma} \cdot \boldsymbol{\gamma}^T \cdot \boldsymbol{\Sigma}_t^*.$$

- We normalize the priors by the measurement probabilities.
- We just update the value of the mean and covariance

Completing the loop

$$\boldsymbol{\mu}_{t+\Delta t}^* = \boldsymbol{\mu}_t \cdot \mathbf{A},$$

$$\boldsymbol{\Sigma}_{t+\Delta t}^* = (\boldsymbol{\mu}_t \cdot \mathbf{A})^{\mathbf{D}} - \mathbf{A}^T \cdot \boldsymbol{\mu}_t^{\mathbf{D}} \cdot \mathbf{A} + \mathbf{A}^T \cdot \boldsymbol{\Sigma}_t \cdot \mathbf{A}.$$

- The mean is updated with the usual formula for single channels

Simulations

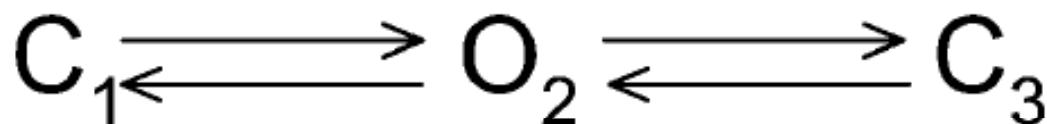
Pet model

$500 \text{ M}^{-1}\text{s}^{-1}$

k_{12}

100 s^{-1}

k_{23}



k_{21}

200 s^{-1}

k_{32}

50 s^{-1}

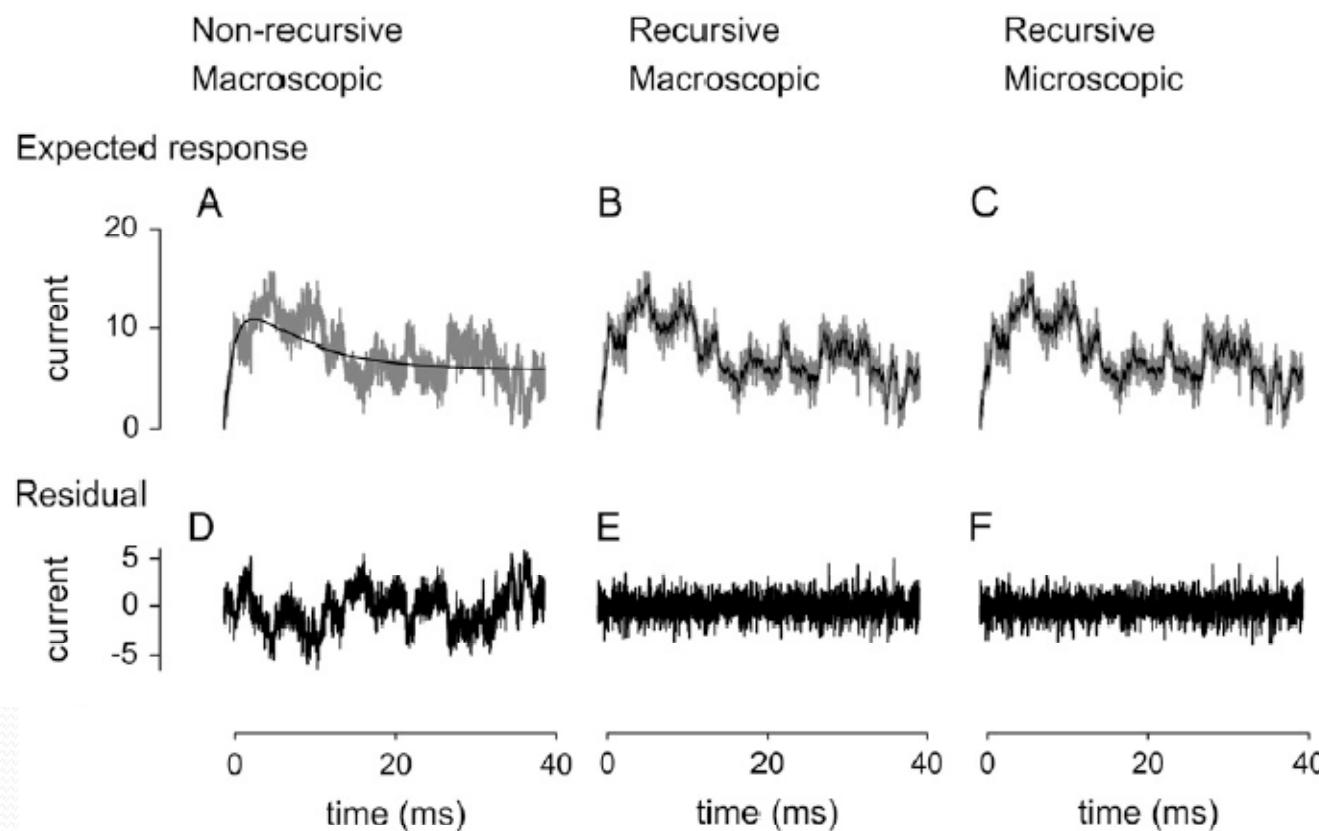
$\varepsilon=1 \text{ pA}; \gamma=1\text{pA}; \text{sample time}=0.02\text{ms}$

Algorithms

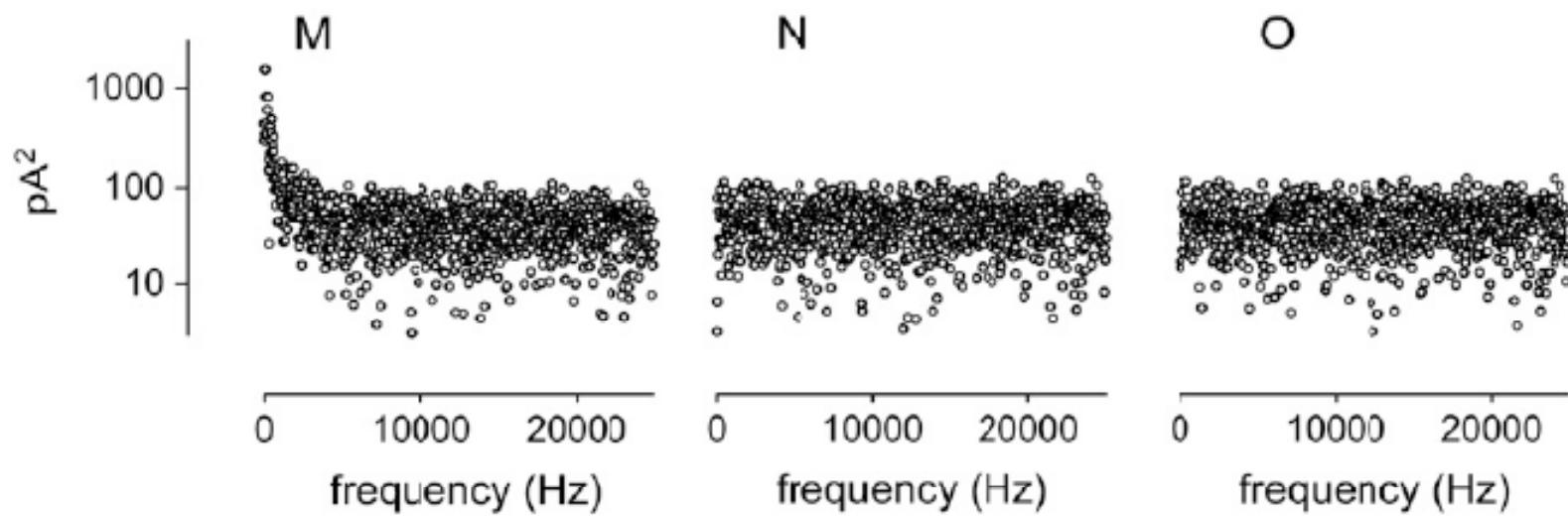
- Microscopic Recursive
- Macroscopic Recursive
- Macroscopic Non Recursive

Non-stationary conditions

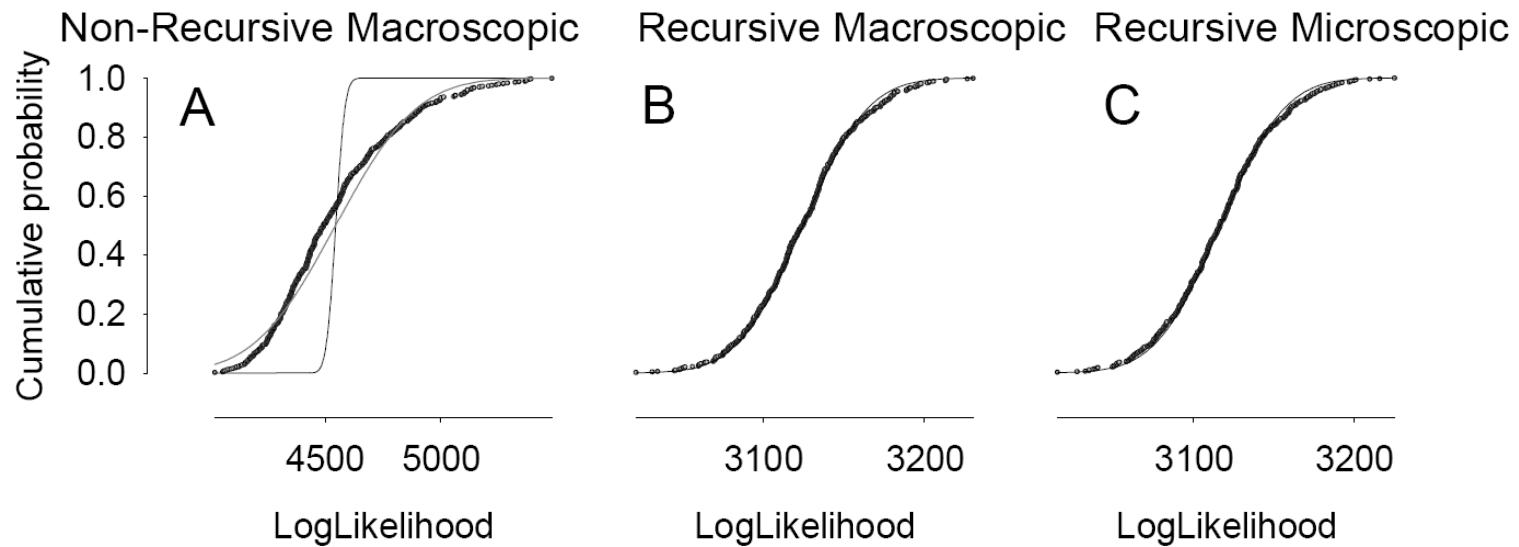
Prediction of the response



FFT of the residuals



Likelihood distribution



- The expected variance of the loglikelihood of a normal is $\frac{1}{2}$, therefore the whole loglikelihood has a $\sigma^2=1/2 n_{\text{samples}}$

Prior probability evolution

Micro R red

Macro R blue

Macro NR black

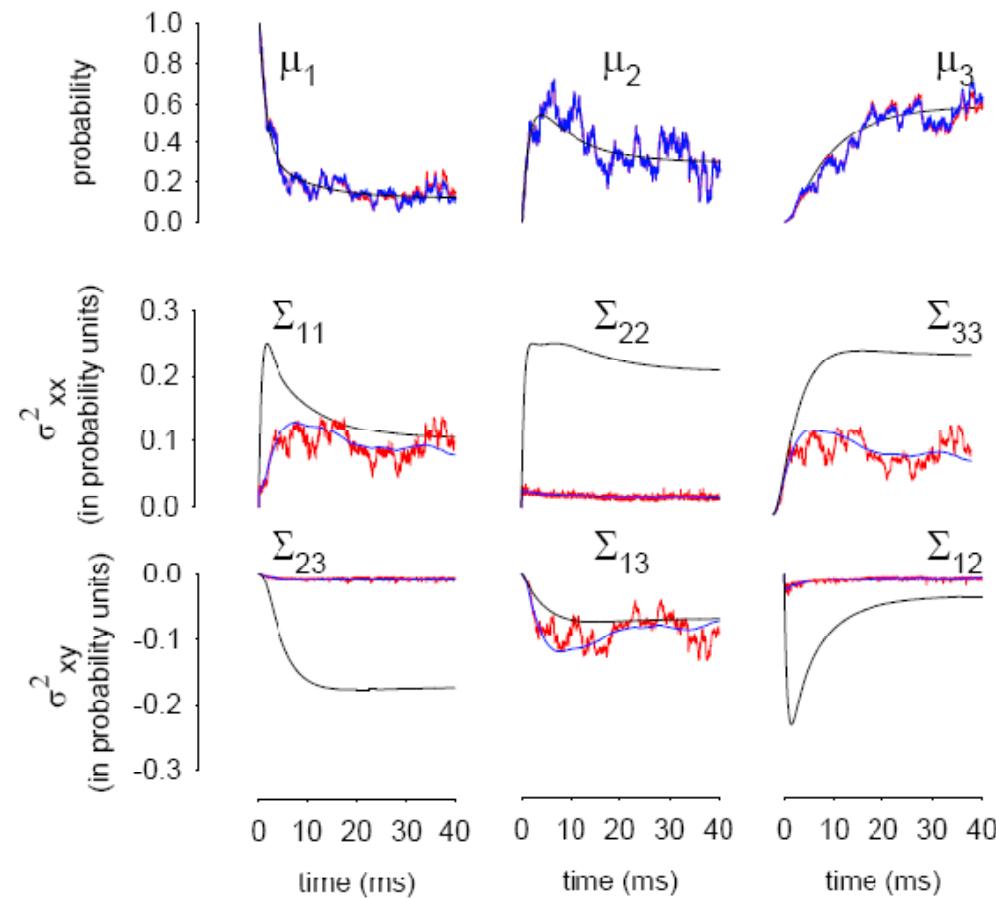
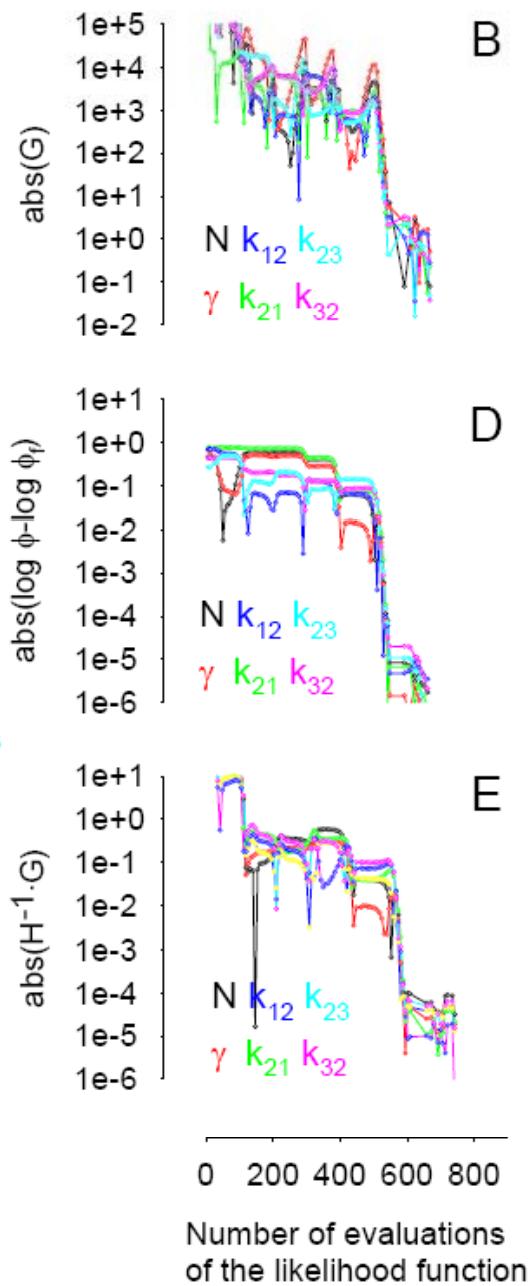
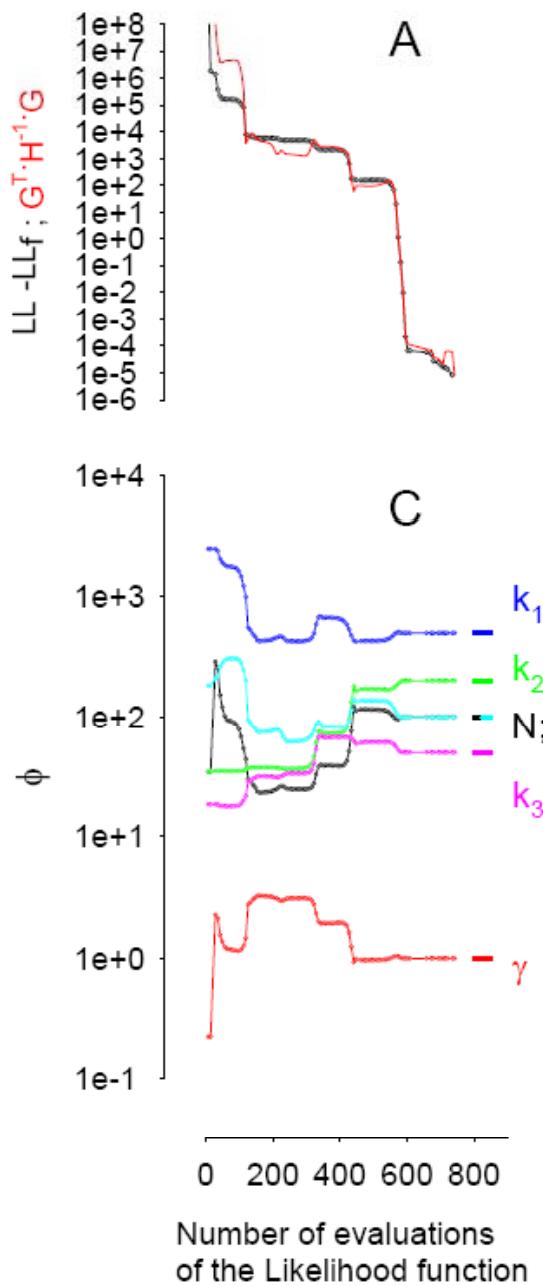
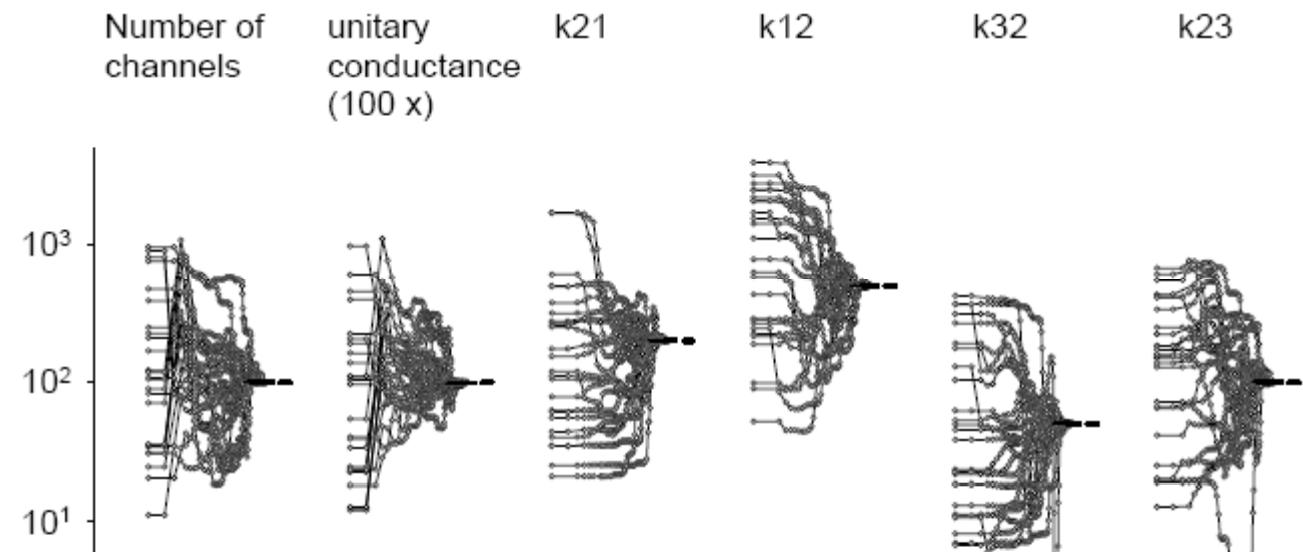


Figure 3

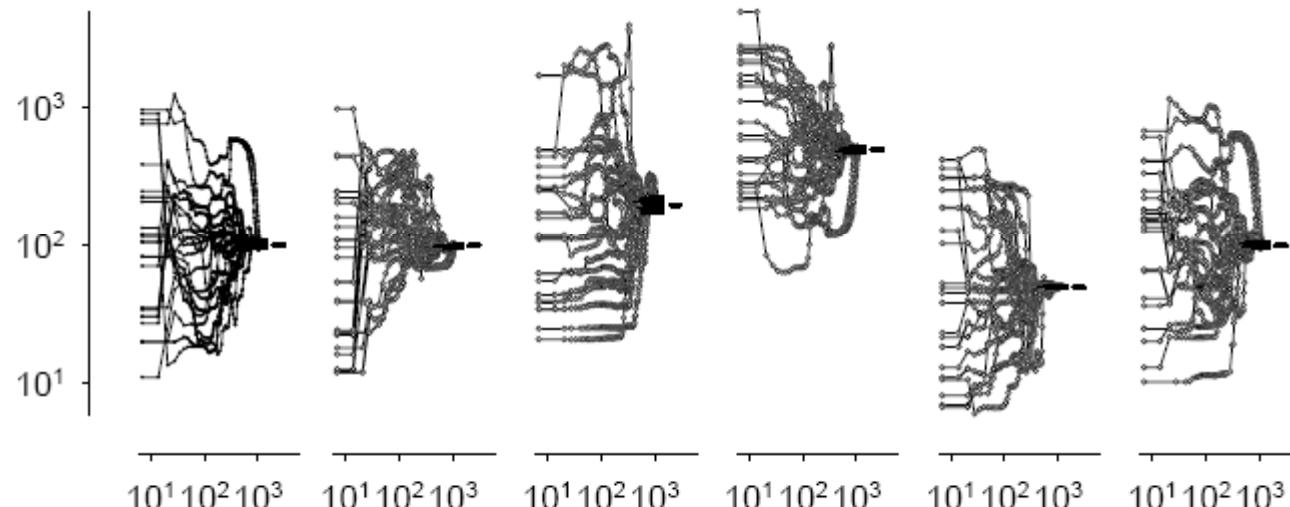


A

Recursive Macroscopic

**B**

Non-Recursive Macroscopic

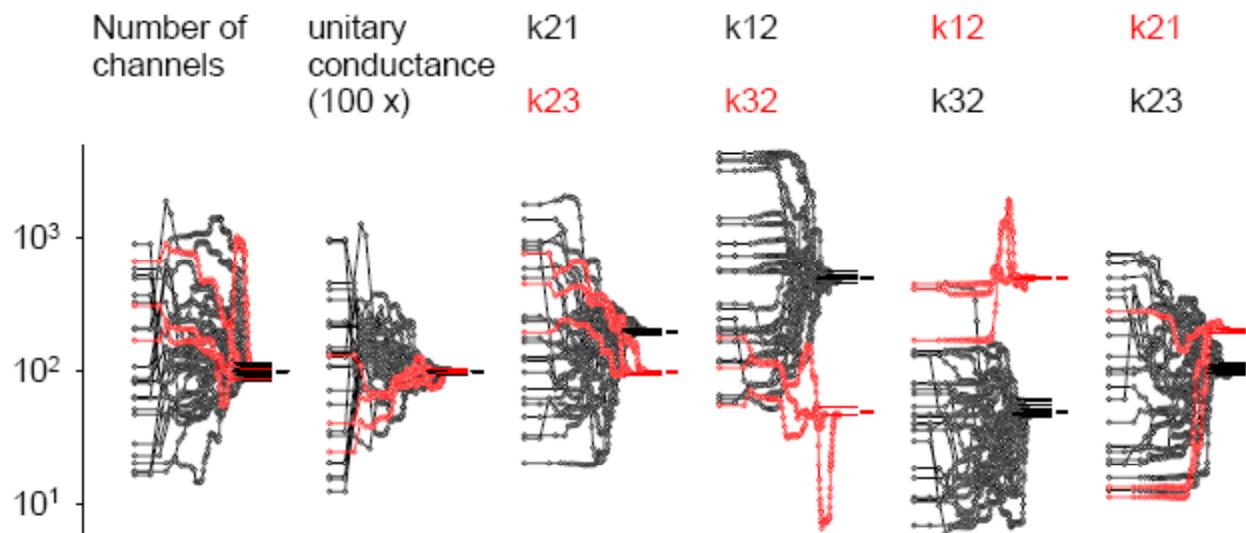


Number of evaluations of the Likelihood function

Stationary conditions

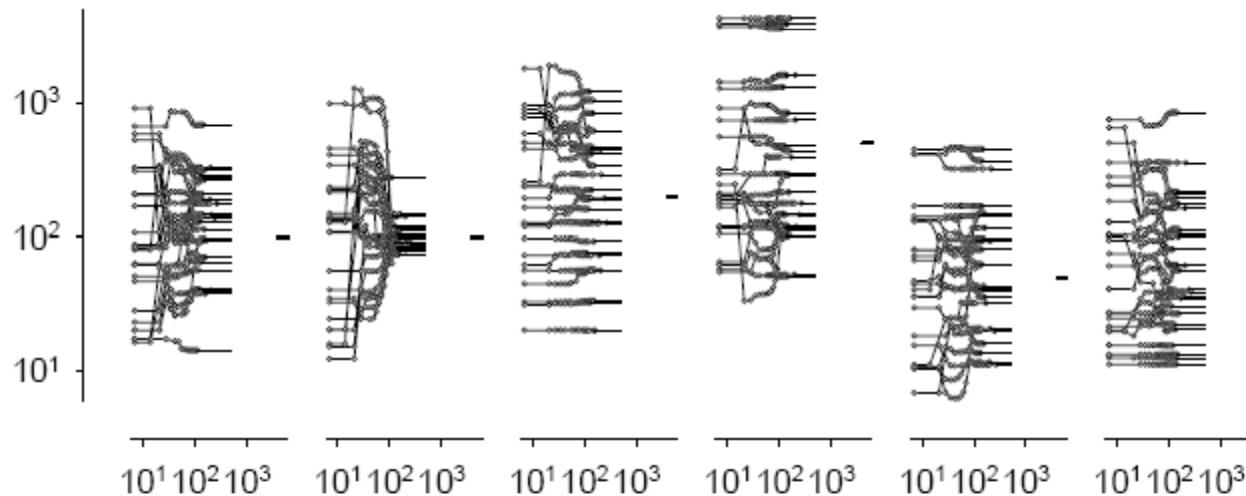
A

Recursive Macroscopic



B

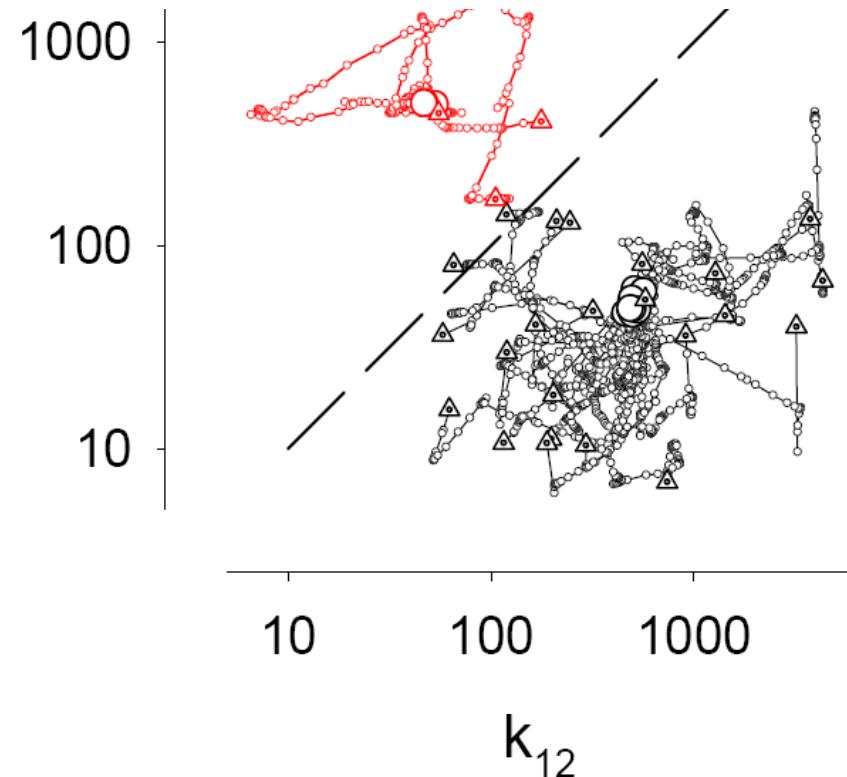
Non recursive Macroscopic

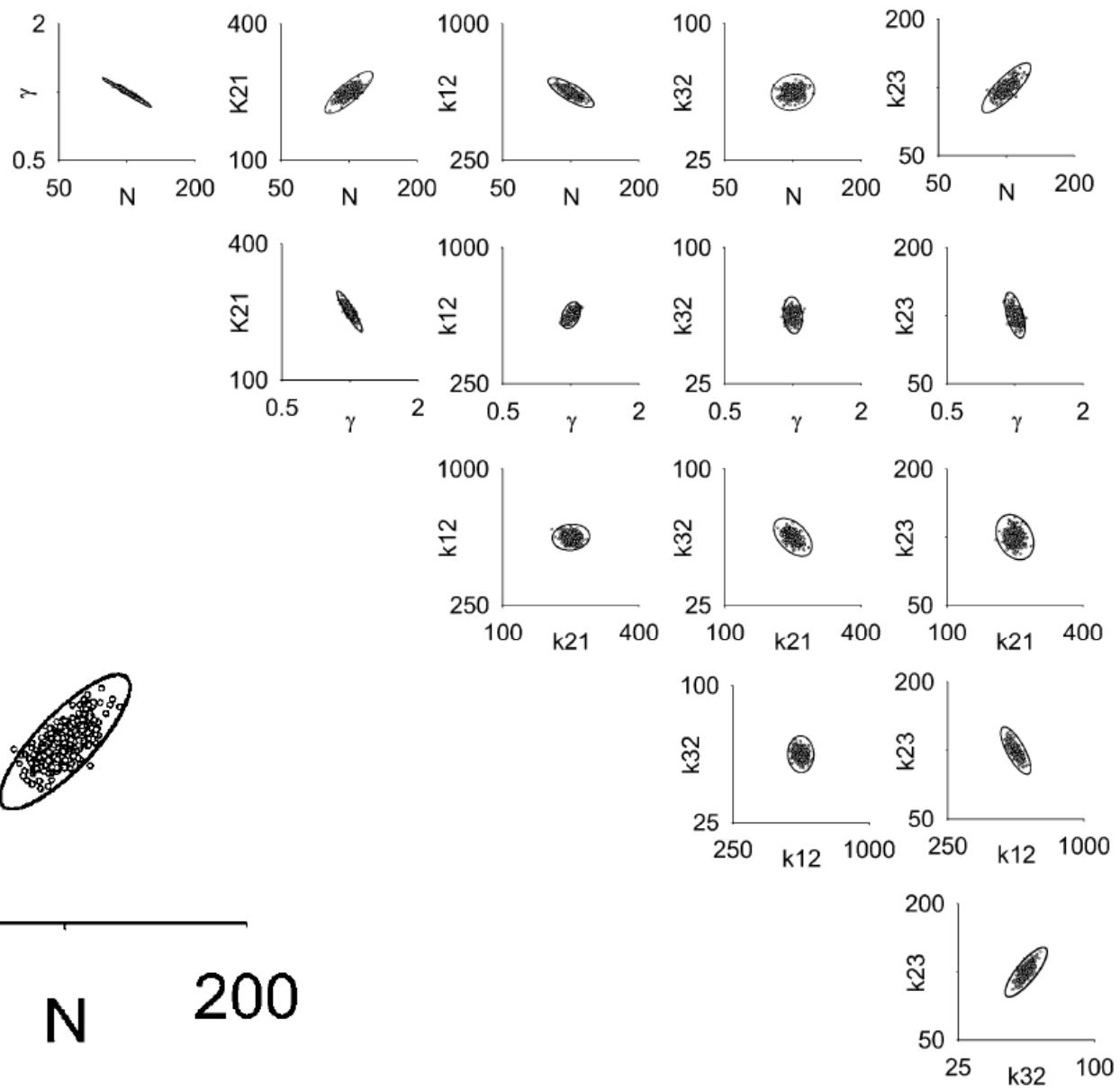
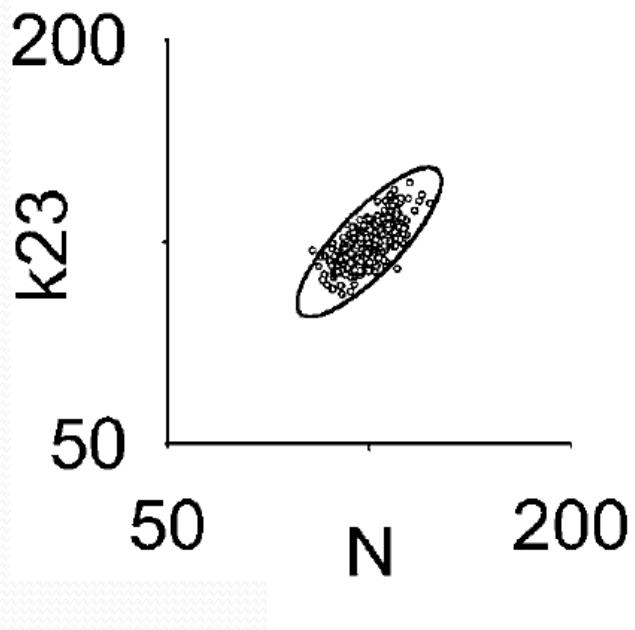


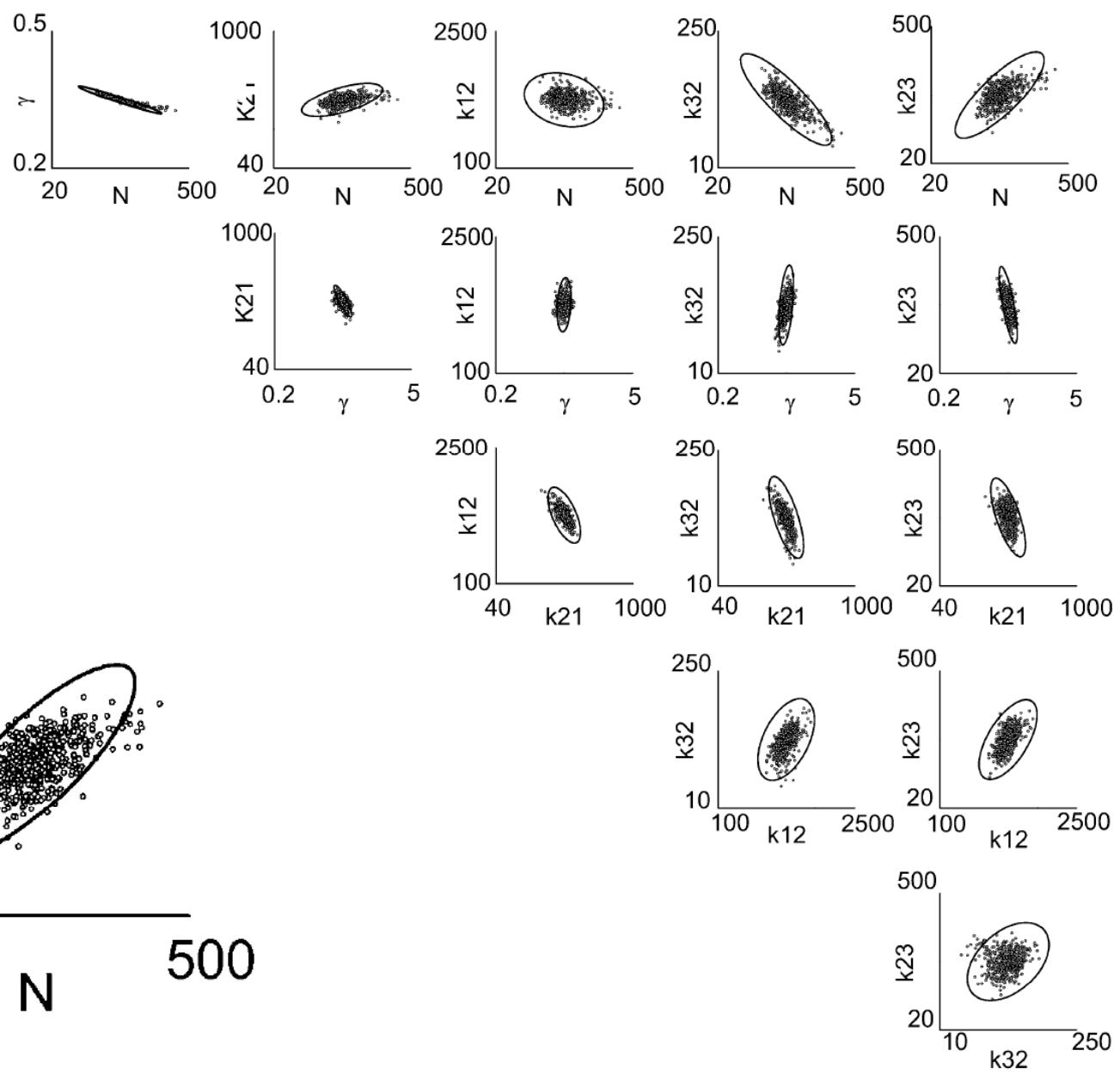
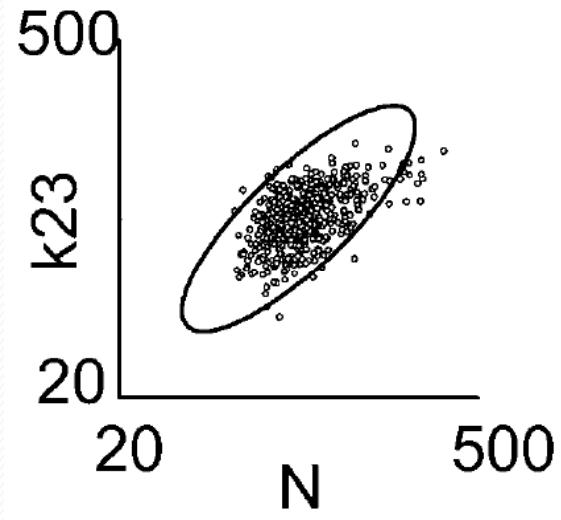
number of Likelihood function calls

When does the algorithm flipped?

When the starting value of k_{32} is larger than the starting value of k_{12} .

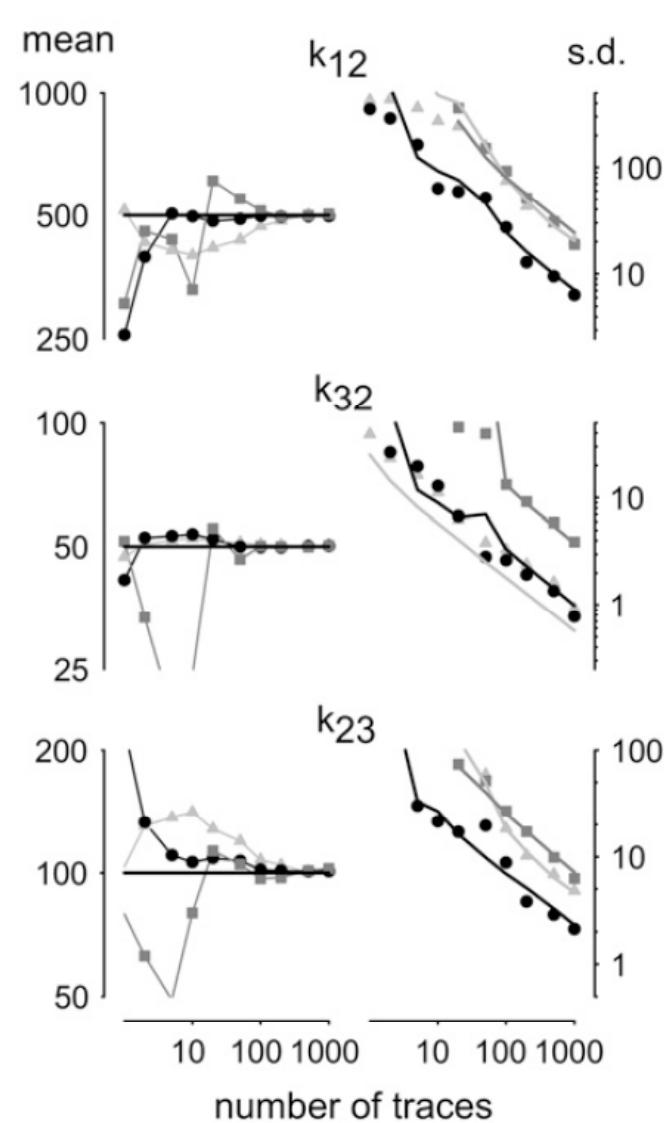
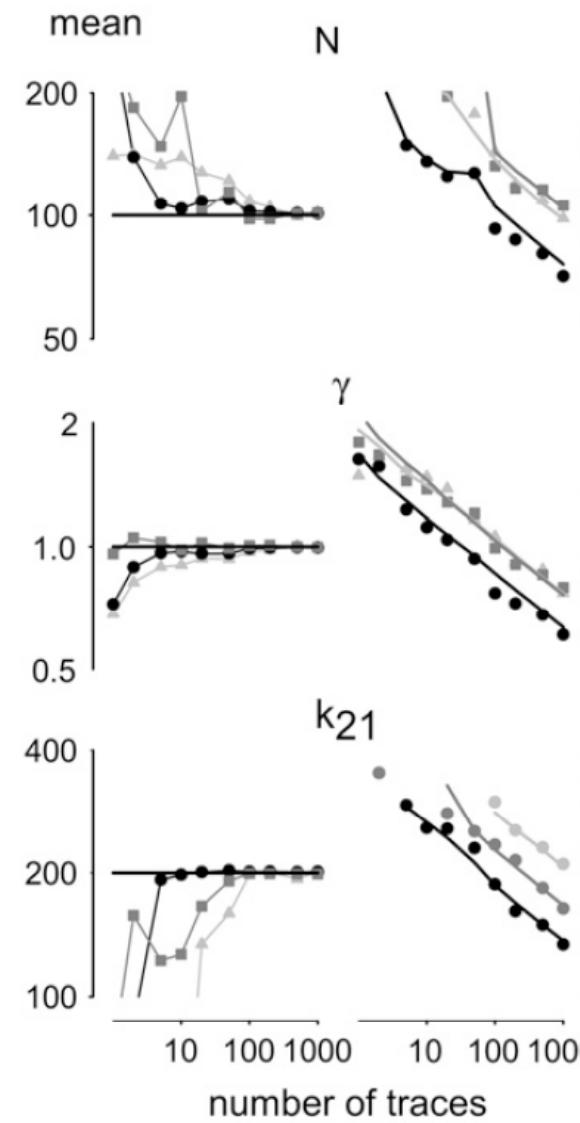






● Recursive, non-stationary
▲ Non-Recursive, non-Stationary,

■ Recursive, Stationary,



Present research

- MacroR-TA, time averaged version of MacroR
- Developing a GPL library than implements MacroR-TA in C++
- Working on electrophysiology experiments with pulses of 20 microseconds.