Maximum Likelihood Inference of Neuronal Dynamics under Noisy and Intermittent Observations using Sequential Monte Carlo EM Algorithms

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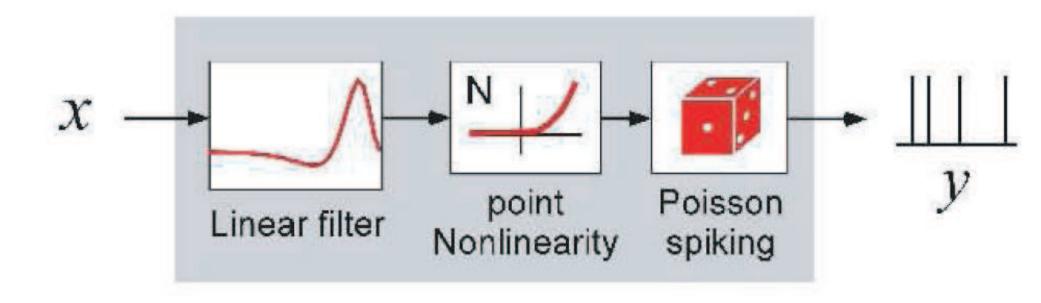
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Motivation

Although technology continues to progress improving the information experimentalists acquire about populations of neurons, the data remains noisy. Because the distribution of noise is, in general, non-Gaussian; and biophysically realistic neural models are nonlinear, sophisticated nonlinear systems identification tools must be developed to infer both (i) the unobserved/hidden states and (ii) the parameters governing the dynamics. By writing models of neurons in terms of observation states (eg, photon or spike counts), and hidden states (eg, voltage or conductance), an Expectation–Maximization (EM) algorithm is a natural framework. By writing the dynamics as Markov Chains, the EM approach is greatly simplified. Because the states are continuous with nonlinear dynamics, exact expectations are not feasible, and we therefore approximate them using Sequential Monte Carlo (SMC) algorithms. This approach is very general and can be applied to a variety of models with increasing complexity.

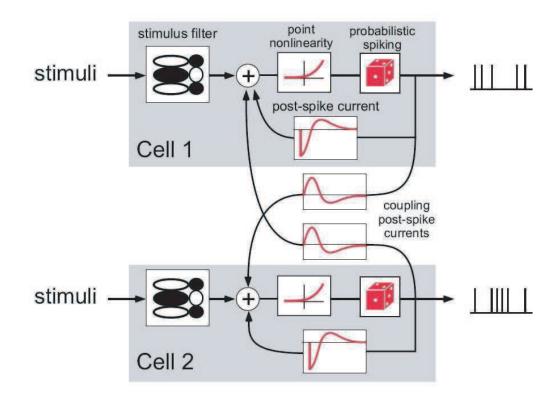
Generalized Linear Models (GLM)

- Cascade Linear-Nonlinear-Poisson (LNP) Models
- Unique global extrema
- Great success in spike train data



Generalizations

- Spike history terms for arbitrary dynamics (e.g., facilitation)
- Spike history terms from other neurons for population effects
- Noisy and intermittent observations
- Maintains properties of GLM (e.g., log concave, etc.)
- Populations of neurons are Hidden Markov Models



Hidden Markov Models (HMM)

Markov Property

$$P[H_t, H_0, \dots, H_{t-1}] = P[H_t | H_{t-1}] \tag{1}$$

Hidden Markov Model

$$P[H_{0:T}, O_{0:T}] = P[O_{0:T}|H_{0:T}]P[H_{0:T}]$$
(2)

$$= P[H_0] \prod_{t=1}^{T} P[O_t|H_t] P[H_t|H_{t-1}]$$
 (3)

Expectation–Maximization (EM)

Expectation Step: Write the expected value of the *joint* likelihood, $Q(\theta, \theta')$

Maximization Step: Maximize $Q(\theta, \theta')$

$$\widehat{\theta} = \max_{\theta} Q(\theta, \theta') \tag{4}$$

$$= \max_{\theta} E_{P(H_{0:T}|O_{0:T})}[\log P_{\theta}(O_{0:T}, H_{0:T})]$$
(5)

$$= \max_{\theta} \sum_{H_0} \cdots \sum_{H_T} \log P_{\theta}(O_{0:T}, H_{0:T}) P_{\theta'}(H_{0:T}|O_{0:T})$$
 (6)

EM for HMM

- Joint likelihood greatly simplifies
- Monotonically increasing
- Recursive algorithms available

$$\widehat{\boldsymbol{\theta}} = \max_{\boldsymbol{\theta}} \sum_{H_0} P_{\boldsymbol{\theta}'} [H_0 \mid O_{0:T}] \times \ln P_{\boldsymbol{\theta}} [H_0]$$

$$+ \sum_{t=1}^{T} \sum_{H_t} \sum_{H_{t-1}} P_{\boldsymbol{\theta}'} [H_t, H_{t-1} \mid O_{0:T}] \times \ln P_{\boldsymbol{\theta}} [H_t \mid H_{t-1}]$$

$$+ \sum_{t=0}^{T} \sum_{H_t} P_{\boldsymbol{\theta}'} [H_t \mid O_{0:T}] \times \ln P_{\boldsymbol{\theta}} [O_t \mid H_t]$$
 (7)

Sequential Monte Carlo (SMC)

- Efficiently (sometimes optimally) sample from HMM
- Many samples approximate the distribution
- SMC–EM Algorithm

$$\widehat{\boldsymbol{\theta}} = \max_{\boldsymbol{\theta}} \sum_{i_0} P_{\boldsymbol{\theta'}} \left[H^{(i_0)} \mid O^{(0:T)} \right] \times \ln P_{\boldsymbol{\theta}} \left[H^{(i_0)} \right]$$

$$+\sum_{t=1}^{T}\sum_{i_{t}}\sum_{j_{t-1}}P_{\boldsymbol{\theta'}}\left[H^{(i_{t})},H^{(j_{t-1})}\mid O^{(0:T)}\right]\times \ln P_{\boldsymbol{\theta}}\left[H^{(i_{t})}\mid H^{(j_{t-1})}\right]$$

$$+ \sum_{t=0}^{T} \sum_{i_{t}} P_{\theta'} \left[H^{(i_{t})} \mid O^{(0:T)} \right] \times \ln P_{\theta} \left[O^{(t)} \mid H^{(i_{t})} \right]$$
 (8)

Forward-Backward Approach

Forward Sampling

$$H_t \sim P[H_t | H_{t-1}, O_{0:t}]$$
 (9)

Backwards Recursion

$$P[H_t, H_{t-1}|O_{0:T}] = P[H_t|H_{t-1}]P[H_{t-1}|O_{0:t-1}] \frac{P[H_t|H_{t-1}]P[H_{t-1}|O_{0:t-1}]}{\int dH_{t-1}P[H_t|H_{t-1}]P[H_{t-1}|O_{0:t-1}]}$$
(10)

Linear-Nonlinear-Poisson Calcium Model

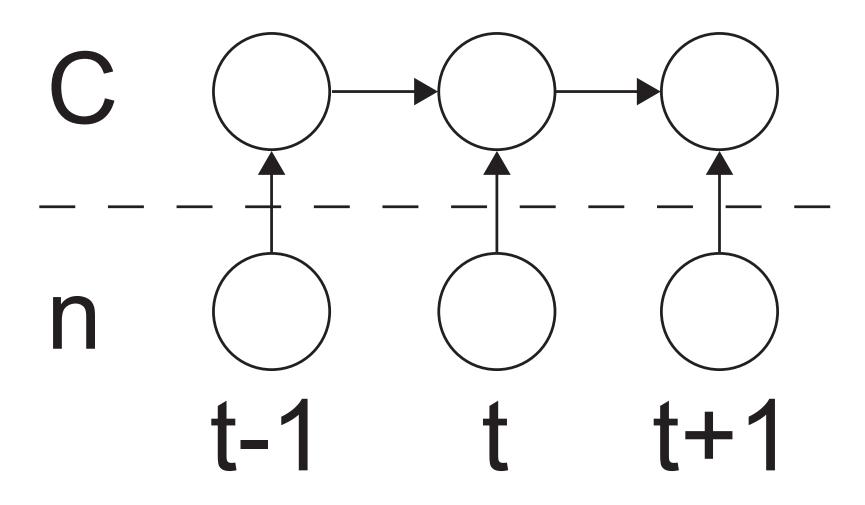
LNP-Ca: Model Specification

$$\lambda(t) = f(b + \mathbf{k}'\mathbf{x}(t)) \tag{11}$$

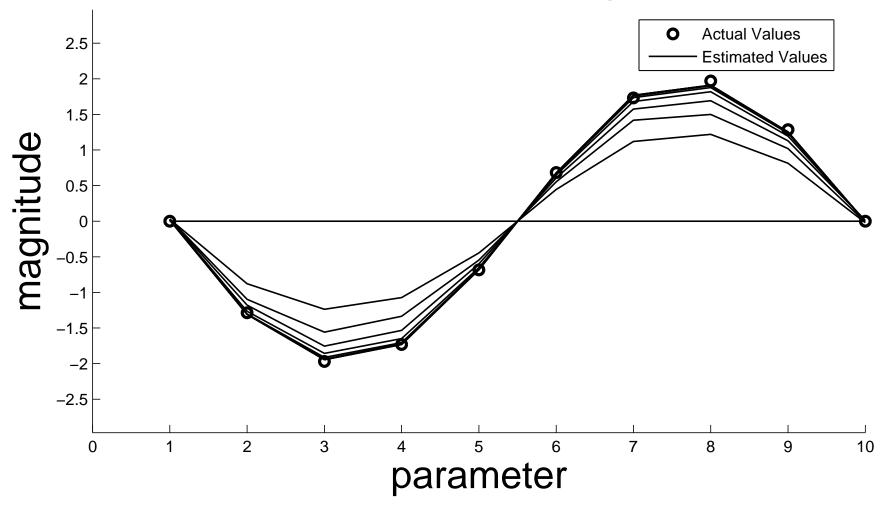
$$n(t) \sim \mathcal{P}[n(t); \lambda(t)dt]$$
 (12)

$$dC(t) = -\alpha C(t)dt + \sigma dB_t + \beta n(t) \tag{13}$$

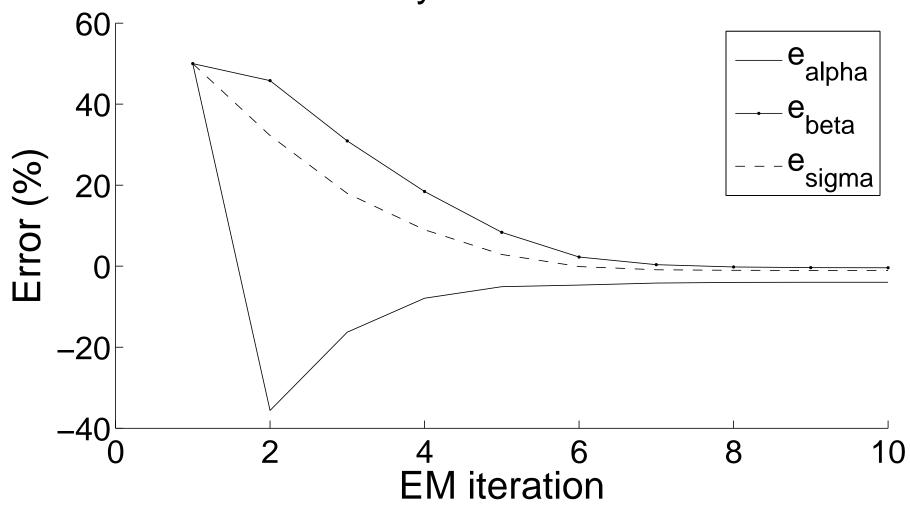
LNP-Ca: Graphical Model

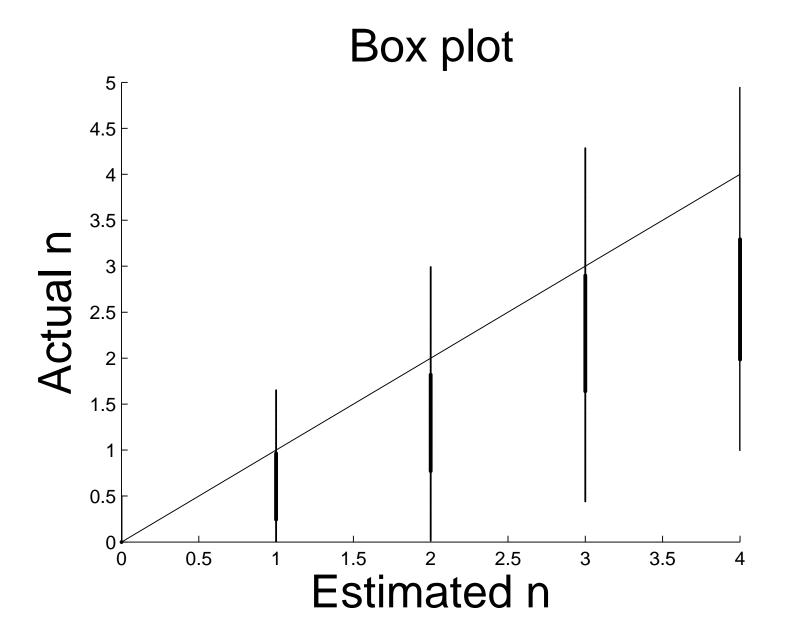


Linear Filter Estimates for Subsequent EM Iterations



Calcium Dynamics Parameters





Linear-Nonlinear-Poisson Intermittent Calcium Observations Model

LNP-ICa: Model Specification

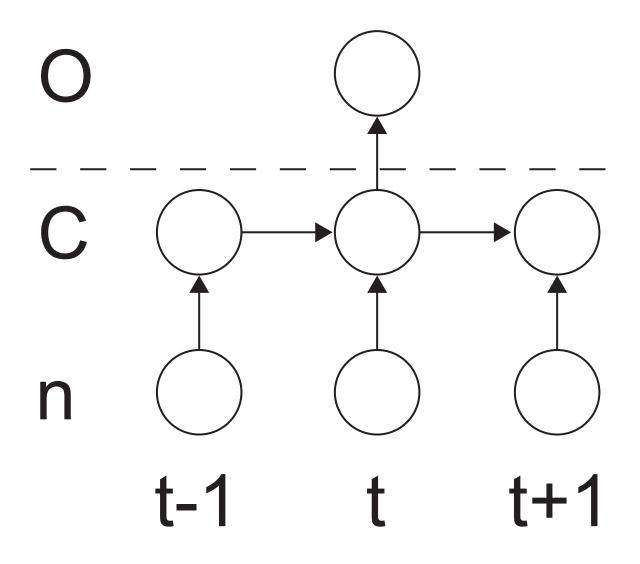
$$\lambda(t) = f(b + \mathbf{k}'\mathbf{x}(t)) \tag{14}$$

$$n(t) \sim \mathcal{P}[n(t); \lambda(t)dt]$$
 (15)

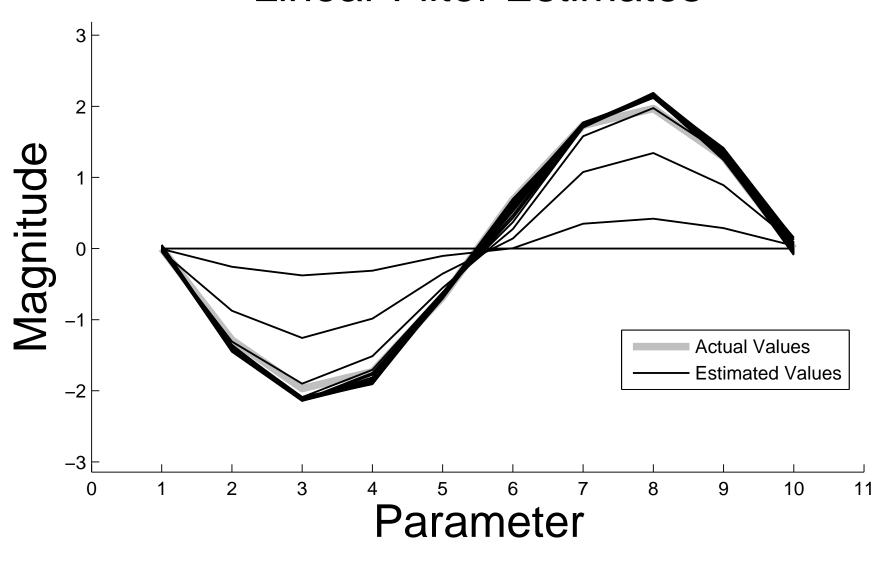
$$dC(t) = -\alpha C(t)dt + \sigma dB_t + \beta n(t) \tag{16}$$

$$O(t) = \eta C(t)$$
 whenever $t/r \in \mathcal{N}$ (17)

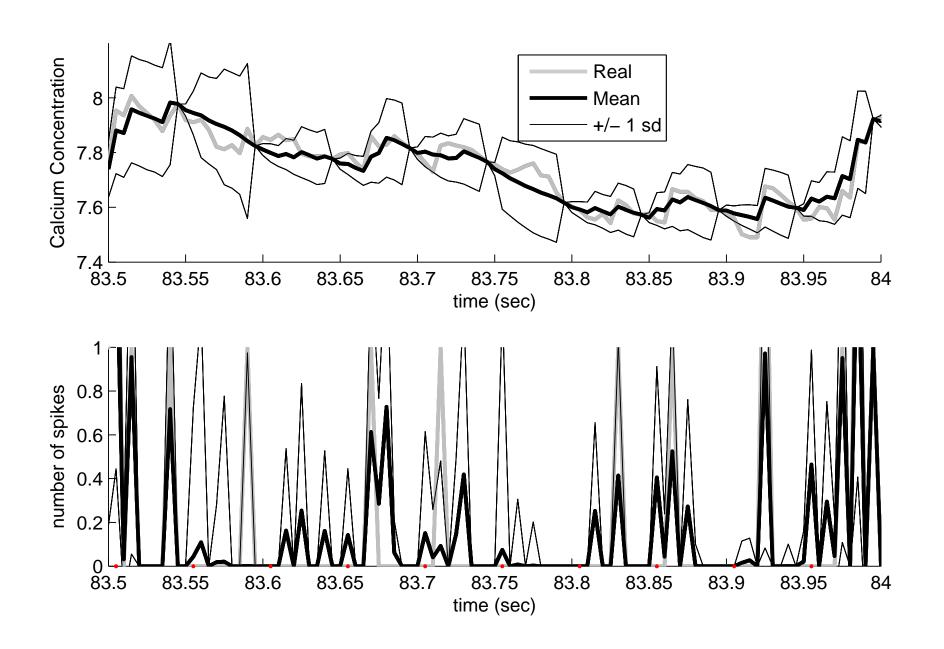
LNP-ICa: Graphical Model



Linear Filter Estimates



LNP-ICa: Density Estimation



Linear-Nonlinear-Poisson Noisy Intermittent Calcium Observations Model

LNP-NICa: Model Specification

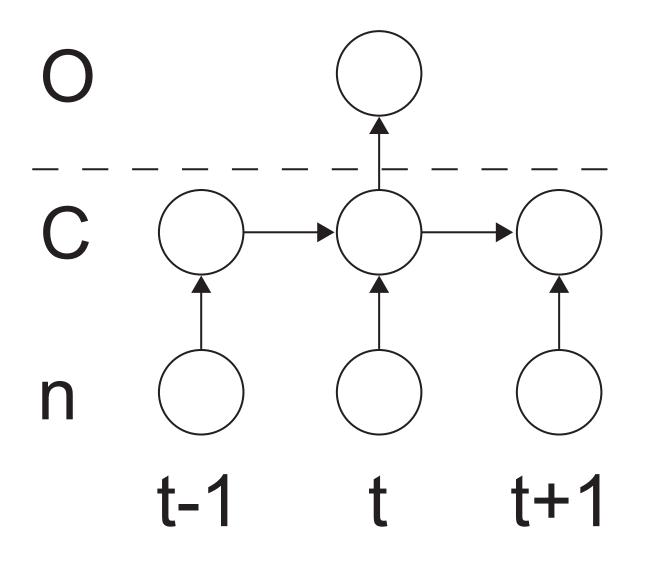
$$\lambda(t) = f(b + \mathbf{k}'\mathbf{x}(t)) \tag{18}$$

$$n(t) \sim \mathcal{P}[n(t); \lambda(t)dt]$$
 (19)

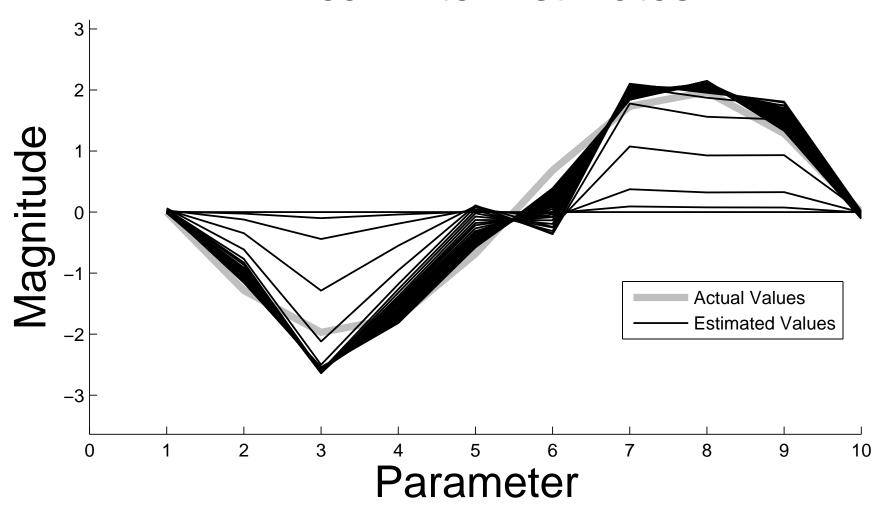
$$dC(t) = -\alpha C(t)dt + \sigma dB_t + \beta n(t)$$
(20)

$$O(t) = \eta C(t) + \varepsilon B_t$$
 whenever $t/r \in \mathcal{N}$ (21)

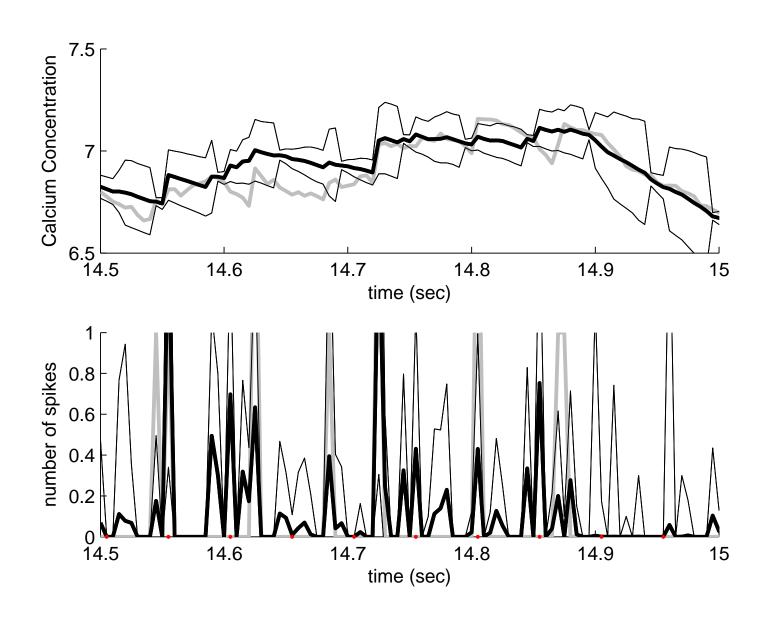
LNP-NICa: Graphical Model



Linear Filter Estimates



LNP-NICa: Density Estimation



Linear-Nonlinear-Poisson with Spike-History and Noisy-Intermittent-Calcium Model

LNP-SH-NICa: Model Specification

$$\lambda(t) = f\left(b + \mathbf{k}'\mathbf{x}(t) + \sum_{l} \omega_{l} h_{l}(t)\right)$$
(22)

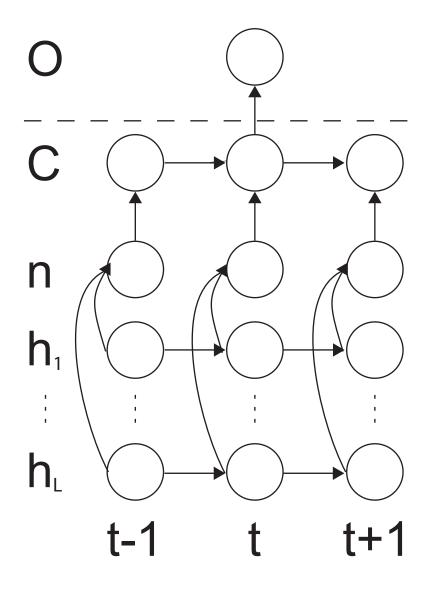
$$dh_l(t) = -\gamma_l h_l(t) + \sigma_h dB_{h,t} + n(t)$$
(23)

$$n(t) \sim \mathcal{P}[n(t); \lambda(t)dt]$$
 (24)

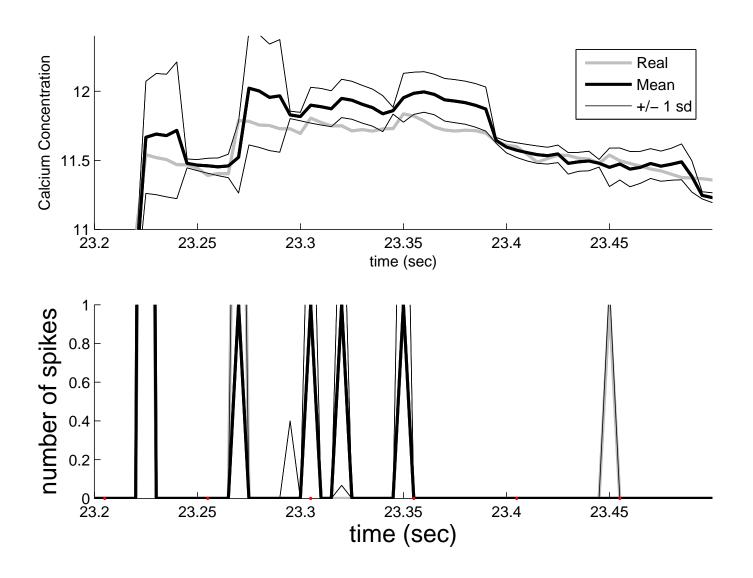
$$dC(t) = -\alpha C(t)dt + \sigma_C dB_{C,t} + \beta n(t)$$
(25)

$$O(t) = \eta C(t) + \varepsilon B_t$$
 whenever $t/r \in \mathcal{N}$ (26)

LNP-SH-NICa: Graphical Model



LNP-SH-NICa: Density Estimation



Discussion

- SMC–EM algorithms efficiently infer hidden states and parameters
- Conceptually straightforward generalization to populations of neurons
- Spike dropping scenario can be dealt with similarly
- Synaptic plasticity easily incorporated

References and Acknowledgements

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