## Approximate inference

## for continuous time Markov processes

Manfred Opper

TU Berlin, Dept of Computer Science

Collaborators:

- Andreas Ruttor & Florian Stimberg (TU Berlin)
- Cédric Archambeau & John Shawe–Taylor (UCL)
- Dan Cornford, Yuan Shen & Michail Vrettas (Aston)
- Guido Sanguinetti & Andrea Ocone (Edinburgh)

### Ito stochastic differential equations

for state  $X_t \in \mathbb{R}^d$ 

$$dX_t = \underbrace{f(X_t)}_{\text{Drift}} dt + \underbrace{\Sigma^{1/2}(X_t)}_{\text{Diffusion}} \times \underbrace{dW_t}_{\text{Wiener process}}$$

Limit of discrete time process  $X_k$ 

$$\Delta X_k \equiv X_{k+1} - X_k = f(X_k) \Delta t + \Sigma^{1/2}(X_k) \sqrt{\Delta t} \epsilon_k .$$

 $\epsilon_k$  i.i.d. Gaussian.

## Overview

- Inference for stochastic dynamics
- Variational approximation in machine learning and physics
- Formulation for probabilities over paths
- Results for low dimensional models
- Hybrid models
- Nonparametric approach to drift estimation

## Motion in double-well potential

$$dX = X(\theta - X^2)dt + \sigma dW.$$





A sample path might look like this



#### **Jump Processes**

Assume that  $X_t$  jumps between discrete states.

Short time behaviour of transition kernel defined by transition rate f:

$$P_{t+\Delta t,t}(x'|x) \simeq f_{\theta}(y|x,t)\Delta t \text{ for } x' \neq x$$
$$P_{t+\Delta t,t}(x|x) \simeq 1 - \sum_{z \neq x} f_{\theta}(z|x,t)\Delta t$$

for  $\Delta t \rightarrow 0$ .



## Simple model of autoregulatory network

On molecular level kinetics is stochastic: Simple autoregulatory network:

2 interacting molecules: mRNA and a Protein

Number of mRNA and Protein molecules:  $X_{mRNA}$ ,  $X_{Prot}$ 

 $\mathbf{X} = (X_{\mathsf{mRNA}}, X_{\mathsf{Prot}})$ 

$$\begin{array}{rcl} X_{\rm Prot} \rightarrow X_{\rm Prot} + 1 & : \mbox{ with Rate } & \gamma X_{\rm mRNA} \,, \\ X_{\rm Prot} \rightarrow X_{\rm Prot} - 1 & : \mbox{ with Rate } & \delta X_{\rm Prot} \,, \\ X_{\rm mRNA} \rightarrow X_{\rm mRNA} - 1 & : \mbox{ with Rate } & \beta X_{\rm mRNA} \,, \\ X_{\rm mRNA} \rightarrow X_{\rm mRNA} + 1 & : \mbox{ with Rate } & \alpha (1 - \alpha_c \Theta (X_{\rm Prot} - \theta_c)) \,, \end{array}$$

where 
$$\Theta(x) = 1$$
 if  $x \ge 0$  and  $\Theta(x) = 0$  for  $x < 0$ .



Simulation of process



(Noisy) observations at discrete times.

#### **Inference Problems**

Given noisy observations  $\{y_i\}_{i=1}^N \equiv y_1, \ldots, y_N$  of hidden process  $X_{t_i}$  at times  $t_i \leq T$  for  $i = 1, \ldots, N$ .

- Estimate  $X_t$  for  $0 \le t \le T$  (smoothing).
- Estimate system parameters  $\theta$  contained in drift f and diffusion  $\Sigma$ .

## **Obvious ? questions**

- Can't we treat this just as a **discrete time** HMM ? Yes, but ....
- Isn't there some simple **forward backward** algorithm ? Yes, but ....
- Can't you just discretize in time and run an MCMC sampler ? Yes, but ....

## What we would like to do

#### • State estimation:

Use **Bayes rule** for conditional distribution over **paths**  $X_{0:T}$  ( $\infty$  dimensional objects)

$$p(X_{0:T}|\{y_i\}_{i=1}^N, \theta) = \frac{p_{prior}(X_{0:T}|\theta)}{p(\{y_i\}_{i=1}^N|\theta)} \prod_{n=1}^N p(y_n|X_{t_n})$$

to compute state prediction  $E[X_t|\{y_i\}_{i=1}^N, \theta]$ 

- Parameter estimation:
  - 1. Maximum Likelihood: Maximise  $p(\{y_i\}_{i=1}^N | \theta)$  with respect to  $\theta$
  - 2. Bayes: Use a prior  $p(\theta)$  to compute  $p(\theta|\{y_i\}_{i=1}^N) \propto p(\{y_i\}_{i=1}^N|\theta)p(\theta)$

#### Path integral representation

of the parameter likelihood (assuming additive noise)

$$p(\{y_i\}_{i=1}^N | \theta) = \int_0^T \left\{ \frac{1}{2\sigma_\theta^2} \left( \frac{dX_t}{dt} - f(X_t) \right)^2 - \sum_n \delta(t - t_n) \ln p(y_n | X_t) \right\} dt \right]$$

λT

with **Onsager–Machlup** type action. One needs to be a bit careful about the correct interpretation of  $\frac{dX_t}{dt}$  and integrals.

#### The variational approximation in statistical physics

(Feynman, Peierls, Bogolubov, Kleinert...)

Let 
$$p(x) = \frac{1}{Z} e^{-H(x)}$$
 and  $q(x) = \frac{1}{Z_0} e^{-H_0(x)}$ 

• The variational bound on the free energy is

$$-\ln Z \leq -\ln Z_0 + \langle H(x) \rangle_0 - \langle H_0(x) \rangle_0 \equiv \mathcal{F}[q]$$

 $\mathcal{F}[q]$  is the variational free energy.

• Approximation for free energies often better than the quality of  $H_0$  suggests.

- Choices for  $H_0$ 
  - 1. Gaussian approximations for path integrals (e.g. Polaron problem)
  - 2. Mean field approximations (factorising distributions)
- Look for a formulation that can easily be applied to a variety of systems without bothering too much about details of path integral formulations.

## The variational approximation (reformulation)

• We would like to approximate intractable distribution

$$p(x|y) = \frac{p(y|x)p_{prior}(x)}{p(y)}$$

by a q(x) which belongs to a family of **simpler tractable** distributions (e.g. factorising = mean field, or Gaussian densities).

• The variational free energy is

$$\mathcal{F}(q) = D[q||p(\cdot|y)] - \ln p(y)$$
  
=  $D[q||p_{prior}] - \int q(x) \ln p(y|x) dx$   
 $\geq -\ln p(y)$ 

• The relative entropy (Kullback–Leibler divergence) is

$$D[q||p] = \int q(x) \ln \frac{q(x)}{p(x|y)} dx$$

## Approximate maximum likelihood estimate

Assume model depends on parameter  $\theta$ . The free energy inherits the dependency.

Let  $q^*(\theta) = \operatorname{argmin}_q \mathcal{F}_{\theta}(q)$ . Since  $-\ln p(y|\theta) \le \mathcal{F}_{\theta}(q^*(\theta))$ 

we can minimise  $\mathcal{F}_{\theta}(q^*)$  wrt  $\theta$  to get an approximate maximum likelihood estimate.

# How to choose the measure q for stochastic differential equations ?

- Process conditioned on data is Markovian!
- It fulfils SDE

$$dX_t = g(X_t, t)dt + \Sigma^{1/2}(X_t) dW_t$$

with a new time dependent drift  $g(X_t, t)$  but the same diffusion  $\Sigma$ .

#### Example

Wiener process with single, noise free observation y = x(t = T) = 0



Posterior drift  $g(x,t) = -\frac{x}{T-t}$  for 0 < t < T.

#### **KL** divergence for path probabilities

Use representation of joint density in term of conditionals and the Markov property (assuming  $q_0(x) = p_0(x)$ ) and work with time discretization  $t_{k+1} - t_k = \Delta t$ .

$$D[q||p] = \int dx_{0:T} q(x_{0:T}) \ln \frac{q(x_{0:T})}{p(x_{0:T})}$$
  

$$\approx \sum_{k=0}^{K-1} \int dx q_{t_k}(x) \int dx' q_{t_{k+1},t_k}(x'|x) \ln \frac{q_{t_{k+1},t_k}(x'|x)}{p_{t_{k+1},t_k}(x'|x)}$$
  

$$= \sum_{k=0}^{K-1} \int dx q_{t_k}(x) D[q_{t_{k+1},t_k}(\cdot|x)||p_{t_{k+1},t_k}(\cdot|x)]$$

in terms of transition and marginal probabilities.

#### We know that short time transition probability

is approximately Gaussian

$$p_{t+\Delta t,t}(x'|x) \propto \exp\left[-\frac{1}{2\Delta t} \left\|x'-x-f(x)\Delta t\right\|^2\right]$$

as  $\Delta t 
ightarrow 0$ ,

with the squared norm  $||F||^2 = F^{\top} \Sigma^{-1} F$ .

Then for small  $\Delta t$ 

$$D\left[q_{t_{k+1},t_k}(\cdot|x) \| p_{t_{k+1},t_k}(\cdot|x)\right] \approx \frac{1}{2} \| g(x,t) - f(x) \|^2 \Delta t$$

# The relative entropy for Stochastic Differential Equations

Let q and p be measures over paths for SDEs with drifts g(X,t) and f(X,t) with same diffusion  $\Sigma(X)$ . Then

$$D[q||p] = \frac{1}{2} \int_0^T dt \left\{ \int dx \ q_t(x) \ \|g(x,t) - f_\theta(x)\|^2 \right\}$$

 $q_t(x)$  is the marginal density of  $X_t$ .

#### Change of measure approach

$$D[Q||P] = E_Q \ln \frac{dQ}{dP}$$

Girsanov's change of measure theorem results in the following Radon-Nikodym derivative:

$$\frac{dQ}{dP} = \exp\left\{-\int_0^T (f-g)^\top \Sigma^{-1/2} \, dB_t + \frac{1}{2} \int_0^T \|f-g\|_{\Sigma}^2 \, dt\right\}$$

where B is a Wiener process with respect to Q.

#### The variational problem (Diffusion)

Minimise variational free energy

$$\mathcal{F}_{\theta}(q) = \frac{1}{2} \int_0^T \int q_t(x) \{ \|g(x,t) - f_{\theta}(x)\|^2 - \sum_i \delta(t-t_i) \ln p(y_i|x) \} dx dt$$

with respect to the posterior drift g(x,t).

The marginal density  $q_t$  and the drift g(x,t) are coupled through the **Fokker - Planck** equation

$$\frac{\partial q_t(x)}{\partial t} = \left\{ -\sum_k \partial_k g_k(x) + \frac{1}{2} \sum_{kl} \partial_k \partial_l \Sigma_{kl}(x) \right\} q_t(x)$$

Variation leads to forward backward PDEs.

## The Variational Gaussian Approximation for SDEs

 Approximate (Gaussian) process over paths X<sub>0:T</sub> induced by linear SDE:

$$dX_t = \{A(t)X_t + b(t)\} dt + \Sigma^{1/2} dW$$

- Diffusion  $\Sigma$  must be independent of X !
- Relative entropy is of the form  $\mathcal{F}_{\theta}[m, S, A, b]$ .
- Constraints are evolution eqs. for marginal mean m(t) and covariance S(t)

$$\frac{dm}{dt} = Am + b$$
$$\frac{dS}{dt} = AS + SA^{\top} + \Sigma.$$

 $\rightarrow$  nonlinear ODEs instead of PDEs !

## Example: Motion in double-well potential

$$dX = X(\theta - X^2)dt + \sigma dW.$$





A trajectory

#### Prediction & comparison with hybrid Monte Carlo

 $T = 20, \ \theta = 1, \ \sigma^2 = 0.8$  with N = 40 observations with noise  $\sigma_o^2 = 0.04$ . Fixed initial conditions.





## Posterior for $\sigma$



## Lorenz 1963

$$dx_t = \sigma(y_t - x_t)dt + \sqrt{\Sigma^x}dW^x$$
  

$$dy_t = (\rho x_t - y_t - x_t z_t)dt + \sqrt{\Sigma_y}dW^y$$
  

$$dz_t = (x_t y_t - \beta z_t)dt + \sqrt{\Sigma_z}dW^z$$









#### More dimensions: Mean field approximation

Approximate further by assuming that processes for different dimensions are independent.

Covariance  $S(t) \rightarrow \text{Diag}(s_1(t), \ldots, s_D(t))$ 

$$\mathcal{F}_{\theta}(q) = \sum_{i=1}^{D} \frac{1}{2\sigma_i^2} \int_0^T E_q \left[ (\dot{m}_i - f_i(X_t))^2 \right] dt + \sum_{i=1}^{D} \frac{1}{2\sigma_i^2} \int_0^T \left\{ \frac{(\dot{s}_i - \sigma_i^2)^2}{4s_i^2} + (\sigma_i^2 - \dot{s}_i) E_q \left[ \frac{\partial f_i(X_t)}{\partial X_t^i} \right] \right\} dt - \sum_{j=1}^{n} E_q \left[ \ln p(y_j | X_{t_j}) \right]$$

Lorenz 1998 model:

 $x=(x^1,\ldots,x^{40})$  with drift  $f_i(x_t)=\left(x_t^{i+1}-x_t^{i-2}\right)x_t^{i-1}-x_t^i+\theta$ 

 $\sigma^2 = 5$  and N = 90 observations.



## Likelihoods



#### The relative entropy for Markov jump processes

Assume transition rates g(x'|x,t) and f(x'|x,t)

$$KL[q||p] = \int_0^T dt \sum_x q_t(x) \sum_{x': x' \neq x} \left\{ g(x'|x,t) \ln \frac{g(x'|x,t)}{f(x'|x,t)} + f(x'|x,t) - g(x'|x,t) \right\}$$

#### Mean field approximation

Multivariate states  $X(t) = (X_1(t), \dots, X_d(t))$ 

**Exact inference**: Linear ODEs in  $S^d$  variables

Variational approximation: Optimise in family of factorising measures, i.e. of the type

 $q(X[0:T]) = \prod_{i=1}^{d} q_i(X_i[0:T])$ 

Linear ODEs in Sd variables.

(Sanguinetti & Opper, 2008, Cohn, El–Hay, Friedman & Kupferman, 2010)

#### Lotka Volterra

#### Comparison with MCMC



Figure 1: Posterior (mean and 90% confidence intervals) over predator paths (observations (circles) only until 1500).

from (V Rao & Y W Teh, 2011)



Figure 2: Posterior (mean and 90% confidence intervals) over prey paths (observations (circles) only until 1500).

# Hybrid models: Inference of transcriptional regulation



- Transcription factors regulate genes by binding to specific sites.
- Hard to measure transcription factor activity directly. Inference must be based on measurement of mRNA concentration of target genes.
- Big networks: Clustering of expression profiles or Factor analysis

## Small subnetworks:

• More detailed dynamical model (Barenco et al)

$$\frac{dx_i}{dt} = -\lambda_i x_i(t) + b_i + A_i \mu(t)$$

which takes sensitivity and degradation into account.

• Try predictions on TF activity  $\mu(t)$  and learn parameters using measurements of mRNA concentration of target genes:

$$y_{ik} = x_i(t_k) + \text{noise}$$

• Assume switching process  $\mu(t) \in \{0, 1\}$  and  $\mu \rightarrow 1 - \mu$  with rates  $f_{\pm}$  modeled by **telegraph process** (Sanguinetti, Ruttor, Archambeau, Opper 2009)

## Multiple (2) transcription factors (toy model)

$$\frac{dx_i}{dt} = -\lambda_i x_i(t) + A_1^i \mu_1(t) + A_2^i \mu_2(t) + A_{12}^i \mu_1(t) \mu_2(t) + b_i$$



Parameter inference for  $A_1^2$  ,  $A_2^2$  and  $A_{12}^2$ .





 $A_1^i$ ,  $A_2^i$  and  $A_{12}^i$  for 5 target genes.

Prediction of activity of transcription factors **FHL1** and **RAP1** (Microarray data from yeast metabolic cycle). Comparison to MCMC



(blue: MCMC, green: Variational upper, red: Var lower bound)

#### Feed-forward-loop

(A Ocone & G Sanginetti, 2011)



**Fig. 3.** p53 network architecture. E2F1 is the master TF, p53 is the target TF and both regulate target genes *DDB2*, *p21*, *BIK*, *PUMA*, *SIVA*, *DRAM*. Target genes of the only E2F1 (*MCM5*, *MCM7*, *LIG1*) have been included.

#### Nonparametric estimation of drift function



Assume that data are generated from  $dX_t = f(X_t)dt + \sigma dW_t$ . Could we directly predict f(x) ?

## Yes, if we use a Gaussian Process prior

distribution  $p(\mathbf{f})$  over functions  $f(\cdot)$ .



If we have 'continuous time' samples → posterior process is Gaussian (Papaspiliopulis et al, 2011).

$$p(\mathbf{f}|X_{0:T}) \propto p(\mathbf{f})L(X_{0:T}|\mathbf{f})$$

with the path 'likelihood'

$$L(X_{0:T}|\mathbf{f}) = \exp\left[-\frac{1}{2\sigma^2}\sum_{t}f^2(X_t)\Delta t + \frac{1}{\sigma^2}\sum_{t}f(X_t)\left(X_{t+\Delta t} - X_t\right)\right]$$

• For sparse samples use EM algorithm which cycles between approximate estimations of latent path  $X_{0:T}$  between observations and recomputing  $f(\cdot)$ .

















## Present & Future work

- Large systems: Simpler classes of approximations, eg. parametric forms for large covariance matrices (projections, low rank representations ?)
- Perturbative corrections (estimate for error)
- State dependent noise.
- Nonparametric estimation of drift f(x) for models with detailed balance
- Combination with optimal stochastic control

## **Publications**

Gaussian Process Approximations of Stochastic Differential Equations, Cédric Archambeau, Dan Cornford, Manfred Opper and John Shawe -Taylor, Journal of Machine Learning Research: Workshop and Conference, Proceedings, 1:1–16. (2007).

Variational Inference for Diffusion Processes, Cedric Archambeau, Manfred Opper, Yuan Shen, Dan Cornford and John Shawe-Taylor, Advances in Neural Information Processing Systems 20 (2008).

*Variational inference for Markov jump processes*, Manfred Opper and Guido Sanguinetti, Advances in Neural Information Processing Systems 20, 1105–1112 (2008).

A comparison of variational and Markov Chain Monte Carlo methods for inference in partially observed stochastic dynamic systems, Yuan Shen, Cédric Archambeau, Dan Cornford, Manfred Opper, John Shawe-Taylor and Remi Barillec. Journal of Signal Processing Systems (2009). *Switching Regulatory Models of Cellular Stress Response*, Guido Sanguinetti, Andreas Ruttor, Manfred Opper and Cédric Archambeau, Bioinformatics, doi: 10.1093/bioinformatics/btp138 (2009).