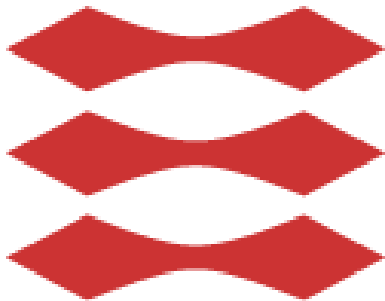


# Well-known shortcomings, advantages and computational challenges in Bayesian modelling: a few case stories

DTU



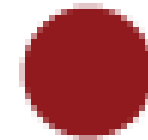
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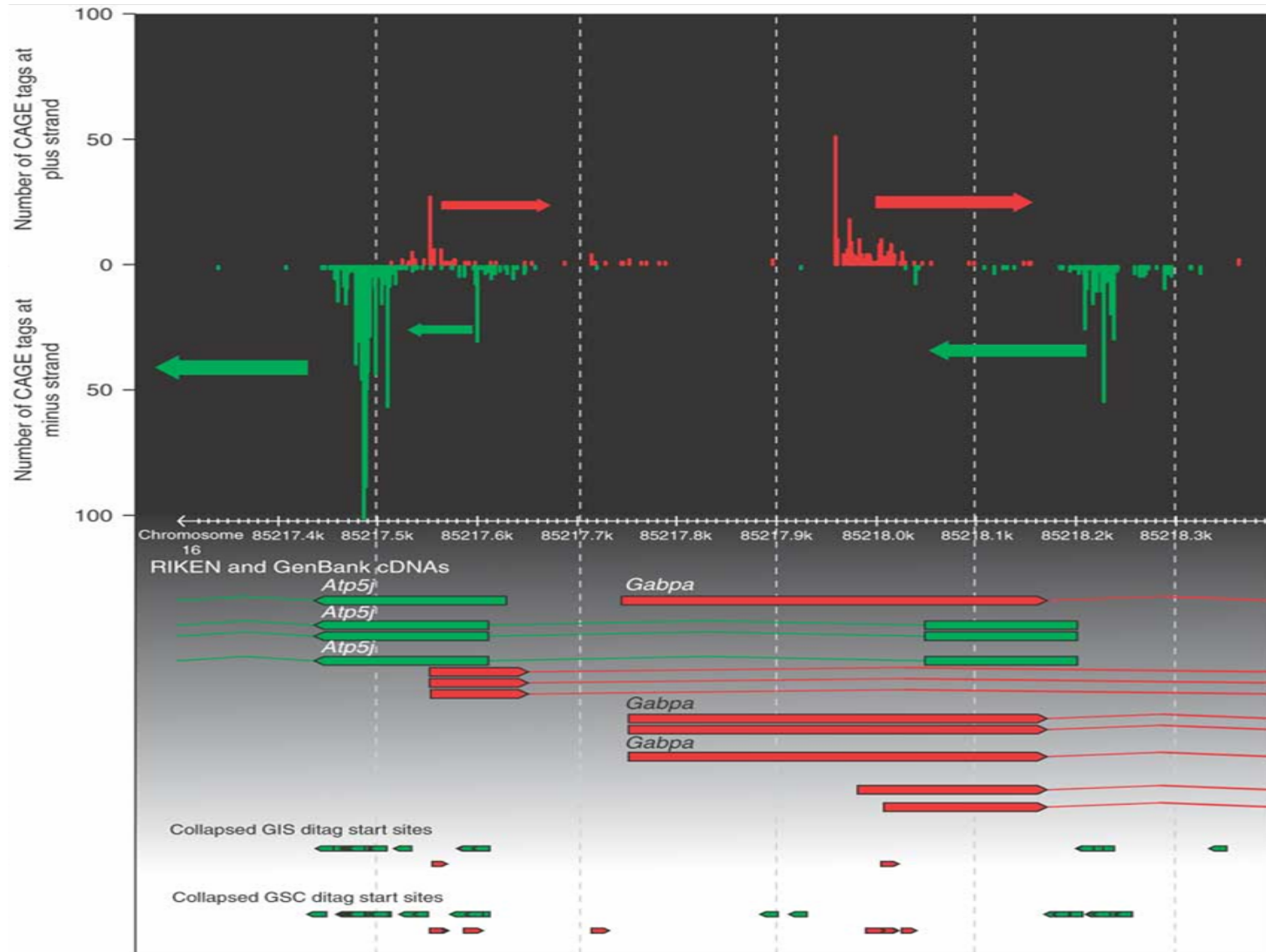
# Overview

1. How many species? predicting sequence tags.
  - Non-parametric Bayes
  - Averaging beats maximum likelihood
  - The model is always wrong (and Bayes can't tell)
2. Computing the marginal likelihood with MCMC
  - Motivation: computing corrections to EP/C
  - The trouble with Gibbs sampling
  - Gaussian process classification

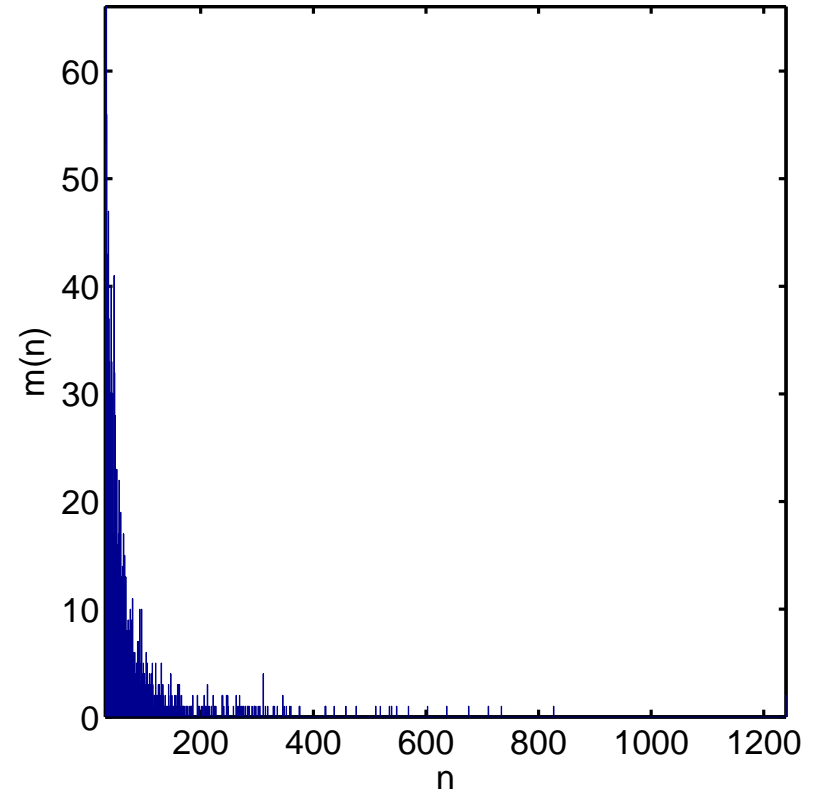
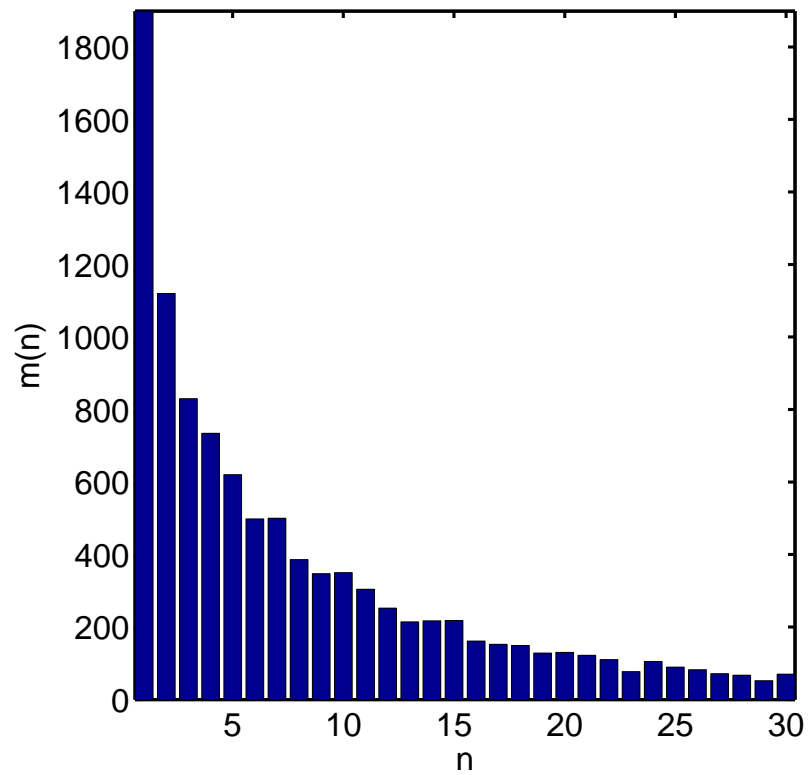
# How many species?



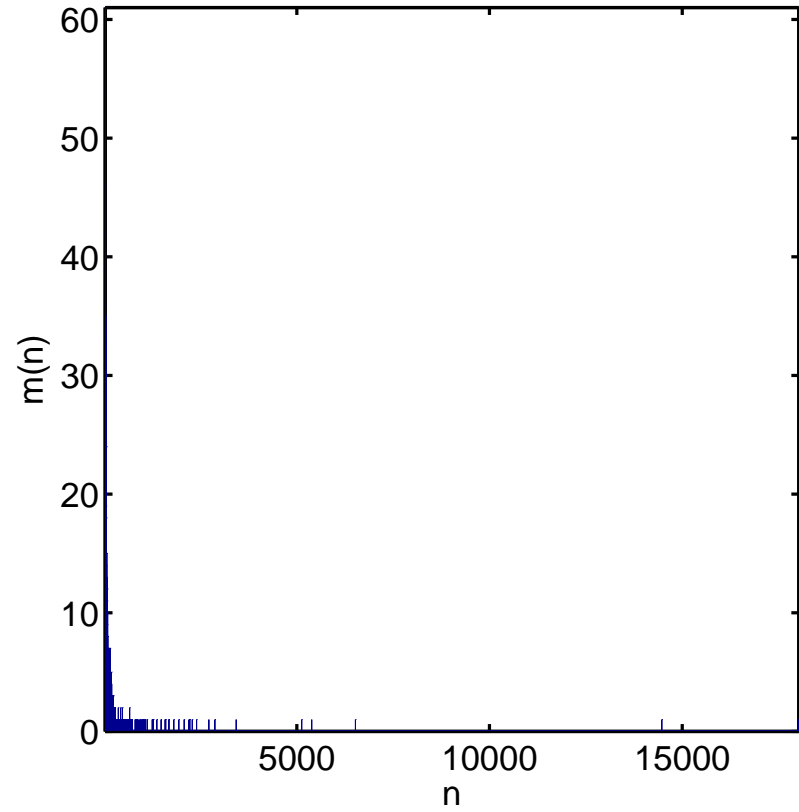
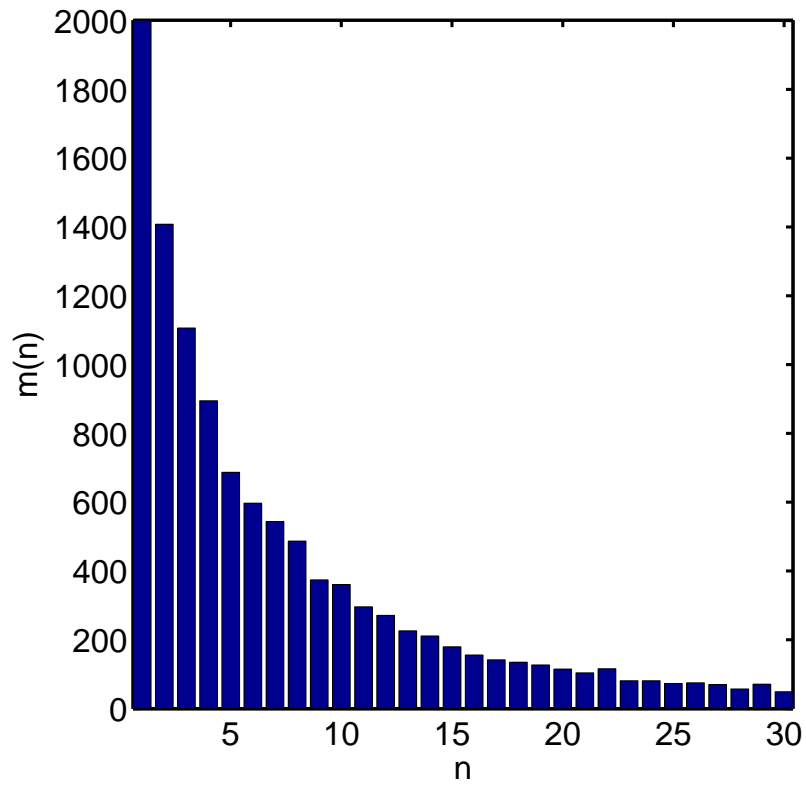
# DNA sequence tags - CAGE



# Look at the data - cerebellum



# Look at the data - embryo



# Chinese restaurant process - Yor-Pitman sampling formula

Observing new species given counts  $\mathbf{n} = n_1, \dots, n_k$  in  $k$  bins:

$$p(n_1, \dots, n_k, 1 | \mathbf{n}, \sigma, \theta) = \frac{\theta + k\sigma}{n + \theta} \quad \text{with} \quad \sum_{i=1}^k n_i = n$$

Re-observing  $j$ :

$$p(n_1, \dots, n_{j-1}, n_j + 1, n_{j+1}, \dots, n_k | \mathbf{n}, \sigma, \theta) = \frac{n_j - \sigma}{n + \theta}$$

Exchangeability – invariant to re-ordering

$$E, E, M, T, T : \quad p_1 = \frac{\theta}{\theta} \frac{1 - \sigma}{1 + \theta} \frac{\theta + \sigma}{2 + \theta} \frac{\theta + 2\sigma}{3 + \theta} \frac{1 - \sigma}{4 + \theta}$$

$$M, T, E, T, E : \quad p_2 = \frac{\theta}{\theta} \frac{\theta + \sigma}{1 + \theta} \frac{\theta + 2\sigma}{2 + \theta} \frac{1 - \sigma}{3 + \theta} \frac{1 - \sigma}{4 + \theta} = \dots = p_1$$

# Inference and prediction

Likelihood function, e.g.  $E, E, M, T, T$

$$\begin{aligned} p(\mathbf{n}|\sigma, \theta) &= \frac{\theta}{\theta} \frac{1-\sigma}{1+\theta} \frac{\theta+\sigma}{2+\theta} \frac{\theta+2\sigma}{3+\theta} \frac{1-\sigma}{4+\theta} \\ &= \frac{1}{\prod_{i=1}^{n-1} (i+\theta)} \prod_{j=1}^{k-1} (\theta+j\sigma) \prod_{i'=1}^k \prod_{j'=1}^{n_{i'}-1} (j'-\sigma) \end{aligned}$$

Flat prior distribution for  $\sigma \in [0, 1]$  and  $\theta$  pseudo-count parameter.

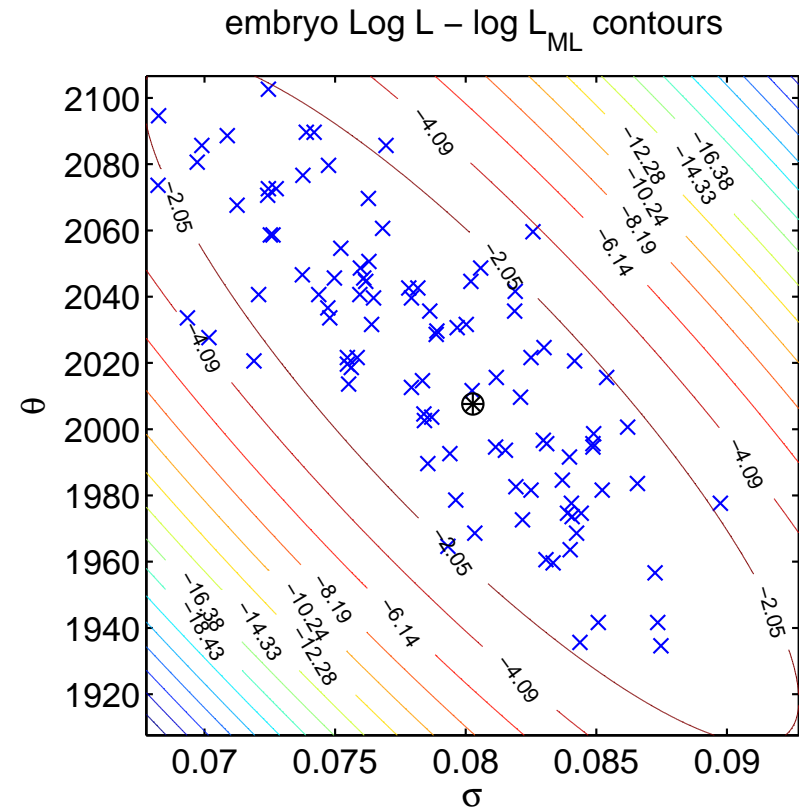
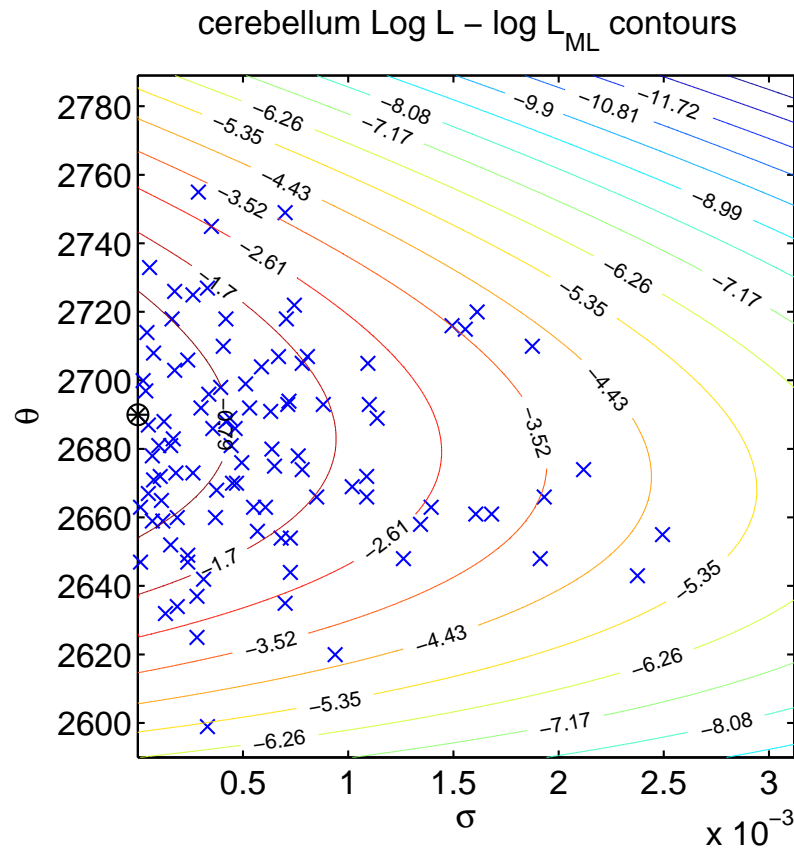
Predictions for new count  $\mathbf{m}$ :

$$p(\mathbf{m}|\mathbf{n}) = \int p(\mathbf{m}|\mathbf{n}, \sigma, \theta) p(\sigma, \theta) d\sigma d\theta$$

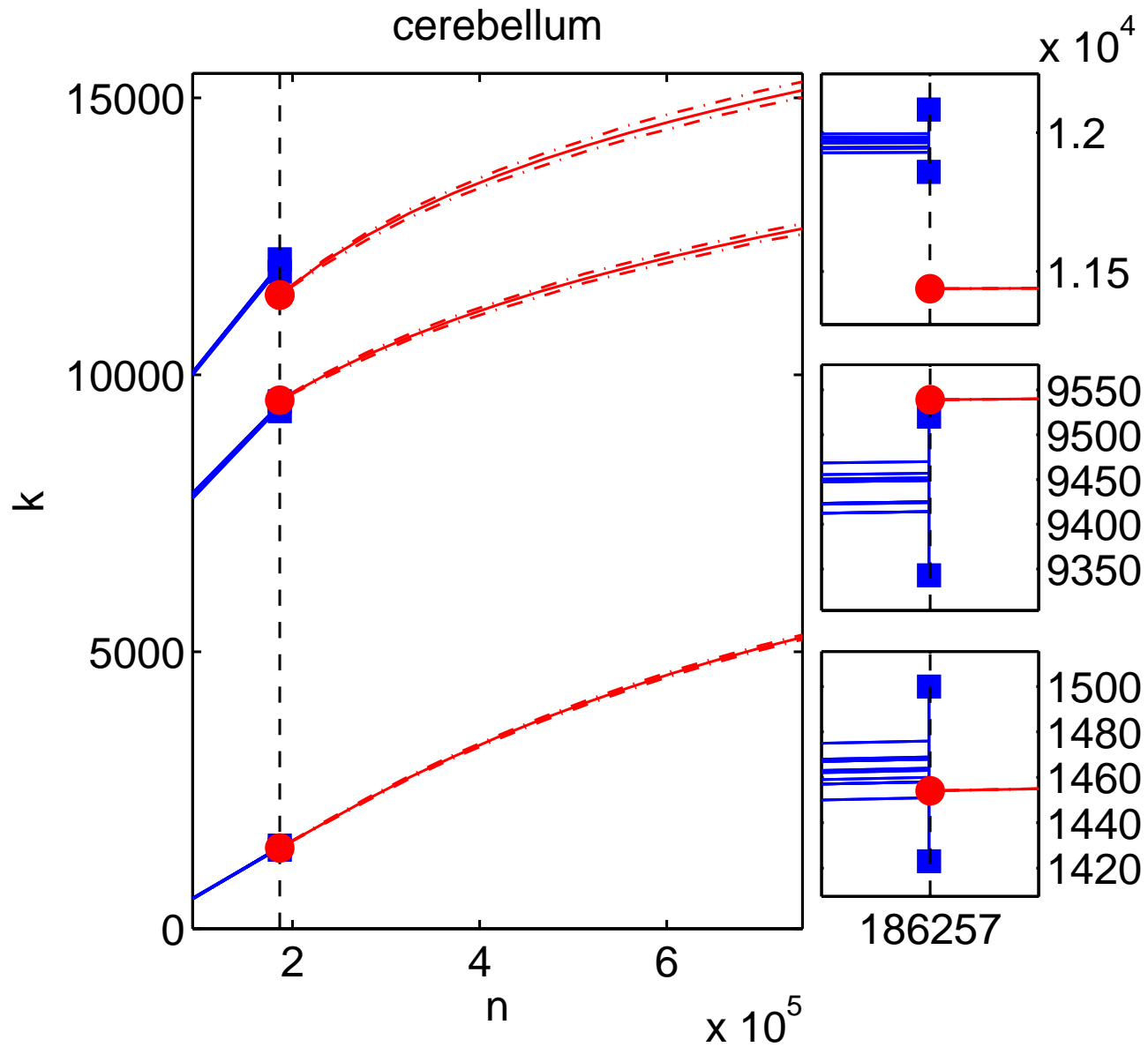
with Gibbs sampling  $(\sigma, \theta)$  and Yor-Pitman sampling for  $\mathbf{m}$ .



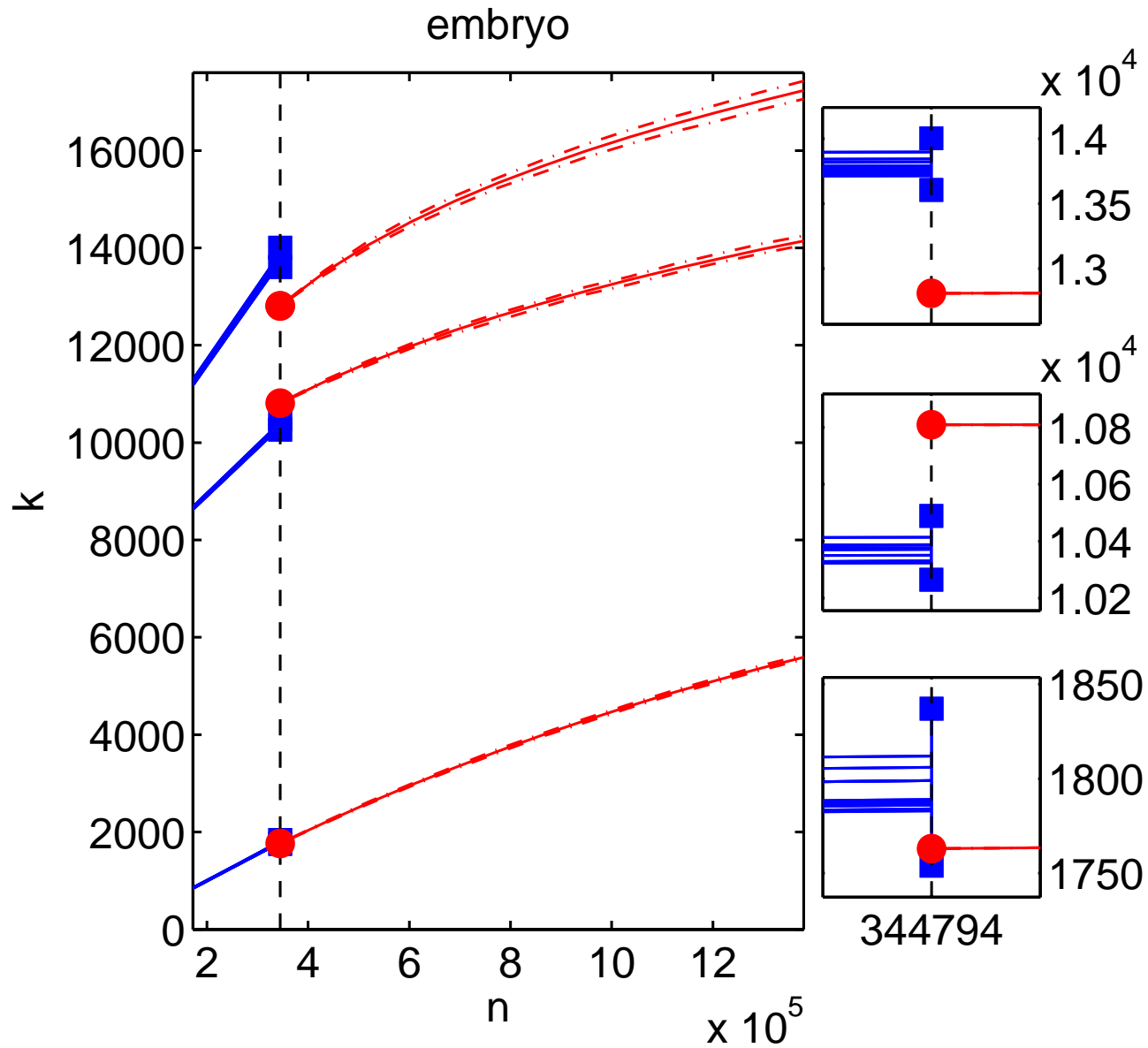
# Averaging versus max. likelihood



# Notice anything funny?

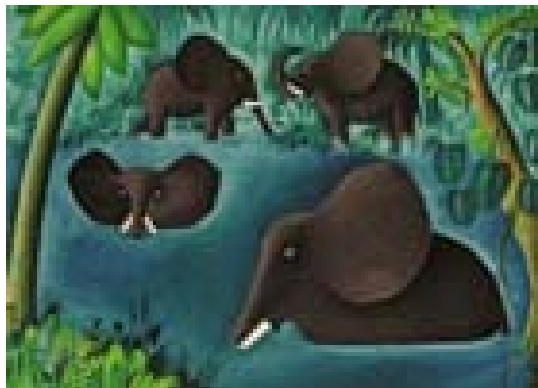


# Notice anything funny? Example 2



# (Well-known) take home messages

- Parameter averaging works!
- The model is always wrong! (as revealed with sufficient data)
- Only one model considered!
- What happened to being Bayesian about model selection?



# Calculating the marginal likelihood

The marginal likelihood:

$$p(\mathcal{D}|\mathcal{H}) = \int p(\mathcal{D}|\mathbf{f}, \mathcal{H}) p(\mathbf{f}|\mathcal{H}) d\mathbf{f}$$

Approximate inference needed!

Monte Carlo: slow mixing, non-trivial to get marginal likelihood estimates

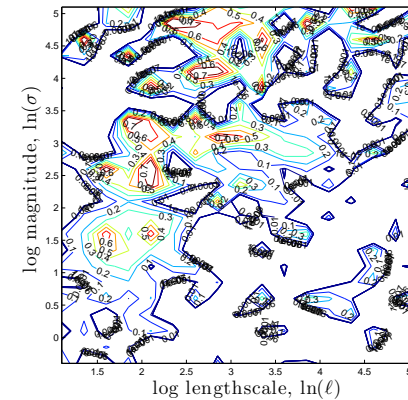
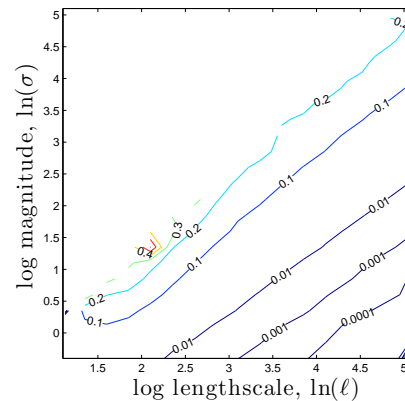
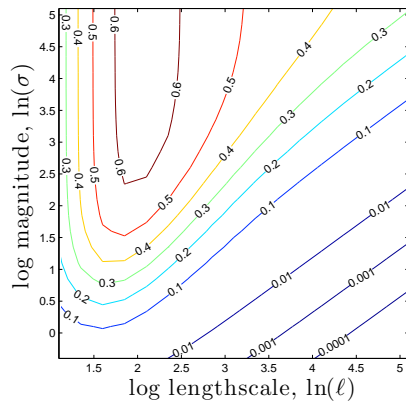
Expectation propagation+, variational Bayes, loopy BP+: sometimes not precise, approximation errors not controllable, not applicable to all models.

# Motivation: validating EP corrections

Kuss-Rasmussen (JMLR 2006)  $N = 767$  3-vs-5 GP USPS digit classification with

$$K(\mathbf{x}, \mathbf{x}') = \sigma^2 \exp\left(-\frac{\|\mathbf{x} - \mathbf{x}'\|^2}{2\ell^2}\right)$$

I:  $\log R = \log Z_{\text{EPC}} - \log Z_{\text{EP}}$  and II+III:  $\log Z_{\text{MCMC}} - \log Z_{\text{EP}}$ .



Thanks to Malte Kuss for making III available.

# Marginal likelihood from importance sampling

Importance sampling

$$p(\mathcal{D}|\mathcal{H}) = \int \frac{p(\mathcal{D}|\mathbf{f}, \mathcal{H}) p(\mathbf{f}|\mathcal{H})}{q(\mathbf{f})} q(\mathbf{f}) d\mathbf{f}$$

Draw samples  $\mathbf{f}_1, \dots, \mathbf{f}_R$  from  $q(\mathbf{f})$  and set

$$p(\mathcal{D}|\mathcal{H}) \approx \frac{1}{R} \sum_{r=1}^R \frac{p(\mathcal{D}|\mathbf{f}_r, \mathcal{H}) p(\mathbf{f}_r|\mathcal{H})}{q(\mathbf{f}_r)}$$

This will usually not work because ratio varies too much.

# Marginal likelihood from thermodynamic integration

Variants: parallel **tempering**, simulated **tempering** and **annealed** importance sampling

$$\begin{aligned}h(\mathbf{f}) &= p(\mathcal{D}|\mathbf{f}, \mathcal{H}) p(\mathbf{f}|\mathcal{H}) \\p(\mathbf{f}|\beta) &= \frac{1}{Z(\beta)} h^\beta(\mathbf{f}) q^{1-\beta}(\mathbf{f}) \\ \log Z(\beta_2) - \log Z(\beta_1) &= \int_{\beta_1}^{\beta_2} \frac{d \log Z(\beta)}{d\beta} d\beta \\ &= \int_{\beta_1}^{\beta_2} \int \log \frac{h(\mathbf{f})}{q(\mathbf{f})} p(\mathbf{f}|\beta) d\mathbf{f} d\beta\end{aligned}$$

Run  $N_\beta$  chains and interpolate

$$\log Z(\beta_2) - \log Z(\beta_1) \approx \frac{\Delta\beta}{R} \sum_{b=1}^{N_\beta} \sum_{r=1}^R \log \frac{h(\mathbf{f}_{rb})}{q(\mathbf{f}_{rb})}$$

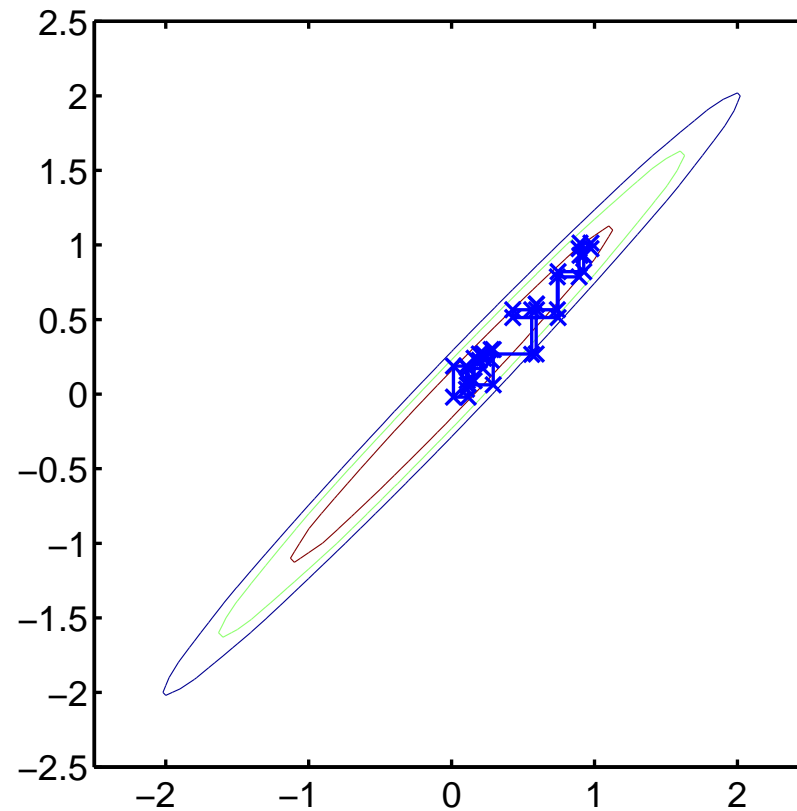
Other things that might work even better: multi-canonical.



# The trouble with Gibbs sampling

Cycle over variables  $f_i$ ,  $i = 1, \dots, N$  and sample conditionals

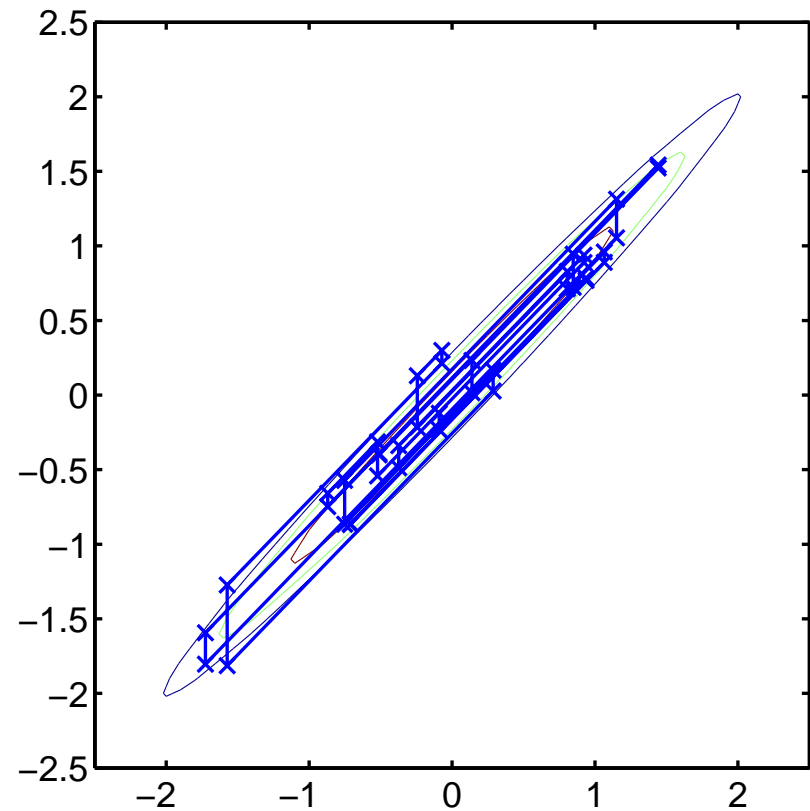
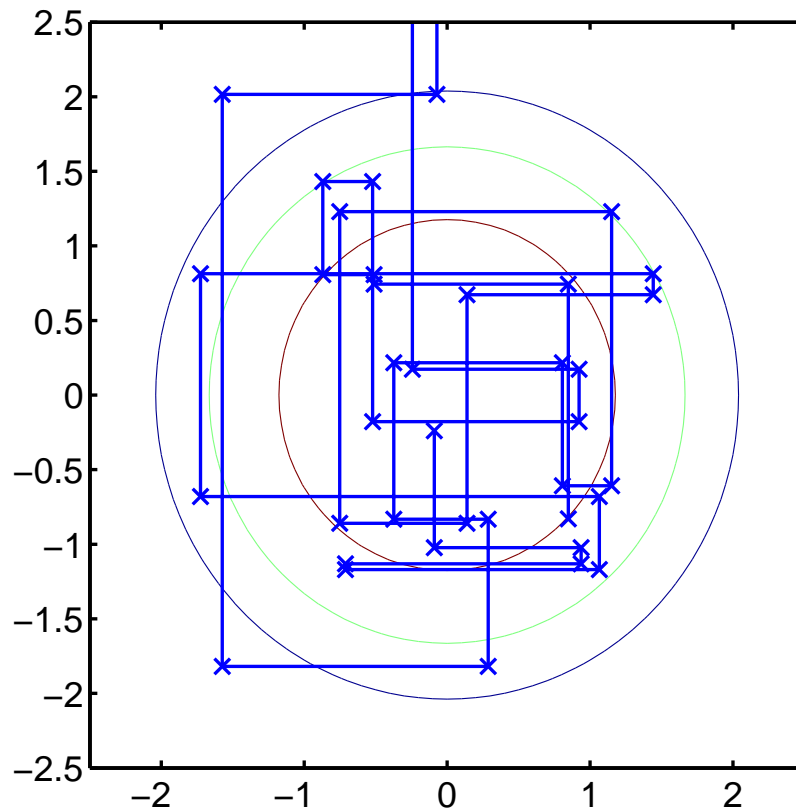
$$p(f_i | \mathbf{f}_{\setminus i}) = \frac{p(\mathbf{f})}{p(\mathbf{f}_{\setminus i})} \propto p(f)$$



# A trivial cure for $\mathcal{N}(\mathbf{f}|0, \mathbf{C})$

Gibbs sample  $z_i, i = 1, \dots, N$  with  $\mathcal{N}(\mathbf{z}|0, \mathbf{I})$

$$\mathbf{f} = \mathbf{Lz} \quad \text{with} \quad \mathbf{C} = \mathbf{L}\mathbf{L}^T$$



# Gaussian process classification (GPC)

$$p(\mathbf{f}|\mathbf{y}, \mathbf{K}, \beta) = \frac{1}{Z(\beta)} \prod_n \phi(y_n f_n) \exp\left(-\frac{\beta}{2} \mathbf{f}^T \mathbf{K}^{-1} \mathbf{f}\right)$$

Noise-free formulation  $\mathbf{f}_{\text{nf}}$

$$\phi(yf) = \int \theta(yf_{\text{nf}}) \mathcal{N}(f_{\text{nf}}|\mathbf{f}, \mathbf{I})$$

Joint distribution

$$p(\mathbf{f}, \mathbf{f}_{\text{nf}}, \mathbf{y}|\mathbf{K}, \beta) = p(\mathbf{y}|\mathbf{f}_{\text{nf}}) p(\mathbf{f}_{\text{nf}}|\mathbf{f}) p(\mathbf{f}|\mathbf{K}, \beta)$$

Marginalize out  $f$

$$p(\mathbf{f}_{\text{nf}}|\mathbf{y}, \mathbf{K}, \beta) \propto \prod_n \theta(y_n f_n) \mathcal{N}(f_{\text{nf}}|0, \mathbf{I} + \mathbf{K}/\beta)$$

Samples of  $\mathbf{f}$  can be recovered from  $p(\mathbf{f}|\mathbf{f}_{\text{nf}})$  (Gaussian)

Efficient sampler of truncated Gaussian needed!

# MCMC for GPC - related work

G. Rodriguez-Yam, R. Davis, and L. Scharf: “Efficient Gibbs Sampling of Truncated Multivariate Normal with Application to Constrained Linear Regression” (preprint 2004)

R. Neal, U. Paquet: Sample joint  $\mathbf{f}$ ,  $\mathbf{f}_{nf}$  and use Adler’s over-relaxation on  $\mathbf{f}$ .

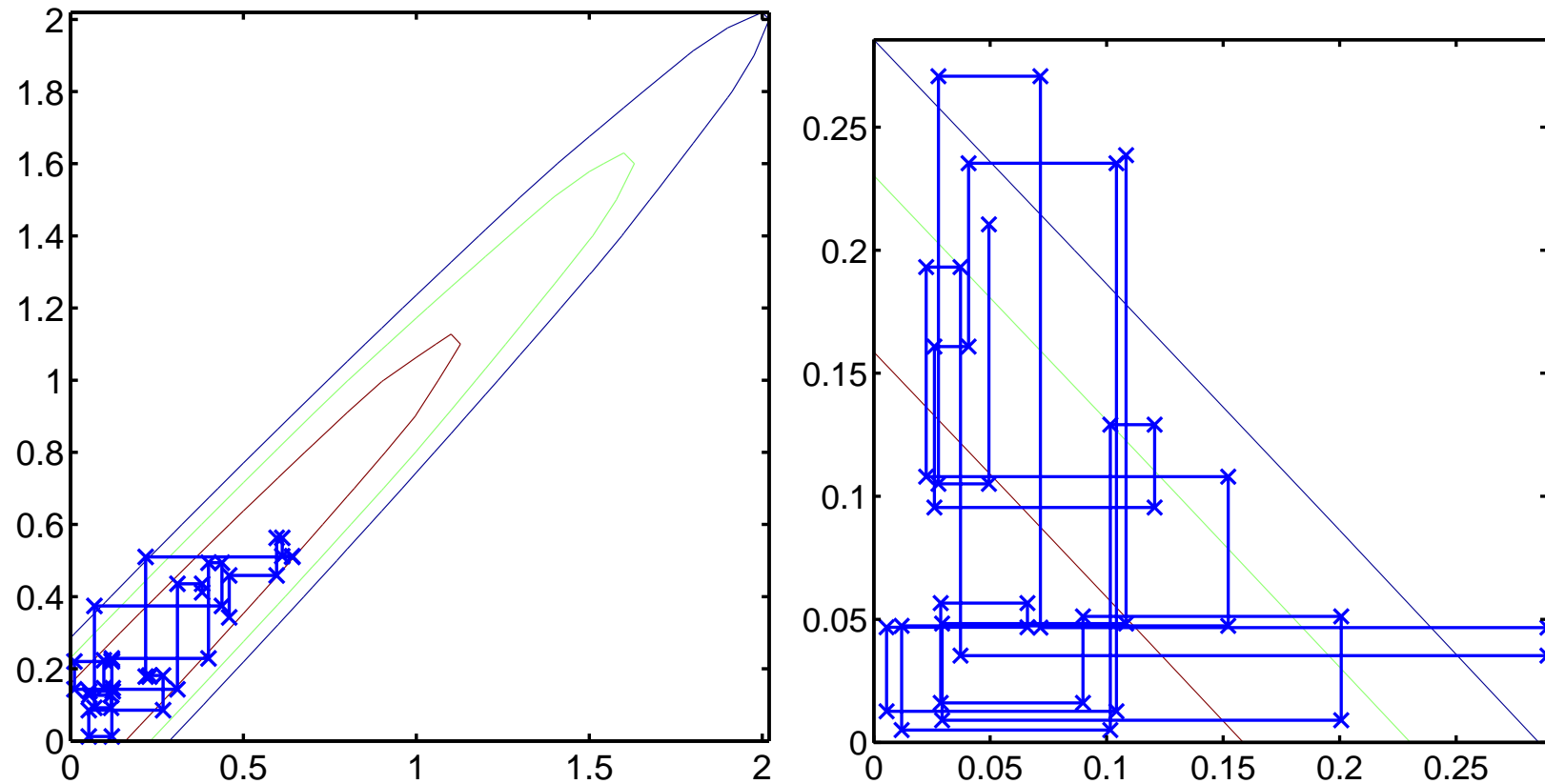
P. Rujan, R. Herbrich: Playing billiards in version space, the Bayes point machine.

M. Kuss+C. Rasmussen: Hybrid Monte Carlo in  $\mathbf{z}$ -space,  $\mathbf{f} = \mathbf{Lz}$  and annealed importance sampling

Y. Qi+T. Minka: Hessian based Metropolis-Hastings - local Gaussian proposals.

# Gibbs sampling - pos/neg covariance

Conditionals are truncated Gaussians! so sampling is easy.



# Efficient Gibbs sampling I

Gibbs sample in whitened space:  $z_i, i = 1, \dots, N$  with  $\mathcal{N}(\mathbf{z}|0, \mathbf{I})$

$$\mathbf{f} = \mathbf{Lz} \quad \text{with} \quad \mathbf{C} = \mathbf{LL}^T$$

What happens to the constraints, say  $\mathbf{f} \geq 0$

$$\mathbf{f} = \mathbf{Lz} \geq 0$$

Just a linear transformation so region in  $\mathbf{z}$ -space is convex and conditional (double) truncated Gaussian.

# Determine limits of conditionals

$j$ th conditional constraints  $i = 1, \dots, N$

$$L_{ij}z_j \geq - \sum_{k \neq i} L_{ik}z_k$$

divide in sets

$$S_{+,j} = \{i | L_{ij} > 0\}$$

$$S_{-,j} = \{i | L_{ij} < 0\}$$

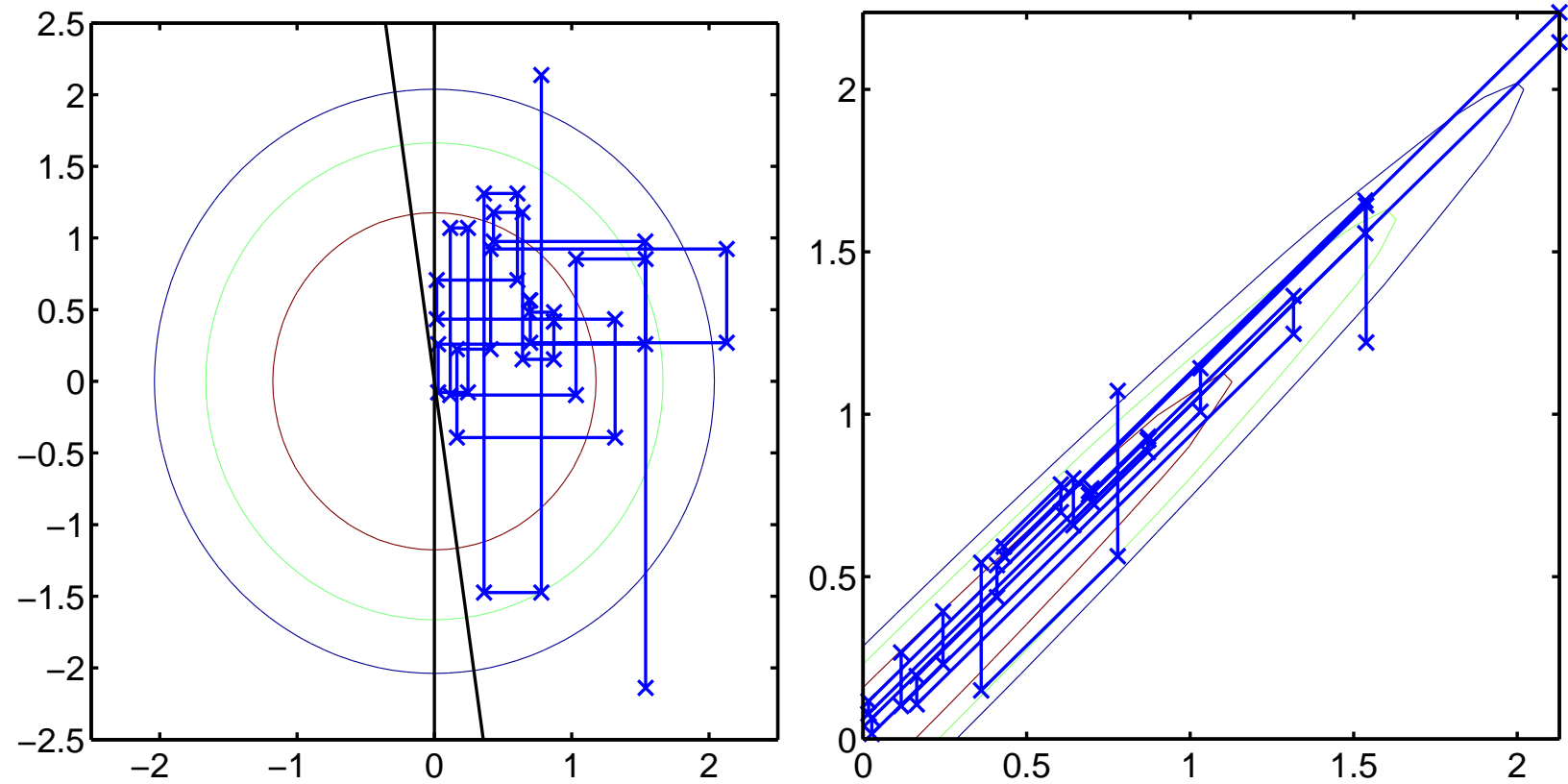
$$S_{0,j} = \{i | L_{ij} = 0\}$$

$$z_{j,\text{lower}} = \max_{i \in S_{+,j}} \frac{- \sum_{k \neq i} L_{ik}z_k}{L_{ij}}$$

$$z_{j,\text{upper}} = \min_{i \in S_{-,j}} \frac{- \sum_{k \neq i} L_{ik}z_k}{L_{ij}}$$

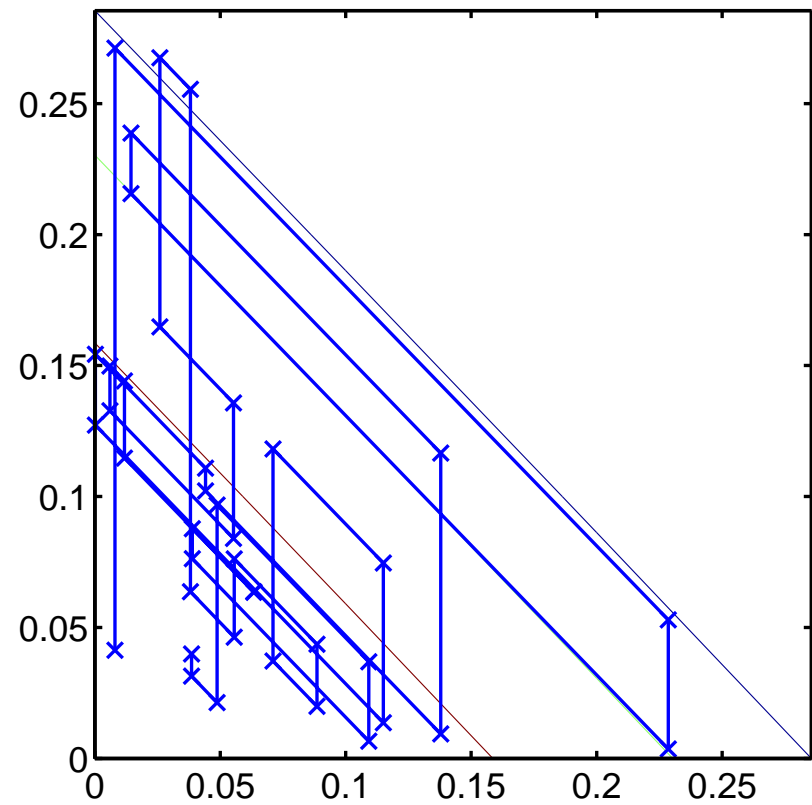
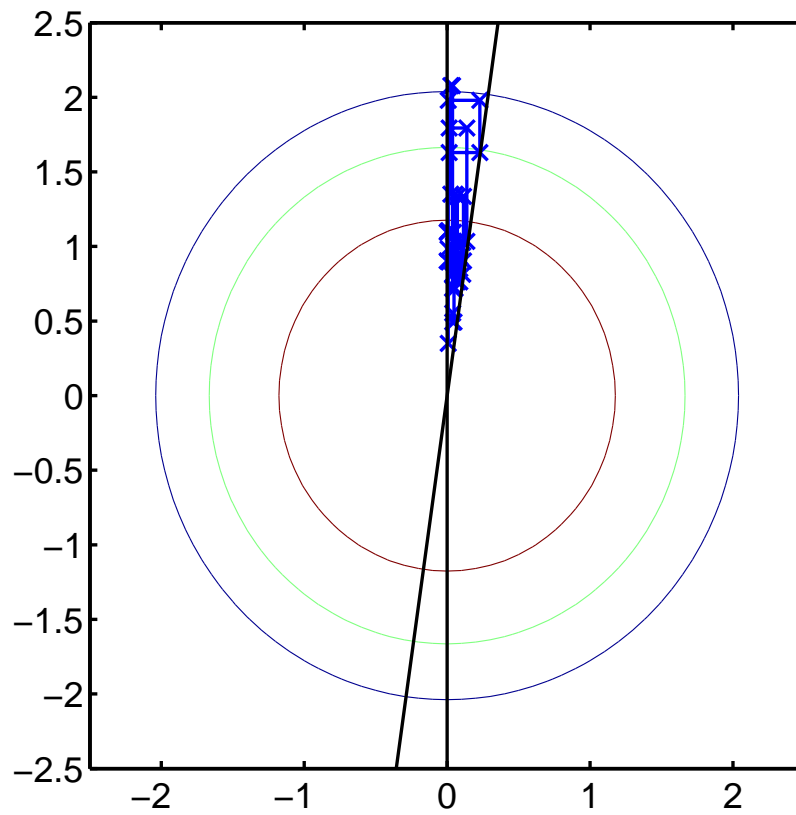
$$z_j = \phi^{-1} \left\{ \phi(z_{j,\text{lower}}) + \text{rand} \left( \phi(z_{j,\text{upper}}) - \phi(z_{j,\text{lower}}) \right) \right\}$$

# Gibbs sampling positive covariance



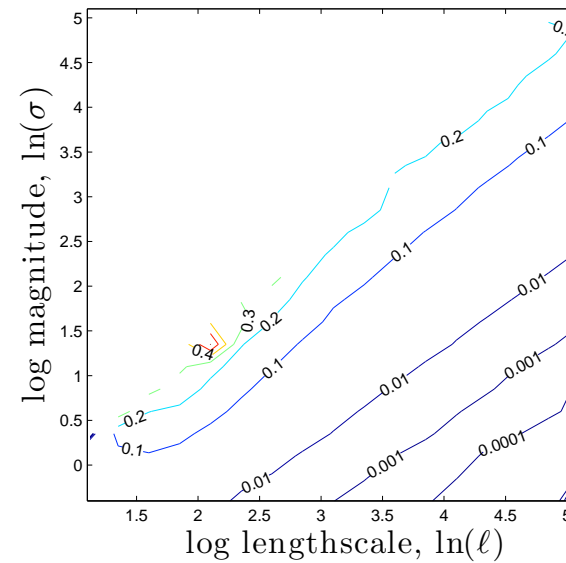
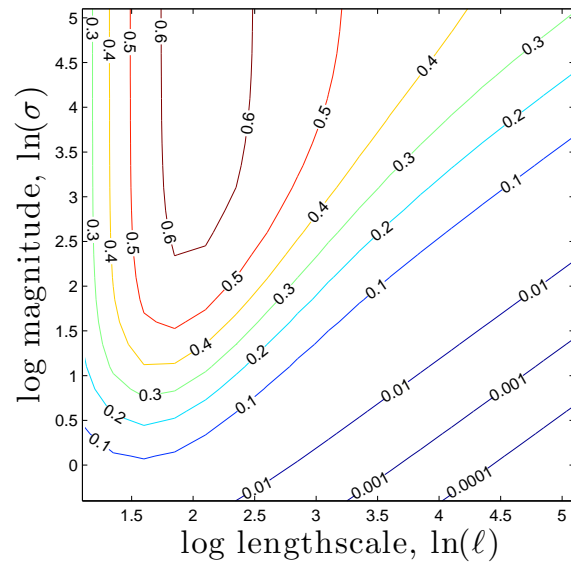


# Gibbs sampling negative covariance



# Kuss+Rasmussen set-up

EP, EP+corrections and MCMC are all very precise!



Details of the EP corrections will (hopefully) come to a conference near you soon!

# Summary

- Averaging works!
- (X-)validation points to model miss-specification!
- How to find better (noise) models?
- Marginal likelihood from sampling
- The trouble with Gibbs sampling and a cure
- Is machine learning becoming (Bayesian) statistics with big data set?

# Acknowledgments

## **Bioinformatics:**

Albin Sandelin

Eivind Valen

## **Machine learning:**

Ulrich Paquet

Manfred Opper

## **Students:** (many shared)

Ricardo Henao (machine learning and bioinformatics)

Morten Hansen (communication)

Carsten Stahlhut (source location in EEG)

Jóan Petur Petersen (machine learning for ship propulsion)

Man-Hung Tang (bioinformatics)

Eivind Valen (bioinformatics)

Troels Marstrand (bioinformatics)