

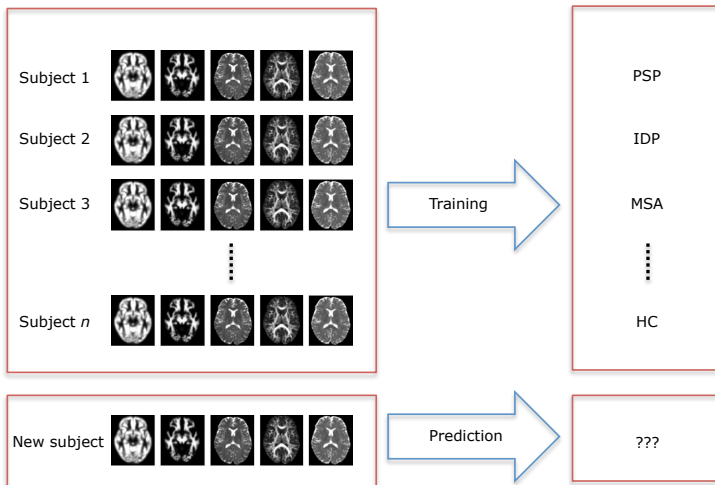
Pseudo-Marginal Bayesian Inference for Gaussian Processes

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Motivating Application



HC - Healthy control

MSA - Multiple system atrophy

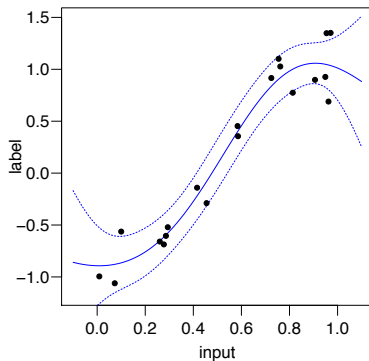
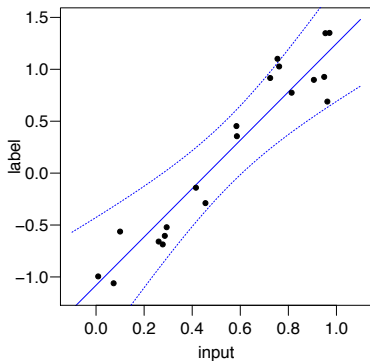
PSP - Progressive Supranuclear Palsy

IDP - Idiopathic Parkinson's disease

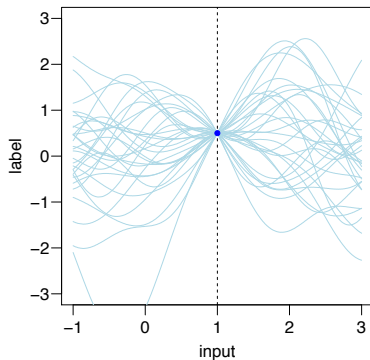
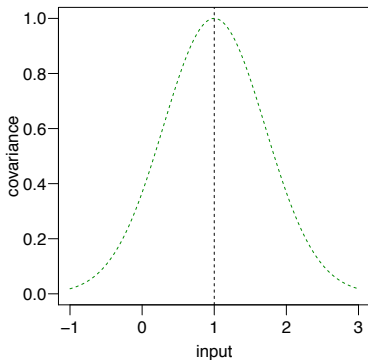
- Mapping input to labels

$$\text{label} = \text{function}(\text{input}, \text{par}, \text{noise})$$

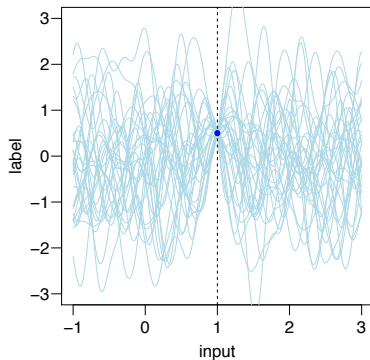
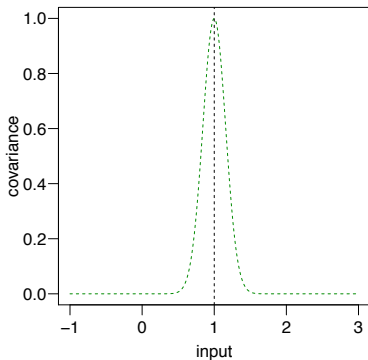
- We may have no clue about function - we need assumptions



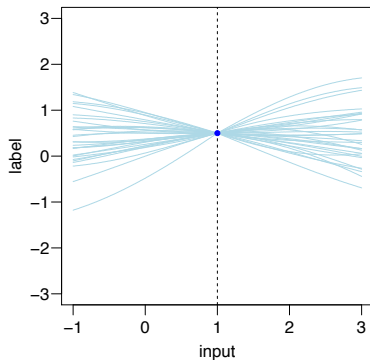
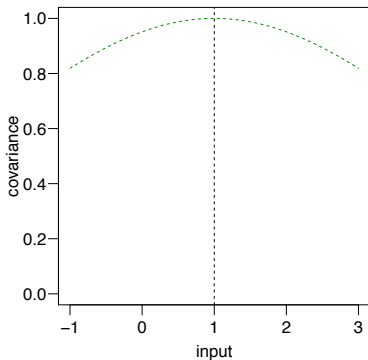
- Gaussians with distance dependent covariance



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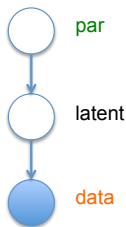


Gaussian Process Models

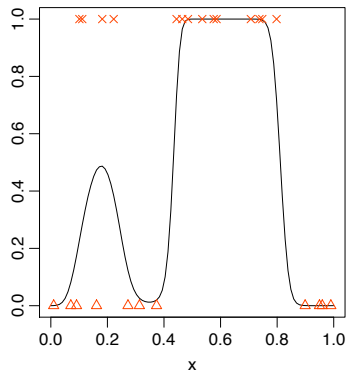
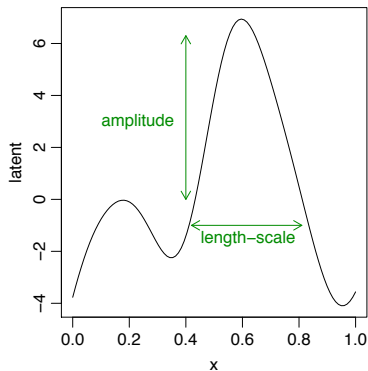
- Class of hierarchical models

$$p(\text{data}|\text{latent}) \quad p(\text{latent}|\text{par}) \quad p(\text{par})$$

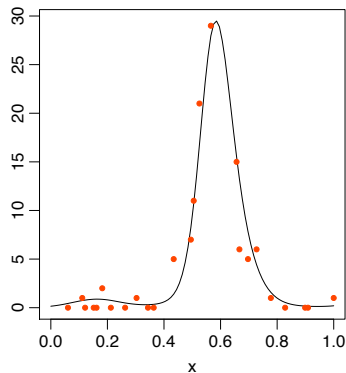
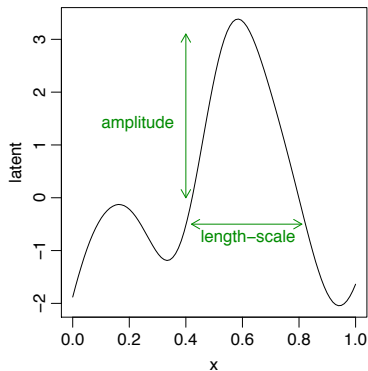
- $p(\text{latent}|\text{par}) = \text{Gaussian Process}$



Gaussian Process Models - Classification example

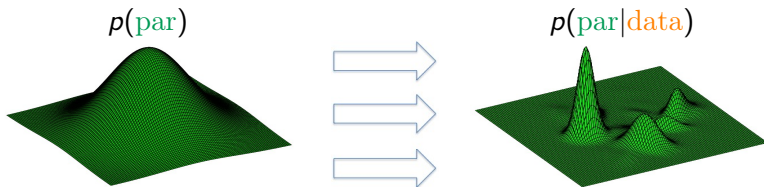


Gaussian Process Models - Count data example

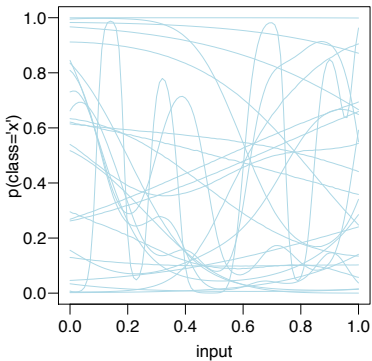
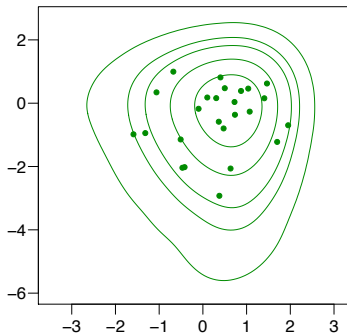


- Inference using Bayes theorem:

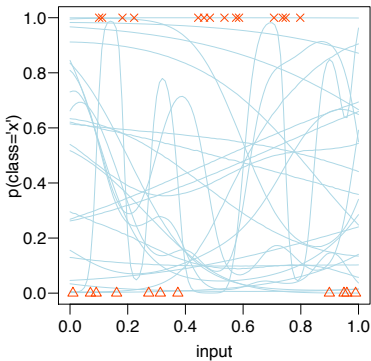
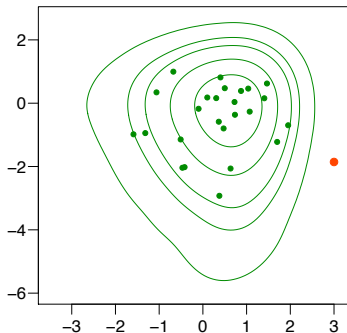
$$p(\text{par}|\text{data}) = \frac{p(\text{data}|\text{par})p(\text{par})}{\int p(\text{data}|\text{par})p(\text{par})d\text{par}}$$



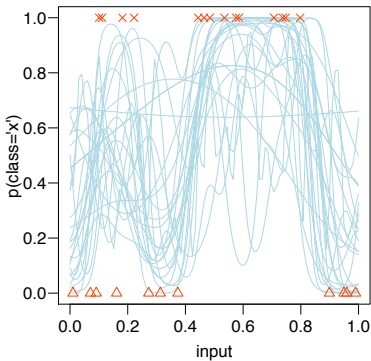
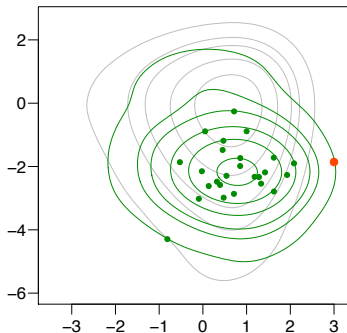
Bayesian Inference - Prior



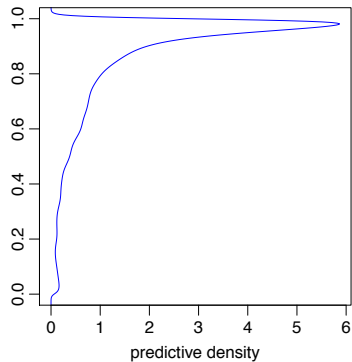
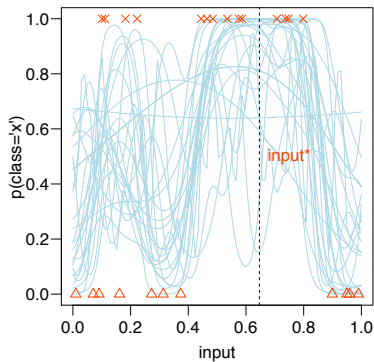
Bayesian Inference - Data



Bayesian Inference - Posterior



Bayesian Inference and Predictions



- Predictions for new data

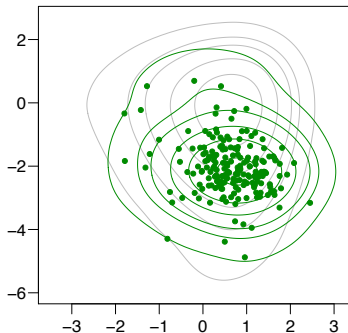
$$p(\text{label}_*|\text{label}) = \int p(\text{label}_*|\text{par})p(\text{par}|\text{label}) d\text{par}$$

- Monte Carlo integration:

$$\int p(\text{label}_*|\text{par})p(\text{par}|\text{label}) d\text{par} \simeq \frac{1}{N} \sum_{i=1}^N p(\text{label}_*|\text{par}^{(i)})$$

with $\text{par}^{(i)}$ drawn from $p(\text{par}|\text{data})$

- Draw samples according to the posterior density



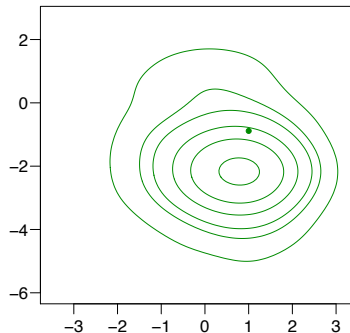
- Bayesian inference

$$p(\text{par}|\text{data}) = \frac{p(\text{data}|\text{par})p(\text{par})}{\int p(\text{data}|\text{par})p(\text{par})d\text{par}}$$

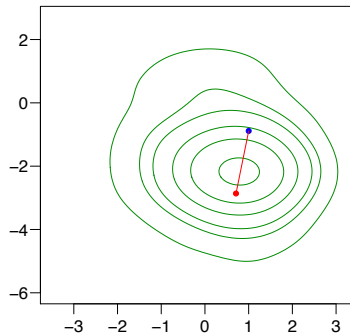
- Random walk sampler - accept a proposal with probability

$$\min \left(1, \frac{p(\text{par}'|\text{data})}{p(\text{par}|\text{data})} \right)$$

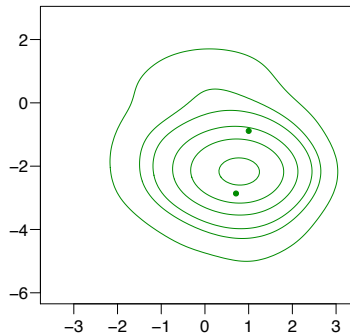
- Explore the parameter space according to the density



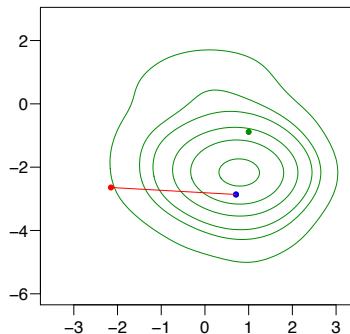
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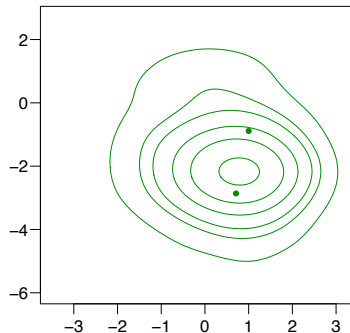
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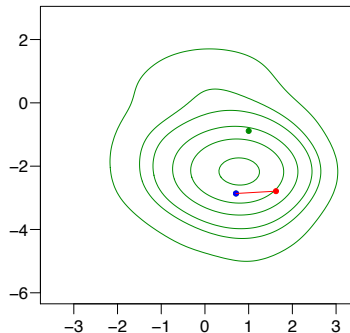
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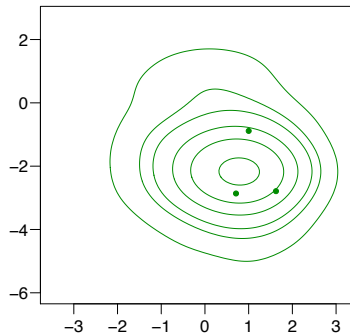
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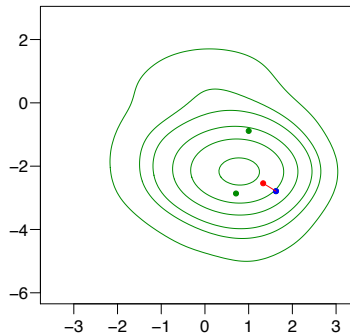
- Explore the parameter space according to the density



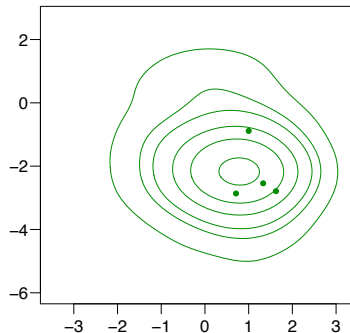
- Explore the parameter space according to the density



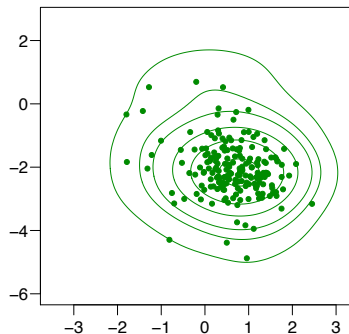
- Explore the parameter space according to the density



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- Explore the parameter space according to the density



How can we draw from $p(\text{par}|\text{data})$ in GPMs?

Marginal likelihood

$$p(\text{data}|\text{par}) = \int p(\text{data}|\text{latent})p(\text{latent}|\text{par})d\text{latent}$$

is unavailable analytically. Options:

- Approximate $p(\text{data}|\text{par})$ within MCMC
- Sample from $p(\text{par}, \text{latent}|\text{data})$
- Pseudo-Marginal MCMC

Gaussian Approximations for marginal likelihood

Gaussian approximation to $p(\text{latent}|\text{data}, \text{par})$

- Laplace Approximation
- Expectation Propagation
- Variational Bayes
- ...

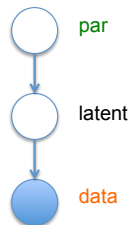
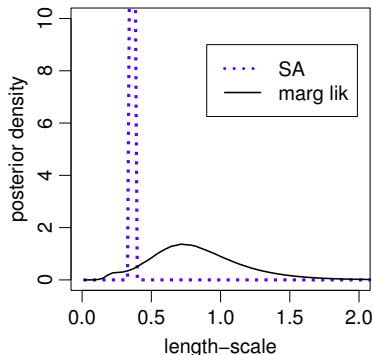
Challenges in MCMC for GPMs - Cost & exploration

- No exact Gibbs steps - need to employ Metropolis within Gibbs steps - waste of computations when rejecting
- Updates of par cost $O(n^3)$
- par can be large dimensional (e.g., Automatic Relevance Determination (ARD) covariance function)
- There are n latent variables (as many as the number of observations)

Challenges in MCMC for GPMs - Structure

Obvious iterative scheme (aka Sufficient Augmentation (SA) scheme). Alternate between:

- Drawing from $p(\text{latent}|\text{par}, \text{data})$
- Drawing from $p(\text{par}|\text{latent})$ (**bad idea** - see figure)

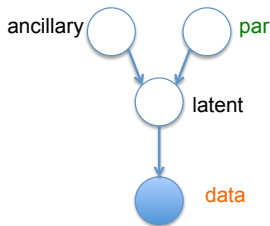
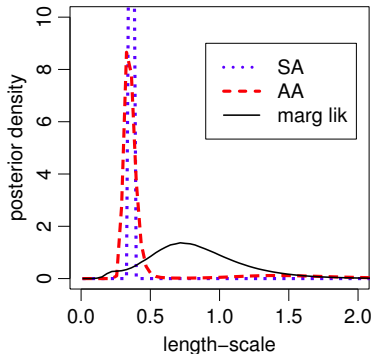


Mitigating coupling effect through reparameterization

Ancillary Augmentation (AA) scheme - reparameterization:

$$K = LL^T \quad \text{ancillary} = L^{-1} \text{latent}$$

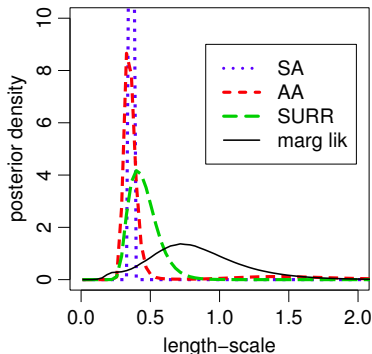
- Replace sampling of **par** with $p(\text{par}|\text{ancillary}, \text{data})$



Mitigating coupling effect through reparameterization

Surrogate data model (SURR):

- Introduce set of auxiliary variables informed by the posterior over latent



$$\text{surrogate} = f(\text{latent}, \text{par})$$

- Replace posterior by unbiased estimate

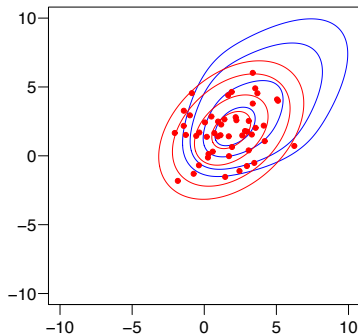
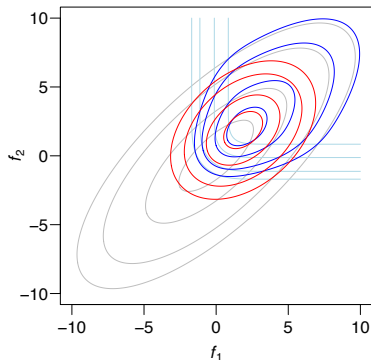
$$\min \left(1, \frac{\tilde{p}(\text{par}'|\text{data})}{\tilde{p}(\text{par}|\text{data})} \right)$$

- Achieved by using an unbiased estimate of $\tilde{p}(\text{data}|\text{par})$

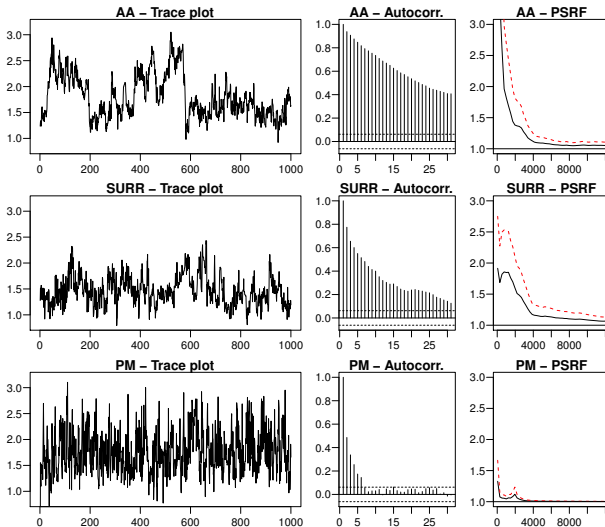
Importance Sampling estimator

- Approximate posterior over latent variables using $q(\text{latent})$
- Then

$$\tilde{p}(\text{data}|\text{par}) = \frac{1}{N} \sum_{i=1}^N \frac{p(\text{data}|\text{latent}^{(i)})p(\text{latent}^{(i)}|\text{par})}{q(\text{latent}^{(i)})}$$

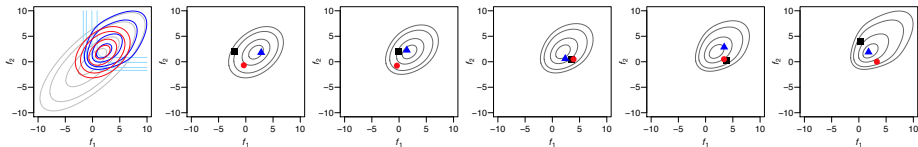


Abalone data set (two classes) $n = 2835$ - inference of length-scale



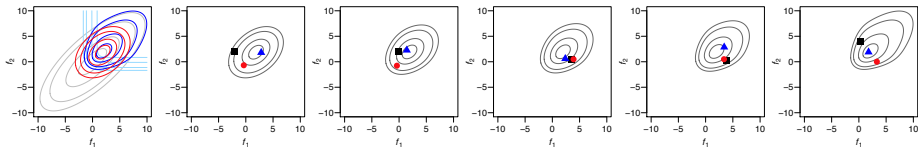
Annealed Importance Sampling estimator

- Annealing from an approximation

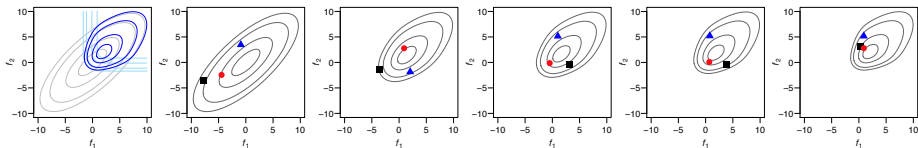


Annealed Importance Sampling estimator

- Annealing from an approximation



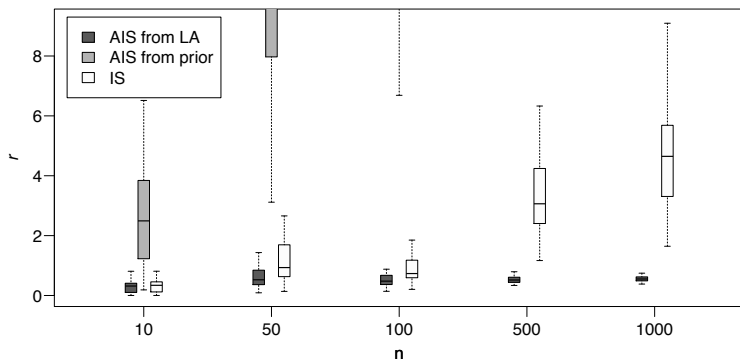
- Annealing from the prior



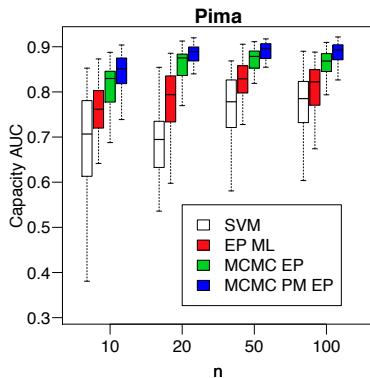
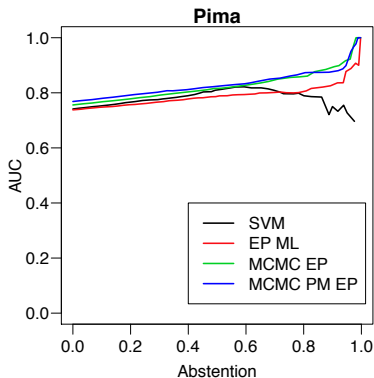
Comparison between AIS with IS

Analysis of the variance of the AIS and IS estimators

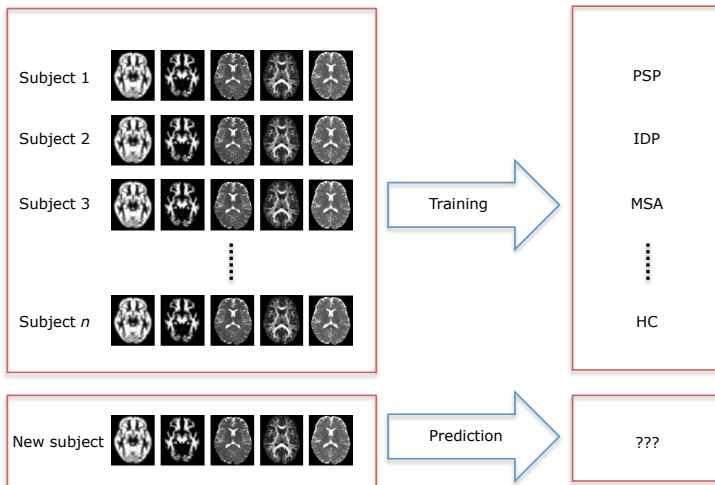
- r is the variance of the \log_{10} marginal likelihood



Some Results



Motivating Application



HC - Healthy control

MSA - Multiple system atrophy

PSP - Progressive Supranuclear Palsy

IDP - Idiopathic Parkinson's disease

- Multiclass classification based on GPs

$$p(\text{disease} = c | \text{sources}) = \text{unknown function}$$

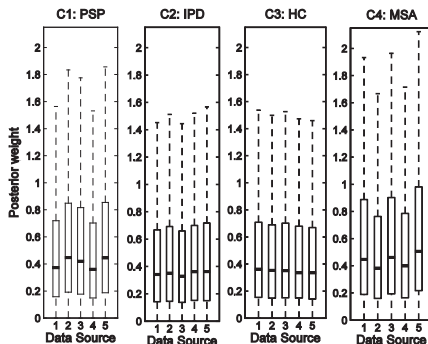
- unknown function modeled using GPs

- Covariance based on source-dependent covariances S_k

$$\sum_{k=1}^K w_{ck} S_k(\text{subject}_i, \text{subject}_j)$$

Parkinson syndromes data - multi source

Method	Accuracy
GP classifier	0.598
SimpleMKL	0.418



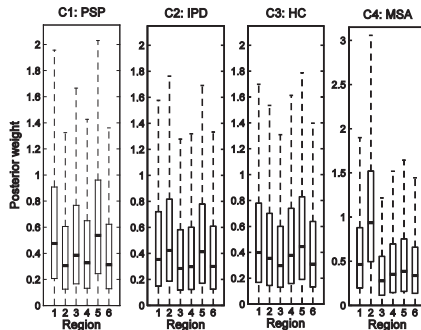
Multiclass classification with multiple regions

Analysis of brain regions

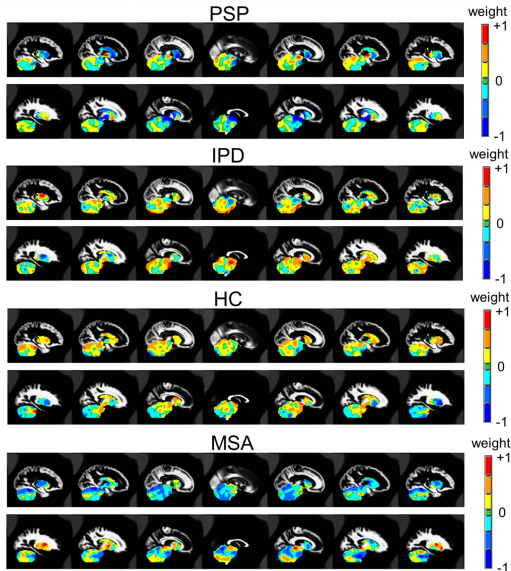
- 1 brainstem
- 2 bilateral cerebellum
- 3 bilateral caudate
- 4 bilateral middle occipital gyrus
- 5 bilateral putamen
- 6 all other regions

Multiclass classification with multiple regions

Method	Accuracy
GP classifier	0.614
SimpleMKL	0.229



Brain Maps



- Gaussian Processes yield flexible and interpretable nonparametric models

Conclusions and ongoing work

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- Gaussian Processes yield flexible and interpretable nonparametric models
- Bayesian inference to accurately quantifying uncertainty in such models
- Pseudo-Marginal MCMC offers a practical way to carry out exact Bayesian computations
- How to make exact Bayesian computations for Gaussian Processes scalable?

- Dr Andre F. Marquand (King's College London)
- Prof Mark Girolami (University College London)
- Dr Guido Sanguinetti (University of Edinburgh)
- Dr Alessandro Vinciarelli (University of Glasgow)

[1] M. Filippone and M. Girolami. Pseudo-Marginal Bayesian inference for Gaussian processes, *IEEE Transactions on Pattern Analysis and Machine Intelligence*, to appear.

[2] M. Filippone. Bayesian inference for Gaussian process classifiers with annealing and pseudo-marginal MCMC, In *ICPR*, 2014.

[3] M. Filippone et al. Probabilistic prediction of neurological disorders with a statistical assessment of neuroimaging data modalities. *Annals of Applied Statistics*, 6(4):1883-1905, 2012.

[4] A. F. Marquand et al. Automated, high accuracy classification of Parkinsonian disorders: a pattern recognition approach. *PLoS ONE*, 2013.

[5] M. Filippone et al. A comparative evaluation of stochastic-based inference methods for Gaussian process models. *Machine Learning*, 93(1):93-114, 2013.