

Quantifying Uncertainty

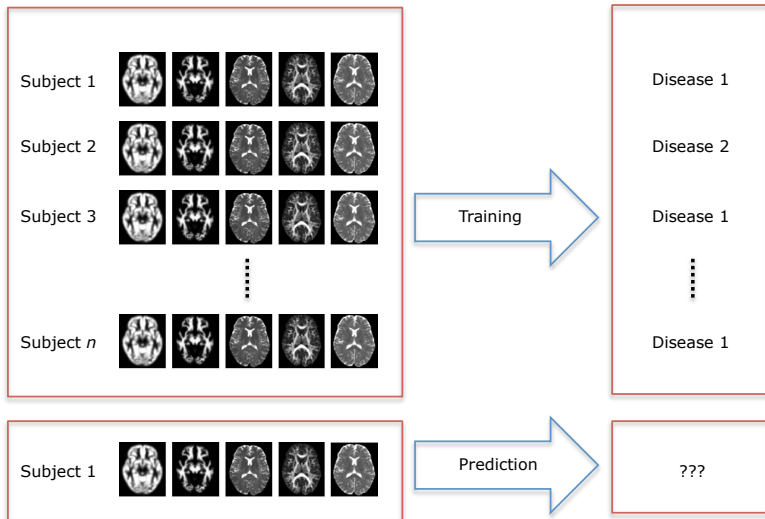
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Motivating Application

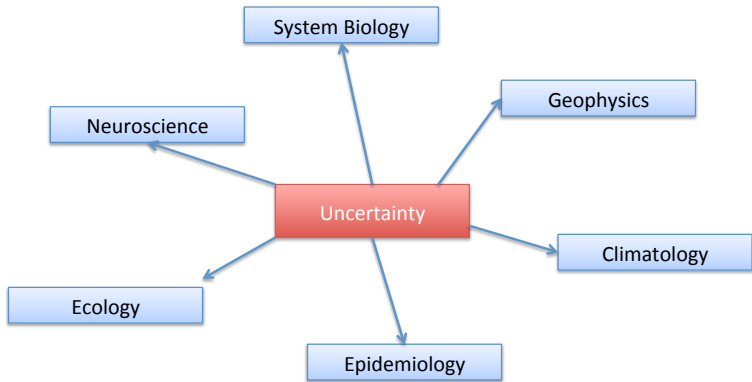
Neurological disorder from neuroimages



Relevance of the problem



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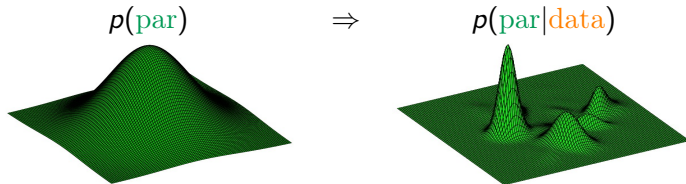


`label = function(input, par, noise)`

- How do we estimate model parameters?
- How do we assess that a model is preferable over another?
- How do we incorporate knowledge by experts?
- How can we attach confidence intervals to our predictions and parameter estimates?

Probabilistic modeling offers an answer to these questions

- Parameters and data are viewed as random variables



- Inference using Bayes theorem:

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

- Predictions for new data

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

- Requires the posterior distribution $p(\text{par}|\text{data})$

Monte Carlo integration

- Predictions for new data $p(\text{data}_*|\text{data})$ is an expectation

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

- Monte Carlo estimation:

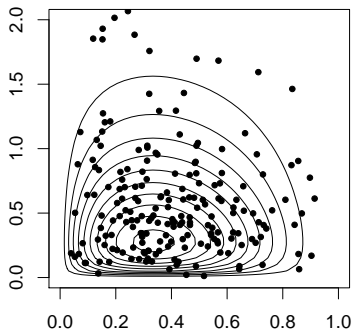
$$\int p(\text{data}_*|\text{par})p(\text{par}|\text{data}) d\text{par} \simeq \frac{1}{N} \sum_{i=1}^N p(\text{data}_*|\text{par}_i)$$

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

- **Good** news: asymptotically correct
- **Bad** news: the “error” $\rightarrow 0$ in $O(1/\sqrt{N})$

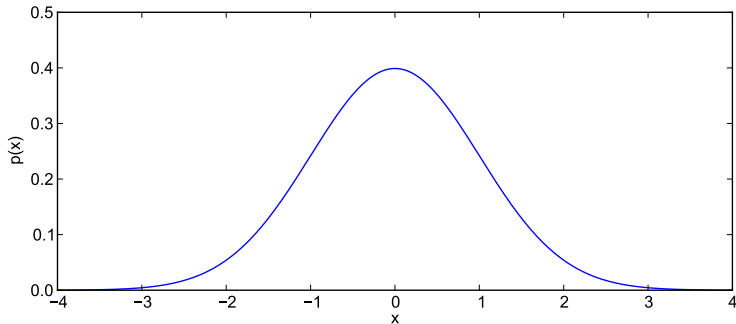
Monte Carlo integration

- Draw samples according to the density



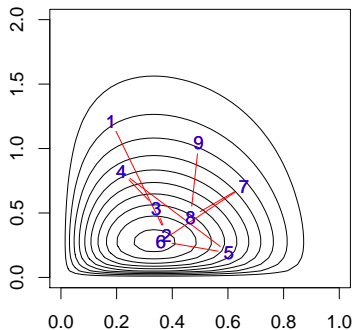
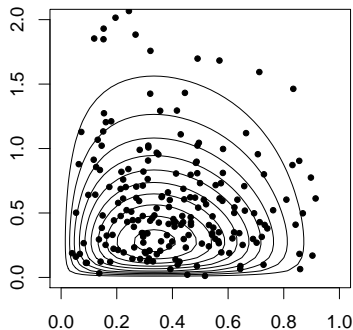
Probability density functions - An important example

- Gaussian probability density function
- We can draw samples from this directly



Markov chain Monte Carlo

- Explore the parameter space according to the density



Often it is not possible to draw samples directly - need to set up a Markov chain

Markov chain Monte Carlo

- MCMC needs the density up to a normalization constant
- Random walk sampler - accept a proposal with probability

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

- by Bayes' theorem

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

- Therefore:

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

A class of hierarchical models

- Models can have more complex structures
- For example:

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

- Note: $p(\text{latent state}|\text{par})$ is actually (for example):

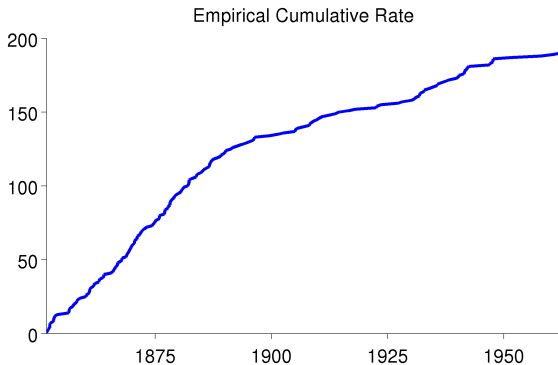
$$p(\text{latent state}|\text{par}, \text{time})$$

or

$$p(\text{latent state}|\text{par}, \text{location})$$

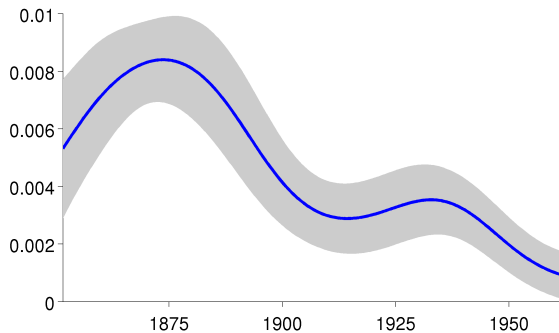
Coal mine disasters data

191 accidents between 1851 and 1962



Coal mine disasters data

191 accidents between 1851 and 1962

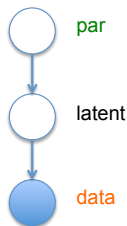


Latent Gaussian Models - LGMs

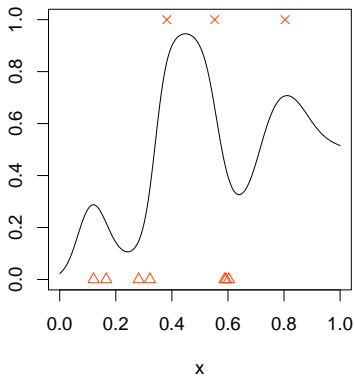
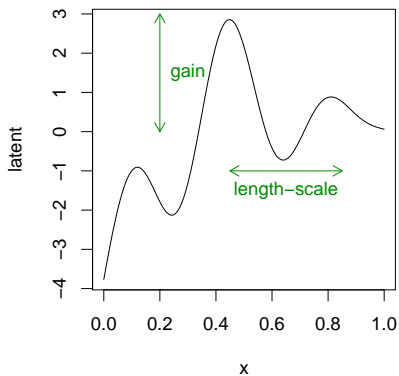
- Class of hierarchical models

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

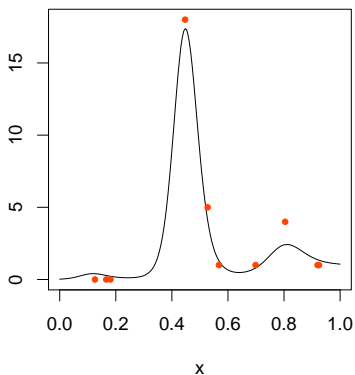
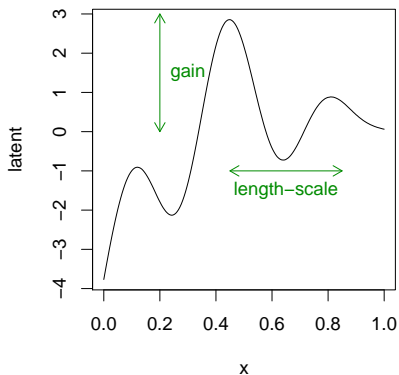
- $p(\text{latent}|\text{par}) = \text{Gaussian}(\text{latent}|\mu(\text{par}), K(\text{par}))$



LGMs - Classification example



LGMs - Count data example



Parkinson syndromes data

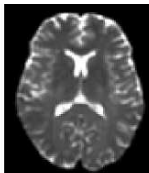
- 62 subjects
- **Early stage** prediction of development of
 - Parkinson Syndromes
 - Multiple System Atrophy
 - Progressive Supranuclear Palsy
- Given neuroimages



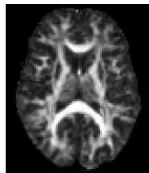
GM



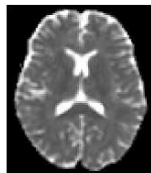
WM



T2



FA



MD

- Multiclass classification based on LGMs

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

- For every pair of subjects x_i, x_j , we can characterize their similarity (**covariance**) $C_s(x_i, x_j)$ as given by the imaging modality s
- We do this by looking at the values of the neuroimages

- Latent variables $f_c(x)$ with LGM prior with covariance

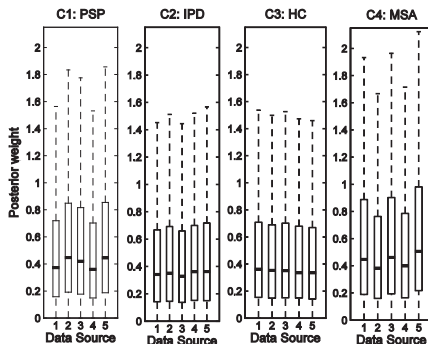
$$\text{cov}(f_c(x_i), f_c(x_j)) = \sum_{s=1}^q w_{cs} C_s(x_i, x_j)$$

- Multinomial likelihood - general case of the logistic function

$$p(\text{disease} = c | \text{latent}, \text{sources}) = \frac{\exp(f_c(x))}{\sum_{r=1}^m \exp(f_r(x))}$$

Parkinson syndromes data - multi source

Method	Accuracy
GP classifier	0.598
SimpleMKL	0.418

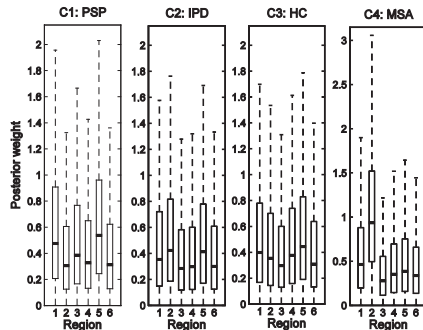


Analysis of brain regions

- For this analysis we used only the GM data source
- We used an anatomical template as in Shattuck et al. 2008 to parcellate the GM images into:
 - brainstem
 - bilateral cerebellum
 - bilateral caudate
 - bilateral middle occipital gyrus
 - bilateral putamen
 - all other regions

Parkinson syndromes data - multi region

Method	Accuracy
GP classifier	0.614
SimpleMKL	0.229



- Bayesian inference offers the possibility to tackle challenging problems by accurately quantifying **uncertainty** in predictions and model parameters
- Computing is playing an important part in bridging the gap between theory and application of Bayesian inference

Thank you!

Questions?