



NTNU
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Introduction to INLA

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Outline

- Introduction
- Bayesian hierarchical models
- Latent Gaussian models
- Deterministic inference
- R-INLA

The screenshot shows the R-bloggers website interface. At the top, there's a navigation bar with links for Home, About, RSS, add your blog!, R jobs, and Contact us. The main content area features a featured article titled "INLA: Bayes goes to Norway" by Luis, dated August 15, 2012. The article has social media sharing buttons for Facebook (14493 likes), Twitter (0 tweets), and Google+ (0). A quote from the article is displayed: "INLA is not the Norwegian answer to ABBA; that would probably be a-ha. INLA is the answer to 'Why do I have enough time to cook a three-course meal while running MCMC analyses?'". Below the quote, the article's abstract is shown: "Integrated Nested Laplace Approximations (INLA) is based on direct numerical integration (rather than simulation as in MCMC) which, according to people 'in the know', allows:

- the estimation of marginal posteriors for all parameters,
- marginal posteriors for each random effect and
- estimation of the posterior for linear combinations of random effects.

 To the right of the article, there are sections for "TOP 3 POSTS FROM THE PAST 2 DAYS" and "TOP 9 ARTICLES OF THE WEEK". A search bar is also visible.

<http://www.r-bloggers.com/inla-bayes-goes-to-norway/>

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What?

The short answer:

INLA is a fast method to do Bayesian inference with latent Gaussian models and R-INLA is an R-package that implements this method with a flexible and simple interface.

A much longer answer:

Rue, Martino, and Chopin (2009) "Approximate Bayesian inference for latent Gaussian models by using integrated nested Laplace approximations." *Journal of the royal statistical society: Series B.* 319–392

INLA - how it all began

- INLA is the result of many years of work, mainly driven by Håvard Rue
- the first “user-friendly” implementation was built on a C-library in 2007
- The first prototype of the R-interface came in Jan/Feb 2008
- today, the complete source-code is about 100 000 lines, in R/C/C++ (view/track/download from inla.googlecode.com)



... and the INLA group increased



(photo 2011)

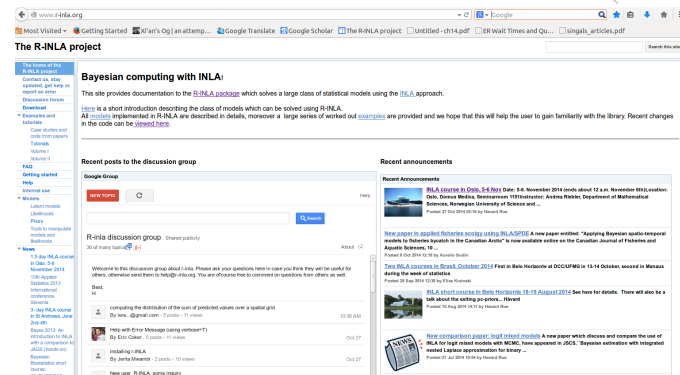
There are more:



...

Where?

The software, information, examples and help can be found at <http://www.r-inla.org>



A complete documentation about INLA is in progress.

So... When can R-INLA be used?

- What type of problems can it solve?
- What type of models can be used?

The core

We have observed something.

We have questions.

We want answers.

How do we find answers?

We need to make choices:

Bayesian or frequentist?

How do we model the data?

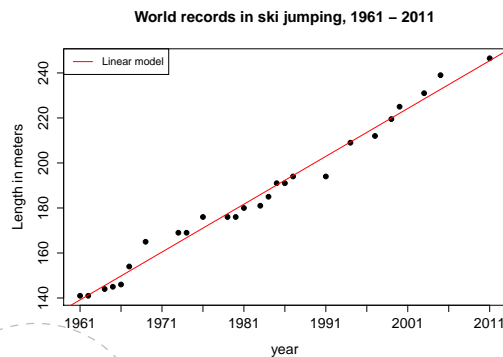
How do we compute the answer?

These questions are *not* independent.

Example: Ski flying records

Assume a simple linear regression model with Gaussian observations $\mathbf{y} = (y_1, \dots, y_n)$, where

$$E(y_i) = \mu + \beta x_i, \quad \text{Var}(y_i) = \tau^{-1}, \quad 1, \dots, n$$

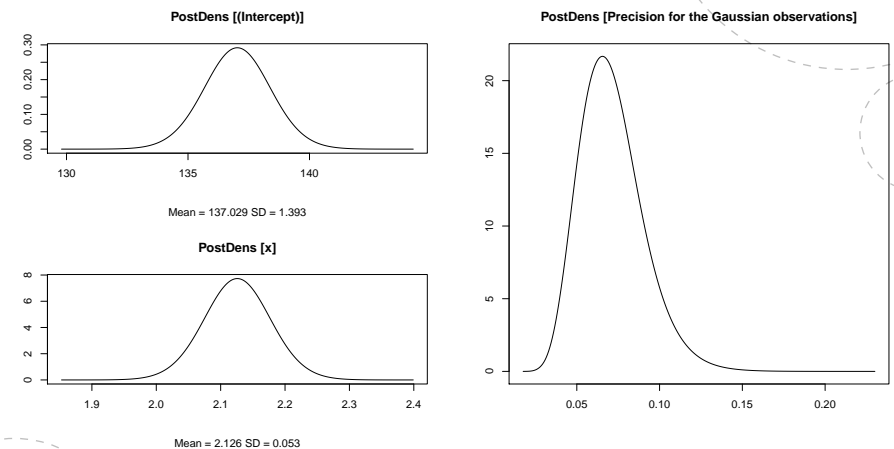


Estimates

Intercept: 137.03 (1.42195),
 β : 2.13 (0.05)

The Bayesian approach

Assign priors to the parameters α , β and τ and calculate posteriors:



Real-world datasets are usually much more complicated!

Using a Bayesian framework:

- Build (hierarchical) models to account for potentially complicated dependency structures in the data.
- Attribute uncertainty to model parameters and latent variables using priors.

Two main challenges:

- Need computationally efficient methods to calculate posteriors.
- Select priors in a sensible way.

Bayesian hierarchical models

INLA can be used with Bayesian hierarchical models where we model in different stages or levels:

Stage 1: What is the distribution of the responses?

Stage 2: What is the distribution of the underlying unobserved (latent) components?

Stage 3: What are our prior beliefs about the parameters controlling the components in the model?

Stage 1

How is our **data** (\mathbf{y}) generated from the underlying components (\mathbf{x}) and hyperparameters (θ) in the model:

- Gaussian response?
- Count data? (E.g. Poisson, negative binomial)
- Spatial point pattern?
- Binary data?
- ...

This information is placed into our *likelihood* $\pi(\mathbf{y}|\mathbf{x}, \theta)$

Stage 2

The underlying **unobserved components** \mathbf{x} are called **latent components** and can be:

- Covariates
- Unstructured random effects (individual effects, group effects)
- Structured random effects, for example an autoregressive process of order 1 (AR(1)-process).

These are linked to the responses in the likelihood through linear predictors.

Stage 3

The likelihood and the latent model typically have hyperparameters that control their behavior. The **hyperparameters** θ can include:

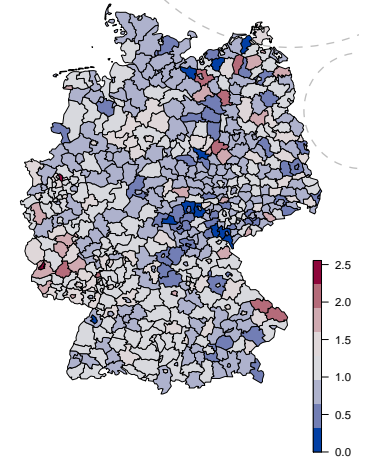
- Variance of observation noise
- Variance of unstructured effects
- Correlation of multivariate effects
- Autocorrelation parameter

Example: Disease mapping in Germany

We observed larynx cancer mortality counts for males in 544 district of Germany from 1986 to 1990 and want to make a model.

Information available:

- y_i : The count at location i .
- E_i : An offset; expected number of cases in district i .
- c_i : A covariate (level of smoking consumption) at location i
- s_i : spatial location i (here, district).



Stage 1: The data

First we decide on the likelihood for our data \mathbf{y}

- Our responses are counts
- We decide to model our responses as

$$y_i | \eta_i \sim \text{Poisson}(E_i \exp(\eta_i))$$

- η_i is a linear function of the latent components

Stage 2: The latent model

The latent field \mathbf{x} consists of two parts:

1. One fixed effect: the intercept μ
2.
 - The spatially structured effect f_s .
 - The unstructured effect \mathbf{u} which accounts for non-observed variability
 - The unknown effect $f(c_i)$ of the exposure covariate which assumes value c_i for district i .

These are combined for each location to give a linear predictor

$$\eta_i = \mu + f_s(s_i) + f(c_i) + u_i$$

The latent field is $\mathbf{x} = (\mu, \{f_s(\cdot)\}, \{f(\cdot)\}, u_1, u_2, \dots, u_n)$

Stage 3: Hyperparameters

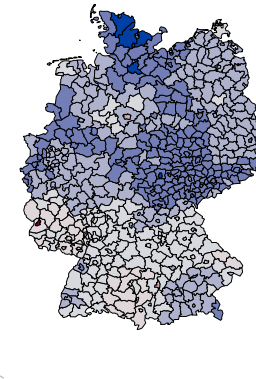
The structured and unstructured spatial effect as well as the smooth covariate effect will be each controlled by one parameter

- $\tau_c, \tau_f, \tau_\eta$: The precisions (inverse variances) of the covariate effect, spatial effect and unstructured effect, respectively.

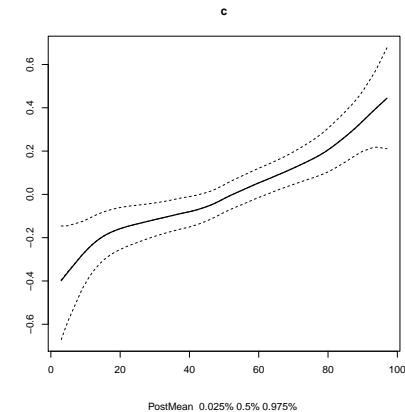
The hyperparameters are $\theta = (\tau_c, \tau_f, \tau_\eta)$, and must be given a prior $\pi(\tau_c, \tau_f, \tau_\eta)$.

Quantities of interest

Structured spatial effect
 $\exp(f_s(s_i))$



Covariate effect $f(c_i)$



Latent Gaussian models

This example is just one example of a very useful class of models called **Latent Gaussian models**.

- The characteristic property is that the **latent part** of the hierarchical model is **Gaussian**, $\mathbf{x}|\theta \sim N(\mathbf{0}, \mathbf{Q}^{-1})$
- The expected value is $\mathbf{0}$
- The *precision* matrix (inverse covariance matrix) is \mathbf{Q}

The general set-up

The set up contains GLMs, GLMMs, GAMs, GAMMs, and more. The mean of the observation i , μ_i , is connected to the linear predictor, η_i , through a link function g ,

$$\eta_i = g(\mu_i) = \mu + \mathbf{z}_i^T \boldsymbol{\beta} + \sum_{\gamma} w_{\gamma,i} f_{\gamma}(c_{\gamma,i}) + u_i, \quad i = 1, 2, \dots, n$$

where

μ : Intercept

$\boldsymbol{\beta}$: Fixed effects of covariates \mathbf{z}

$\{f_{\gamma}(\cdot)\}$: Non-linear/smooth effects of covariates \mathbf{c}

$\{w_{\gamma,i}\}$: Known weights defined for each observed data point

\mathbf{u} : Unstructured error terms

Specification of the latent field

- Collect all parameters (random variables) in the linear predictor in a **latent field** $\mathbf{x} = \{\mu, \beta, \{f_\gamma(\cdot)\}, \eta\}$.
- A latent Gaussian model is obtained by assigning Gaussian priors to all elements of \mathbf{x} .
- Very flexible due to many different forms of the unknown functions $\{f_\lambda(\cdot)\}$:
- **Hyperparameters** account for variability and length/strength of dependence

Flexibility through f -functions

The functions $\{f_\gamma\}$ in the linear predictor make it possible to capture very different types of random effects in the same framework:

- $f(\text{time})$: For example, an AR(1) process, a random walk of first or second order (RW1 or RW2)
- $f(\text{spatial location})$: For example, a Matérn field
- $f(\text{covariate})$: For example, a RW1 or RW2 on the covariate values
- $f(\text{time, spatial location})$ can be a spatio-temporal effect
- And much more

Additivity

- One of the most useful features of the framework is the additivity.
- Effects can easily be removed and added without difficulty.
- Each component might add a new latent part and might add new hyperparameters, but the modelling framework and computations stay the same.

A small point to think about

From a Bayesian point of view fixed effects and random effects are all the same.

- Fixed effects are also random
- They only differ in the prior we put on them

Example: Smoothing binary time-series

- Have observed a sequence y_1, y_2, \dots, y_n of 0s and 1s
- Each time t has an associated covariate x_t
- We want to smooth the time series by inferring the sequence ρ_t , for $t = 1, 2, \dots, n$, of probabilities for 1s at each time step

Example: Smoothing time series

Stage 1: We choose a Bernoulli distribution for the responses, so that

$$y_t | \eta_t \sim \text{Bernoulli} \left(\frac{\exp(\eta_t)}{1 + \exp(\eta_t)} \right)$$

Stage 2: Covariates, AR(1) component, i.e. $a_t = \rho a_{t-1} + \epsilon_t$, and random noise are connected to likelihood by

$$\eta_t = \beta_0 + \beta_1 x_t + a_t + v_t$$

Stage 3: ρ : Dependence parameter in AR(1) process

σ_a^2 : Marginal variance in AR(1) process

σ_v^2 : Variance of unstructured term

Loads of examples

- Generalized linear and additive (mixed) models
- Disease mapping
- Survival analysis
- Log-Gaussian Cox-processes
- Spatio and spatio-temporal models
- Stochastic volatility models
- Measurement error models
- And more!

Computations

So...

Now we have a modelling framework

But how do we get our answers?

What do we care about?

It depends on the problem!

- A single element of the latent field (e.g. the sign or quantiles of a fixed effect)
- A linear combination of elements from the latent field (the average over an area of a spatial effect, the difference of two effects)
- A single hyperparameter (the correlation)
- Predictions at unobserved locations

What do we need to compute?

Often we are interested in the posterior probability density of an element of the latent field

$$\pi(x_i|\mathbf{y})$$

or the posterior probability density of an element of the hyperparameters

$$\pi(\theta_j|\mathbf{y})$$

or some other statistics

$$\pi(f(\mathbf{x}, \boldsymbol{\theta})|\mathbf{y})$$

But, as always in Bayesian statistics, we need to do high-dimensional integrals that cannot be computed analytically.

Traditional approach: MCMC*

Based on sampling. Construct Markov chains with the target posterior as stationary distribution.

- Extensively used within Bayesian inference since the 1980's.
- Flexible and general, sometimes the only thing we can do!
- A generic tool is available with JAGS/OpenBUGS.
- Tools for specific models are of course available, e.g. BayesX and stan.

* Markov chain Monte Carlo

Approximate inference

Bayesian inference can (almost) never be done exactly. Some form of approximation must always be done.

- MCMC “works” for everything, but it can be incredibly slow
- Is it possible to make a quicker, more specialized inference scheme which only needs to work for this limited class of models?

Recall: What is our model framework?

Latent Gaussian models

$$\mathbf{y}|\mathbf{x}, \boldsymbol{\theta} \sim \prod_i \pi(y_i|\eta_i, \boldsymbol{\theta})$$

$$\mathbf{x}|\boldsymbol{\theta} \sim \pi(\mathbf{x}|\boldsymbol{\theta}) \sim \mathcal{N}(\mathbf{0}, \mathbf{Q}(\boldsymbol{\theta})^{-1})$$

$$\boldsymbol{\theta} \sim \pi(\boldsymbol{\theta})$$

Gaussian!

Not Gaussian

where the precision matrix $\mathbf{Q}(\boldsymbol{\theta})$ is sparse. Generally these “sparse” Gaussian distributions are called **Gaussian Markov random fields** (GMRFs).

The sparseness can be exploited for very quick computations for the Gaussian part of the model through numerical algorithms for sparse matrices.

The INLA idea

Use the posterior distribution

$$\pi(\mathbf{x}, \boldsymbol{\theta} | \mathbf{y}) \propto \pi(\boldsymbol{\theta})\pi(\mathbf{x} | \boldsymbol{\theta})\pi(\mathbf{y} | \mathbf{x}, \boldsymbol{\theta})$$

to approximate the posterior marginals

$$\pi(x_i | \mathbf{y}) \quad \text{and} \quad \pi(\theta_j | \mathbf{y})$$

directly.

Let us consider a toy example to illustrate the ideas.

Smoothing noisy observations (I)

Observations

$$y_i = m(i) + \epsilon_i, \quad i = 1, \dots, n$$

for Gaussian iid noise ϵ_j with *known* precision.

Will assume $m(i)$ is a smooth function wrt i

How does INLA work?

Observations

$$y_i = m(i) + \epsilon_i, \quad i = 1, \dots, n$$

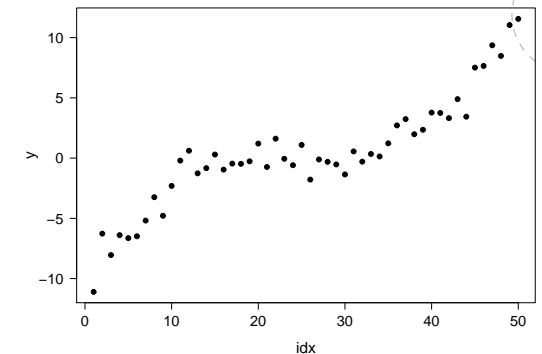
Here, we assume that $m(i)$ is a smooth function wrt i and

$\epsilon_j \stackrel{\text{iid}}{\sim} \mathcal{N}(0, \tau_0)$ with *known* precision τ_0 .

```

1 n = 50
2 idx = 1:n
3 fun = 100*((idx-n/2)/n)^3
4 y = fun + rnorm(n, mean
5   =0, sd=1)
6 plot(idx, y)

```



Assumed hierarchical model

1. **Data:** Gaussian observations with known precision

$$y_i | x_i, \theta \sim \mathcal{N}(x_i, \tau_0)$$

2. **Latent model:** A Gaussian model for the smooth function¹

$$\pi(\mathbf{x} | \theta) \propto \theta^{(n-2)/2} \exp\left(-\frac{\theta}{2} \sum_{i=2}^n (x_i - 2x_{i-1} + x_{i-2})^2\right)$$

3. **Hyperparameter:** The smoothing parameter θ which we assign a $\Gamma(a, b)$ prior

$$\pi(\theta) \propto \theta^{a-1} \exp(-b\theta), \quad \theta > 0$$

¹model="rw2"

Derivation of posterior marginals (I)

Since

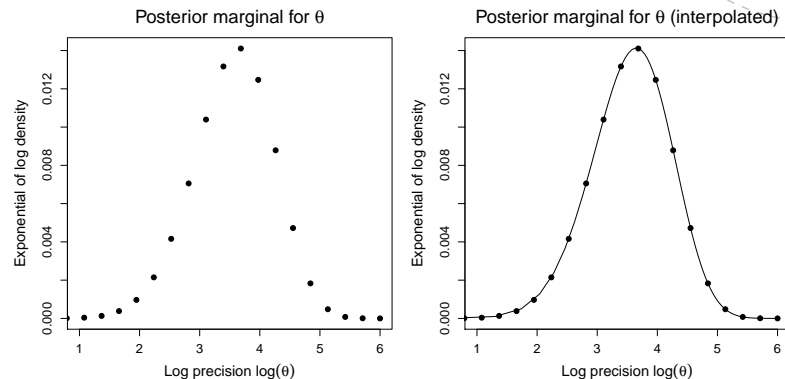
$$\mathbf{x}, \mathbf{y} | \theta \sim \mathcal{N}(\cdot, \cdot)$$

(derived using $\pi(\mathbf{x}, \mathbf{y} | \theta) \propto \pi(\mathbf{y} | \mathbf{x}, \theta) \pi(\mathbf{x} | \theta)$), we can compute (numerically) all marginals, using that

$$\pi(\theta | \mathbf{y}) \propto \frac{\overbrace{\pi(\mathbf{x}, \mathbf{y} | \theta)}^{\text{Gaussian}} \pi(\theta)}{\underbrace{\pi(\mathbf{x} | \mathbf{y}, \theta)}_{\text{Gaussian}}}$$

Posterior marginal for hyperparameter

Select a grid of points to represent the density $\theta | \mathbf{y}$. (Here, they points are chosen to be equi-distant.)



Derivation of posterior marginals (II)

From

$$\mathbf{x} | \mathbf{y}, \theta \sim \mathcal{N}(\cdot, \cdot)$$

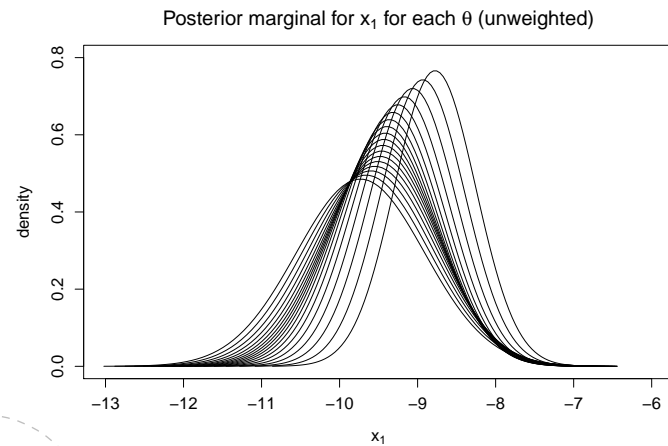
we can compute

$$\begin{aligned} \pi(x_i | \mathbf{y}) &= \int \underbrace{\pi(x_i | \theta, \mathbf{y})}_{\text{Gaussian}} \pi(\theta | \mathbf{y}) d\theta \\ &\approx \sum_k \pi(x_i | \theta_k, \mathbf{y}) \pi(\theta_k | \mathbf{y}) \Delta_k \end{aligned}$$

where $\theta_k, k = 1, \dots, K$, correspond to representative points of $\theta | \mathbf{y}$ and Δ_k are the corresponding weights.

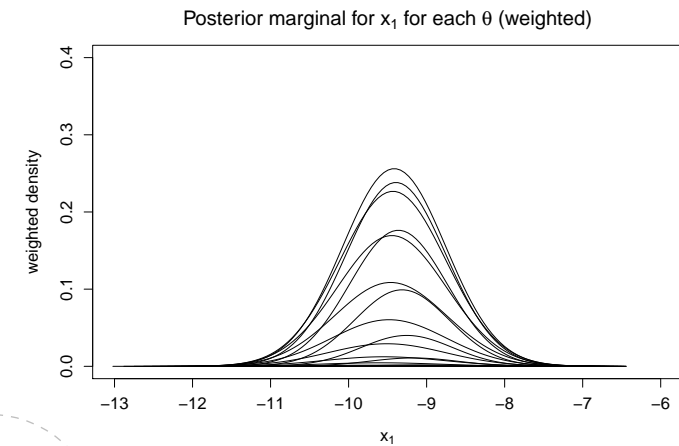
Posterior marginal for latent parameters

Compute the conditional posterior marginal for each x_i given each θ_k .



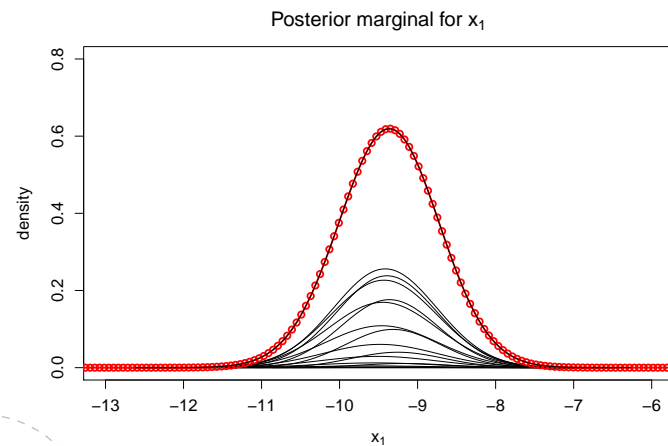
Posterior marginal for latent parameters

Weigh the resulting (conditional) marginal posteriors by the density associated with each θ_k .



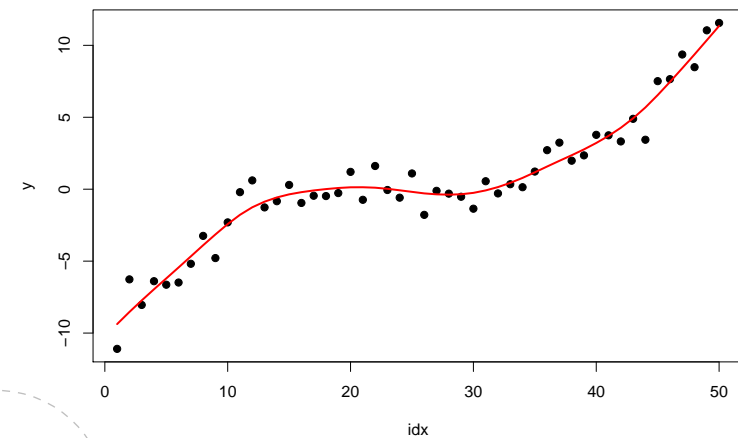
Posterior marginal for latent parameters

Numerically sum over all conditional densities to obtain the posterior marginal for each x_i .



Fitted spline

The posterior marginals are used to calculate summary statistics, like means, variances and credible intervals:



Extensions

This is the basic idea behind INLA. It is quite simple.

However, we need to extend this basic idea so we can deal with

- More than one hyperparameter
- Non-Gaussian observations

The non-Gaussian part of the model

- In many cases $\pi(\mathbf{x} | \mathbf{y}, \theta)$ is very close to a Gaussian distribution, and can be replaced with a Laplace approximation
- This means that all the really hard, high-dimensional integrals with respect to the latent field are easy, and only the integrals with respect to the hyperparameters remain
- If the number of hyperparameters is low, these integrals can be done efficiently numerically

Limitations

- The dimension of the latent field \mathbf{x} can be large (10^2 – 10^6)
- But the dimension of the hyperparameters θ must be small (≤ 9)

In other words, each random effect can be big, but there cannot be too many random effects unless they share parameters.

How to use INLA?

INLA is implemented through the package R-INLA in the R software which

- has a very user friendly `formula` interface

```
linear_model <- lm(weight ~ group)
```

Fits the linear model

$$\text{weight}_i = \mu + \text{group}_i + \epsilon_i$$

Example: Ski flying records

Steps to run INLA

1. Make an object to store responses and covariates

```
data = list(y = y, x = x)
```

2. Make a formula specifying the model

```
formula = y~x
```

3. Call INLA

```
res=inla(formula, data=data, family="gaussian")
```

Example: Summary

Call:

```
"inla(formula = formula, family = \"gaussian\", data = data)"
```

Time used:

Pre-processing	Running inla	Post-processing	Total
0.0581	0.0161	0.0181	0.0924

Fixed effects:

	mean	sd	0.025quant	0.5quant	0.975quant	mode	kld
(Intercept)	137.0288	1.3929	134.2798	137.0288	139.7741	137.0288	0
x	2.1259	0.0526	2.0221	2.1259	2.2295	2.1259	0
...							

Summary of INLA

Three main ingredients in INLA

- Latent Gaussian models
- Laplace approximations
- Gaussian Markov random fields

These ingredients give a very nice tool for Bayesian inference which is

- fast
- accurate
- scales well for moderate sizes