# Bayesian multivariate density estimation for observables and random effects 

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#### Abstract

Multivariate density estimation is approached using Bayesian nonparametric mixture of normals models. Two models are developed which are both centred over a multivariate normal distribution but make different prior assumptions about how the unknown distribution departs from a normal distribution. The priors are applied to density estimation of both observables and random effects (or other unobservable random quantities). Markov chain Monte Carlo methods are described for estimation of all models. The models are applied to density estimation for observables and the application of a nonparametric linear mixed model to repeated cholesterol measurements from the Framingham study.


Keywords: Dirichlet process mixture models, Mixtures of normals, Adaptive Monte Carlo, Centring.

## 1 Introduction

The use of Dirichlet process mixture (DPM) models in density estimation has become increasingly popular in Bayesian statistics, following the seminal work of Lo (1984)

[^0]and Ferguson (1983). The mixture of normals has become the main example and is often combined with a conjugate prior as described by Escobar and West (1995). Suppose that $y_{1}, y_{2}, \ldots, y_{n}$ are a sample of $p$-dimensional vectors. The model has the following hierarchical form for multivariate observations $y_{1}, y_{2}, \ldots, y_{n}$
\[

$$
\begin{gathered}
y_{i} \sim \mathrm{~N}\left(\theta_{i}, \Sigma_{i}\right) \\
\left(\theta_{i}, \Sigma_{i}\right) \sim G \\
G \sim \operatorname{DP}(M, H)
\end{gathered}
$$
\]

where $\mathrm{N}(\theta, \Sigma)$ represents a multivariate normal distribution with mean $\theta$ and covariance $\Sigma, \mathrm{DP}(M, H)$ represents a Dirichlet process (Ferguson, 1973) with mass parameter $M$ and centring distribution $H$. In this case, the centring distribution $H$ is chosen to be $\mathrm{N}\left(\theta \mid \theta_{0}, n_{0} \Sigma\right) \operatorname{IW}\left(\Sigma \mid \alpha_{\Sigma}, \beta_{\Sigma}\right)$ where IW represents an inverse Wishart distribution with shape parameter $\alpha_{\Sigma}$ and mean $\beta_{\Sigma} /\left(\alpha_{\Sigma}-p-1\right)$. The marginal distribution of $y_{i}$ is then an infinite mixture of normals distribution. The hyperparameter $\theta_{0}$ is straightforward to choose since it is the prior mean of $y_{i}$. However, it is less clear how $\alpha_{\Sigma}, \beta_{\Sigma}$ and $n_{0}$ should be chosen. In practice, their value has an important effect on the form of the density estimate is often chosen by a process of "trial-and-error".

In univariate density estimation with mixtures of normals, Griffin (2010) suggests a simplified version of the conjugate DPM model where

$$
\begin{gather*}
y_{i} \sim \mathrm{~N}\left(\theta_{i}, a \sigma^{2}\right), \quad 0<a<1  \tag{1}\\
\theta_{i} \sim G \\
G \sim \operatorname{DP}(M, H)
\end{gather*}
$$

and now $H=\mathrm{N}\left(\theta \mid \mu,(1-a) \sigma^{2}\right)$. This model implies that the prior predictive distribution of $y_{i}$ is normal with mean $\mu$ and variance $\sigma^{2}$. The parameter $a$ can be interpreted as a smoothness parameter with larger values of $a$ leading to a prior which places increasing mass on unimodal distributions which look increasingly normally shaped. The model does not allow different variance for each normal component, unlike the model of Escobar and West (1995), but the model approximate any distribution of $y_{i}$ since it has full support. Griffin (2010) shows that this model leads to better performance, in terms of out-of-sample prediction, than previously proposed "default" priors with mixing on both the mean and variance.

This paper is concerned with extending the model of Griffin (2010) to problems of multivariate density estimation. A useful property of the univariate model is that $\mu$ and $\sigma^{2}$ can be given standard priors (including the standard non-informative prior for the normal model) so that the nonparametric analysis will tend towards a parametric
analysis, assuming a normal distribution for $y_{i}$, as $a$ tends to 1 . This allows us to centre semiparametric models, such as linear mixed models, defined using nonparametric priors for some distributions directly over their parametric counterparts. This theme is developed in this paper.

The paper is organized as follows: Section 2 discusses two extensions of the univariate model to multivariate problems, Section 3 describes the Markov chain Monte Carlo algorithms needed to fit the models, Section 4 presents applications of the models to density estimation and semiparametric linear mixed effect models, Section 5 is a discussion.

## 2 Models

Suppose that the density of $y_{i}$ is $f(\cdot)$ then this density is our parameter of interest in density estimation. The model in (1) implies that

$$
f\left(y_{i}\right)=\sum_{i=1}^{\infty} w_{i} \mathbf{N}\left(y_{i} \mid \theta_{i}, a \sigma^{2}\right)
$$

where $w_{i}$ are the sizes of the jumps of a Dirichlet process and $\theta_{i} \sim \mathbf{N}\left(\mu,(1-a) \sigma^{2}\right)$, which defines a prior on $f(\cdot)$. The prior mean of $f(\cdot)$ is

$$
\mathrm{E}\left[f\left(y_{i}\right) \mid a, \mu, \sigma^{2}\right]=\mathrm{N}\left(y_{i} \mid \mu, \sigma^{2}\right)
$$

Therefore, the prior mean does not depend on $a$ and the prior for $f(\cdot)$ is centred over a normal distribution with mean $\mu$ and variance $\sigma^{2}$. The parameter $a$ controls the proportion of the overall variation, $\sigma^{2}$, allocated to within-component variation (given by $a \sigma^{2}$ ) and between-component variation (given by $(1-a) \sigma^{2}$ ). Larger values of $a$ lead to more variation placed into the within-component variation and lead to realized densities $f(\cdot)$ that look increasingly like a normal distribution. Therefore, the parameter can be interpret as a measure of the departure of the density of $y_{i}$ from a normal distribution with larger $a$ representing a density closer to a normal density.

The parameterisation of the model in terms of a smoothness parameter, $a$, and a fixed prior mean, $\mu$, for $f(\cdot)$ which does not depend on the smoothness parameter defines a useful model for univariate density estimation. Extending these ideas to a multivariate model, with $p$-dimensional observations $y_{i}$, would imply that the prior should have the property that

$$
\begin{equation*}
\mathrm{E}\left[f\left(y_{i}\right) \mid a, \mu, \Sigma\right]=\mathrm{N}\left(y_{i} \mid \mu, \Sigma\right) \tag{2}
\end{equation*}
$$

where $\mu$ is a $p$-dimensional mean vector, $\Sigma$ is a $p \times p$-dimensional covariance matrix and $a$ is a vector of remaining smoothness parameters. A direct extension of the model
in (1) would define $y_{i} \sim \mathrm{~N}\left(\theta_{i}, a \Sigma\right)$ and $H=\mathrm{N}(\theta \mid \mu,(1-a) \Sigma)$ leading to the required centring property in (2) but would imply that the prior for the marginal distribution of $y_{i j}$ is the model in (1) with $\sigma^{2}$ replaced by $\Sigma_{j j}, \theta$ replaced by $\theta_{j}$ and smoothness parameter $a$. The departures from normality would be the same for all variables. This will often be unrealistically simple in practice since there may be dimensions or directions in which the marginal distributions are close to normality and other dimensions or directions in which the marginal distributions are far from normal, e.g. multi-modal.

Two models will be defined which allow different levels of smoothing for different dimensions of direction of $y_{i}$. Both models have the centring property that $\mathrm{E}\left[f\left(y_{i}\right) \mid a, \mu, \Sigma\right]=\mathrm{N}\left(y_{i} \mid \mu, \Sigma\right)$. The idea that there may be some dimensions of $y_{i}$ whose marginal distribution is closer to normality than others will underlie Model 1 and the idea that there may be some directions of $y_{i}$ in which the marginal distribution is close to normality will underlie Model 2.

Model 1 defines smoothness parameters $a_{1}, a_{2}, \ldots, a_{p}$ and assumes that

$$
\begin{gathered}
y_{i} \sim \mathrm{~N}\left(y_{i} \mid \theta_{i}, A^{1 / 2} \Sigma A^{1 / 2}\right), \\
\theta_{i} \sim G, \\
G \sim \operatorname{DP}(M, H)
\end{gathered}
$$

where

$$
H=\mathrm{N}\left(\theta \mid \mu, \Sigma-A^{1 / 2} \Sigma A^{1 / 2}\right)
$$

and $A^{1 / 2}$ is a $p \times p$-dimensional matrix of the form

$$
A^{1 / 2}=\left(\begin{array}{ccc}
\sqrt{a_{1}} & & 0  \tag{3}\\
& \ddots & \\
0 & & \sqrt{a_{p}}
\end{array}\right)
$$

The model is only well-defined if both covariance matrices are positive-definite which is true if $0<a_{j}<1$ for all $j$. The covariance matrix $A^{1 / 2} \Sigma A^{1 / 2}$ is clearly positive definite (since it is a product of positive definite matrices) and so $f\left(y_{i} \mid \theta_{i}, a, \Sigma\right)$ is welldefined. The positive-definiteness of $\Sigma-A^{1 / 2} \Sigma A^{1 / 2}$ can be shown in the following way. Clearly

$$
\Sigma-A^{1 / 2} \Sigma A^{1 / 2}=\left(I-A^{1 / 2}\right) \Sigma\left(I+A^{1 / 2}\right)-\Sigma A^{1 / 2}+A^{1 / 2} \Sigma
$$

and so

$$
x^{T}\left(\Sigma-A^{1 / 2} \Sigma A^{1 / 2}\right) x=x^{T}\left(I-A^{1 / 2}\right) \Sigma\left(I+A^{1 / 2}\right) x
$$

for any vector $x$ of appropriate length. The matrix $\left(I-A^{1 / 2}\right) \Sigma\left(I+A^{1 / 2}\right)$ is clearly positive-definite since it is the product of positive-definite matrices and so $\Sigma-A^{1 / 2} \Sigma A^{1 / 2}$ is also positive-definite.

The model implies that the components of the mixture have the same correlation matrix as the correlation matrix of the data since the $(j, k)$-th term of $A^{1 / 2} \Sigma A^{1 / 2}$ is $\sqrt{a_{j} a_{k}} \Sigma_{j k}$ and so the correlation between $y_{i j}$ and $y_{i k}$ conditional on $\theta_{i}$ is $\frac{\Sigma_{i j}}{\sqrt{\Sigma_{i i} \Sigma_{j j}}}$. The prior for the marginal distribution of $y_{i j}$ follows the model in (1) with $\mu$ replaced by $\mu_{j}, \sigma^{2}$ replaced by $\Sigma_{j j}$ and $a$ replaced by $a_{j}$. Therefore, the departure from normality in dimension $j$ is controlled by $a_{j}$.


Figure 1: Four realisations of the density under Model 1 for different value of $a_{1}$ and $a_{2}$ with $M=5$ and zero correlation.

Figure 1 shows realizations from the model when $p=2$ with different values of $a_{1}$ and $a_{2}$. Clearly, the interpretation of the smoothness parameters is the same as the univariate models: larger values of $a_{k}$ imply that the distribution of $y_{i k}$ is increasingly normal and, small values of $a_{k}$ tend to lead to multi-modal marginal distributions of $y_{i k}$. The multi-modality of the joint distribution depends on the value of both $a_{1}$ and $a_{2}$. In the Figure, small values for both $a_{1}$ and $a_{2}$ lead to a distribution with many modes whereas the combination of a small value for $a_{1}$ and a large value for $a_{2}$ leads to a few modes (and similarly for large $a_{1}$ and small $a_{2}$ ). The effect of introducing correlation between $a_{1}$ and $a_{2}$ is shown in Figure 2. The distribution is


Figure 2: Four realisations of the density under Model 1 for different value of $a_{1}$ and $a_{2}$ with $M=5$ and correlation of 0.5 .
close to a normal distribution if both $a_{1}$ and $a_{2}$ are large. Small values of $a_{1}$ and $a_{2}$ leads to distributions with many modes where each component distribution have the same correlation. The effect of a small $a_{1}$ and a large $a_{2}$ is interesting and illustrates an important consequence of choosing this model. The marginal distribution of $y_{1}$ is multi-modal and the marginal distribution of $y_{2}$ is close to normality. Consequently, the joint distribution is constructed from several components with small variances in the first dimension and large variances in the second dimension.

The idea underlying Model 2 is that there are linear transforms of the data for which the marginal distribution is relatively normal and other linear transformations for which the marginal distribution is far from normal. This is achieved by defining $z_{i}=B^{-1}\left(y_{i}-\mu\right)$ and assuming that $z_{i 1}, z_{i 2}, \ldots, z_{i p}$ are independent and follow the model in (1) with $\mu=0, \sigma^{2}=1$ and smoothness parameter $a_{j}$. This implies that $z_{i}$ is centred over a multivariate normal distribution with mean $(0,0, \ldots, 0)$ and the identity matrix $I_{p}$ as its covariance matrix. Therefore, $y_{i}$ is centred over $\mathrm{N}\left(\mu, B B^{T}\right)$ and a natural choice for $B$, which will be followed in this paper, is the Cholesky


Figure 3: Four realisations of the density under Model 2 for different value of $a_{1}$ and $a_{2}$ with $M=5$ and correlation of 0.5 .
decomposition of the covariance matrix $\Sigma$. The model for $y_{i}$ is then

$$
\begin{aligned}
y_{i} & \sim \mathrm{~N}\left(\theta_{i}, B A^{\star} B^{T}\right) \\
\theta_{i} & \sim \mathrm{DP}(M H)
\end{aligned}
$$

where $H=\mathrm{N}\left(\theta \mid \mu, B B^{T}-B A^{\star} B^{T}\right)$ and $A^{*}$ is a diagonal matrix with non-zero elements $a_{1}, a_{2}, \ldots, a_{p}$.

Model 1 and Model 2 will be the same if the correlation between $y_{i 1}$ and $y_{i 2}$ is zero. However, they differ in terms of how correlation is introduced. Unlike Model 1, Model 2 rotates the distribution if the correlation is non-zero. Realisations of this prior with a correlation of 0.5 are shown in Figure 3. Clearly small values of $a_{1}$ and $a_{2}$ lead to distributions with many modes and typically well-seperated components. The value of $a=0.2$ and larger give rise to distribution with less modes and a more cohesive distribution.

## 3 Density estimation in random effect models

The distribution of unobserved quantities, such as random effects, will often have an unknown form. Bayesian approaches to density estimation can be naturally extended to this problem by assuming a nonparametric prior for the unknown distribution and so defines a semiparametric model. The Linear Mixed Model (LMM) will be the main focus of interest in this section although the ideas extend naturally to other models with random effects. The usual Normal LMM assumes that responses $y_{i 1}, \ldots, y_{i T}$ are observed for the $i$-th individual with regressors $X_{i 1}, \ldots, X_{i T}$ and $Z_{i 1}, \ldots, Z_{i T}$ where

$$
\begin{gathered}
y_{i t}=\alpha+X_{i t} \beta+Z_{i t} \gamma_{i}+\epsilon_{i t}, \\
\gamma_{i} \stackrel{i . i . d .}{\sim} \mathrm{N}(\mu, \Sigma)
\end{gathered}
$$

and $\epsilon_{i t}$ are mutually independent and $\epsilon_{i t} \sim \mathrm{~N}\left(0, \sigma^{2}\right)$. The model makes the distinction that $\beta$ are the same for all individuals and so $X_{i t}$ has the same effect for all individuals whereas $\gamma_{i}$ varies from individual-to-individual and so allows for different effects of $Z_{i t}$ across individuals. The parameters of the model are then given priors which often have the form: $\beta \sim \mathrm{N}\left(0, \Sigma_{\beta}\right), \alpha \sim \mathrm{N}\left(0, \sigma_{\alpha}\right), \mu \sim \mathrm{N}\left(0, \Sigma_{\mu}\right)$ and $\Sigma \sim \operatorname{IW}(\eta, \Upsilon)$. The main focus of interest is usually the regression parameters $\beta$ and the hyperparameters $\mu$. A nonparametric specification for the distribution of $\gamma_{i}$ defines a semiparametric model and leads to robust estimates of the regression coefficients.Initial work on Bayesian nonparametric estimation in these models is described by Bush and MacEachern (1996) and Kleinman and Ibrahim (1998). These assume that the distribution of $\gamma_{i}$ follows a Dirichlet process mixture model. In this paper, specific forms of Dirichlet process mixture model (Model 1 and Model 2) are used to define LMM-Model 1 and LMM-Model 2. LMM-Model 1 assumes

$$
\begin{gathered}
\gamma_{i} \sim \mathrm{~N}\left(\theta_{i}, A^{1 / 2} \Sigma A^{1 / 2}\right) \\
\theta_{i} \sim \operatorname{DP}(M H)
\end{gathered}
$$

where $H=\mathrm{N}\left(\theta \mid \mu, \Sigma-A^{1 / 2} \Sigma A^{1 / 2}\right), A$ is defined in (3) and $\beta, \alpha, \mu$ and $\Sigma$ are given the priors for the Normal LMM. The model conditional on $\beta, \alpha, \mu$ and $\Sigma$ is centred over the normal model and the nonparametric model is the same as the parametric model model if $a_{1}, a_{2}, \ldots, a_{p}$ are all equal to one. Similarly, LMM-Model 2 can be defined by assuming that

$$
\begin{gathered}
\gamma_{i} \sim \mathrm{~N}\left(\theta_{i}, B A^{\star} B^{T}\right) \\
\theta_{i} \sim \mathrm{DP}(M H)
\end{gathered}
$$

where $H=\mathrm{N}\left(\theta \mid \mu, B B^{T}-B A^{\star} B^{T}\right), A^{\star}$ is a diagonal matrix with non-zero elements $a_{1}, a_{2}, \ldots, a_{p}$ and $\beta, \alpha, \mu$ and $\Sigma$ are given the priors for the Normal LMM. Once again
the model conditional on $\beta, \alpha, \mu$ and $\Sigma$ is centred over the normal model with mean $\mu$ and variance-covariance matrix $\Sigma=B B^{T}$ and the nonparametric model is the same as the parametric model model if $a_{1}, a_{2}, \ldots, a_{p}$ are all equal to one.

In both models inference about $a_{1}, a_{2}, \ldots, a_{p}$ provides information about the dimensions in which the distribution departs from normality. This is indicated by small values of $a_{i}$. Larger values of $a_{i}$ will lead to an analysis which is closer to the parametric analysis and so allows us to take advantage of the normal assumption if it is supported by the data.

## 4 Computational Methods

Both Model 1 and Model 2 are conjugate Dirichlet process mixture models and so can be fitted using standard Pólya urn scheme methods described in MacEachern (1998). The choice of a Dirichlet process prior implies there are $K$ distinct values of $\theta_{1}, \theta_{2}, \ldots, \theta_{n}$ which will be denoted $\theta_{(1)}, \theta_{(2)}, \ldots, \theta_{(K)}$. Latent variables $s_{1}, s_{2}, \ldots, s_{n}$ are introduced to link these values so that $\theta_{i}=\theta_{\left(s_{i}\right)}$.

### 4.1 Model 1

The parameter $\mu$ and $\Sigma$ are assumed to have the priors $\mu \sim \mathrm{N}\left(\mu_{0}, \Sigma_{\mu}\right)$ and $\Sigma \sim$ $\operatorname{IW}(\eta, \Upsilon)$. Let $\Sigma_{j}=A^{1 / 2} \Sigma A^{1 / 2}, \Sigma_{0}=\Sigma-A^{1 / 2} \Sigma A^{1 / 2}$ and $n_{j}=\#\left\{i \mid s_{i}=j\right\}$. The full conditional distributions are as follows.

## Updating $s$

Let $K^{-}$be the number of distinct values for $\theta_{1}, \ldots \theta_{i-1}, \theta_{i+1}, \ldots, \theta_{n}$ and $n_{j}^{-}=$ $\#\left\{k \mid s_{k}=j, k \neq i\right\}$. The full conditional distribution of $s_{i}$ is

$$
p\left(s_{i}=j\right) \propto n_{j} \mathrm{~N}\left(y_{i} \mid \theta_{j}^{\star}, \Sigma_{j}^{\star}\right), \quad j=1,2, \ldots, K^{-}
$$

and

$$
p\left(s_{i}=K^{-}+1\right) \propto M \mathrm{~N}\left(y_{i} \mid \mu, \Sigma\right)
$$

where

$$
\theta_{j}^{\star}=\left(n_{j}^{-} \Sigma_{k}^{-1}+\Sigma_{0}^{-1}\right)^{-1}\left(\Sigma_{k}^{-1} \sum_{\left\{k \mid s_{k}=j, k \neq i\right\}} y_{k}+\Sigma_{0}^{-1} \mu_{0}\right)
$$

and

$$
\Sigma_{j}^{\star}=\left(n_{j}^{-} \Sigma_{k}^{-1}+\Sigma_{0}^{-1}\right)^{-1}
$$

After updating $s_{1}, s_{2}, \ldots, s_{n}$, the values of $\theta_{(1)}, \theta_{(2)}, \ldots, \theta_{(K)}$ are sampled from their full conditional distribution $\theta_{(j)} \sim \mathrm{N}\left(\theta_{j}^{\star}, \Sigma_{j}^{\star}\right)$ where

$$
\theta_{j}^{\star}=\left(n_{j} \Sigma_{k}^{-1}+\Sigma_{0}^{-1}\right)^{-1}\left(\Sigma_{k}^{-1} \sum_{\left\{k \mid s_{k}=j\right\}} y_{k}+\Sigma_{0}^{-1} \mu_{0}\right)
$$

and

$$
\Sigma_{j}^{\star}=\left(n_{j} \Sigma_{k}^{-1}+\Sigma_{0}^{-1}\right)^{-1}
$$

## Updating $\mu$

The full conditional distribution of $\mu$ is $\mathrm{N}\left(\mu \mid \mu^{\dagger}, \Sigma^{\dagger}\right)$ where

$$
\begin{gathered}
\mu^{\dagger}=\left(K \Sigma_{0}^{-1}+\Sigma_{\mu}^{-1}\right)^{-1}\left(\Sigma_{0}^{-1} \sum_{i=1}^{K} \theta_{(i)}+\Sigma_{\mu}^{-1} \mu_{0}\right) \\
\Sigma^{\dagger}=\left(K \Sigma_{0}^{-1}+\Sigma_{\mu}^{-1}\right)^{-1}
\end{gathered}
$$

## Updating $\Sigma$ and $a$

The covariance matrix $\Sigma$ and $a$ can be updated in the following way. We reparametrise $\Sigma$ to $\left(P, \sigma_{1}^{2}, \ldots, \sigma_{p}^{2}\right)$ where $P$ is the $(p \times p)$-dimensional correlation matrix with terms

$$
P_{i j}=\frac{\Sigma_{i j}}{\Sigma_{11}^{1 / 2} \Sigma_{22}^{1 / 2}}
$$

and $\sigma_{j}^{2}=\Sigma_{j j}$ and update these parameters using a Metropolis-Hastings random walk. The full conditional distribution is $p(\Sigma) \propto \Sigma_{k}^{-n / 2} \Sigma_{0}^{K / 2} \exp \left\{-\frac{1}{2}\left[\sum_{i=1}^{n}\left(y_{i}-\theta_{i}\right)^{T} \Sigma_{k}^{-1}\left(y_{i}-\theta_{i}\right)^{T}+\sum_{j=1}^{K}\left(\theta_{(j)}-\mu\right)^{T} \Sigma_{0}^{-1}\left(\theta_{(j)}-\mu\right)\right]\right\}$.

The value of $\sigma_{j}^{2}$ is updated with $a_{j}$. The new value of $\sigma_{j}^{2}$ is proposed as $\sigma_{j}^{2 \prime}=$ $\sigma_{j}^{2} \exp \left\{\phi_{j}^{2} \epsilon\right\}$ where $\epsilon \sim \mathrm{N}(0,1)$ and the new value of $a_{j}$ is proposed as $a_{j}^{\prime}=\frac{\left(1+a_{j}\right) /\left(1-a_{j}\right) \exp \left\{\nu_{j} \epsilon\right\}-1}{\left(1+a_{j}\right) /\left(1-a_{j}\right) \exp \left\{\nu_{j} \epsilon\right\}+1}$.
Let $\Sigma^{\prime}$ be the values of the covariance matrix with $\sigma_{j}^{2}$ and $a_{j}$ replaced by $\sigma_{j}^{2 \prime}$ and $a_{j}^{\prime}$.
The new values is accepted with probability
$\max \left\{1, \frac{p\left(a_{j}^{\prime}\right) p\left(\Sigma^{\prime}\right)\left|\Sigma^{\prime}\right|^{(\eta+p+1) / 2} \exp \left\{-\operatorname{tr}\left(\Upsilon \Sigma^{\prime-1}\right) / 2\right\} \sigma_{j}^{2^{\prime}(p+1) / 2}\left(\frac{1}{1+a_{j}^{\prime}}+\frac{1}{1-a_{j}^{\prime}}\right)}{p\left(a_{j}\right) p(\Sigma)|\Sigma|^{(\eta+p+1) / 2} \exp \left\{-\operatorname{tr}\left(\Upsilon \Sigma^{-1}\right) / 2\right\} \sigma_{j}^{2(p+1) / 2}\left(\frac{1}{1+a_{j}}+\frac{1}{1-a_{j}}\right)}\right\}$.

The new value of $P_{i j}$ is proposed as $P_{i j}^{\prime}=\frac{\left(1+P_{i j}\right) /\left(1-P_{i j}\right) \exp \left\{\zeta_{i j} \epsilon\right\}-1}{\left(1+P_{i j}\right) /\left(1-P_{i j}\right) \exp \left\{\zeta_{i j} \epsilon\right\}+1}$. Let $\Sigma^{\prime}$ be the values of the covariance matrix with $P_{i j}$ replaced by $P_{i j}^{\prime}$. The new value is accepted with probability

$$
\max \left\{1, \frac{p\left(\Sigma^{\prime}\right)\left|\Sigma^{\prime}\right|^{(\eta+p+1) / 2} \exp \left\{-\operatorname{tr}\left(\Upsilon \Sigma^{\prime-1}\right) / 2\right\}\left(\frac{1}{1+P_{i j}^{\prime}}+\frac{1}{1-P_{i j}^{\prime}}\right)}{p(\Sigma)|\Sigma|^{(\eta+p+1) / 2} \exp \left\{-\operatorname{tr}\left(\Upsilon \Sigma^{-1}\right) / 2\right\}\left(\frac{1}{1+P_{i j}}+\frac{1}{1-P_{i j}}\right)}\right\}
$$

The variance of the increments $\zeta_{i j}, \nu_{j}$ and $\phi_{j}$ are tuned using the adaptive scheme of Atchadé and Rosenthal (2005) with a target acceptance rate of 0.3.

## Updating $M$

The full conditional distribution for $M$ is

$$
p(M) \frac{\Gamma(M)}{\Gamma(M+n)} M^{K}
$$

and can be updated using a Metropolis-Hastings random walk step. The variance of the increments is tuned using the adaptive scheme of Atchade and Rosenthal (2005).

### 4.2 Model 2

We define $\Sigma=B B^{T}$ and assume the priors $\mu \sim \mathrm{N}\left(\mu_{0}, \Sigma_{\mu}\right)$ and $\Sigma \sim \operatorname{IW}\left(\eta, \Sigma_{0}\right)$. Model 2 can be updated in the same way as Model 1 with $\Sigma_{k}=B A^{1 / 2} A^{1 / 2} B^{T}$ and $\Sigma_{0}=B B^{T}-B A^{1 / 2} A 1 / 2 B^{T}$.

### 4.3 LMM-Model 1

The model in Section 3 can be fitted using a Gibbs sampler with the updating scheme as follows. The parameters $s$ and $\mu$ are updated marginalizing over $\gamma$ which lead to much better mixing of the sampler than the Gibbs sampler including $\gamma$ at all steps.

## Updating $\beta$

The full conditional distribution of $\beta$ is

$$
\mathrm{N}\left(\left(\sigma^{-2} \sum_{i=1}^{n} \sum_{t=1}^{T} X_{i t}^{\prime} X_{i t}+\Lambda_{\beta}\right)^{-1} \sigma^{-2} \sum_{i=1}^{n} \sum_{t=1}^{T} X_{i t}^{\prime}\left(y_{i t}-Z_{i t} \gamma_{i}\right),\left(\sigma^{-2} \sum_{i=1}^{n} \sum_{t=1}^{T} X_{i t}^{\prime} X_{i t}+\Lambda_{\beta}\right)^{-1}\right)
$$

## Updating $\sigma^{2}$

The full conditional distribution of $\sigma^{2}$ is

$$
\mathrm{Ga}\left(\alpha+n T, \beta+\sum_{i=1}^{n} \sum_{t=1}^{T}\left(y_{i}-X_{i t} \beta-Z_{i t} \gamma_{i}\right)^{2}\right)
$$

## Updating $s$

The full conditional distribution of $s_{i}$ is

$$
p\left(s_{i}=j\right) \propto n_{j} \mathrm{~N}\left(y_{i}^{\star} \mid Z_{i} \theta_{(j)}, Z_{i} A^{1 / 2} \Sigma A^{1 / 2} Z_{i}^{T}+\sigma^{2}\right), \quad j=1,2, \ldots, K^{-}
$$

and

$$
p\left(s_{i}=K^{-}+1\right) \propto M \mathrm{~N}\left(y_{i}^{\star} \mid Z_{i} \mu, Z_{i} \Sigma Z_{i}^{T}+\sigma^{2}\right)
$$

where

$$
y_{i k}^{\star}=y_{i k}-X_{i k} \beta
$$

If $s_{i}=K^{-}+1$ then $\theta_{\left(K^{-}+1\right)}$ is sampled from $\mathrm{N}\left(\theta^{\star}, \Sigma^{\star}\right)$ where

$$
\theta^{\star}=\left(Z_{i}^{\prime}\left(Z_{i} \Sigma_{k} Z_{k}^{T}+\sigma^{-2} I_{T}\right)^{-1} Z_{i}+\Sigma_{0}^{-1}\right)^{-1}\left(Z_{i}^{\prime}\left(Z_{i} \Sigma_{k} Z_{k}^{T}+\sigma^{-2} I_{T}\right)^{-1} y_{i}+\Sigma_{0}^{-1} \mu\right)
$$

and

$$
\Sigma^{\star}=\left(Z_{i}^{\prime}\left(Z_{i} \Sigma_{k} Z_{k}^{T}+\sigma^{-2} I_{T}\right)^{-1} Z_{i}+\Sigma_{0}^{-1}\right)^{-1}
$$

where $\Sigma_{k}=A^{1 / 2} \Sigma A^{1 / 2}$ and $\Sigma_{0}=\Sigma-A^{1 / 2} \Sigma A^{1 / 2}$

## Updating $\theta$

The full conditional distribution of $\theta_{(k)}$ is $\mathrm{N}\left(\theta^{\star}, \Sigma^{\star}\right)$ where

$$
\theta^{\star}=\left(\sum_{i \mid s_{i}=k} Z_{i}^{\prime}\left(Z_{i} \Sigma_{k} Z_{k}^{T}+\sigma^{-2} I_{T}\right)^{-1} Z_{i}+\Sigma_{0}^{-1}\right)^{-1}\left(\sum_{i \mid s_{i}=k} Z_{i}^{\prime}\left(Z_{i} \Sigma_{k} Z_{k}^{T}+\sigma^{-2} I_{T}\right)^{-1} y_{i}+\Sigma_{0}^{-1} \mu\right)
$$

and

$$
\Sigma^{\star}=\left(\sum_{i \mid s_{i}=k} Z_{i}^{\prime}\left(Z_{i} \Sigma_{k} Z_{k}^{T}+\sigma^{-2} I_{T}\right)^{-1} Z_{i}+\Sigma_{0}^{-1}\right)^{-1}
$$

## Updating $\gamma$

The full conditional distribution for $\gamma_{i}$ is

$$
\mathrm{N}\left(\left(\sigma^{-2} \sum_{t=1}^{T} Z_{i t}^{\prime} Z_{i t}+\Sigma_{k}^{-1}\right)^{-1}\left(\sigma^{-2} \sum_{t=1}^{T} Z_{i t}\left(y_{i t}-X i t \beta\right)+\Sigma_{k}^{-1} \mu_{s_{i}}\right),\left(\sigma^{-2} \sum_{t=1}^{T} Z_{i t}^{\prime} Z_{i t}+\Sigma_{k}^{-1}\right)^{-1}\right)
$$

## Updating $\Sigma$ and $a$

The covariance matrix $\Sigma$ and $a$ can be updated in the following way. We reparametrise $\Sigma$ to $\left(P, \sigma_{1}^{2}, \ldots, \sigma_{p}^{2}\right)$ where $P$ is the $(p \times p)$-dimensional correlation matrix with terms

$$
P_{i j}=\frac{\Sigma_{i j}}{\Sigma_{11}^{1 / 2} \Sigma_{22}^{1 / 2}}
$$

and $\sigma_{j}^{2}=\Sigma_{j j}$ and update these parameters using a Metropolis-Hastings random walk. The full conditional distribution is $p(\Sigma) \propto \Sigma_{k}^{-n / 2} \Sigma_{0}^{K / 2} \exp \left\{-\frac{1}{2}\left[\sum_{i=1}^{n}\left(\gamma_{i}-\theta_{i}\right)^{T} \Sigma_{k}^{-1}\left(\gamma_{i}-\theta_{i}\right)^{T}+\sum_{j=1}^{K}\left(\theta_{(j)}-\mu\right)^{T} \Sigma_{0}^{-1}\left(\theta_{(j)}-\mu\right)\right]\right\}$.

The value of $\sigma_{j}^{2}$ is updated with $a_{j}$. The new value of $\sigma_{j}^{2}$ is proposed as $\sigma_{j}^{2 \prime}=$ $\sigma_{j}^{2} \exp \left\{\phi_{j}^{2} \epsilon\right\}$ where $\epsilon \sim \mathrm{N}(0,1)$ and the new value of $a_{j}$ is proposed as $a_{j}^{\prime}=\frac{\left(1+a_{j}\right) /\left(1-a_{j}\right) \exp \left\{\nu_{j} \epsilon\right\}-1}{\left(1+a_{j}\right) /\left(1-a_{j}\right) \exp \left\{\nu_{j} \epsilon\right\}+1}$. Let $\Sigma^{\prime}$ be the values of the covariance matrix with $\sigma_{j}^{2}$ and $a_{j}$ replaced by $\sigma_{j}^{2 \prime}$ and $a_{j}^{\prime}$.
The new values is accepted with probability
$\max \left\{1, \frac{p\left(a_{j}^{\prime}\right) p\left(\Sigma^{\prime}\right)\left|\Sigma^{\prime}\right|^{(\eta+p+1) / 2} \exp \left\{-\operatorname{tr}\left(\Upsilon \Sigma^{\prime-1}\right) / 2\right\} \sigma_{j}^{2^{\prime}(p+1) / 2}\left(\frac{1}{1+a_{j}^{\prime}}+\frac{1}{1-a_{j}^{\prime}}\right)}{p\left(a_{j}\right) p(\Sigma)|\Sigma|^{(\eta+p+1) / 2} \exp \left\{-\operatorname{tr}\left(\Upsilon \Sigma^{-1}\right) / 2\right\} \sigma_{j}^{2(p+1) / 2}\left(\frac{1}{1+a_{j}}+\frac{1}{1-a_{j}}\right)}\right\}$.
The new value of $P_{i j}$ is proposed as $P_{i j}^{\prime}=\frac{\left(1+P_{i j}\right) /\left(1-P_{i j}\right) \exp \left\{\zeta_{i j} \epsilon\right\}-1}{\left(1+P_{i j}\right) /\left(1-P_{i j}\right) \exp \left\{\left\{_{i j} \epsilon\right\}+1\right.}$. Let $\Sigma^{\prime}$ be the values of the covariance matrix with $P_{i j}$ replaced by $P_{i j}^{\prime}$. The new value is accepted with probability

$$
\max \left\{1, \frac{p\left(\Sigma^{\prime}\right)\left|\Sigma^{\prime}\right|^{(\eta+p+1) / 2} \exp \left\{-\operatorname{tr}\left(\Upsilon \Sigma^{\prime-1}\right) / 2\right\}\left(\frac{1}{1+P_{i j}^{\prime}}+\frac{1}{1-P_{i j}^{\prime}}\right)}{p(\Sigma)|\Sigma|^{(\eta+p+1) / 2} \exp \left\{-\operatorname{tr}\left(\Upsilon \Sigma^{-1}\right) / 2\right\}\left(\frac{1}{1+P_{i j}}+\frac{1}{1-P_{i j}}\right)}\right\} .
$$

The variance of the increments $\zeta_{i j}, \nu_{j}$ and $\phi_{j}$ are tuned using the adaptive scheme of Atchadé and Rosenthal (2005) with a target acceptance rate of 0.3.

## Updating $M$

The full conditional distribution for $M$ is

$$
p(M) \frac{\Gamma(M)}{\Gamma(M+n)} M^{K}
$$

and can be updated using a Metropolis-Hastings random walk step. The variance of the increments is tuned using the adaptive scheme of Atchadé and Rosenthal (2005).

## Updating $\mu$

The full conditional distribution for $\mu$ is

$$
\mathrm{N}\left(\left(K \Sigma_{0}^{-1}+\Sigma_{\mu}^{-1}\right)^{-1}\left(\Sigma_{0}^{-1} \sum_{k=1}^{K} \theta_{(k)}+\Sigma_{\mu}^{-1} \mu_{0}\right),\left(K \Sigma_{0}^{-1}+\Sigma_{\mu}^{-1}\right)^{-1}\right) .
$$

### 4.4 LMM-Model 2

The sampler for LMM-Model 1 can be used with $\Sigma_{k}=B A^{1 / 2} A^{1 / 2} B^{T}$ and $\Sigma_{0}=$ $B B^{T}-B A^{1 / 2} A^{1 / 2} B^{T}$.

## 5 Examples

The models for Bayesian density estimation are applied to two problems in bivariate density estimation for observables, and a simulated and a real data example for unobservable random effects in a linear mixed model.

### 5.1 Bivariate density estimation: Old faithfull data

Data measuring the waiting time between eruptions and the duration of the eruption of the Old Faithful geyser in Yellowstone National Park are available in the R package 'datasets' and have become a popular data set for bivariate density estimation. Model 1 and Model 2 were fitted with two prior settings for $a_{i}$. The first prior setting assumes that $a_{i}$ follows a uniform distribution, which is a natural default choice for parameters on $(0,1)$, and will be written as $\operatorname{Be}(1,1)$, a Beta distribution with both parameters equal to ones. The second choice is $a_{i} \sim \operatorname{Be}(1,10)$, a Beta distribution with parameters 1 and 10 , which places more mass on small values of $a_{i}$ and so more mass is placed on distributions which are far from a normal distribution (Griffin, 2010). The parameters $\mu$ and $\Sigma$ are given the priors $\mu \sim \mathrm{N}\left(\bar{y}, 10^{4}\right)$ and $\Sigma \sim \operatorname{IW}(p+2, \hat{\Sigma})$ where $\bar{y}$ and $\hat{\Sigma}$ are the sample mean and variance-covariance matrix of the data. This choice ensures that the prior mean of $\Sigma$ is $\hat{\Sigma}$.

The posterior mean density of the observations and the data are shown in Figure 4 For all models and all priors, the estimated density is bimodal with one mode around $(2,55)$ and another mode around $(4.5,80)$ and follows the data. The density around $(4.5,80)$ is clearly skewed. The posterior distribution of $a_{1}$ and $a_{2}$ for Model 1 are summarized in Table 1. The posterior median of $a_{1}$ is much smaller than those of $a_{2}$ for both priors. The results indicate that the marginal density of duration is close to normal but the marginal density of waiting time is far from normal. Results for $a_{1}$ and


Figure 4: Contour plots of the posterior mean density under Models 1 and 2 with two different priors for $a_{1}$ and $a_{2}$ for the Old Faithfull data (observations are shown as dots).

|  | $a_{1}$ | $a_{2}$ |
| :---: | :---: | :---: |
| $\operatorname{Be}(1,1)$ | $0.12(0.04,0.24)$ | $0.31(0.14,0.51)$ |
| $\operatorname{Be}(1,10)$ | $0.09(0.03,0.19)$ | $0.20(0.08,0.37)$ |

Table 1: The posterior distribution of $a_{1}$ and $a_{2}$ summarized as the posterior median and 95\% credible interval for the Old Faithfull data using Model 1 with two different priors for $a_{1}$ and $a_{2}$.
$a_{2}$ are harder to interpret in Model 2 since the prior and likelihood is invariant to the permutation of $a_{1}$ and $a_{2}$ and no results are presented.

### 5.2 Bivariate density estimation: Aircraft data

Bowman and Azzalini (1997) describe bivariate density estimation for data related to a study of the development of aircraft technology originally analysed by Saviotti and Bowman (1984). This will be re-analyzed using the methods described in this paper. The data contain six characteristics (total engine power, wing span, length, maximum take-off weight, maximum speed and range) of aircraft designs. The first two principal
components are shown in figure and can be interpreted as "size" and "speed adjusted for size". Further details are given in the reference. The priors for $a_{i}, \mu$ and $\Sigma$ were chosen in the same way as in the example in Section 5.1.


Figure 5: Contour plots of the posterior mean density under Models 1 and 2 with two different priors for $a_{1}$ and $a_{2}$ for the aircraft data (observations are shown as dots).

Figure 5 shows a scatterplot of the data and the posterior mean density of the observations across both models and prior settings for $a_{i}$. The results are robust to these choices. The posterior mean gives a good description of the data. In particular, the higher density of points around -4 for the first dimension $x_{1}$ is well captured. The posterior distributions of $a_{1}$ and $a_{2}$ for Model 1 under the two priors for $a_{i}$ are

|  | $a_{1}$ | $a_{2}$ |
| :---: | :---: | :---: |
| $\operatorname{Be}(1,1)$ | $0.11(0.06,0.24)$ | $0.10(0.05,0.20)$ |
| $\operatorname{Be}(1,10)$ | $0.09(0.04,0.19)$ | $0.08(0.04,0.15)$ |

Table 2: The posterior distribution of $a_{1}$ and $a_{2}$ summarized as the posterior median and $95 \%$ credible interval for the aircraft data using Model 1 with two different priors for $a_{1}$ and $a_{2}$.
summarised in Table 2. The posterior medians for both $a_{1}$ and $a_{2}$ are close to 0.1
indicating that there are similar levels of non-normality in both variables, as illustrated by the posterior mean densities.

### 5.3 Nonparametric linear mixed model: Simulated Data

The nonparametric linear mixed model of Section 3 was applied to a simulated data set. The design of the simulated data sets follows from Ghidey et al. (2004). It is assumed that observations, $y_{i 1}, y_{i 2}, \ldots, y_{i 6}$ are made at 6 time points $t_{i}=(0,2,4,6,8,10)$ and that

$$
Y_{i j}=\beta_{0}+\beta_{1} t_{i j}+\gamma_{0 i}+\gamma_{1 i} t_{i j}+\epsilon_{i j}
$$

where $\epsilon_{i j} \sim \mathrm{~N}(0,0.04)$. This linear growth model where the intercept and slope are assumed to be different for each individual. Data were generated from this model with $\beta_{0}=2.35$ and $\beta_{1}=0.28$. Two distributions were assumed for the random effect $\gamma=\left(\gamma_{0}, \gamma_{1}\right):$

- Example 1

$$
\gamma \sim \mathrm{N}\left(\binom{0}{0},\left(\begin{array}{cc}
0.15 & 0.02 \\
0.02 & 0.04
\end{array}\right)\right)
$$

- Example 2

$$
\gamma \sim 0.5 \times \mathrm{N}\left(\binom{-1}{0},\left(\begin{array}{cc}
0.15 & 0.02 \\
0.02 & 0.04
\end{array}\right)\right)+0.5 \times \mathrm{N}\left(\binom{1}{0},\left(\begin{array}{cc}
0.15 & 0.02 \\
0.02 & 0.04
\end{array}\right)\right)
$$

Example 1 assumes a normal distribution for the random effects whereas Example 2 assumes that they are drawn from a mixture of two normal distributions. The marginal distribution of $b_{0}$ in Example 2 is bi-modal whereas the marginal distribution of $b_{1}$ is normal. A sample was generated from the two examples with $n=200$.

The posterior mean density under Models 1 and 2 for Example 1, shown in Figure 6 , is a good estimate of the true random effects distribution. In Model 1, the posterior medians of $a_{1}$ and $a_{2}$ are 0.21 and 0.17 respectively indicating that both marginal distributions are close to normal. The estimates of the regression parameters $\beta_{0}$ and $\beta_{1}$, shown in Table 3, are very similar for the parametric model (which is correctly specified) and both nonparametric models. Therefore, the two nonparametric models can replicate the parametric analysis when it is supported by the data.

The posterior means of the unknown density under Models 1 and 2 for Example 2 are shown in Figure 7. Both estimates show the bi-modal random effects distribution generating the data. The posterior medians of $a_{1}$ and $a_{2}$ are 0.09 and 0.21 indicating that the marginal distribution of $b_{1}$ is less normally distributed than $b_{2}$ and that $b_{2}$ is close to normally distributed which both match the properties of the model used to


Figure 6: Contour plots of the posterior mean density of $\gamma$ under Models 1 and 2 with two different priors for $a_{1}$ and $a_{2}$ for Example 1.
generate the data. The esimated regression coefficients for Example 2 are shown in Table 4. Both two nonparametric models and the parametric models show very similar results for the posterior median. However, the parametric model underestimates the uncertainty in the estimate of $\beta_{0}$, since it assumes a normal distribution for $b_{0}$ rather than the true bi-modal distribution.

### 5.4 Nonparametric linear mixed model: Cholesterol Data

The use of the model for density estimation in a linear mixed model are illustrated using repeated cholesterol measurements for a sample of 200 subjects from the Framingham study. The data was originally analysed by Zhang and Davidian (2001) and has been subsequently re-analysed by Ghidey et al. (2004) and Ho and Hu (2008). The measurements for taken every 2 years for a period of 10 years. There are two fixed effects, age and sex, and the intercept and the effect of time are assumed to be random effects drawn from a bivariate distribution. Assuming a nonparametric specification for the distribution leads to a semiparametric model. The full model can be written

$$
Y_{i j}=\beta_{0 i}+\beta_{1} \operatorname{age}_{i}+\beta_{2} \operatorname{sex}_{i}+\beta_{3 i} t_{i j}+\epsilon_{i j}
$$

| Model 1 |  |  |
| :---: | :---: | :---: |
|  $\operatorname{Be}(1,10)$ $\operatorname{Be}(1,1)$ <br> $\beta_{0}$ $2.34(2.28,2.40)$ $2.34(2.29,2.40)$ <br> $\beta_{1}$ $0.28(0.25,0.31)$ $0.28(0.25,0.31)$ <br> $a_{1}$ $0.209(0.055,0.464)$ $0.706(0.240,0.980)$ <br> $a_{2}$ $0.175(0.046,0.423)$ $0.612(0.158,0.972)$ |  |  |


| Model 2 |  |  |
| :---: | :---: | :---: |
|  | $\operatorname{Be}(1,10)$ | $\operatorname{Be}(1,1)$ |
| $\beta_{0}$ | $2.34(2.29,2.40)$ | $2.34(2.29,2.40)$ |
| $\beta_{1}$ | $0.28(0.25,0.31)$ | $0.28(-0.25,0.31)$ |


| Parametric |  |
| :--- | :--- |
| $\beta_{0}$ | $2.35(2.29,2.40)$ |
| $\beta_{1}$ | $0.28(0.25,0.31)$ |

Table 3: The posterior distribution of the regression coefficients summarized as the posterior median and $95 \%$ credible interval for Example 1 using parametric Normal model and Models 1 and Model 2 with two different priors for $a_{1}$ and $a_{2}$.

$$
\binom{\beta_{0 i}}{\beta_{3 i}} \sim F
$$

where $Y_{i j}$ is the $j$-th observation for the $i$-th subject, $t_{i j}$ is the time of the measurement, $\operatorname{age}_{i}$ and sex ${ }_{i}$ are the measured covariates for the $i$-th subject. It is assumed that $\epsilon_{i j} \stackrel{i . i . d .}{\sim} \mathrm{N}\left(0, \sigma^{2}\right)$ and $F$ is assumed to follow Model 1 or Model 2. The posterior means of $F$ under the different models and different prior settings are shown in Figure 8 . The estimated distribution show a positive correlation between the intercept and the slope. The distribution is not close to normally distributed with the iso-contour not so tightly packed for larger values of the intercept. The posterior means of the marginal distributions for the intercept and slope are shown in Figure 9. The intercept has a clear positive skewness for all models and all priors whereas the slope has a negative skewness which becomes more pronounced for the $\operatorname{Be}(1,10)$ prior.

The estimated regression coefficients for the data under the two models with the two beta priors are shown in Table 5. The estimates for the slope under the two models for both prior setting are similar and are relatively robust. The posterior median of the intercept parameter varies more across the different models and differ from the value for the parametric analysis. The $\operatorname{Be}(1,1)$ prior for both models led to results closer to the value for the parametric analysis. A little surprisingly, the posterior median of


Figure 7: Contour plots of the posterior mean density of $\gamma$ under Models 1 and 2 with two different priors for $a_{1}$ and $a_{2}$ for Example 2.
$a_{1}$ is larger than $a_{2}$ showing that the marginal distribution of the intercept is closer to normality than the marginal distribution of the slope. This is due to heavier than normal tails in the posterior mean distributions of $a_{2}$. The effect of age has very similar posterior median values across all models fitted to the data. However, there is difference between the estimate of the effect of sex under the nonparametric models, which have posterior medians close to -0.06 , and the parametric model, which has a median of -0.041 .

## 6 Discussion

This paper has presented methods for Bayesian nonparametric density estimation for multivariate data and a semiparametric linear mixed model with an unknown random effects distribution. The prior for the unknown distribution is centred over a normal distribution, in the sense that the prior predictive distribution of an observation $y$ conditional on $\mu$ and $\Sigma$ is a normal distribution with mean $\mu$ and variance $\Sigma$. Two specifications are considered. One assumes parameters that measure the departure from normalty in each dimensions. The second assumes that a rotated version of $y$ is

| Model 1 |  |  |
| :---: | :---: | :---: |
|  $\operatorname{Be}(1,10)$ $\operatorname{Be}(1,1)$ <br> $\beta_{0}$ $2.34(2.13,2.55)$ $2.34(2.13,2.55)$ <br> $\beta_{1}$ $0.29(0.26,0.32)$ $0.29(0.26,0.32)$ <br> $a_{1}$ $0.093(0.025,0.210)$ $0.119(0.040,0.270)$ <br> $a_{2}$ $0.208(0.064,0.445)$ $0.659(0.189,0.960)$ |  |  |

Model 2

| Model 2 |  |  |
| :---: | :---: | :---: |
|  | $\operatorname{Be}(1,10)$ | $\operatorname{Be}(1,1)$ |
| $\beta_{0}$ | $2.34(2.13,2.55)$ | $2.34(2.13,2.55)$ |
| $\beta_{1}$ | $0.29(0.26,0.32)$ | $0.29(0.26,0.32)$ |


|  | Parametric |
| :--- | :--- |
| $\beta_{0}$ | $2.34(2.13,2.55)$ |
| $\beta_{1}$ | $0.29(0.26,0.32)$ |

Table 4: The posterior distribution of the regression coefficients summarized as the posterior median and $95 \%$ credible interval for Example 2 using parametric Normal model and Models 1 and Model 2 with two different priors for $a_{1}$ and $a_{2}$.
centred over a normal distribution with covariance given by the identity matrix. Semiparametric linear mixed models can be defined by assuming that the distribution of the random effects is modelled using one of these nonparametric models. The semiparametric model is then centred over the semiparametric model and the inference can be "shrunk" towards the parametric model when the data supports the parametric model.

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Figure 8: Contour plots of the posterior mean density under Models 1 and 2 with two different priors for $a_{1}$ and $a_{2}$ for the chlosterol data.

$$
\operatorname{Be}(1,1)
$$


Model 1



Model 2
+


$\operatorname{Be}(1,10)$



Intercept

Figure 9: Plots of the posterior mean marginal densities for intercept and slope under Models 1 and 2 with two different priors for $a_{1}$ and $a_{2}$ for the chlosterol data.

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Model 1

|  | $\operatorname{Be}(1,10)$ | $\operatorname{Be}(1,1)$ |
| :---: | :---: | :---: |
| $\beta_{1}$ (age) | $0.015(0.008,0.022)$ | $0.017(0.010,0.022)$ |
| $\beta_{2}$ (sex) | $-0.064(-0.158,0.031)$ | $-0.056(-0.153,0.039)$ |
| intercept | $1.73(1.43,2.04)$ | $1.67(1.42,2.01)$ |
| slope | $0.28(0.23,0.33)$ | $0.28(0.23,0.33)$ |
| $a_{1}$ | $0.140(0.048,0.325)$ | $0.283(0.097,0.635)$ |
| $a_{2}$ | $0.094(0.013,0.286)$ | $0.305(0.065,0.857)$ |

Model 2

|  | $\operatorname{Be}(1,10)$ | $\operatorname{Be}(1,1)$ |
| :---: | :---: | :---: |
| $\beta_{1}$ (age) | $0.016(0.010,0.022)$ | $0.016(0.010,0.022)$ |
| $\beta_{2}$ (sex) | $-0.063(-0.157,0.031)$ | $-0.060(-0.154,0.037)$ |
| intercept | $1.70(1.42,1.97)$ | $1.68(1.42,1.97)$ |
| slope | $0.28(0.23,0.33)$ | $0.28(0.23,0.33)$ |

Parametric

| $\beta_{1}$ (age) | $0.017(0.010,0.026)$ |
| :---: | :---: |
| $\beta_{2}$ (sex) | $-0.041(-0.160,0.076)$ |
| intercept | $1.64(1.26,1.96)$ |
| slope | $0.28(0.22,0.34)$ |

Table 5: The posterior distribution of the regression coefficients and the smoothness parameters summarized as the posterior median and $95 \%$ credible interval for Example 1 using parametric Normal model and Models 1 and Model 2 with two different priors for $a_{1}$ and $a_{2}$.

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