

Appendix

Appendix

The following paragraphs contain additional information about the **simulation studies** that were covered in chapter ??.

Section ??: Correlated Predictor Variables

The simulation design was chosen in the following way:

- The design matrix $\mathbf{X} = (\mathbf{x}_1 \quad \mathbf{x}_2 \quad \mathbf{x}_3)$ is simulated from a three dimensional normal distribution $\mathcal{N}_3(\boldsymbol{\mu}, \boldsymbol{\Sigma})$ with mean vector $\boldsymbol{\mu} = (-5 \quad 2 \quad 0)^T$ and covariance matrix $\begin{pmatrix} 1 & \rho & \rho \\ \rho & 3 & \rho \\ \rho & \rho & 5 \end{pmatrix}$. Hence, the dependence among the regressors is fully determined by the parameter ρ .
- The design matrix $\mathbf{Z} = (\mathbf{z}_1 \quad \mathbf{z}_2)$ consists of linear combinations of the regressors \mathbf{x}_1 up to \mathbf{x}_3 , more specifically $\mathbf{z}_1 = 0.8 \cdot \mathbf{x}_1 + 0.2 \cdot \mathbf{x}_2$ and $\mathbf{z}_2 = \mathbf{x}_2 - 0.5 \cdot \mathbf{x}_3$. In both design matrices intercept columns are added for estimation purposes.
- The true coefficient vectors are given by $\boldsymbol{\beta} = (\beta_0 \quad \beta_1 \quad \beta_2 \quad \beta_3)^T = (0 \quad 3 \quad -1 \quad 1)^T$ and $\boldsymbol{\gamma} = (\gamma_0 \quad \gamma_1 \quad \gamma_2)^T = (0 \quad 2 \quad 0)^T$.
- The outcome variable \mathbf{y} is generated according to the correctly specified location-scale model $y_i \stackrel{iid}{\sim} \mathcal{N}(\mathbf{x}_i^T \boldsymbol{\beta}, \exp(\mathbf{z}_i^T \boldsymbol{\gamma})^2)$ for $i = 1, \dots, n$ with sample size $n = 50$. Before data generation all columns in \mathbf{X} and \mathbf{Z} are standardized, i.e. mean-centered around 0 and scaled to unit variance.
- Three different values were chosen for $\rho \in \{0, -0.5, 0.9\}$ to compare the ‘nice’ case of uncorrelated predictors with the performance for negative and positive dependence. For each covariance structure the three models `mcmc_ridge()`, `mcmc()` and `lmls()` are fitted to the standardized covariates, where each Posterior Mean estimate from both of the Markov Chain Monte Carlo samplers is based on 10.000 samples.

Section ??: Sample Size

The simulation study is based on the following conditions:

- The design matrix $\mathbf{X} = (\mathbf{1}_n \quad \mathbf{x}_1 \quad \mathbf{x}_2)$ contains two independently sampled regressor variables plus one intercept column:
 - $\mathbf{x}_1 \stackrel{iid}{\sim} \mathcal{N}(1, 1)$,
 - $\mathbf{x}_2 \stackrel{iid}{\sim} \mathcal{N}(2, 1)$.
- The design matrix $\mathbf{Z} = (\mathbf{1}_n \quad \mathbf{z}_1 \quad \mathbf{z}_2)$ is structured in the same way with the regressor variables:
 - $\mathbf{z}_1 \stackrel{iid}{\sim} \mathcal{N}(1, 1)$,
 - $\mathbf{z}_2 \stackrel{iid}{\sim} \mathcal{N}(2, 1)$.
- After sampling the matrices \mathbf{X} and \mathbf{Z} , the columns of $\mathbf{x}_1, \mathbf{x}_2, \mathbf{z}_1$ and \mathbf{z}_2 are standardized.

- The true coefficient vectors are given by $\beta = (\beta_0 \ \beta_1 \ \beta_2)^T = (1 \ -1 \ 4)^T$ and $\gamma = (\gamma_0 \ \gamma_1 \ \gamma_2)^T = (0 \ -0.5 \ 1)^T$.
- The posterior means are analyzed with respect to 6 different sample sizes $n \in \{0, 50, 100, 200, 300, 500\}$.
- In the next step, the outcome vector $y \in \mathbb{R}^n$ is simulated and passed to the `mcmc_ridge()` function with `nsim` = 500 simulations.
- To make the results more stable, the above procedure is repeated 100 times. For each coefficient, the mean value of the Posterior Mean estimates of each coefficient is calculated as well as the Mean Absolute Error (MAE) with respect to the true values of β and γ .

Section ??: Redundant Covariates

We again state the conditions that the simulation study is based on:

- The design matrix $\mathbf{X} = (\mathbf{1}_n \ \mathbf{x}_1 \ \dots \ \mathbf{x}_{20})$ consists of one intercept column plus 10 *pairs* of successive regressors, starting with the pair $(\mathbf{x}_1, \mathbf{x}_2)$. Each pair $(\mathbf{x}_i, \mathbf{x}_{i+1})$ for $i \in \{1, 3, \dots, 19\}$ is (independently from all remaining pairs) drawn from a bivariate normal distribution with mean vector $\mu = (0 \ 0)^T$ and correlation matrix $\begin{pmatrix} 1 & 0.9 \\ 0.9 & 1 \end{pmatrix}$.
- The design matrix $\mathbf{Z} = (\mathbf{1}_n \ \mathbf{x}_1 \ \mathbf{x}_3)$ is of minor interest in this case and consists of an intercept column plus two uncorrelated columns chosen from \mathbf{X} .
- The true coefficients of β are determined by the pattern $\beta_i = 0$, if i is even and $\beta_i = 1$, if i is odd. Thus, all covariates with even subscripts are redundant, whereas those with odd subscripts contribute to \mathbf{y} . The true γ , again of minor interest here, is given by $\gamma = (0 \ 1 \ 1)^T$.
- The outcome variable \mathbf{y} is generated according to the correctly specified location-scale model $y_i \stackrel{iid}{\sim} \mathcal{N}(\mathbf{x}_i^T \beta, \exp(\mathbf{z}_i^T \gamma)^2)$ for $i = 1, \dots, n$, where the covariates are used on their original scale.
- The sample size $n = 50$ is deliberately chosen small compared to the number of regressors. Before fitting each of the three models, all columns of \mathbf{X} except the intercept column are standardized to zero mean and unit variance. Both of the Bayesian models generate 10.000 simulated values for each coefficient.

Section ??: Challenging the Model Assumptions

The data for this simulation study is generated by the following conventions:

- The design matrix $\mathbf{X} = (\mathbf{1}_n \ \mathbf{x}_1 \ \mathbf{x}_2 \ \mathbf{x}_3 \ \mathbf{x}_4)$ contains four independently sampled regressor variables plus one intercept column:
 - $\mathbf{x}_1 \stackrel{iid}{\sim} \mathcal{N}(5, 16)$,
 - $\mathbf{x}_2 \stackrel{iid}{\sim} \text{Exp}(5)$,
 - $\mathbf{x}_3 \stackrel{iid}{\sim} \mathcal{U}([-2, 12])$,
 - $\mathbf{x}_4 \stackrel{iid}{\sim} \text{Ber}(0.3)$.
- The design matrix $\mathbf{Z} = (\mathbf{1}_n \ \mathbf{x}_1 \ \mathbf{x}_2 \ \mathbf{z}_3)$ contains the additional regressor variable $\mathbf{z}_3 \stackrel{iid}{\sim} t_{10}$, which is independently sampled from all other columns.
- The true coefficient vectors are given by $\beta = (\beta_0 \ \beta_1 \ \beta_2 \ \beta_3 \ \beta_4)^T = (0 \ -10 \ -5 \ -3 \ -1)^T$ and $\gamma = (\gamma_0 \ \gamma_1 \ \gamma_2 \ \gamma_3)^T = (0 \ 3 \ 5 \ 10)^T$.
- Three different specifications for the outcome distribution were chosen:
 - $y_i \sim \mathcal{N}(\mu, \sigma^2)$,
 - $y_i \sim \mu + \left(\sigma \cdot \sqrt{\frac{3}{5}}\right) T$, where $T \sim t_5$,

– $y_i \sim \mu + \sigma \cdot U$, where $U \sim \mathcal{U}([0, 1])$.

In all cases, the outcome vectors are generated with the covariates on their original (unstandardized) scale.

- In order to isolate the impact of the different shapes of the three probability distributions from the effect of varying moment structures, the mean $\mu = \mathbf{x}_i^T \boldsymbol{\beta}$ and the variance $\sigma^2 = \exp(\mathbf{z}_i^T \boldsymbol{\gamma})^2$ are held constant across the models.
- All three models `mcmc_ridge()`, `mcmc()` and `lmls()` are fitted with standardized covariates. The sample size is set to $n = 50$ and the result of both Bayesian samplers are based on 10.000 simulations.

Section ??: Hyperparameters - Impact on Estimation Accuracy

The simulation study of the impact of the Hyperparameters on the estimated coefficients is constructed as follows:

- The design matrix $\mathbf{X} = (\mathbf{x}_1 \ \mathbf{x}_2)$ is simulated from a two dimensional normal distribution $\mathcal{N}_2(\boldsymbol{\mu}, \boldsymbol{\Sigma})$ with mean vector $\boldsymbol{\mu} = (1 \ 2)^T$ and identity covariance matrix $\boldsymbol{\Sigma} = \mathbf{I}_2$. The same holds true for the design matrix $\mathbf{Z} = (\mathbf{z}_1 \ \mathbf{z}_2)$ with mean vector $\boldsymbol{\mu} = (5 \ 3)^T$ and identity covariance matrix. However, after simulating \mathbf{X} and \mathbf{Z} , both are standardized to zero mean and unit variance.
- In both design matrices intercept columns are added for estimation purposes. The true coefficient vectors are given by $\boldsymbol{\beta} = (\beta_0 \ \beta_1 \ \beta_2)^T = (0 \ -1 \ 4)^T$ and $\boldsymbol{\gamma} = (\gamma_0 \ \gamma_1 \ \gamma_2)^T = (0 \ -2 \ 1)^T$.
- The outcome variable \mathbf{y} is generated according to the correctly specified location-scale model $y_i \stackrel{iid}{\sim} \mathcal{N}(\mathbf{x}_i^T \boldsymbol{\beta}, \exp(\mathbf{z}_i^T \boldsymbol{\gamma})^2)$ for $i = 1, \dots, n$ with sample size $n = 50$ as well as standardized data matrices \mathbf{X} and \mathbf{Z} .
- For sampling the location parameter, the full conditional multivariate normal distribution of $\boldsymbol{\beta}$ is chosen, i.e. `mcmc_ridge(..., mh_location = FALSE)` is used. Therefore, the location estimate is directly affected by the hyperparameters.
- For simulating the influence of the hyperparameters, nine different values are chosen: $a_\tau, b_\tau, a_\xi, b_\xi \in \{-1, 0, 0.5, 1, 2, 10, 50, 100, 200\}$. Since for statistical properties like the mean of an Inverse Gamma distribution $\frac{b}{a-1}$ the condition $a > 1$ is required, particular attention is given to larger values. However, it is an aim to inspect the performance of the sampler for hyperparameter values smaller than 1 as well.

Section ??: Hyperparameters - Impact on Ridge Penalty

This simulation study is conducted in the following way:

- The column vectors of the design matrices $\mathbf{X} = (\mathbf{x}_1 \ \mathbf{x}_2)$ and $\mathbf{Z} = (\mathbf{z}_1 \ \mathbf{z}_2)$ are independently drawn from normal distributions with variance $\sigma^2 = 1$ and varying means $\mu \in (1, 2, 5, 3)$. Then both design matrices are standardized and intercept columns are added.
- The true coefficient vectors are given by $\boldsymbol{\beta} = (\beta_0 \ \beta_1 \ \beta_2)^T = (0 \ 8 \ 2)^T$ and $\boldsymbol{\gamma} = (\gamma_0 \ \gamma_1 \ \gamma_2)^T = (0 \ 3 \ 3)^T$.
- The outcome variable \mathbf{y} is generated based on the standardized covariates according to the correctly specified location-scale model $y_i \stackrel{iid}{\sim} \mathcal{N}(\mathbf{x}_i^T \boldsymbol{\beta}, \exp(\mathbf{z}_i^T \boldsymbol{\gamma})^2)$ for $i = 1, \dots, n$ with sample size $n = 50$.
- The hyperparameter pairs (a_τ, b_τ) and (a_ξ, b_ξ) take values on a grid, which is constructed by all combinations of the sequence $(\frac{1}{16}, \frac{1}{8}, \frac{1}{4}, \frac{1}{2}, 1, 2, 4, 8, 16, 32, 64, 128, 256)$. While one pair is varied, the other pair is fixed on the values $(1, 1)$.
- The `mcmc_ridge()` function does not use the `lmls()` function as basis in this case, but rather works with the standardized data matrices \mathbf{X} and \mathbf{Z} directly. The number of simulations `num_sim` is chosen as 1000 and the starting values `beta_start` and `gamma_start` are set to $(1 \ 1)^T$, respectively.