Bayesian Ridge Regression - Second Report

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Introduction

1 Correlated Predictor Variables

Up to this point, we have often illustrated the usage and results of the mcmc_ridge() sampler with simulated data from the built-in toy_data set. There, each regressor variable is independently sampled from a normal distribution and the outcome variable is simulated based on a correctly specified location-scale regression model $y_i \sim \mathcal{N}\left(\mathbf{x}_i^T \boldsymbol{\beta}, \exp\left(\mathbf{z}_i^T \boldsymbol{\gamma}\right)^2\right)$. All these conditions lead to an excellent performance of the MCMC sampler, but might arguably not represent the most challenging task.

The next two sections analyze the sampler's performance on simulated data, which might be closer to data found in the real world. First, we will induce correlation among the predictor variables, whereas in the following section the distributional assumptions are considerably changed. Further, the mcmc_ridge() performance is compared to the Maximum Likelihood based lmls() estimator and the Markov Chain Monte Carlo mcmc() sampler from the lmls package.

1.1 Simulation Setting

- The design matrix $\mathbf{X} = \begin{pmatrix} \mathbf{x}_1 & \mathbf{x}_2 & \mathbf{x}_3 \end{pmatrix}$ is simulated from a three dimensional normal distribution $\mathcal{N}_3(\boldsymbol{\mu}, \boldsymbol{\Sigma})$ with mean vector $\boldsymbol{\mu} = \begin{pmatrix} -5 & 2 & 0 \end{pmatrix}^T$ 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 $\mathbf{Z} = \begin{pmatrix} \mathbf{z}_1 & \mathbf{z}_2 \end{pmatrix}$ 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 $\boldsymbol{\beta} = \begin{pmatrix} \beta_0 & \beta_1 & \beta_2 & \beta_4 \end{pmatrix}^T = \begin{pmatrix} 0 & 3 & -1 & 1 \end{pmatrix}^T$ and $\boldsymbol{\gamma} = \begin{pmatrix} \gamma_0 & \gamma_1 & \gamma_2 \end{pmatrix}^T = \begin{pmatrix} 0 & 2 & 0 \end{pmatrix}^T$.
- Three different values (0, -0.5 and 0.9) were chosen for ρ 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() were fitted, where each Posterior Mean estimate from both of the Markov Chain Monte Carlo samplers is based on 1000 samples.
- Moreover, we compared the performance of the usual $mcmc_ridge()$ implementation, which draws β from the closed form full conditional (multivariate normal) distribution, with an alternative sampling process that uses a Metropolis-Hastings approach for both, the location parameter β as well as the scale parameter γ . The latter is initiated by the $mcmc_ridge()$ argument $ml_location = TRUE$. The variance of the corresponding proposal distribution is set to a carefully chosen default value, but can be manually changed by means of the $prop_var_loc$ argument.

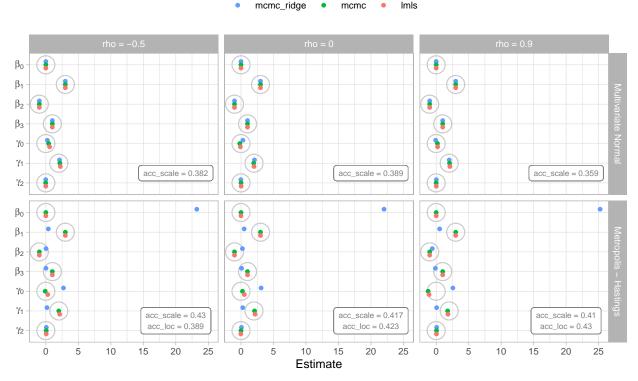
1.2 Simulation Results

The following plot displays the posterior mean estimates for the MCMC samplers and the Maximum Likelihood estimates for the lmls() function of one complete iteration of the simulation. For a better visual comparison

the true values for each coefficient are indicated by grey circles, whereas the acceptance rate(s) of the Metropolis-Hastings sampling process are provided in grey boxes:

Model Performance for different Predictor Correlation Structures

True coefficient values are indicated by grey circles



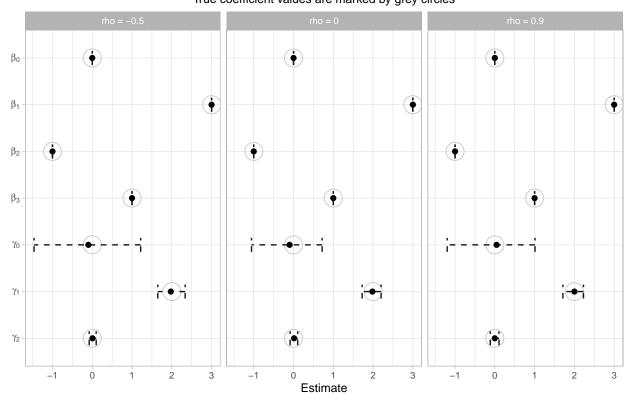
The scaling of the x- axis is dominated by one outlier in the lower panel for each correlation structure. While the Metropolis-Hastings approach for β performs moderately well for most of the coefficients, it massively overestimates the intercept β_0 . This observation can be made across many different data sets: In some special cases the performance is close to (but never better) than sampling directly from a multivariate normal distribution, however, most of the time the performance is significantly worse and the samples show (obviously) much larger correlation requiring a higher number of simulations for stable estimation. For that reason, we limit the Metropolis-Hastings sampling process for β to this one illustration and will focus on the classical mcmc_ridge() implementation in the remaining parts of the report.

The upper panel indicates a very good performance by all three estimation procedures in consideration. Further, all acceptance probabilities are in a reasonable range supporting a fast convergence of all Markov Chains.

It is important to remember, that each point in the plot only represents exactly one measurement. In order to make any conclusions about bias and variance of the estimation procedures, the above procedure is repeated 50 times. The black points represent the mean of these 50 posterior mean estimates. Since we cannot rely on distributional theory for the standard errors, the variability of the estimates is displayed by nonparametric 'confidence' intervals, which are simply given by the range from the empirical 0.05 quantile to the 0.95 quantile of the 50 estimated values.

Further investigations have shown that the mcmc_ridge(), the mcmc() and the lmls() functions perform very similar for each correlation structure. For that reason only the results of the mcmc_ridge() sampler are shown in the following plot:

Empirical 90% Confidence Intervals for Posterior Mean Estimates True coefficient values are marked by grey circles



There are three conclusions from this first simulation study:

- 1. The correlation structure does not have a significant impact of the sampling results. The three plot facets look almost identical.
- 2. The mcmc_ridge() sampler (as well as the mcmc() and lmls() functions) are very robust towards correlated data and perform extremely well. In particular, all three approaches (visually) provide close to unbiased estimates.
- 3. The variability among the β estimates is almost nonexistent, such that results from a single simulation are already reliable and representative. While the estimates for the γ vector are still correct on average, the variability across different simulations is significant (particularly for γ_0). Thus, averaging the results from multiple repetitions of the sampling process is advisable.

2 Challenging the Model Assumptions

2.1 Simulation Setting

2.2 Technical Aspects

As outlined in the previous paragraph, a total of $50 \cdot 3 \cdot 3 = 450$ models were fitted to analyze the performance differences. In order to speed up the involved computations of this specific and some of the other simulation studies illustrated in this report, we used the *parallel computing* capabilities of R.

There are many options from various packages to choose from. We decided to use the furrr package which is built on top of the future package specialized on parallel processing. As the name suggests furrr provides a convenient way to use many functions from the popular purrr package using multiple cores at the same time. This functional programming based approach (similar to the apply() family in 'base R') is particularly well

suited for simulation studies and provides some structural as well as minor performance advantages compared to the classical for-loop approach that are beyond the scope of this report.

The following (slightly modified) code snippet provides a brief insight into the implementation:

```
plan(multisession, workers = 8)

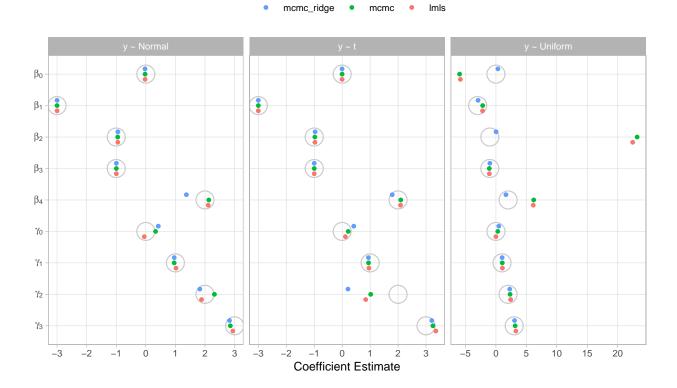
full_results <- tibble(id = 1:50) %>%
  mutate(samples = future_map(
    .x = id,
    .f = ~ show_results(n = 50, num_sim = 1000),
    .options = furrr_options(seed = 1)
  ))
```

The plan() function borrowed from the future package initializes the parallel computing process and the number of cores available for computation. The show_results() helper function fits all three models mcmc_ridge(), mcmc() and lmls() for each outcome distribution in a single iteration. This procedure is repeated 50 times in parallel using the future_map() function from the furr package, where the results of all 450 models are saved in a well organized structure inside of a list column. This column of the data frame contains complete information such that all required elements for the further analysis can be easily extracted and post processed.

Finally, the .options() argument allows the specification of a random seed. Random number generation in the context of parallel computing is slightly more involved than in the sequential approach. This additional complexity is automatically handled by the future_map() function, such that all results are sampled in a statistically valid and fully reproducible manner.

2.3 Simulation Results

Posterior Means / MLE for (misspecified) Regression Models True coefficient values are marked by grey circles



Empirical 90% Confidence Intervals for Posterior Mean Estimates

True coefficient values are marked by grey circles

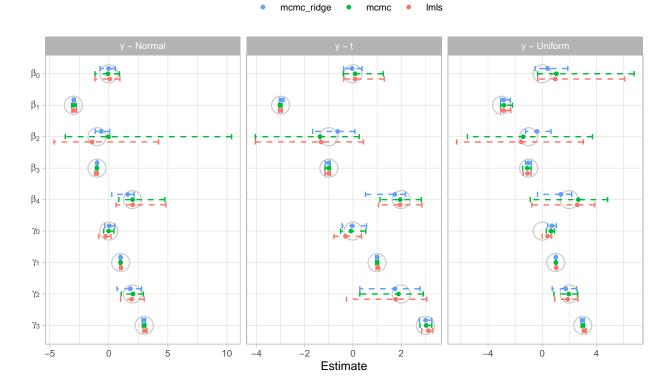


Table 1: Bias of Coefficient Estimates

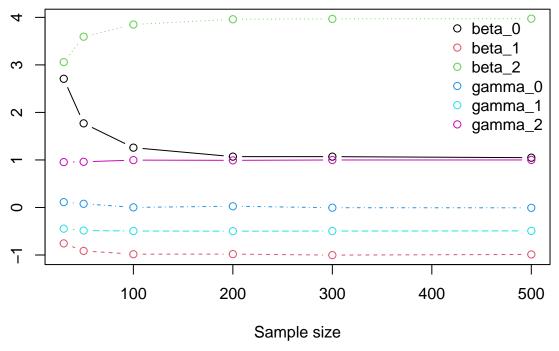
	Normal			t			Uniform		
	lmls	mcmc	mcmc_ridge	lmls	mcmc	mcmc_ridge	lmls	mcmc	mcmc_ridge
β_0	0.07	-0.07	-0.02	0.09	0.09	-0.04	0.97	1.05	0.40
β_2	-0.43	0.97	0.35	-0.32	-0.37	0.37	-0.55	-0.39	0.61
β_4	0.04	0.01	-0.40	-0.05	-0.04	-0.26	0.59	0.67	-0.62
γ_0	-0.27	-0.02	0.03	-0.31	-0.08	-0.03	0.40	0.66	0.72
γ_2	-0.06	0.04	-0.18	-0.23	-0.11	-0.26	-0.11	-0.04	-0.25

Table 2: Standard Errors of Coefficient Estimates

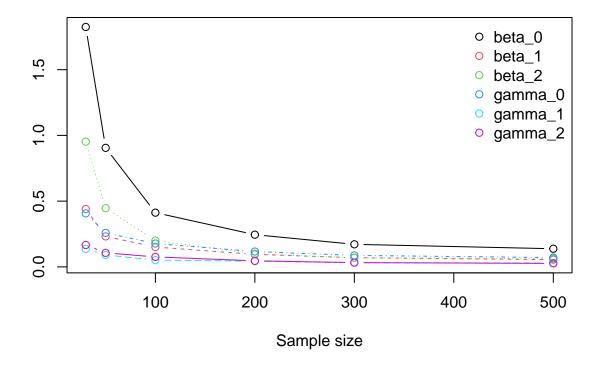
	Normal			t			Uniform		
	lmls	mcmc	mcmc_ridge	lmls	mcmc	mcmc_ridge	lmls	mcmc	mcmc_ridge
β_0	0.83	0.69	0.47	0.62	0.62	0.49	2.15	2.37	1.47
β_2	7.11	4.41	0.40	2.54	2.58	0.54	6.48	6.48	0.67
β_4	1.32	1.33	0.68	0.57	0.56	0.53	4.10	4.30	0.81
γ_0	0.33	0.28	0.27	0.35	0.33	0.34	0.22	0.19	0.21
γ_2	0.78	0.65	0.62	1.01	0.93	0.89	0.55	0.54	0.56

3 Sample Size

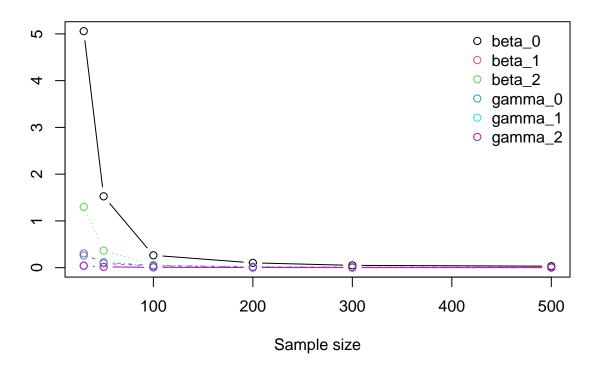
Mean of Posterior Means



MAE of Posterior Means

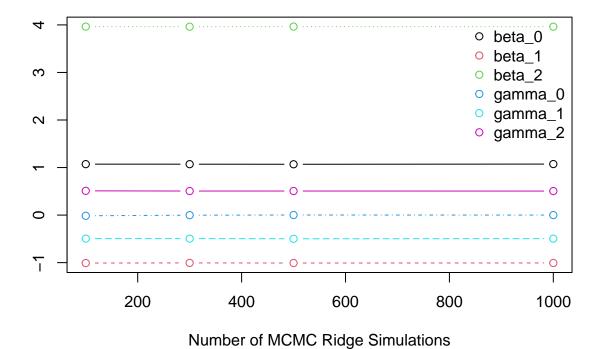


MSE of Posterior Means

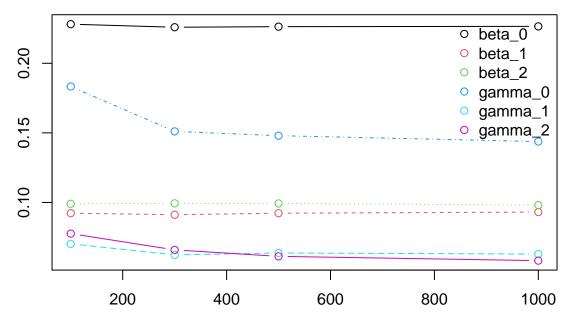


4 Number of Simulations

Mean of Posterior Means

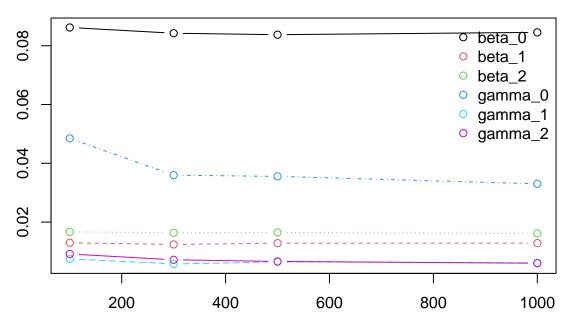


MAE of Posterior Means



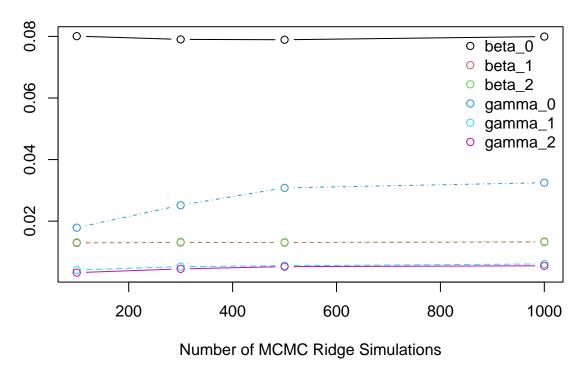
Number of MCMC Ridge Simulations

MSE of Posterior Means



Number of MCMC Ridge Simulations

Mean of Variances within the Samples



5 Hyperparameters

Next Steps

In the process of writing the first and second report, the main work of developing both the mathematical foundation as well as the code base of the asp21bridge package has been done. Thus, the following weeks will focus on filling remaining gaps and working out details that were left open due to time constraints. This includes taking a closer look at code efficiency and possible code refactoring for the sake of modularity.

The final step consists of combining all pieces of the project into a single structured and consistent final report. Aside of the work shown in the first two reports, some new elements like a brief discussion of design and implementation choices during the development stage will be added with the aim of providing the reader a solid understanding of the underlying ideas as well as the practical usage of the package.