Half Title

Introduction to Bayesian Inference: A GUIded tour using R

Title Page

Introduction to Bayesian Inference: A GUIded tour using R

by Andrés Ramírez-Hassan, PhD. Statistical Science.

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Introduction



FIGURE 1
Supposedly portrait of Thomas Bayes.

Since late 90's Bayesian inference has gained a lot of popularity among researchers due to the computational revolution and availability of algorithms to solve complex integrals. However, many researchers, students and practitioners still lack understanding and application of this inferential approach. The main reason is the requirement of good programming skills.

Introduction to Bayesian inference: A GUIded tour using R mainly targets those who want to apply Bayesian regression analysis having a good conceptual and formal understanding, but not having time to develop programming skills. Thus, this book provides a graphical user interface (GUI) to carry out Bayesian regression in a very friendly environment. The book also provides the basic theory, and its code implementation using R software [61], some econometric/statistical applications to highlight the potential of Bayesian regression, and theory and computational exercises, for those who are interested in developing more complex models. In particular, this book contains the mathematical proofs step by step of the basic model, which are the base for obtaining the most relevant mathematical results of more complex models.

Our GUI is based on an interactive web application using shiny [15], and some packages in **R**. Users can estimate univariate, multivariate, time series, panel data (longitudinal) and Bayesian model average models using our GUI.

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In addition, it gives basic summaries and formal and graphical diagnostics of the posterior chains. Our GUI can be run in any operating system, and is freely available at https://github.com/besmarter/BSTApp.

Users can get simulated and real datasets in the folders **DataSim**, and **DataApp**, respectively. The former folder also includes the files that were used to simulate different processes, so, the population parameters are available, and as a consequence, these files can be used as a pedagogical tool to show some statistical properties. The latter folder contains the datasets used in our applications. Users should use these datasets as templates to structuring their own datasets. Simply type **shiny::runGitHub("besmarter/BSTApp""**, **launch.browser=T)** in the **R** package console or any **R** code editor to run our GUI.¹

This book has three parts. The first one is about theory (conceptual and mathematical), programming and simulation foundations (chapters 1 to 5), the second part is about applications of regression analysis (chapters 6 to 10), and the third part is about advanced methods in Bayesian inference (chapters 11 to 15). I show in some detail the mathematical deductions in the first part of the book, whereas I do not show any proof in the second and third parts. However, same mathematical steps can be used to find the results of parts two and three of the book. I also show three levels regarding computational implementation in the second part of the book: programming ourselves the algorithms, using Bayesian R packages, and using our GUI.

Chapter 1 begins with an introduction to formal concepts in Bayesian inference starting with the Bayes' rule, all its components with their formal definitions and basic examples. Then, it presents the basics of Bayesian inference based on decision theory under uncertainty. Chapter 2 presents conceptual differences between Bayesian and Frequentist statistical approaches, and a historical and philosophical perspective about Bayesian statistics and econometrics highlighting differences compared to the Frequentist approach. Chapter 3 presents the differences between the objective and subjective schools in Bayesian inference. Particular attention is put to elicitation techniques, that is, how to transform expert knowledge into prior probabilistic statements. In Chapter 4 I introduce conjugate families in basic statistical models, solving them analytically and computationally. Simulation based methods are shown in Chapter 5, these algorithms are very important in modern Bayesian inference as most realistic models do not have standard forms or analytical solutions. Univariate and multivariate regression models are presented in chapters 6 and 7, respectively. Chapter 8 presents the state-space representation of time series models, and Chapter 9 presents Bayesian panel data (longitudinal) models. Chapter 10 introduces Bayesian model averaging. In the third part, there are Chapter 11 introducing hierarchical models, Chapter 12 shows causal inference, Chapter 13 shows Bayesian methods in machine learning algorithms, Chapter 14 introduces spatial models, and Chapter 15 describes

¹I strongly recommend to type the code line rather than copy and paste it.

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some recent methodological developments such as approximate Bayesian computation (ABC), synthetic likelihood (SL), variational Bayes (VB), and integrated nested Laplace approximations (INLA).

About me

My name is Andrés Ramírez-Hassan, I am an applied and theory econometrician working as a Distinguished Professor in the School of Finance, Economics and Government at Universidad EAFIT (Medellín, Colombia). I got a PhD in Statistical Science, a masters degree in Finance, and another in Economics, and also a bachelor's degree in Economics. I was a research fellow at the Department of Econometrics and Business Statistics at Monash University, and a visiting Professor in the Department of Economics at the University of Melbourne and the University of Glasgow. Having completed my PhD degree, much of my research has been in the area of Bayesian Econometrics with applications in crime, finance, health, sports and utilities. My work has been published (or is forthcoming) in the International Journal of Forecasting, Journal of Applied Econometrics, Econometric Reviews, Journal of Computational and Graphical Statistics, The R Journal, Economic Modelling, Spatial Economic Analysis, Economic Inquiry, World Development, Journal of Sport Economics, Empirical Economics, Australian and New Zealand Journal of Statistics, Brazilian Journal of Probability and Statistics, and other highly regarded international research outlets.

I founded **BEsmarter** –**B**ayesian **E**conometrics: simulations, **m**odels and **a**pplications to **research**, **teaching** and **e**ncoding with **r**esponsibility–. This is a research group whose **mission** is to lead and excel in the generation and dissemination of Bayesian Econometric knowledge through research, teaching and software. We envision worldwide econometric research, teaching and applications based on the Bayesian framework that:

- Inspires new econometric ideas
- Creates a user friendly environment for applications of Bayesian econometrics
- Transforms classic econometric research, teaching and applications
- And where one of the main concerns of science is to solve social problems
 mail: aramir21@gmail.com / aramir21@eafit.edu.co
 website: http://www.besmarter-team.org / https://sites.google.com/view/arh-



FIGURE 2

bayesian

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Foreword

Preface

The main goal of this book is to make more approachable the Bayesian inferential framework to students, researchers and practitioners who want to understand and apply this statistical/econometric approach, but who do not have time to develop programming skills. I tried to have a balance between applicability and theory. Then, this book comes with a very friendly graphical user interface (GUI) to implement the most common regression models, but also contains the basic mathematical developments, as well as their code implementation, for those who are interested in advancing in more complex models.

To instructors and students

This book is divided in three parts, foundations (chapters 1 to 5), regression analysis (chapters 6 to 10), and Advanced methods (chapters 11 to 15). Our graphical user interface (GUI) targets the second part. This can be download at https://github.com/besmarter/BSTApp. Instructors and students can have all codes, simulated and real data sets are also there. To install our GUI just type shiny::runGitHub("besmarter/BSTApp", launch.browser=T) in the R package console or any R code editor, and execute it.

Students should have some basic knowledge in probability theory and statistics, particularly, regression analysis. It is strongly recommended to have some familiarity with standard univariate and multivariate probability distributions.

I included some formal and computational exercises at the end of each chapter. This would help students to have a better understanding of the material shown in each chapter. A manual with the solutions of exercises accompanies this book.

Instructors can use this book as a text in a course of introduction to Bayesian Econometrics/Statistics with a high emphasis on implementation and applications. This book is complementary, rather than substitute, of excellent books in the topic such as [30, 66, 37, 34, 46] and [45].

Acknowledgments

I started our GUI in the 2016 after being diagnosed with cervical dystonia. I used to work in this side project on weekends, I named this time "nerd weekends", and it was a kind of release from my health condition. Once I got better, I invited Mateo Graciano, my former student, business partner and friend, to be part of the project, he helped me a lot developing our GUI, and I am enormously thankful to Mateo. I would also like to thank members

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of the BEsmarter research group from Universidad EAFIT, and NUMBATs members from Monash University for your comments and recommendations to improve our GUI.

This book is an extension of the paper A GUIded tour of Bayesian regression [65], which is a brief user guide of our GUI. So, I decided to write this book to show the underlying theory and codes in our GUI, and use it as a text book in my course in Bayesian econometrics/statistics. I acknowledge and offer my gratitude to my students in this subject, their insight and thoughtful questions have helped me to get a better understanding of this material.

I also thank Chris Parmeter for your suggestions about how to present our user guide, Professor Raul Pericchi and Juan Carlos Correa who introduced me to Bayesian statistics, Liana Jacobi and Chun Fung Kwok (Jackson) from the University of Melbourne and David Frazier from Monash University for nice talks and amazing collaborations in Bayesian Econometrics/statistics, Professor Peter Diggle to support my career, and particularly, Professor Gael Martin, who gave me a chance to work with her, she is an inspiring intellectual figure. Finally, my colleagues and staff from Universidad EAFIT have always given me their support.

To my parents, Orlando and Nancy, who have given me their unconditional support. They have taught me that the primary aspect of the human being's spiritual evolution is humility. I am in my way to learn this.

Symbols

Symbol Description

\neg	Negation symbol	argmax	Argument of the maximum
\propto	Proportional symbol	argmin	Argument of the minimum
\perp	Independence symbol	tr	Trace operator
$\mathcal R$	The Real set	vec	Vectorization operator
Ø	Empty set	\lim	Limit
1	Indicator function	\otimes	Kronecker product
P	Probability measure	$\operatorname{diag}\{\cdot\}$	Diagonal matrix
:=	Is defined as	$dim\{\cdot\}$	Dimension of an object

Part I

Foundations: Theory, simulation methods and programming

Basic formal concepts

We introduce formal concepts in Bayesian inference starting with the Bayes' rule, all its components with their formal definitions and basic examples. In addition, we present some nice features of Bayesian inference such as Bayesian updating, and asymptotic sampling properties, and the basics of Bayesian inference based on decision theory under uncertainty, presenting important concepts like loss function, risk function and optimal rules.

1.1 The Bayes' rule

As expected the point of departure to perform Bayesian inference is the Bayes' rule, which is the Bayes' solution to the inverse probability of causes, this rule combines prior beliefs with objective probabilities based on repeatable experiments. In this way, we can move from observations to probable causes.

Formally, the conditional probability of A_i given B is equal to the conditional probability of B given A_i times the marginal probability of A_i over the marginal probability of B,

$$P(A_i|B) = \frac{P(A_i, B)}{P(B)}$$

$$= \frac{P(B|A_i) \times P(A_i)}{P(B)},$$
(1.1)

where by the law of total probability $P(B) = \sum_{i} P(B|A_i)P(A_i) \neq 0$, $\{A_i, i = 1, 2, ...\}$ is a finite or countably infinite partition of a sample space.

In the Bayesian framework, B is sample information that updates a probabilistic statement about an unknown object A_i following probability rules. This is done by means of the Bayes' rule using prior "beliefs" about A_i , that is, $P(A_i)$, sample information relating B to the particular state of the nature A_i through a probabilistic statement, $P(B|A_i)$, and the probability of observing that specific sample information P(B).

¹Observe that I use the term "Bayes' rule" rather than "Bayes' theorem". It was Laplace [48] who actually generalized the Bayes' theorem [5]. His generalization is named the Bayes' rule.

Let's see a simple example, the base rate fallacy:

Assume that the sample information comes from a positive result from a test whose true positive rate (sensitivity) is 98%, P(+|disease) = 0.98. On the other hand, the prior information regarding being infected with this disease comes from a base incidence rate that is equal to 0.002, that is P(disease) = 0.002. Then, what is the probability of being actually infected?

This is an example of the base rate fallacy, where having a positive test result from a disease whose base incidence rate is tiny gives a low probability of actually having the disease.

The key to answer the question is based on understanding the difference between the probability of having the disease given a positive result, P(disease|+), versus the probability of a positive result given the disease, P(+|disease). The former is the important result, and the Bayes' rule help us to get the answer. Using the Bayes' rule (equation 1.1):

$$\begin{split} P(\text{disease}|+) &= \frac{P(+|\text{disease}) \times P(\text{disease})}{P(+)} \\ &= \frac{0.98 \times 0.002}{0.98 \times 0.002 + (1 - 0.98) \times (1 - 0.002)} \\ &= 0.09, \end{split}$$

where $P(+) = P(+|\text{disease}) \times P(\text{disease}) + P(+|\neg \text{disease}) \times P(\neg \text{disease})$.

R code. The base rate fallacy

We observe that despite of having a positive result, the probability of having the disease is low. This due to the base rate being tiny.

Another interesting example, which is at the heart of the origin of the Bayes' theorem [5], is related to the existence of God [74]. The Section X of David Hume's "An Inquiry concerning Human Understanding, 1748" is named *Of Miracles*. There, Hume argues that when someone claims to have seen a

 $^{^2\}neg$ is the negation symbol. In addition, we have that $P(B|A)=1-P(B|A^c)$ in this example, where A^c is the complement of A. However, it is not always the case that $P(B|A)\neq 1-P(B|A^c)$.

The Bayes' rule 5

miracle, this is poor evidence it actually happened, since it goes against what we see every day. Then, Richard Price, who actually finished and published "An essay towards solving a problem in the doctrine of chances" in 1763 after Bayes died in 1761, argues against Hume saying that there is a huge difference between *impossibility* as used commonly in conversation and *physical impossibility*. Price used an example of a dice with a million sides, where *impossibility* is getting a particular side when throwing this dice, and *physical impossibility* is getting a side that does not exist. In millions throws, the latter case never would occur, but the former eventually would.

Let's say that there are two cases of resurrection (Res), Jesus Christ and Elvis, and the total number of people who have ever lived is 108.5 billion,³ then the prior base rate is $2/(108.5 \times 10^9)$. On the other hand, let's say that the sample information comes from a very reliable witness whose true positive rate is 0.9999999. Then, what is the probability of this miracle?⁴

Using the Bayes' rule:

```
\begin{split} P(\text{Res}|\text{Witness}) &= \frac{P(\text{Witness}|\text{Res}) \times P(\text{Res})}{P(\text{Witness})} \\ &= \frac{2/(108.5*10^9) \times 0.9999999}{2/(108.5*10^9) \times 0.9999999 + (1-2/(108.5*10^9)) \times (1-0.9999999)} \\ &= 0.000184297806959661 \end{split}
```

where $P(\text{Witness}) = P(\text{Witness}|\text{Res}) \times P(\text{Res}) + (1 - P(\text{Witness}|\text{Res})) \times (1 - P(\text{Res}))$.

Thus, 1.843×10^{-4} is the probability of a resurrection given a very reliable witness.

Observe that we can condition on many events in the Bayes' rule. Let's have two conditioning events B and C, then equation 1.1 becomes

 $^{^3} https://www.wolframalpha.com/input/?i=number+of+people+who+have+ever+lived+on+Earth\ ^4 https://www.r-bloggers.com/2019/04/base-rate-fallacy-or-why-no-one-is-justified-to-believe-that-jesus-rose/$



FIGURE 1.1
The Monty Hall problem.

$$P(A_i|B,C) = \frac{P(A_i,B,C)}{P(B,C)} = \frac{P(B|A_i,C) \times P(A_i|C) \times P(C)}{P(B|C)P(C)}.$$
 (1.2)

Let's use this rule in one of the most intriguing statistical puzzles, the Monty Hall problem, to illustrate how to use equation 1.2 [70, 71]. This was the situation faced by a contestant in the American television game show Let's Make a Deal. There, the contestant was asked to choose a door where behind one door there is a car, and behind the others, goats. Let's say that the contestant picks door No. 1, and the host (Monty Hall), who knows what is behind each door, opens door No. 3, where there is a goat (see Figure 1.1). Then, the host asks the tricky question to the contestant, do you want to pick door No. 2?

Let's name P_i the event **contestant picks door No.** i, which stays close, H_i the event **host picks door No.** i, which is open, and there is a goat, and C_i the event **car is behind door No.** i. In this particular setting, the contestant is interested in the probability of the event $P(C_2|H_3, P_1)$. A naive answer would be that it is irrelevant as initially $P(C_i) = 1/3$, i = 1, 2, 3, and now $P(C_i|H_3) = 1/2$, i = 1, 2 as the host opened door No. 3. So, why bothering changing the initial guess if the odds are the same (1:1)? The important point here is that the host knows what is behind each door, and picks a door where there is a goat given contestant choice. In this particular

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setting, $P(H_3|C_3, P_1) = 0$, $P(H_3|C_2, P_1) = 1$ and $P(H_3|C_1, P_1) = 1/2$. Then, using equation 1.2

$$\begin{split} P(C_2|H_3,P_1) &= \frac{P(C_2,H_3,P_1)}{P(H_3,P_1)} \\ &= \frac{P(H_3|C_2,P_1)P(C_2|P_1)P(P_1)}{P(H_3|P_1)\times P(P_1)} \\ &= \frac{P(H_3|C_2,P_1)P(C_2)}{P(H_3|P_1)} \\ &= \frac{1\times 1/3}{1/2}, \end{split}$$

where the third equation uses the fact that C_i and P_i are independent events, and $P(H_3|P_1) = 1/2$ due to this depending just on P_1 (not on C_2).

Therefore, changing the initial decision increases the probability of getting the car from 1/3 to 2/3! Thus, it is always a good idea to change the door.

Let's see a simulation exercise to check this answer:

R code. The Monty Hall problem

```
set.seed(0101) # Set simulation seed
2 S <- 100000 # Simulations
  Game <- function(switch = 0){</pre>
     # switch = 0 is not change
     # switch = 1 is to change
    opts <- 1:3
     car <- sample(opts, 1) # car location</pre>
     guess1 <- sample(opts, 1) # Initial guess</pre>
     if(car != guess1) {
     host <- opts[-c(car, guess1)]
    } else {
12
     host <- sample(opts[-c(car, guess1)], 1)
    win1 <- guess1 == car # Win no change
15
     guess2 <- opts[-c(host, guess1)]</pre>
     win2 <- guess2 == car # Win change
     if(switch == 0){
      win <- win1
    } else {
      win <- win2
21
22
23
     return(win)
24 }
26 #Win probabilities not changing
        <- mean(replicate(S, Game(switch = 0)))</pre>
28 Prob
29 0.3334
31 #Win probabilities changing
32 Prob <- mean(replicate(S, Game(switch = 1)))</pre>
33 Prob
34 0.6654
```

1.2 Bayesian framework: A brief summary of theory

For two random objects θ and y, the Bayes' rule may be analogously used,⁵

⁵From a Bayesian perspective θ is fixed, but unknown. Then, it is treated as a random object despite the lack of variability (see Chapter 2).

$$\pi(\boldsymbol{\theta}|\mathbf{y}) = \frac{p(\mathbf{y}|\boldsymbol{\theta}) \times \pi(\boldsymbol{\theta})}{p(\mathbf{y})},$$
(1.3)

where $\pi(\boldsymbol{\theta}|\mathbf{y})$ is the posterior density function, $\pi(\boldsymbol{\theta})$ is the prior density, $p(\mathbf{y}|\boldsymbol{\theta})$ is the likelihood (statistical model), and

$$p(\mathbf{y}) = \int_{\mathbf{\Theta}} p(\mathbf{y}|\boldsymbol{\theta}) \pi(\boldsymbol{\theta}) d\boldsymbol{\theta} = \mathbb{E} \left[p(\mathbf{y}|\boldsymbol{\theta}) \right]$$
 (1.4)

is the marginal likelihood or prior predictive. Observe that for this expected value to be meaningful the prior should be a proper density, that is, integrates to one, otherwise, it does not make sense.

Observe that $p(\mathbf{y}|\boldsymbol{\theta})$ is not a density in $\boldsymbol{\theta}$. In addition, $\pi(\boldsymbol{\theta})$ does not have to integrate to 1, that is, $\pi(\boldsymbol{\theta})$ can be an improper density function, $\int_{\Theta} \pi(\boldsymbol{\theta}) d\boldsymbol{\theta} = \infty$. However, $\pi(\boldsymbol{\theta}|\mathbf{y})$ is a proper density function, that is, $\int_{\Theta} \pi(\boldsymbol{\theta}|\mathbf{y}) d\boldsymbol{\theta} = 1$. For instance, set $\pi(\boldsymbol{\theta}) = c$, where c is a constant, then $\int_{\Theta} cd\boldsymbol{\theta} = \infty$. However, $\int_{\Theta} \pi(\boldsymbol{\theta}|\mathbf{y}) d\boldsymbol{\theta} = \int_{\Theta} \frac{p(\mathbf{y}|\boldsymbol{\theta}) \times c}{\int_{\Theta} p(\mathbf{y}|\boldsymbol{\theta}) \times cd\boldsymbol{\theta}} d\boldsymbol{\theta} = 1$ where c cancels out.

 $\pi(\boldsymbol{\theta}|\mathbf{y})$ is a sample updated "probabilistic belief" version of $\pi(\boldsymbol{\theta})$, where $\pi(\boldsymbol{\theta})$ is a prior probabilistic belief which can be constructed from previous empirical work, theory foundations, expert knowledge and/or mathematical convenience (see Chapters 3 and 4). This prior usually depends on parameters, which are named *hyperparameters*. In addition, the Bayesian approach implies using a probabilistic model about \mathbf{y} given $\boldsymbol{\theta}$, that is, $p(\mathbf{y}|\boldsymbol{\theta})$, where its integral over $\boldsymbol{\Theta}$, $p(\mathbf{y})$ is named the model evidence due to being a measure of model fit to the data.

Observe that the Bayesian inferential approach is conditional, that is, what can we learn about an unknown object $\boldsymbol{\theta}$ given that we already observed \mathbf{y} ? The answer is also conditional on the probabilistic model, that is $p(\mathbf{y}|\boldsymbol{\theta})$. So, what if we want to compare different models, let's say \mathcal{M}_m , $m = \{1, 2, ..., M\}$. Then, we should make explicit this in the Bayes' rule formulation,

$$\pi(\boldsymbol{\theta}|\mathbf{y}, \mathcal{M}_m) = \frac{p(\mathbf{y}|\boldsymbol{\theta}, \mathcal{M}_m) \times \pi(\boldsymbol{\theta}|\mathcal{M}_m)}{p(\mathbf{y}|\mathcal{M}_m)}.$$
 (1.5)

The posterior model probability is

$$\pi(\mathcal{M}_m|\mathbf{y}) = \frac{p(\mathbf{y}|\mathcal{M}_m) \times \pi(\mathcal{M}_m)}{p(\mathbf{y})},$$
(1.6)

where $p(\mathbf{y}|\mathcal{M}_m) = \int_{\mathbf{\Theta}} p(\mathbf{y}|\boldsymbol{\theta}, \mathcal{M}_m) \times \pi(\boldsymbol{\theta}|\mathcal{M}_m) d\boldsymbol{\theta}$ due to equation 1.5, and $\pi(\mathcal{M}_m)$ is the prior model probability.

Calculating $p(\mathbf{y})$ in equations 1.3 and 1.6 is very demanding most of the realistic cases. Fortunately, it is not required when performing inference about $\boldsymbol{\theta}$ as this is integrated out from it. Then, all what you need to know about the

shape of $\boldsymbol{\theta}$ is in $p(\mathbf{y}|\boldsymbol{\theta}, \mathcal{M}_m) \times \pi(\boldsymbol{\theta}|\mathcal{M}_m)$ or without explicitly conditioning on \mathcal{M}_m ,

$$\pi(\boldsymbol{\theta}|\mathbf{y}) \propto p(\mathbf{y}|\boldsymbol{\theta}) \times \pi(\boldsymbol{\theta}).$$
 (1.7)

Equation 1.7 is a very good shortcut to perform Bayesian inference about θ .

We also can avoid calculating $p(\mathbf{y})$ when performing model selection (hypothesis testing) using posterior odds ratio, that is, comparing models \mathcal{M}_1 and \mathcal{M}_2 ,

$$PO_{12} = \frac{\pi(\mathcal{M}_1|\mathbf{y})}{\pi(\mathcal{M}_2|\mathbf{y})}$$
$$= \frac{p(\mathbf{y}|\mathcal{M}_1)}{p(\mathbf{y}|\mathcal{M}_2)} \times \frac{\pi(\mathcal{M}_1)}{\pi(\mathcal{M}_2)}, \tag{1.8}$$

where the first term in equation 1.8 is named the Bayes factor, and the second term is the prior odds. Observe that the Bayes factor is a ratio of ordinates for \mathbf{y} under different models. Then, the Bayes factor is a measure of relative sample evidence in favor of model 1 compared to model 2.

However, we still need to calculate $p(\mathbf{y}|\mathcal{M}_m) = \int_{\Theta} p(\mathbf{y}|\boldsymbol{\theta}, \mathcal{M}_m) \pi(\boldsymbol{\theta}|\mathcal{M}_m) d\boldsymbol{\theta} = \mathbb{E}\left[p(\mathbf{y}|\boldsymbol{\theta}, \mathcal{M}_m)\right]$. For this integral to be meaningful, the prior must be proper. Using improper prior has unintended consequences when comparing models, for instance, parsimonious models are favored by posterior odds or Bayes factors depend on units of measure (see Chapter 4).

A nice feature of comparing models using posterior odds is that if we have an exhaustive set of competing models such that $\sum_{m=1}^{M} \pi(\mathcal{M}_m|\mathbf{y}) = 1$, then we can recover $\pi(\mathcal{M}_m|\mathbf{y})$ without calculating $p(\mathbf{y})$. In particular, given two models \mathcal{M}_1 and \mathcal{M}_2 such that $\pi(\mathcal{M}_1|\mathbf{y}) + \pi(\mathcal{M}_2|\mathbf{y}) = 1$. Then, $\pi(\mathcal{M}_1|\mathbf{y}) = \frac{PO_{12}}{1+PO_{12}}$ and $\pi(\mathcal{M}_2|\mathbf{y}) = 1 - \pi(\mathcal{M}_1|\mathbf{y})$. In general, $\pi(\mathcal{M}_m|\mathbf{y}) = \frac{p(\mathbf{y}|\mathcal{M}_m) \times \pi(\mathcal{M}_m)}{\sum_{l=1}^{M} p(\mathbf{y}|\mathcal{M}_l) \times \pi(\mathcal{M}_l)}$. These posterior model probabilities can be used to perform Bayesian model averaging.

Table 1.1 shows guidelines for the interpretation of $2 \log(PO_{12})$ [44]. This transformation is done to replicate the structure of the likelihood ratio test statistic. However, posterior odds do not require nested models as the likelihood ratio test does.

Observe that the posterior odds ratio is a relative criterion, that is, we specify an exhaustive set of competing models, and compare them. However, we may want to check the performance of a model in its own or use a non-informative prior. In this case, we can use the posterior predictive p-value [28, 29].⁶

 $^{^{6}}$ [2] show potential issues due to using data twice in the construction of the predictive p values. They also present alternative proposals, for instance, the partial posterior predictive p value.

TABLE 1.1Kass and Raftery guidelines.

$2 \times \log(PO_{12})$	PO_{12}	Evidence against \mathcal{M}_2
0 to 2	1 to 3	Not worth more than a bare mention
2 to 6	3 to 20	Positive
6 to 10	20 to 150	Strong
> 10	> 150	Very strong

The intuition behind the predictive p-value is simple: analyze discrepancy between model's assumptions and data by checking a potential extreme tailarea probability. Observe that this approach does not check if a model is true, its focus is on potential discrepancies between a model and the data at hand.

This is done simulating pseudo-data from our sampling model $(\mathbf{y}^{(s)}, s = 1, 2, ..., S)$ using draws from the posterior distribution, and then calculating a discrepancy measure, $D(\mathbf{y}^{(s)}, \boldsymbol{\theta})$, to estimate the posterior predictive p-value, $p_D(\mathbf{y}) = P[D(\mathbf{y}^{(s)}, \boldsymbol{\theta}) \geq D(\mathbf{y}, \boldsymbol{\theta})]$ using the proportion of the S draws for which $D(\mathbf{y}^{(s)}, \boldsymbol{\theta}^{(s)}) \geq D(\mathbf{y}, \boldsymbol{\theta}^{(s)})$. Extreme tail probabilities $(p(D_{\mathbf{y}}) \leq 0.05 \text{ or } p(D_{\mathbf{y}}) \geq 0.95)$ suggest potential discrepancy between the data and the model. [29] also suggest the posterior predictive p-value based on the minimum discrepancy, $D_{min}(\mathbf{y}) = \min_{\boldsymbol{\theta}} D(\mathbf{y}, \boldsymbol{\theta})$, and the average discrepancy statistic $D(\mathbf{y}) = \mathbb{E}[D(\mathbf{y}, \boldsymbol{\theta})] = \int_{\boldsymbol{\Theta}} D(\mathbf{y}, \boldsymbol{\theta}) \pi(\boldsymbol{\theta}|\mathbf{y}) d\boldsymbol{\theta}$. These alternatives can be more computational demanding.

The Bayesian approach is also suitable to get probabilistic predictions, that is, we can obtain a posterior predictive density

$$\pi(\mathbf{Y}_0|\mathbf{y}, \mathcal{M}_m) = \int_{\mathbf{\Theta}} \pi(\mathbf{Y}_0, \boldsymbol{\theta}|\mathbf{y}, \mathcal{M}_m) d\boldsymbol{\theta}$$
$$= \int_{\mathbf{\Theta}} \pi(\mathbf{Y}_0|\boldsymbol{\theta}, \mathbf{y}, \mathcal{M}_m) \pi(\boldsymbol{\theta}|\mathbf{y}, \mathcal{M}_m) d\boldsymbol{\theta}. \tag{1.9}$$

Observe that equation 1.9 is again an expectation $\mathbb{E}[\pi(\mathbf{Y}_0|\boldsymbol{\theta},\mathbf{y},\mathcal{M}_m)]$, this time using the posterior distribution. Therefore, the Bayesian approach takes estimation error into account when performing prediction.

As we have shown many times, expectation (integration) is a common feature in Bayesian inference. That is why the remarkable relevance of computation based on *Monte Carlo integration* in the Bayesian framework (see Chapter 5).

Bayesian model average (BMA) allows considering model uncertainty in prediction or any unknown probabilistic object. In the prediction case,

$$\pi(\mathbf{Y}_0|\mathbf{y}) = \sum_{m=1}^{M} \pi(\mathcal{M}_m|\mathbf{y})\pi(\mathbf{Y}_0|\mathbf{y}, \mathcal{M}_m), \tag{1.10}$$

and parameters case,

$$\pi(\boldsymbol{\theta}|\mathbf{y}) = \sum_{m=1}^{M} \pi(\mathcal{M}_m|\mathbf{y})\pi(\boldsymbol{\theta}|\mathbf{y}, \mathcal{M}_m), \qquad (1.11)$$

where

$$\mathbb{E}(\boldsymbol{\theta}|\mathbf{y}) = \sum_{m=1}^{M} \hat{\boldsymbol{\theta}}_{m} \pi(\mathcal{M}_{m}|\mathbf{y}), \tag{1.12}$$

and

$$Var(\boldsymbol{\theta}|\mathbf{y}) = \sum_{m=1}^{M} \pi(\mathcal{M}_m|\mathbf{y}) \widehat{Var}(\boldsymbol{\theta}|\mathbf{y}, \mathcal{M}_m) + \sum_{m=1}^{M} \pi(M_m|\mathbf{y}) (\hat{\boldsymbol{\theta}}_m - \mathbb{E}[\boldsymbol{\theta}|\mathbf{y}])^2,$$
(1.13)

 $\hat{\boldsymbol{\theta}}_m$ and $\widehat{Var}(\boldsymbol{\theta}|\mathbf{y}, \mathcal{M}_m)$ are the posterior mean and variance under model m, respectively.

Observe how the variance in equation 1.13 encloses extra variability due to potential differences between mean posterior estimates associated with each model, and the posterior mean involving model uncertainty in equation 1.12.

A nice advantage of the Bayesian approach, which is very useful in *state* space representations (see Chapter 8), is the way that the posterior distribution updates with new sample information. Given $\mathbf{y} = \mathbf{y}_{1:t+1}$ a sequence of observations from 1 to t+1, then

$$\pi(\boldsymbol{\theta}|\mathbf{y}_{1:t+1}) \propto p(\mathbf{y}_{1:t+1}|\boldsymbol{\theta}) \times \pi(\boldsymbol{\theta})$$

$$= p(y_{t+1}|\mathbf{y}_{1:t},\boldsymbol{\theta}) \times p(\mathbf{y}_{1:t}|\boldsymbol{\theta}) \times \pi(\boldsymbol{\theta})$$

$$\propto p(y_{t+1}|\mathbf{y}_{1:t},\boldsymbol{\theta}) \times \pi(\boldsymbol{\theta}|\mathbf{y}_{1:t}). \tag{1.14}$$

We observe that the new prior is just the posterior distribution using the previous observations. This is particular useful under the assumption of conditional independence, that is, $y_{t+1} \perp \mathbf{y}_{1:t}|\boldsymbol{\theta}$, then $p(y_{t+1}|\mathbf{y}_{1:t},\boldsymbol{\theta}) = p(y_{t+1}|\boldsymbol{\theta})$ such that the posterior can be recovered recursively [60]. This facilities online updating due to all information up to t being in $\boldsymbol{\theta}$. Then, $\pi(\boldsymbol{\theta}|\mathbf{y}_{1:t+1}) \propto p(y_{t+1}|\boldsymbol{\theta}) \times \pi(\boldsymbol{\theta}|\mathbf{y}_{1:t}) \propto \prod_{h=1}^{t+1} p(y_h|\boldsymbol{\theta}) \times \pi(\boldsymbol{\theta})$. This recursive expression can be calculated faster at some specific point in time t compared to a batch mode algorithm, which requires processing simultaneously all information up to t.

It is also important to wonder about the sampling properties of "Bayesian estimators". This topic has attracted attention of statisticians and econometricians long time ago. For instance, asymptotic posterior concentration at the population parameter vector is discussed by [11]. Convergence of posterior distributions is stated by the Bernstein-von Mises theorem [49], which

creates a link between credible intervals (sets) and confidence intervals (sets), where a credible interval is an interval in the domain of the posterior distribution within which an unknown parameter falls with a particular probability. Credible intervals treat bounds as fixed and parameters as random, whereas confidence intervals reverse this. There are many settings in parametric models where Bayesian credible intervals with α level converge asymptotically to confidence intervals at α level. This suggests that Bayesian inference is asymptotically correct from a sampling perspective in these settings.

A heuristic approach to show this in the simplest case where we assume random sampling and $\theta \in \mathcal{R}$ is the following: $p(\mathbf{y}|\theta) = \prod_{i=1}^N p(y_i|\theta)$ such that the log likelihood is $l(\mathbf{y}|\theta) \equiv \log p(\mathbf{y}|\theta) = \sum_{i=1}^N \log p(y_i|\theta) = N \times \bar{l}(\mathbf{y}|\theta)$ where $\bar{l} \equiv \frac{1}{N} \sum_{i=1}^N \log p(y_i|\theta)$ is the mean likelihood. Then, the posterior distribution is proportional to

$$\pi(\theta|\mathbf{y}) \propto p(\mathbf{y}|\theta) \times \pi(\theta)$$

$$= \exp\left\{N \times \bar{l}(\mathbf{y}|\theta)\right\} \times \pi(\theta). \tag{1.15}$$

Observe that as the sample size gets large, that is, $N \to \infty$, the exponential term should dominate the prior distribution as long as this does not depend on N such that the likelihood determines the posterior distribution asymptotically.

Maximum likelihood theory shows that $\lim_{N\to\infty} \bar{l}(\mathbf{y}|\theta) \to \bar{l}(\mathbf{y}|\theta_0)$ where θ_0 is the population parameter of the data generating process. In addition, doing a second order Taylor expansion of the log likelihood at the Maximum likelihood estimator,

$$l(\mathbf{y}|\theta) \approx l(\mathbf{y}|\hat{\theta}) + \frac{dl(\mathbf{y}|\theta)}{d\theta} \Big|_{\hat{\theta}} (\theta - \hat{\theta}) + \frac{1}{2} \frac{d^2 l(\mathbf{y}|\theta)}{d\theta^2} \Big|_{\hat{\theta}} (\theta - \hat{\theta})^2$$

$$= l(\mathbf{y}|\hat{\theta}) + \frac{1}{2} \sum_{i=1}^{N} \frac{d^2 l(y_i|\theta)}{d\theta^2} \Big|_{\hat{\theta}} (\theta - \hat{\theta})^2$$

$$= l(\mathbf{y}|\hat{\theta}) - \frac{1}{2} N \left[-\bar{l}''|_{\hat{\theta}} \right] (\theta - \hat{\theta})^2$$

$$= l(\mathbf{y}|\hat{\theta}) - \frac{N}{2\sigma^2} (\theta - \hat{\theta})^2$$

where
$$\frac{dl(\mathbf{y}|\theta)}{d\theta}\Big|_{\hat{\theta}} = 0$$
, $\bar{l}'' \equiv \frac{1}{N} \sum_{i=1}^{N} \frac{d^{2}l(y_{i}|\theta)}{d\theta^{2}}\Big|_{\hat{\theta}}$ and $\sigma^{2} := \left[-\bar{l}''|_{\hat{\theta}}\right]^{-1}$. 8 Then,

⁷Take into account that in the likelihood function the argument is θ . However, we keep the notation for facility in exposition.

⁸The last definition follows from standard theory in maximum likelihood estimation.

$$\pi(\theta|\mathbf{y}) \propto \exp\left\{l(\mathbf{y}|\theta)\right\} \times \pi(\theta)$$

$$\approx \exp\left\{l(\mathbf{y}|\hat{\theta}) - \frac{N}{2\sigma^2}(\theta - \hat{\theta})^2\right\} \times \pi(\theta)$$

$$\propto \exp\left\{-\frac{N}{2\sigma^2}(\theta - \hat{\theta})^2\right\} \times \pi(\theta)$$

Observe that we have that the posterior density is proportional to the kernel of a normal density with mean $\hat{\theta}$ and variance σ^2/N as long as $\pi(\hat{\theta}) \neq 0$. This kernel dominates as the sample size gets large due to N in the exponential term. Observe that the prior should not exclude values of θ that are logically possible, such as $\hat{\theta}$.

1.2.1 Example: Health insurance

Suppose that you are analyzing to buy a health insurance next year. To make a better decision you want to know what is the probability that you visit your Doctor at least once next year? To answer this question you have records of the number of times that you have visited your Doctor the last 5 years, $y = \{0, 3, 2, 1, 0\}$. How to proceed?

Assuming that this is a random sample⁹ from a data generating process (statistical model) that is Poisson, that is, $Y_i \sim P(\lambda)$, and your probabilistic prior beliefs about λ are well described by a Gamma distribution with shape and scale parameters α_0 and β_0 , $\lambda \sim G(\alpha_0, \beta_0)$, then, you are interested in calculating the probability $P(Y_0 > 0|\mathbf{y})$. You need to calculate the posterior predictive density $\pi(Y_0|\mathbf{y})$ to answer this question in a Bayesian way.

In this example, $p(\mathbf{y}|\lambda)$ is Poisson, and $\pi(\lambda)$ is Gamma. Then, using 1.9

$$\pi(Y_0|\mathbf{y}) = \int_0^\infty \frac{\lambda^{y_0} \exp\left\{-\lambda\right\}}{y_0!} \times \pi(\lambda|\mathbf{y}) d\lambda,$$

where the posterior distribution is $\pi(\lambda|\mathbf{y}) \propto \lambda^{\sum_{i=1}^{N} y_i + \alpha_0 - 1} \exp\left\{-\lambda\left(\frac{\beta_0 N + 1}{\beta_0}\right)\right\}$ by equation 1.3.

Observe that the last expression is the kernel of a Gamma distribution with parameters $\alpha_n = \sum_{i=1}^N y_i + \alpha_0$ and $\beta_n = \frac{\beta_0}{\beta_0 N+1}$. Given that $\int_0^\infty \pi(\lambda|\mathbf{y}) d\lambda = 1$, then the constant of proportionality in the last expression is $\Gamma(\alpha_n)\beta_n^{\alpha_n}$, where $\Gamma(\cdot)$ is the gamma function. Thus, the posterior density function $\pi(\lambda|\mathbf{y})$ is $G(\alpha_n, \beta_n)$.

Observe that

⁹Independent and identically distributed draws.

$$\mathbb{E}[\lambda|\mathbf{y}] = \alpha_n \beta_n$$

$$= \left(\sum_{i=1}^N y_i + \alpha_0\right) \left(\frac{\beta_0}{\beta_0 N + 1}\right)$$

$$= \bar{y} \left(\frac{N\beta_0}{N\beta_0 + 1}\right) + \alpha_0 \beta_0 \left(\frac{1}{N\beta_0 + 1}\right)$$

$$= w\bar{y} + (1 - w)\mathbb{E}[\lambda],$$

where \bar{y} is the sample mean, which is the maximum likelihood estimator of λ in this example, $w = \left(\frac{N\beta_0}{N\beta_0+1}\right)$ and $\mathbb{E}[\lambda] = \alpha_0\beta_0$ is the prior mean. The posterior mean is a weighted average of the maximum likelihood estimator (sample information) and the prior mean. Observe that $\lim_{N\to\infty} w = 1$, that is, the sample information asymptotically dominates.

The predictive distribution is

$$\begin{split} \pi(Y_0|\mathbf{y}) &= \int_0^\infty \frac{\lambda^{y_0} \exp\left\{-\lambda\right\}}{y_0!} \times \frac{1}{\Gamma(\alpha_n)\beta_n^{\alpha_n}} \lambda^{\alpha_n - 1} \exp\left\{-\lambda/\beta_n\right\} d\lambda \\ &= \frac{1}{y_0! \Gamma(\alpha_n)\beta_n^{\alpha_n}} \int_0^\infty \lambda^{y_0 + \alpha_n - 1} \exp\left\{-\lambda \left(\frac{1 + \beta_n}{\beta_n}\right)\right\} d\lambda \\ &= \frac{\Gamma(y_0 + \alpha_n) \left(\frac{\beta_n}{\beta_n + 1}\right)^{y_0 + \alpha_n}}{y_0! \Gamma(\alpha_n)\beta_n^{\alpha_n}} \\ &= \left(\frac{y_0 + \alpha_n - 1}{y_0}\right) \left(\frac{\beta_n}{\beta_n + 1}\right)^{y_0} \left(\frac{1}{\beta_n + 1}\right)^{\alpha_n}. \end{split}$$

The third equality follows from the kernel of a Gamma density, and the fourth from $\binom{y_0+\alpha_n-1}{y_0}=\frac{(y_0+\alpha_n-1)(y_0+\alpha_n-2)...\alpha_n}{y_0!}=\frac{\Gamma(y_0+\alpha_n)}{\Gamma(\alpha_n)y_0!}$ using a property of the Gamma function.

Observe that this is a Negative Binomial density, that is $Y_0|\mathbf{y} \sim NB(\alpha_n, p_n)$ where $p_n = \frac{\beta_n}{\beta_n + 1}$.

Up to this point, we have said nothing about the hyperparameters, which are required to give a concrete response to this exercise. Thus, we show two approaches to set them. First, we set $\alpha_0 = 0.001$ and $\beta_0 = 1/0.001$ which imply vague prior information about λ due to having a large degree of variability compared to the mean information.¹⁰ In particular, $\mathbb{E}[\lambda] = 1$ and $\mathbb{V}ar[\lambda] = 1000$.

In this setting, $P(Y_0 > 0|\mathbf{y}) = 1 - P(Y_0 = 0|\mathbf{y}) \approx 0.67$. That is, the probability of visiting the Doctor at least once next year is approximately 0.67.

¹⁰We should be aware that there may be technical problems using this king of hyperparameters in this setting [31].

Another approach is using *Empirical Bayes*, where we set the hyperparameters maximizing the logarithm of the marginal likelihood, that is, $\left[\hat{\alpha}_0 \ \hat{\beta}_0\right]^{\top} = \underset{\alpha_0,\beta_0}{\operatorname{argmax}} \, \ln p(\mathbf{y})$ where

$$p(\mathbf{y}) = \int_0^\infty \left\{ \frac{1}{\Gamma(\alpha_0)\beta_0^{\alpha_0}} \lambda^{\alpha_0 - 1} \exp\left\{-\lambda/\beta_0\right\} \prod_{i=1}^N \frac{\lambda^{y_i} \exp\left\{-\lambda\right\}}{y_i!} \right\} d\lambda$$

$$= \frac{\int_0^\infty \lambda^{\sum_{i=1}^N y_i + \alpha_0 - 1} \exp\left\{-\lambda \left(\frac{\beta_0 N + 1}{\beta_0}\right)\right\} d\lambda}{\Gamma(\alpha_0)\beta_0^{\alpha_0} \prod_{i=1}^N y_i!}$$

$$= \frac{\Gamma(\sum_{i=1}^N y_i + \alpha_0) \left(\frac{\beta_0}{N\beta_0 + 1}\right)^{\sum_{i=1}^N y_i} \left(\frac{1}{N\beta_0 + 1}\right)^{\alpha_0}}{\Gamma(\alpha_0) \prod_{i=1}^N y_i}$$

Using the empirical Bayes approach, we get $\hat{\alpha}_0 = 51.8$ and $\hat{\beta}_0 = 0.023$, then $P(Y_0 > 0|\mathbf{y}) = 1 - P(Y_0 = 0|\mathbf{y}) \approx 0.70$.

Observe that we can calculate the posterior odds comparing the model using an Empirical Bayes prior (model 1) versus the vague prior (model 2). We assume that $\pi(\mathcal{M}_1) = \pi(\mathcal{M}_2) = 0.5$, then

$$PO_{12} = \frac{p(\mathbf{y}|\text{Empirical Bayes})}{p(\mathbf{y}|\text{Vague prior})}$$

$$= \frac{\frac{\Gamma(\sum_{i=1}^{N} y_i + 51.807) \left(\frac{0.023}{N \times 0.023 + 1}\right)^{\sum_{i=1}^{N} y_i} \left(\frac{1}{N \times 0.023 + 1}\right)^{51.807}}{\Gamma(\sum_{i=1}^{N} y_i + 0.001) \left(\frac{1/0.001}{N/0.001 + 1}\right)^{\sum_{i=1}^{N} y_i} \left(\frac{1}{N/0.001 + 1}\right)^{0.001}}{\Gamma(0.001)}$$

$$\approx 919.$$

Then, $2 \times \log(PO_{12}) = 13.64$, there is very strong evidence against the vague prior model (see Table 1.1). In particular, $\pi(\text{Empirical Bayes}|\mathbf{y}) = \frac{919}{1+919} = 0.999$ and $\pi(\text{Vague prior}|\mathbf{y}) = 1 - 0.999 = 0.001$. These probabilities can be used to perform Bayesian model average (BMA). In particular,

$$\mathbb{E}(\lambda|\mathbf{y}) = 1.2 \times 0.999 + 1.2 \times 0.001 = 1.2$$

$$Var(\lambda|\mathbf{y}) = 0.025 \times 0.999 + 0.24 \times 0.001$$

$$+ (1.2 - 1.2)^2 \times 0.999 + (1.2 - 1.2)^2 \times 0.001 = 0.025.$$

The BMA predictive distribution is a mix of negative binomial distributions, that is, $y_0|\mathbf{y} \sim 0.999 \times NB(57.8, 0.02) + 0.001 \times NB(6.001, 0.17)$.

R code. Health insurance, predictive distribution using vague hyperparameters

```
set.seed(010101)
2 y <- c(0, 3, 2, 1, 0) # Data
3 N <- length(y)
4 ProbBo <- function(y, a0, b0){
    N <- length(y)
    #sample size
    an \leftarrow a0 + sum(y)
     # Posterior shape parameter
    bn <- b0 / ((b0 * N) + 1)
# Posterior scale parameter
    p <- bn / (bn + 1)
     # Probability negative binomial density
     Pr <- 1 - pnbinom(0, size=an, prob=(1 - p))</pre>
     # Probability of visiting the Doctor at least once next
     # Observe that in R there is a slightly different
       parametrization.
     return(Pr)
16
17 }
18 # Using a vague prior:
19 a0 <- 0.001 # Prior shape parameter
_{20} b0 <- 1 / 0.001 # Prior scale parameter
PriMeanV <- a0 * b0 # Prior mean
PriVarV <- a0 * b0^2 # Prior variance
^{23} Pp <- ProbBo(y, a0 = 0.001, b0 = 1 / 0.001)
24 # This setting is vague prior information.
25 Pp
26 0.67
```

R code. Health insurance, predictive distribution using empirical Bayes

```
1 # Using Empirical Bayes
2 LogMgLik <- function(theta, y){</pre>
3 N <- length(y)
#sample size
a0 <- theta[1]</pre>
6 # prior shape hyperparameter
7 b0 <- theta[2]
_{8} # prior scale hyperparameter
   an <- sum(y) + a0
   # posterior shape parameter
   if(a0 <= 0 || b0 <= 0){
    #Avoiding negative values
    lnp <- -Inf
13
14
    }else{
    lnp <- lgamma(an) + sum(y)*log(b0/(N*b0+1)) - a0*log(N*b0
       +1) - lgamma(a0)
16 }
17 # log marginal likelihood
   return(-lnp)
19 }
20 theta0 \leftarrow c(0.01, 1/0.1)
_{21} # Initial values
22 control <- list(maxit = 1000)</pre>
23 # Number of iterations in optimization
24 EmpBay <- optim(theta0, LogMgLik, method = "BFGS", control =
       control , hessian = TRUE , y = y)
25 # Optimization
26 EmpBay $convergence
28 aOEB <- EmpBay par [1]
29 # Prior shape using empirical Bayes
30 a0EB
31 51.81
32 bOEB <- EmpBay par [2]
33 # Prior scale using empirical Bayes
34 b0EB
35 0.023
36 PriMeanEB <- a0EB * b0EB
37 # Prior mean
38 PriVarEB <- a0EB * b0EB^2
39 # Prior variance
40 PpEB \leftarrow ProbBo(y, a0 = a0EB, b0 = b0EB)
41 # This setting is using emprical Bayes.
42 PpEB
43 0.70
```

R code. Health insurance, density plots

```
1 # Density figures:
2 # This code helps plotting densities
_3 lambda <- seq(0.01, 10, 0.01)
4 # Values of lambda
5 VaguePrior <- dgamma(lambda,shape=a0,scale = b0)</pre>
6 EBPrior <- dgamma(lambda, shape=a0EB, scale = b0EB)
7 PosteriorV <- dgamma(lambda, shape = a0 + sum(y), scale = b0</pre>
       / ((b0 * N) + 1))
8 PosteriorEB <- dgamma(lambda, shape = a0EB+sum(y), scale =</pre>
      bOEB / ((bOEB * N) + 1))
9 # Likelihood function
10 Likelihood <- function(theta, y){</pre>
11 LogL <- dpois(y, theta, log = TRUE)</pre>
12 Lik <- prod(exp(LogL))</pre>
13 return(Lik)
14 }
15 Liks <- sapply(lambda, function(par) {Likelihood(par, y = y)
      })
16 Sc <- max(PosteriorEB)/max(Liks)</pre>
17 #Scale for displaying in figure
18 LiksScale <- Liks * Sc
19 data <- data.frame(cbind(lambda, VaguePrior, EBPrior,</pre>
      PosteriorV, PosteriorEB, LiksScale)) #Data frame
20 require(ggplot2) # Cool figures
21 require(latex2exp) # LaTeX equations in figures
require(ggpubr) # Multiple figures in one page
23 fig1 <- ggplot(data = data, aes(lambda, VaguePrior)) + geom_
      line() + xlab(TeX("$\\lambda$")) + ylab("Density") +
      ggtitle("Prior: Vague Gamma")
24 fig2 <- ggplot(data = data, aes(lambda, EBPrior)) + geom_
      line() + xlab(TeX("$\\lambda$")) + ylab("Density") +
      ggtitle("Prior: Empirical Bayes Gamma")
25 fig3 <- ggplot(data = data, aes(lambda, PosteriorV)) + geom_
      line() + xlab(TeX("$\\lambda$")) + ylab("Density") +
      ggtitle("Posterior: Vague Gamma")
26 fig4 <- ggplot(data = data, aes(lambda, PosteriorEB)) + geom
       _{	t line()} + {	t xlab(TeX("$ \ \ )} + {	t ylab("Density")} +
      ggtitle("Posterior: Empirical Bayes Gamma")
27 FIG <- ggarrange(fig1, fig2, fig3, fig4, ncol = 2, nrow = 2)
28 annotate_figure(FIG, top = text_grob("Vague versus Empirical
       Bayes: Poisson-Gamma model", color = "black", face = "
      bold", size = 14))
29 dataNew <- data.frame(cbind(rep(lambda, 3), c(EBPrior,
      PosteriorEB, LiksScale), rep(1:3, each = 1000))) #Data
30 colnames(dataNew) <- c("Lambda", "Density", "Factor")
31 dataNew$Factor <- factor(dataNew$Factor, levels=c("1", "3",</pre>
      "2"),
32 labels=c("Prior", "Likelihood", "Posterior"))
33 ggplot(data = dataNew, aes_string(x = "Lambda", y = "Density
       ", group = "Factor")) + geom_line(aes(color = Factor)) +
       xlab(TeX("$\\lambda$")) + ylab("Density") + ggtitle("
      Prior, likelihood and posterior: Empirical Bayes Poisson
      -Gamma model") + guides(color=guide_legend(title="
      Information")) + scale_color_manual(values = c("red", "
      yellow", "blue"))
```



FIGURE 1.2 Vague versus Empirical Bayes: Poisson-Gamma model.



FIGURE 1.3 Prior, likelihood and posterior: Empirical Bayes Poisson-Gamma model.

Figure 1.2 displays prior and posterior densities based on vague and Empirical Bayes hyperparameters. We see that prior and posterior densities using the latter are more informative as expected.

Figure 1.3 shows the prior, scaled likelihood and posterior densities of λ based on the hyperparameters of the Empirical Bayes approach. The posterior density is a compromise between prior and sample information.

R code. Health insurance, Predictive density

```
# Predictive distributions
  PredDen <- function(y, y0, a0, b0){</pre>
    N <- length(y)
    #sample size
    an \leftarrow a0 + sum(y)
    # Posterior shape parameter
    bn \leftarrow b0 / ((b0 * N) + 1)
    # Posterior scale parameter
    p <- bn / (bn + 1)
    # Probability negative binomial density
    Pr <- dnbinom(y0, size=an, prob=(1 - p))
    # Predictive density
12
    # Observe that in R there is a slightly different
      parametrization.
     return(Pr)
15 }
16 y0 <- 0:10
17 PredVague <- PredDen(y=y, y0=y0, a0=a0, b0=b0)
18 PredEB <- PredDen(y=y, y0=y0, a0=a0EB, b0=b0EB)
19 dataPred <- as.data.frame(cbind(y0, PredVague, PredEB))</pre>
20 colnames(dataPred) <- c("y0", "PredictiveVague",</pre>
       PredictiveEB")
ggplot(data = dataPred) + geom_point(aes(y0, PredictiveVague
       , color = "red")) +
22 xlab(TeX("$y_0$")) + ylab("Density") + ggtitle("Predictive
      density: Vague and Empirical Bayes priors") + geom_point
(aes(y0, PredictiveEB, color = "yellow")) +
guides(color = guide_legend(title="Prior")) + scale_color_
      manual(labels = c("Vague", "Empirical Bayes"), values =
       c("red", "yellow")) + scale_x_continuous(breaks=seq
       (0,10,by=1))
```

Figure 1.4 displays the predictive probability mass of not having any visits to a physician the next year, having one, two, and so on using Empirical Bayes and vague hyperparameters. The predictive probability of not having any visits are approximately equal to 30% and 33% based on the Empirical Bayes and vague hyperparameters.



FIGURE 1.4
Predictive density: Vague and Empirical Bayes.

R code. Health insurance, Bayesian model average 1 # Posterior odds: Vague vs Empirical Bayes 2 PO12 <- exp(-LogMgLik(c(aOEB, bOEB), y = y))/exp(-LogMgLik(c (a0, b0), y = y))3 PO12 4 919 5 PostProMEM <- P012/(1 + P012)</pre> 6 PostProMEM 8 # Posterior model probability Empirical Bayes 9 PostProbMV <- 1 - PostProMEM 10 PostProbMV 11 0.002 12 # Posterior model probability vague prior 13 # Bayesian model average (BMA) 14 PostMeanEB \leftarrow (a0EB + sum(y)) * (b0EB / (b0EB * N + 1)) 15 # Posterior mean Empirical Bayes 16 PostMeanV \leftarrow (a0 + sum(y)) * (b0 / (b0 * N + 1)) 17 # Posterior mean vague priors 18 BMAmean <- PostProMEM * PostMeanEB + PostProbMV * PostMeanV 19 BMAmean 20 1.2 21 # BMA posterior mean 22 PostVarEB <- (a0EB + sum(y)) * (b0EB/(b0EB * N + 1))^2 23 # Posterior variance Empirical Bayes 24 PostVarV \leftarrow (a0 + sum(y)) * (b0 / (b0 * N + 1))^2 25 # Posterior variance vague prior 26 BMAVar <- PostProMEM * PostVarEB + PostProbMV*PostVarV + PostProMEM * (PostMeanEB - BMAmean)^2 + PostProbMV * (PostMeanV - BMAmean)^2 27 # BMA posterior variance 28 BMAVar 29 0.025

$ig| R \ code. \ Health \ insurance, \ Bayesian \ model \ average$

Predictive density: BMA



FIGURE 1.5 Bayesian model average: Predictive density.

Figure 1.5 displays the predictive density using Bayesian model average based on the vague and Empirical Bayes hyperparameters. This figure essentially resembles the predictive probability mass function based on the Empirical Bayes framework, as the posterior model probability for that setting is nearly one.

Figure 1.6 displays how the posterior distribution updates given new sample information based on an initial non-informative prior (iteration 1). We



FIGURE 1.6 Bayesian updating: Posterior densities.

see that iteration 5 is based on all the sample information in our example, as a consequence, the posterior density in iteration 5 is equal to the posterior density in Figure 1.3.

R code. Health insurance, Bayes updating

```
1 # Bayesian updating
2 BayUp <- function(y, lambda, a0, b0){</pre>
    N <- length(y)
    #sample size
    an \leftarrow a0 + sum(y)
    # Posterior shape parameter
    bn <- b0 / ((b0 * N) + 1)
     # Posterior scale parameter
    p <- dgamma(lambda, shape = an, scale = bn)
    # Posterior density
    return(list(Post = p, a0New = an, b0New = bn))
12 }
13 PostUp <- NULL
14 for(i in 1:N){
   if(i == 1){
15
       PostUpi \leftarrow BayUp(y[i], lambda, a0 = 0.001, b0 = 1/0.001)
     else{
      PostUpi <- BayUp(y[i], lambda, a0 = PostUpi$a0New, b0 =
18
       PostUpi$b0New)
19
    PostUp <- cbind(PostUp, PostUpi$Post)</pre>
20
21 }
22 DataUp <- data.frame(cbind(rep(lambda, 5), c(PostUp), rep</pre>
       (1:5, each = 1000))) #Data frame
23 colnames(DataUp) <- c("Lambda", "Density", "Factor")</pre>
24 DataUp$Factor <- factor(DataUp$Factor, levels=c("1", "2", "3
       ", "4", "5"),
25 labels=c("Iter 1", "Iter 2", "Iter 3", "Iter 4", "Iter 5"))
26 ggplot(data = DataUp, aes_string(x = "Lambda", y = "Density"
       , group = "Factor")) + geom_line(aes(color = Factor)) +
       xlab(TeX("$\\lambda$")) + ylab("Density") + ggtitle("
       Bayesian updating: Poisson-Gamma model with vague prior"
       ) + guides(color=guide_legend(title="Update")) + scale_
       color_manual(values = c("red", "purple", "blue", "yellow
       ", "black"))
27 S <- 100000 # Posterior draws
28 PostMeanLambdaUps <- sapply(1:N, function(i) {mean(sample())</pre>
       lambda, S, replace = TRUE, prob = PostUp[ , i])))) #
       Posterior mean update i
29 paste("Posterior means using all information and sequential
       updating are:", round(PostMeanV, 2), "and", round(
       PostMeanLambdaUps[5], 2), sep = " ")
30 Posterior means using all information and sequential
       updating are: 1.2 and 1.2
31 PostVarLambdaUps <- sapply(1:N, function(i) {var(sample(</pre>
       lambda, S, replace = TRUE, prob = PostUp[ , i])))) #
       Posterior variance update i
32 paste("Posterior variances using all information and
       sequential updating are:", round(PostVarV, 2), "and",
round(PostVarLambdaUps[5], 2), sep = " ")
33 Posterior variances using all information and sequential
       updating are: 0.24 and 0.24
```

1.3 Bayesian reports: Decision theory under uncertainty

The Bayesian framework allows reporting the full posterior distributions. However, some situations demand to report a specific value of the posterior distribution (point estimate), an informative interval (set), point or interval predictions and/or selecting a specific model. Decision theory offers an elegant framework to make a decision regarding what are the optimal posterior values to report [9].

The point of departure is a loss function, which is a non-negative real value function whose arguments are the unknown state of nature (Θ) , and a set of actions to be made (A), that is,

$$L(\boldsymbol{\theta}, a) : \boldsymbol{\Theta} \times \mathcal{A} \to \mathcal{R}^+.$$

This function is a mathematical expression of the loss of making mistakes. In particular, selecting action $a \in \mathcal{A}$ when $\theta \in \Theta$ is the true. In our case, the unknown state of nature can be parameters, functions of them, future or unknown realizations, models, etc.

From a Bayesian perspective, we should choose the action that minimizes the posterior expected loss $(a^*(\mathbf{y}))$, that is, the posterior risk function $(\mathbb{E}[L(\boldsymbol{\theta}, a)|\mathbf{y}])$,

$$a^*(\mathbf{y}) = \underset{a \in \mathcal{A}}{\operatorname{argmin}} \ \mathbb{E}[L(\boldsymbol{\theta}, a) | \mathbf{y}],$$

where $\mathbb{E}[L(\boldsymbol{\theta}, a)|\mathbf{y}] = \int_{\boldsymbol{\Theta}} L(\boldsymbol{\theta}, a) \pi(\boldsymbol{\theta}|\mathbf{y}) d\boldsymbol{\theta}$. 11

Different loss functions imply different optimal decisions. We illustrate this assuming $\theta \in \mathcal{R}$.

• The quadratic loss function, $L(\theta, a) = [\theta - a]^2$, gives as optimal decision the posterior mean, $a^*(\mathbf{y}) = \mathbb{E}[\theta|\mathbf{y}]$, that is

$$\mathbb{E}[\theta|\mathbf{y}] = \underset{a \in \mathcal{A}}{\operatorname{argmin}} \ \int_{\Theta} [\theta - a]^2 \pi(\theta|\mathbf{y}) d\theta.$$

To get this results, let's use the first condition order, differentiate the risk function with respect to a, interchange differential and integral order, and set this equal to zero, $-2\int_{\Theta}[\theta-a^*]\pi(\theta|\mathbf{y})d\theta=0$ implies that $a^*\int_{\Theta}\pi(\theta|\mathbf{y})d\theta=a^*(\mathbf{y})=\int_{\Theta}\theta\pi(\theta|\mathbf{y})d\theta=\mathbb{E}[\theta|\mathbf{y}]$, that is, the posterior mean is the Bayesian optimal action. This means that we should report the posterior mean as a point estimate of θ when facing the quadratic loss function.

¹¹[16] propose Laplace type estimators (LTE) based on the *quasi-posterior*, $p(\theta) = \frac{\exp\{L_n(\theta)\}\pi(\theta)}{\int_{\Theta} \exp\{L_n(\theta)\}\pi(\theta)d\theta}$ where $L_n(\theta)$ is not necessarily a log-likelihood function. The LTE minimizes the *quasi-posterior risk*.

- The generalized quadratic loss function, $L(\theta, a) = w(\theta)[\theta a]^2$, where $w(\theta) > 0$ is a weighting function, gives as optimal decision rule the weighted mean. We should follow same steps as the previous result to get $a^*(\mathbf{y}) = \frac{\mathbb{E}[w(\theta) \times \theta|\mathbf{y}]}{\mathbb{E}[w(\theta)|\mathbf{y}]}$. Observe that the weighted average is driven by the weighting function $w(\theta)$.
- The absolute error loss function, $L(\theta, a) = |\theta a|$, gives as optimal action the posterior median (Exercise 5).
- The generalized absolute error function,

$$L(\theta, a) = \begin{cases} K_0(\theta - a), \theta - a \ge 0 \\ K_1(a - \theta), \theta - a < 0 \end{cases}, K_0, K_1 > 0,$$

implies the following risk function,

$$\mathbb{E}[L(\theta, a)|\mathbf{y}] = \int_{-\infty}^{a} K_1(a - \theta)\pi(\theta|\mathbf{y})d\theta + \int_{a}^{\infty} K_0(\theta - a)\pi(\theta|\mathbf{y})d\theta.$$

Differentiating with respect to a, interchanging differentials and integrals, and equating to zero,

$$K_1 \int_{-\infty}^{a^*} \pi(\theta|\mathbf{y}) d\theta - K_0 \int_{a^*}^{\infty} \pi(\theta|\mathbf{y}) d\theta = 0,$$

then, $\int_{-\infty}^{a^*} \pi(\theta|\mathbf{y}) d\theta = \frac{K_0}{K_0 + K_1}$, that is, any $K_0/(K_0 + K_1)$ -percentile of $\pi(\theta|\mathbf{y})$ is an optimal Bayesian estimate of θ .

We can also use decision theory under uncertainty in hypothesis testing. In particular, testing $H_0: \theta \in \Theta_0$ versus $H_1: \theta \in \Theta_1$, $\Theta = \Theta_0 \cup \Theta_1$ and $\emptyset = \Theta_0 \cap \Theta_1$, there are two actions of interest, a_0 and a_1 , where a_j denotes no rejecting H_j , $j = \{0, 1\}$.

Given the $0 - K_i$ loss function,

$$L(\theta, a_j) = \left\{ \begin{matrix} 0, & \theta \in \Theta_j \\ K_j, & \theta \in \Theta_j, j \neq i \end{matrix} \right\},$$

where there is no loss if the right decision is made, for instance, no rejecting H_0 when $\theta \in \Theta_0$, and the loss is K_j when an error is made, for instance, type I error, rejecting the null hypothesis (H_0) when it is true $(\theta \in \Theta_0)$, implies a loss equal to K_1 due to picking a_1 , no rejecting H_1 .

The posterior expected loss associated with decision a_j , that is, no rejecting H_j , is $\mathbb{E}[L(\theta, a_j)|\mathbf{y}] = 0 \times P(\Theta_j|\mathbf{y}) + K_jP(\Theta_i|\mathbf{y}) = K_jP(\Theta_i|\mathbf{y}), j \neq i$. Therefore, the Bayes optimal decision is the one that gives the smallest posterior expected loss, that is, the null hypothesis is rejected $(a_1 \text{ is not rejected})$, when

 $K_0P(\Theta_1|\mathbf{y}) > K_1P(\Theta_0|\mathbf{y})$. Given our framework $(\Theta = \Theta_0 \cup \Theta_1, \emptyset = \Theta_0 \cap \Theta_1)$, then $P(\Theta_0|\mathbf{y}) = 1 - P(\Theta_1|\mathbf{y})$, and as a consequence, $P(\Theta_1|\mathbf{y}) > \frac{K_1}{K_1 + K_0}$, that is, the rejection region of the Bayesian test is $R = \left\{\mathbf{y} : P(\Theta_1|\mathbf{y}) > \frac{K_1}{K_1 + K_0}\right\}$.

Decision theory also helps to construct interval (region) estimates. Let $\Theta_{C(\mathbf{y})} \subset \Theta$ a credible set for θ , and $L(\theta, \Theta_{C(\mathbf{y})}) = 1 - \mathbb{1} \{ \theta \in \Theta_{C(\mathbf{y})} \}$, where

$$\mathbb{1}\left\{\theta \in \Theta_{C(\mathbf{y})}\right\} = \left\{\begin{matrix} 1, & \theta \in \Theta_{C(\mathbf{y})} \\ 0, & \theta \notin \Theta_{C(\mathbf{y})} \end{matrix}\right\}.$$

Then,

$$L(\theta,\Theta_{C(\mathbf{y})}) = \begin{cases} 0, & \theta \in \Theta_{C(\mathbf{y})} \\ 1, & \theta \notin \Theta_{C(\mathbf{y})} \end{cases},$$

where the 0–1 loss function is equal to zero if $\theta \in \Theta_{C(\mathbf{y})}$, and one if $\theta \notin \Theta_{C(\mathbf{y})}$. Then, the risk function is $1 - P(\theta \in \Theta_{C(\mathbf{y})})$.

Given a measure of credibility $(\alpha(\mathbf{y}))$ that defines the level of trust that $\theta \in \Theta_{C(\mathbf{y})}$; then, we can measure the accuracy of the report by $L(\theta, \alpha(\mathbf{y})) = [\mathbbm{1}\{\theta \in \Theta_{C(\mathbf{y})}\} - \alpha(\mathbf{y})]^2$. This loss function could be used to suggest a choice of the report $\alpha(\mathbf{y})$. Given that this is a quadratic loss function, the optimal action is the posterior mean, that is $\mathbb{E}[\mathbbm{1}\{\theta \in \Theta_{C(\mathbf{y})}\}|\mathbf{y}] = P(\theta \in \Theta_{C(\mathbf{y})}|\mathbf{y})$. This probability can be calculated given the posterior distribution, that is, $P(\theta \in \Theta_{C(\mathbf{y})}|\mathbf{y}) = \int_{\Theta_{C(\mathbf{y})}} \pi(\theta|\mathbf{y})d\theta$. This is a measure of the belief that $\theta \in \Theta_{C(\mathbf{y})}$ given the prior beliefs and sample information.

The set $\Theta_{C(\mathbf{y})} \in \Theta$ is a $100(1-\alpha)\%$ credible set with respect to $\pi(\theta|\mathbf{y})$ if $P(\theta \in \Theta_{C(\mathbf{y})}|\mathbf{y}) = \int_{\Theta_{C(\mathbf{y})}} \pi(\theta|\mathbf{y}) = 1-\alpha$.

Two alternatives to report credible sets are the symmetric credible set and the highest posterior density set (HPD). The former is based on $\frac{\alpha}{2}\%$ and $(1-\frac{\alpha}{2})\%$ percentiles of the posterior distribution, and the latter is a $100(1-\alpha)\%$ credible interval for θ with the property that it has the smallest distance compared to any other $100(1-\alpha)\%$ credible interval for θ based on the posterior distribution. That is, $C(\mathbf{y}) = \{\theta : \pi(\theta|\mathbf{y}) \ge k(\alpha)\}$, where $k(\alpha)$ is the largest number such that $\int_{\theta:\pi(\theta|\mathbf{y})\ge k(\alpha)} \pi(\theta|\mathbf{y})d\theta = 1-\alpha$. The HPD set can be a collection of disjoint intervals when working with multimodal posterior densities. In addition, they have the limitation of not necessary being invariant under transformations.

Decision theory can also be used to perform prediction (point, sets or probabilistic). Suppose that there is a loss function $L(Y_0, a)$ involving the prediction of Y_0 . Then, $\mathbb{E}_{Y_0}[L(Y_0, a)] = \int_{\mathcal{Y}_0} L(Y_0, a)\pi(Y_0|\mathbf{y})dY_0$, where $\pi(Y_0|\mathbf{y})$ is the predictive density function. Thus, we make an optimal choice for prediction that minimizes the risk function given a specific loss function.

Although BMA allows incorporating model uncertainty in a regression framework, sometimes it is desirable to select just one model. A compelling alternative is the model with the highest posterior model probability. This model is the best alternative for prediction in the case of a 0–1 loss function [19].

1.3.1 Example: Health insurance continues

We show some optimal rules in the health insurance example. In particular, the best point estimates of λ given the quadratic, absolute and generalized absolute loss functions. For the third, we assume that underestimating λ is twice as costly as overestimating it, that is, $K_0 = 2$ and $K_1 = 1$.

Taking into account that the posterior distribution of λ is $G(\alpha_0 + \sum_{i=1}^{N} y_i, \beta_0/(\beta_0 N + 1))$, using the hyperparameters from empirical Bayes, we have that $\mathbb{E}[\lambda|\mathbf{y}] = \alpha_n \beta_n = 1.2$, the median is 1.19, and the 2/3-th quantile is 1.26. Those are the optimal point estimates for the quadratic, absolute and generalized absolute loss functions.

In addition, we test the null hypothesis H_0 . $\lambda \in [0,1)$ versus H_1 . $\lambda \in [1,\infty)$ setting $K_0 = K_1 = 1$ we should reject the null hypothesis due to $P(\lambda \in [0,1)) = 0.9 > K_1/(K_0 + K_1) = 0.5$.

We get that the 95% symmetric credible interval is (0.91, 1.53), and the highest posterior density interval is (0.9, 1.51). Finally, the optimal point prediction under a quadratic loss function is 1.2, which is the mean value of the posterior predictive distribution, and the optimal model assuming a 0-1 loss function is the model using the hyperparameters from the empirical Bayes procedure due to the posterior model probability of this model being approximately 1, whereas the posterior model probability of the model using vague hyperparameters is approximately 0.

R code. Health insurance, Bayesian reports

```
1 an \leftarrow sum(y) + a0EB
2 # Posterior shape parameter
3 bn <- b0EB / (N*b0EB + 1)
4 # Posterior scale parameter
5 S <- 1000000
6 # Number of posterior draws
7 Draws <- rgamma (1000000, shape = an, scale = bn)
8 # Posterior draws
9 ##### Point estimation #######
10 OptQua <- an*bn
11 # Mean: Optimal choice quadratic loss function
12 OptQua
13 1.200952
14 OptAbs <- qgamma(0.5, shape = an, scale = bn)
# Median: Optimal choice absolute loss function
16 OptAbs
17 1.194034
_{\rm 18} # Setting KO = 2 and K1 = 1, that is, to underestimate
      lambda is twice as costly as to overestimate it.
19 KO <- 2; K1 <- 1
20 OptGenAbs <- quantile(Draws, KO/(KO + K1))</pre>
21 # Median: Optimal choice generalized absolute loss function
22 OptGenAbs
23 66.66667%
24 1.262986
25 ##### Hypothesis test #######
26 # HO: lambda in [0,1) vs H1: lambda in [1, Inf]
27 KO <- 1; K1 <- 1
28 ProbHO <- pgamma(1, shape = an, scale = bn)
29 ProbHO # Posterior probability HO
30 0.09569011
31 ProbH1 <- 1 -ProbH0
32 ProbH1 # Posterior probability H1
33 0.9043099
34 # We should reject HO given ProbH1 > K1 / (KO + K1)
35 ##### Credible intervals #######
36 LimInf <- qgamma(0.025, shape = an, scale = bn) # Lower
      bound
37 LimInf
38 0.9114851
39 LimSup <- qgamma(0.975, shape = an, scale = bn) # Upper
      bound
40 LimSup
41 1.529724
42 HDI <- HDInterval::hdi(Draws, credMass = 0.95) # Highest
      posterior density credible interval
43 HDI
      lower
44
45 0.8971505 1.5125911
46 attr(,"credMass")
47 [1] 0.95
48 ##### Predictive optimal choices #######
_{
m 49} p <- bn / (bn + 1) # Probability negative binomial density
50 OptPred <- p/(1-p)*an # Optimal point prediction given a
      quadratic loss function in prediction
51 OptPred
52 1.200952
```

1.4 Summary

We introduce the Bayes' rule to update probabilistic statements using funny examples. Then, we study the three probabilistic objects of main relevance in Bayesian inference: the posterior distribution, the marginal likelihood and the predictive density. The first allows performing inference regarding parameters, the second is required to perform hypothesis test for model selection using the Bayes factor, and the third to perform probabilistic predictions. We also review some sampling properties of Bayesian estimators, and Bayes update. All those concepts were developed using a simple example in R software. Finally, we introduce some concepts of decision theory that can be used to report summary statistics minimizing posterior expected losses.

1.5 Exercises

1. The court case: the blue or green cap

A cab was involved in a hit and run accident at night. There are two cab companies in the town: blue and green. The former has 150 cabs, and the latter 850 cabs. A witness said that a blue cab was involved in the accident; the court tested his/her reliability under the same circumstances, and got that 80% of the times the witness correctly identified the color of the cab. What is the probability that the color of the cab involved in the accident was blue given that the witness said it was blue?

2. The Monty Hall problem

What is the probability of winning a car in the *Monty Hall problem* switching the decision if there are four doors, where there are three goats and one car? Solve this problem analytically and computationally. What if there are n doors, n-1 goats and one car?

- 3. Solve the health insurance example using a Gamma prior in the rate parametrization, that is, $\pi(\lambda) = \frac{\beta_0^{\alpha_0}}{\Gamma(\alpha_0)} \lambda^{\alpha_0 1} \exp{\{-\lambda \beta_0\}}$.
- 4. Suppose that you are analyzing to buy a car insurance next year. To make a better decision you want to know what is the probability that you have a car claim next year? You have the records of your car claims in the last 15 years, $\mathbf{y} = \{0, 1, 0, 1, 0, 1, 1, 0, 0, 1, 0, 0, 1, 1, 0\}$. Assume that this is a random sample from a data generating process (statistical model) that is Bernoulli, $Y_i \sim Ber(p)$, and your probabilistic prior beliefs about p are well described by a beta distribution with parameters α_0 and β_0 , $p \sim B(\alpha_0, \beta_0)$, then, you are

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interested in calculating the probability of a claim the next year $P(Y_0 = 1|\mathbf{y})$.

Solve this using an empirical Bayes approach and a non-informative approach where $\alpha_0=\beta_0=1$ (uniform distribution).

5. Show that given the loss function, $L(\theta, a) = |\theta - a|$, then the optimal decision rule minimizing the risk function, $a^*(\mathbf{y})$, is the median.

Conceptual differences of the Bayesian and Frequentist approaches

We give some of the conceptual differences between the Bayesian and Frequentist inferential approaches. We emphasize in the Bayesian concepts as most of the readers can be familiarized with the Frequentist statistical framework.

2.1 The concept of probability

Let's begin with the following thought experiment: Assume that you are watching the international game show "Who wants to be a millionaire?", the contestant is asked to answer a very simple question: What is the last name of the brothers who are credited with inventing the world's first successful motor-operated airplane?

- What is the probability that the contestant answers this question correctly?
 - Unless you have:
 - watched this particular contestant participating in this show many times.
 - 2. seen him asked this same question each time,
 - 3. and computed the relative frequency with which he gives the correct answer,

you need to answer this question as a Bayesian!

Uncertainty about the event answer this question needs to be expressed as a "degree of belief" informed both by information coming from data on the skill of the particular participant, and how much he knows about inventors, and possibly prior knowledge on his performance in other game shows. Of course, your prior knowledge of the contestant may be minimal, or it may be very informed. Either way, your final answer remains a degree of belief held about an uncertain, and inherently unrepeatable state of nature.

The point of this hypothetical, light-hearted scenario is simply to highlight that a key distinction between the Frequentist and Bayesian approaches to inference is not the use (or nature) of prior information, but simply the manner in which probability is used. To the Bayesian, probability is the mathematical construct used to quantify uncertainty about an unknown state of nature, conditional on observed data and prior knowledge about the context in which that state of nature occurs. To the Frequentist, probability is linked intrinsically to the concept of a repeated experiment, and the relative frequency with which a particular outcome occurs, conditional on that unknown state. This distinction remains key whether the Bayesian chooses to be *informative or subjective* in the specification of prior information, or chooses to be non-informative or objective.

Frequentists consider probability as a physical phenomenon, like mass or wavelength, whereas Bayesians stipulate that probability lives in the mind of scientists as any scientific construct [58].

It seems that the understanding of the concept of probability for the common human being is more associated with "degrees of belief" rather than relative frequency. Peter Diggle, President of The Royal Statistical Society between 2014 and 2016, was asked in an interview "A different trend which has surged upwards in statistics during Peter's career is the popularity of Bayesian statistics. Does Peter consider himself a Bayesian?", and he replied "... you can't not believe in Bayes' theorem because it's true. But that doesn't make you a Bayesian in the philosophical sense. When people are making personal decisions – even if they don't formally process Bayes' theorem in their mind – they are adapting what they think they should believe in response to new evidence as it comes in. Bayes' theorem is just the formal mathematical machinery for doing that."

However, we should say that psychological experiments suggest that human beings suffer from *anchoring*, that is, a cognitive bias that causes us to rely too heavily on the previous information (prior) such that the updating process (posterior) due to new information (likelihood) being low compared to the Bayes' rule [42].

2.2 Subjectivity is not the key

The concepts of *subjectivity* and *objectivity* indeed characterize both statistical paradigms in differing ways. Among Bayesians there are those who are immersed in *subjective* rationality [64, 21, 67, 50], but others who adopt *objective* prior distributions such as Jeffreys', reference, empirical or robust [4, 47, 40, 8] to operationalize Bayes' rule, and thereby weight quantitative (data-based) evidence. Among Frequentists, there are choices made about significance levels which, if not explicitly subjective, are typically not grounded in any objective and documented assessment of the relative losses of Type I and Type II er-

rors.¹ In addition, both Frequentist and Bayesian statisticians make decisions about the form of the data generating process, or "model", which – if not subject to rigorous diagnostic assessment – retains a subjective element that potentially influences the final inferential outcome. Although we all know that by definition a model is a schematic and simplified approximation to reality,

"Since all models are wrong the scientist cannot obtain a *correct* one by excessive elaboration. On the contrary following William of Occam he should seek an economical description of natural phenomena." [12].

We also know that "All models are wrong, but some are useful" [13], that is why model diagnostics are important. This task can be performed in both approaches. Particularly, the Bayesian framework can use predictive p-values for absolute testing [28, 2] or posterior odds ratios for relative statements [39, 44]. This is because the marginal likelihood, conditional on data, is interpreted as the evidence of the prior distribution [7].

In addition, what is objectivity in a Frequentist approach? For example, why should we use a 5% or 1% significance level rather than any other value? As someone said, the apparent objectivity is really a consensus [50]. In fact "Student" (William Gosset) saw statistical significance at any level as being "nearly valueless" in itself [80]. But, this is not just a situation in the Frequentist approach. The cut–offs given to "establish" scientific evidence against a null hypothesis in terms of log_{10} scale [40] or log_e scale [44] in Table 1.1 are also $ad\ hoc$.

Although the true state of nature in Bayesian inference is expressed in "degrees of belief", the distinction between the two paradigms does not reside in one being more, or less, *subjective* than the other. Rather, the differences are philosophical, pedagogical, and methodological.

2.3 Estimation, hypothesis testing and prediction

All what is required to perform estimation, hypothesis testing (model selection) and prediction in the Bayesian approach is to apply the Bayes' rule. This means coherence under a probabilistic view. But, there is no free lunch, coherence reduces flexibility. On the other hand, the Frequestist approach may be not coherent from a probabilistic point of view, but it is very flexible. This approach can be seen as a tool kit that offers inferential solutions under the umbrella of understanding probability as relative frequency. For instance, a point estimator in a Frequentist approach is found such that satisfies good sampling properties like unbiasness, efficiency, or a large sample property as consistency.

¹Type I error is rejecting the null hypothesis when this is true, and the Type II error is not rejecting the null hypothesis when this is false.

A remarkable difference is that optimal Bayesian decisions are calculated minimizing the expected value of the loss function with respect to the posterior distribution, that is, it is conditional on observed data. On the other hand, Frequentist "optimal" actions are base on the expected values over the distribution of the estimator (a function of data) conditional on the unknown parameters, that is, it considers sampling variability.

The Bayesian approach allows to obtain the posterior distribution of any unknown object such as parameters, latent variables, future or unobserved variables or models. A nice advantage is that prediction can take into account estimation error, and predictive distributions (probabilistic forecasts) can be easily recovered.

Hypothesis testing (model selection) is based on *inductive logic* reasoning (*inverse probability*); on the basis of what we see, we evaluate what hypothesis is most tenable, and is performed using posterior odds, which in turn are based on Bayes factors that evaluate evidence in favor of a null hypothesis taking explicitly the alternative [44], following the rules of probability [50], comparing how well the hypothesis predicts data [36], minimizing the weighted sum of type I and type II error probabilities [22, 59], and taking the implicit balance of losses [40, 10] into account. Posterior odds allows to use the same framework to analyze nested and non-nested models and perform model average. However, Bayes factors cannot be based on improper or vague priors [45], the practical interplay between model selection and posterior distributions is not as easy as it maybe in the Frequentist approach, and the computational burden can be more demanding due to solving potentially difficult integrals.

On the other hand, the Frequentist approach establishes most of its estimators as the solution of a system of equations. Observe that optimization problems reduce to solve systems. We can potentially get the distribution of these estimators, but most of the time it is needed asymptotic arguments or resampling techniques. Hypothesis testing requires pivotal quantities and/or also resampling, and prediction most of the time is based on a plug-in approach, which means not taking estimation error into account. In addition, ancillary statistics can be used to build prediction intervals. Comparing models depends on their structure, for instance, there are different Frequentist statistical approaches to compare nested and non-nested models. A nice feature in some situations is that there is a practical interplay between hypothesis testing and confidence intervals, for instance in the normal population mean hypothesis framework you cannot reject at α significance level (Type I error) any null hypothesis H_0 . $\mu = \mu^0$ if μ^0 is in the $1 - \alpha$ confidence interval $P(\mu \in [\hat{\mu} - |t_{N-1}^{\alpha/2}| \times \hat{\sigma}_{\hat{\mu}}, \hat{\mu} + |t_{N-1}|^{\alpha/2} \times \hat{\sigma}_{\hat{\mu}}]) = 1 - \alpha$, where $\hat{\mu}$ and $\hat{\sigma}_{\hat{\mu}}$ are the maximum likelihood estimators of the mean and standard error, and $t_{N-1}^{\alpha/2}$ is

²A pivot quantity is a function of unobserved parameters and observations whose probability distribution does not depend on the unknown parameters.

³An ancillary statistic is a pivotal quantity that is also a statistic.

the quantile value of the Student's t distribution at $\alpha/2$ probability and N-1 degrees of freedom, N is the sample size.

A remarkable difference between the Bayesian and the Frequentist inferential frameworks is the interpretation of credible/confidence intervals. Observe that once we have estimates, such that for example the previous interval is [0.2,0.4] given a 95% confidence level, we cannot say that $P(\mu \in [0.2,0.4]) = 0.95$ in the Frequentist framework. In fact, this probability is 0 or 1 under this approach, as μ can be there or not, the problem is that we will never know in applied settings. This due to that $P(\mu \in [\hat{\mu} - |t_{N-1}^{0.025}| \hat{\times} \sigma_{\hat{\mu}}, \hat{\mu} + |t_{N-1}^{0.025}| \times \hat{\sigma}_{\hat{\mu}}]) = 0.95$ being in the sense of repeated sampling. On the other hand, once we have the posterior distribution, we can say that $P(\mu \in [0.2, 0.4]) = 0.95$ under the Bayesian framework.

Following common practice, most of researchers and practitioners do hypothesis testing based on the p-value in the Frequentist framework. But, **what** is a p-value? Most of the users do not know the answer due to many time statistical inference is not performed by statisticians [8].⁴ A p-value is the probability of obtaining a statistical summary of the data equal to or *more extreme* than what was actually observed, assuming that the null hypothesis is true.

Therefore, p-value calculations involve not just the observed data, but also more extreme hypothetical observations. So,

"What the use of p implies, therefore, is that a hypothesis that may be true may be rejected because it has not predicted observable results that have not occurred." [40]

It seems that common Frequentist inferential practice intertwined two different logic reasoning arguments: the p-value [25] and $significance\ level$ [57]. The former is an informal short-run criterion, whose philosophical foundation is $reduction\ to\ absurdity$, which measures the discrepancy between the data and the null hypothesis. So, the p-value is not a direct measure of the probability that the null hypothesis is false. The latter, whose philosophical foundations is deduction, is based on a long-run performance such that controls the overall number of incorrect inferences in the repeated sampling without care of individual cases. The p-value fallacy consists in interpreting the p-value as the strength of evidence against the null hypothesis, and using it simultaneously with the frequency of type I error under the null hypothesis [36].

The American Statistical Association has several concerns regarding the use of the p-value as a cornerstone to perform hypothesis testing in science. This concern motivates the ASA's statement on p-values [78], which can be summarized in the following principles:

• "P-values can indicate how incompatible the data are with a specified statistical model."

⁴https://fivethirtyeight.com/features/not-even-scientists-can-easily-explain-p-values/

- "P-values do not measure the probability that the studied hypothesis is true, or the probability that the data were produced by random chance alone."
- "Scientific conclusions and business or policy decisions should not be based only on whether a p-value passes a specific threshold."
- "Proper inference requires full reporting and transparency."
- "A *p*-value, or statistical significance, does not measure the size of an effect or the importance of a result."
- "By itself, a *p*-value does not provide a good measure of evidence regarding a model or hypothesis."

To sum up, Fisher proposed the p-value as a witness rather than a judge. So, a p-value lower than the significance level means more inspection of the null hypothesis, but it is not a final conclusion about it.

Another difference between the Frequentists and the Bayesians is the way how scientific hypothesis are tested. The former use the p-value, whereas the latter use the Bayes factor. Observe that the p-value is associated with the probability of the data given the hypothesis, whereas the Bayes factor is associated with the probability of the hypothesis given the data. However, there is an approximate link between the t statistic and the Bayes factor for regression coefficients [62]. In particular, $|t| > (log(N)+6)^{1/2}$, corresponds to strong evidence in favor of rejecting the not relevance of a control in a regression. Observe that in this setting the threshold of the t statistic, and as a consequence the significant level, depends on the sample size. Observe that this setting agrees with the idea in experimental designs of selecting the sample size such that we control Type I and Type II errors. In observational studies we cannot control the sample size, but we can select the significance level.

See also [69] and [6] for nice exercises to reveal potential flaws of the p-value (p) due to $p \sim U[0,1]$ under the null hypothesis,⁵ and calibrations of the p-value to interpret them as the odds ratio and the error probability. In particular, $B(p) = -e \times p \times \log(p)$ when $p < e^{-1}$, and interpret this as the Bayes factor of H_0 to H_1 , where H_1 denotes the unspecified alternative to H_0 , and $\alpha(p) = (1 + [-e \times p \times \log(p)]^{-1})^{-1}$ as the error probability α in rejecting H_0 . Take into account that B(p) and $\alpha(p)$ are lower bounds.

Logic of argumentation in the Frequentist approach is based on *deductive logic*, this means that it starts from a statement about the true state of nature (null hypothesis), and predicts what should be seen if this statement were true. On the other hand, the Bayesian approach is based on *inductive logic*, this means that it defines what hypothesis is more consistent with what is seen. The former inferential approach establishes that the true of the premises implies the true of the conclusion, that is why we reject or not reject hypothesis. The latter establishes that the premises supply some evidence, but not

 $^{^5}$ https://joyeuserrance.wordpress.com/2011/04/22/proof-that-p-values-under-the-null-are-uniformly-distributed/ for a simple proof.

full assurance, of the true of the conclusion, that is why we get probabilistic statements.

Here, there is a difference between effects of causes (forward causal inference) and causes of effects (reverse causal inference) [32, 20]. To illustrate this point, imagine that a firm increases the price of a specific good, then economic theory would say that its demand decreases. The premise (null hypothesis) is a price increase, and the consequence is a demand reduction. Another view would be to observe a demand reduction, and try to identify which cause is more tenable. For instance, demand reduction can be caused by any positive supply shocks or any negative demand shocks. The Frequentist logic sees the first view, and the Bayesian reasoning gives the probability associated with possible causes.

2.4 The likelihood principle

The **likelihood principle** states that in making inference or decisions about the state of the nature all the relevant *experimental* information is given by the *likelihood function*. The Bayesian framework follows this statement, that is, it is conditional on observed data.

We follow [7], who in turns followed [51], to illustrate the likelihood principle. We are given a coin such that we are interested in the probability, θ , of having it come up heads when flipped. It is desired to test H_0 . $\theta = 1/2$ versus H_1 . $\theta > 1/2$. An experiment is conducted by flipping the coin (independently) in a series of trials, the results of which is the observation of 9 heads and 3 tails.

This is not yet enough information to specify $p(y|\theta)$, since the series of trials was not explained. Two possibilities:

- 1. The experiment consisted of a predetermine 12 flips, so that Y = [Heads] would be $B(12, \theta)$, then $p_1(y|\theta) = \binom{n}{y} \theta^y (1-\theta)^{n-y} = 220 \times \theta^9 (1-\theta)^3$.
- 2. The experiment consisted of flipping the coin until 3 tails were observed (r=3). Then, Y, the number of failures (heads) until getting 3 tails, is $NB(3, 1-\theta)$. Then, $p_2(y|\theta) = {y+r-1 \choose r-1}(1-(1-\theta)^y(1-\theta)^r = 55 \times \theta^9(1-\theta)^3$.

Using a Frequentist approach, the significance level of y=9 using the Binomial model against $\theta=1/2$ would be:

$$\alpha_1 = P_{1/2}(Y \ge 9) = p_1(9|1/2) + p_1(10|1/2) + p_1(11|1/2) + p_1(12|1/2) = 0.073.$$

R code. The likelihood principle: Binomial model

```
1 success <- 9
2 # Number of observed success in n trials
3 n <- 12
4 # Number of trials
5 siglevel <- sum(sapply(9:n,function(y)dbinom(y,n,0.5)))
6 siglevel
7 0.073</pre>
```

For the Negative Binomial model, the significance level would be:

```
\alpha_2 = P_{1/2}(Y \ge 9) = p_2(9|1/2) + p_2(10|1/2) + \dots = 0.0327.
```

$R\ code.\ The\ likelihood\ principle:\ Negative\ Binomial\ model$

```
success <- 3
2 # Number of target success (tails)
3 failures <- 9
4 # Number of failures
5 siglevel <- 1 - pnbinom((failures - 1), success, 0.5)
6 siglevel
7 0.0327</pre>
```

We arrive to different conclusions using a significance level equal to 5%, whereas we obtain the same outcomes using a Bayesian approach because the kernels of both distributions are the same $(\theta^9 \times (1-\theta)^3)$.

2.5 Why is not the Bayesian approach that popular?

At this stage, we may wonder why the Bayesian statistical framework is not the dominant inferential approach despite that it has its historical origin in 1763 [5], whereas the Frequentist statistical framework was largely developed in the early 20th century. The scientific battle over the Bayesian inferential approach lasted for 150 years, and this maybe explained by some of the following facts.

There is an issue regarding apparent subjectivity as the Bayesian inferential approach runs counter the strong conviction that science demands objectivity, and Bayesian probability is a measure of degrees of belief, where the initial prior maybe just a guess; this was not accepted as objective and rigorous science. Initial critics said that Bayes was quantifying ignorance as he set equal probabilities to any potential result. As a consequence, prior distributions were damned [55].

Bayes himself seemed not to have believed in his idea. Although, it seems that Bayes achieved his breakthrough during the late 1740s, he did not send it off to the Royal Society for publication. It was his friend, Richard Price, another Presbyterian minister, who rediscovered Bayes' idea, polished it and published.

However, it was Laplace who independently generalized Bayes' theorem in 1781. He used it initially in gambling problems, and soon after in astronomy, mixing different sources of information in order to leverage research in specific situations where data was scarce. Then, he wanted to use his discovery to find the probability of causes, and thought that this required large data sets, and turned into demography. In this field, he had to perform large calculations that demanded to develop smart approximations, creating the Laplace's approximation and the central limit theorem [47]; although, at the cost of apparently leaving his research on Bayesian inference.

Once Laplace was gone in 1827, the Bayes' rule disappeared from the scientific spectrum for almost a century. In part, personal attacks against Laplace made the rule be forgotten, and also, the old fashion thought that statistics does not have to say anything about causation, and that the prior is very subjective to be compatible with science. Although, practitioners used it to solve problems in astronomy, communication, medicine, military and social issues with remarkable results.

Thus, the concept of degrees of belief to operationalize probability was gone in name of scientific objectivity, and probability as the frequency an event occurs in many repeatable trials became the rule. Laplace critics argued that those concepts were diametric opposites, although, Laplace considered them as basically equivalent when large sample sizes are involved [55].

The era of the Frequentists or sampling theorists began, lead by Karl Pearson, and his nemesis, Ronald Fisher, both brilliant, against the inverse probability approach, persuasive and dominant characters that made impossible to argue against their ideas. Karl Pearson legacy was taken by his son Egon, and Egon's friend Neyman, both inherited the anti-Bayesian and anti-Fisher legacy.

Despite the anti-Bayesian campaign among statisticians, there were some independent characters developing Bayesian ideas, Borel, Ramsey and de Fineti, all of them isolated in different countries, France, England and Italy. However, the anti-Bayesian trio of Fisher, Neyman and Egon Person got all the attention during the 1920s and 1930s. Only, a geophysicist, Harold Jeffreys, kept alive Bayesian inference in the 1930s and 1940s. Jeffreys was a

very quiet, shy, uncommunicative gentleman working at Cambridge in the astronomy department. He was Fisher's friend thanks to his character, although they were diametric opposites regarding the Bayesian inferential approach, facing very high intellectual battles. Unfortunately for the Bayesian approach, Jeffreys lost, he was very technical using confusing high level mathematics, worried about inference from scientific evidence, not guiding future actions based on decision theory, which was very important in that era for mathematical statistics due to the Second World War. On the other hand, Fisher was a very dominant character, persuasive in public and a master of practice, his techniques were written in a popular style with minimum mathematics.

However, Bayes' rule achieved remarkable results in applied settings like the AT&T company or the social security system in USA. Bayesian inference also had a relevant role during the second World War and the Cold War. Alan Turing used inverse probability at Bletchley Park to crack German messages called Enigma code used by U-boats, Andrei Kolmogorov used it to improved firing tables of Russia's artillery, Bernard Koopman applied it for searching targets in the open sea and the RAND Corporation used it in the Cold War. Unfortunately, these Bayesian developments were top secrets for almost 40 years that keep classified the contribution of inverse probability in modern human history.

During 1950s and 1960s three mathematicians lead the rebirth of the Bayesian approach, Good, Savage and Lindley. However, it seems that they were unwilling to apply their theories to real problems, and despite that the Bayesian approach proved its worth, for instance, in business decisions, navy search, lung cancer, etc, it was applied to simple models due to its mathematical complexity and requirement of large computations. But, there were some breakthrough that change this. First, hierarchical models introduced by Lindley and Smith, where a complex model is decomposed into many easy to solve models, and second, Markov chain Monte Carlo methods developed by Hastings in the 1970s [38] and the Geman brothers in the 1980s [33]. These methods were introduced into the Bayesian inferential framework in the 1990s by Gelfand and Smith [26], and Tierney [76], when desktop computers got enough computational power to solve complex models. Since then, the Bayesian inferential framework has gained increasing popularity among practitioners and scientists.

A simple working example

We will illustrate some conceptual differences between the Bayesian and Frequentist statistical approaches performing inference given a random sample $\mathbf{y} = [y_1, y_2, \dots, y_N], \text{ where } y_i \stackrel{iid}{\sim} N(\mu, \sigma^2), i = 1, 2, \dots, N.$ In particular, we set $\pi(\mu, \sigma) = \pi(\mu)\pi(\sigma) \propto \frac{1}{\sigma}$. This is a standard non-

informative improper prior (Jeffreys prior, see Chapter 4), that is, this prior is perfectly compatible with sample information. In addition, we are assuming independent priors for μ and σ . Then,

$$\pi(\mu, \sigma | \mathbf{y}) \propto \frac{1}{\sigma} \times (\sigma^2)^{-N/2} \exp\left\{-\frac{1}{2\sigma^2} \sum_{i=1}^N (y_i - \mu)^2\right\}$$

$$= \frac{1}{\sigma} \times (\sigma^2)^{-N/2} \exp\left\{-\frac{1}{2\sigma^2} \sum_{i=1}^N ((y_i - \bar{y}) - (\mu - \bar{y}))^2\right\}$$

$$= \frac{1}{\sigma} \exp\left\{-\frac{N}{2\sigma^2} (\mu - \bar{y})^2\right\} \times (\sigma)^{-N} \exp\left\{-\frac{1}{2\sigma^2} \sum_{i=1}^N (y_i - \bar{y})^2\right\}$$

$$= \frac{1}{\sigma} \exp\left\{-\frac{N}{2\sigma^2} (\mu - \bar{y})^2\right\} \times (\sigma)^{-(\alpha_n + 1)} \exp\left\{-\frac{\alpha_n \hat{\sigma}^2}{2\sigma^2}\right\},$$

where $\bar{y} = \frac{\sum_{i=1}^{N}}{N}$, $\alpha_n = N - 1$ and $\hat{\sigma}^2 = \frac{\sum_{i=1}^{N} (y_i - \bar{y})^2}{N - 1}$. The first term in the last expression is the kernel of a normal den-

The first term in the last expression is the kernel of a normal density, $\mu|\sigma, \mathbf{y} \sim N(\bar{y}, \sigma^2/N)$. The second term is the kernel of an inverted gamma density [79], $\sigma|\mathbf{y} \sim IG(\alpha_n, \hat{\sigma}^2)$. Therefore, $\pi(\mu|\sigma, \mathbf{y}) = (2\pi\sigma^2/N)^{-1/2} \exp\left\{\frac{-N}{2\sigma^2}(\mu-\bar{y})^2\right\}$ and $\pi(\sigma|\mathbf{y}) = \frac{2}{\Gamma(\alpha_n/2)} \left(\frac{\alpha_n\hat{\sigma}^2}{2}\right)^{\alpha_n/2} \frac{1}{\sigma^{\alpha_n+1}} \times \exp\left\{-\frac{\alpha_n\hat{\sigma}^2}{2\sigma^2}\right\}$.

Observe that $\mathbb{E}[\mu|\sigma,\mathbf{y}]=\bar{y}$, this is also the maximum likelihood (Frequentist) point estimate of μ in this setting. In addition, the Frequentist $(1-\alpha)\%$ confidence interval and the Bayesian $(1-\alpha)\%$ credible interval have exactly the same form, $\bar{y}\pm|z_{\alpha/2}|\frac{\sigma}{\sqrt{N}}$, where $z_{\alpha/2}$ is the $\alpha/2$ percentile of a standard normal distribution. However, the interpretations are totally different. The confidence interval has a probabilistic interpretation under sampling variability of \bar{Y} , that is, in repeated sampling $(1-\alpha)\%$ of the intervals $\bar{Y}\pm|z_{\alpha/2}|\frac{\sigma}{\sqrt{N}}$ would include μ , but given an observed realization of \bar{Y} , say \bar{y} , the probability of $\bar{y}\pm|z_{\alpha/2}|\frac{\sigma}{\sqrt{N}}$ including μ is 1 or 0, that is why we say a $(1-\alpha)\%$ confidence interval. On the other hand, $\bar{y}\pm|z_{\alpha/2}|\frac{\sigma}{\sqrt{N}}$ has a simple probabilistic interpretation in the Bayesian framework, there is a $(1-\alpha)\%$ probability that μ lies in this interval.

If we want to get the marginal posterior density of μ ,

$$\pi(\mu|\mathbf{y}) = \int_0^\infty \pi(\mu, \sigma|\mathbf{y}) d\sigma$$

$$\propto \int_0^\infty \frac{1}{\sigma} \times (\sigma^2)^{-N/2} \exp\left\{-\frac{1}{2\sigma^2} \sum_{i=1}^N (y_i - \mu)^2\right\} d\sigma$$

$$= \int_0^\infty \left(\frac{1}{\sigma}\right)^{N+1} \exp\left\{-\frac{N}{2\sigma^2} \frac{\sum_{i=1}^N (y_i - \mu)^2}{N}\right\} d\sigma$$

$$= \left[\frac{2}{\Gamma(N/2)} \left(\frac{N \sum_{i=1}^N (y_i - \mu)^2}{2N}\right)^{N/2}\right]^{-1}$$

$$\propto \left[\sum_{i=1}^N (y_i - \mu)^2\right]^{-N/2}$$

$$= \left[\sum_{i=1}^N ((y_i - \bar{y}) - (\mu - \bar{y}))^2\right]^{-N/2}$$

$$= \left[\alpha_n \hat{\sigma}^2 + N(\mu - \bar{y})^2\right]^{-N/2}$$

$$\propto \left[1 + \frac{1}{\alpha_n} \left(\frac{\mu - \bar{y}}{\hat{\sigma}/\sqrt{N}}\right)^2\right]^{-(\alpha_n + 1)/2}.$$

The fourth line is due to having the kernel of a inverted gamma density with N degrees of freedom in the integral [79].

The last expression is the kernel of a Student's t density function with $\alpha_n = N-1$ degrees of freedom, expected value equal to \bar{y} , and variance $\frac{\hat{\sigma}^2}{N}\left(\frac{\alpha_n}{\alpha_n-2}\right)$. Then, $\mu|\mathbf{y}\sim t\left(\bar{y},\frac{\hat{\sigma}^2}{N}\left(\frac{\alpha_n}{\alpha_n-2}\right),\alpha_n\right)$.

Observe that a $(1-\alpha)\%$ confidence interval and $(1-\alpha)\%$ credible interval have exactly the same expression, $\bar{y}\pm|t_{\alpha/2}^{\alpha_n}|\frac{\hat{\sigma}}{\sqrt{N}}$, where $t_{\alpha/2}^{\alpha_n}$ is the $\alpha/2$ per-

centile of a Student's t distribution. But again, the interpretations are totally different.

The mathematical similarity between the Frequentist and Bayesian expressions in this example is due to using an improper prior.

2.6.1 Example: Math test

You have a random sample of math scores of size N = 50 from a normal distribution, $Y_i \sim N(\mu, \sigma)$. The sample mean and variance are equal to 102 and 10, respectively. Assuming an improper prior equal to $1/\sigma$,

- Get 95% confidence and credible intervals for μ .
- What is the posterior probability that $\mu > 103$?

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Using $\mu|\mathbf{y} \sim t\left(\bar{y}, \frac{\hat{\sigma}^2}{N}\left(\frac{\alpha_n}{\alpha_n-2}\right), \alpha_n\right)$, which implies that $\bar{y} \pm |t_{\alpha/2}^{\alpha_n}| \frac{\hat{\sigma}}{\sqrt{N}}$, where $\bar{y} = 102$, $\hat{\sigma}^2 = 10$ and $\alpha_n = 49$, the 95% confidence and credible intervals for μ are the same (101.1, 102.9), and $P(\mu > 103) = 1.49\%$ given the sample information.

R code. Example: Math test

```
1 N <- 50 # Sample size
2 y_bar <- 102 # Sample mean
3 s2 <- 10 # Sample variance
4 alpha <- N - 1
5 serror <- (s2/N)^0.5
6 LimInf <- y_bar - abs(qt(0.025, alpha)) * serror
7 LimInf
8 101.101
9 # Lower bound
10 LimSup <- y_bar + abs(qt(0.025, alpha)) * serror
1 LimSup
11 102.898
13 # Upper bound
14 y.cut <- 103
15 P <- 1-metRology::pt.scaled(y.cut, df = alpha, mean = y_bar, sd = serror)
16 P
17 0.0149
18 # Probability of mu greater than y.cut
```

2.7 Summary

The differences between the Bayesian and Frequentist inferential approaches are philosophical, including as pertains to the role of probability; pedagogical, in particular as relates to the use of inference to inform decision making; and methodological, as having differences in their mathematical and computational frameworks. Although at methodological level, the debate has become considerably muted, except for some aspects of inference, with the recognition that each approach has a great deal to contribute to statistical practice [35, 3, 43]. As Bradley Efron said "Computer-age statistical inference at its most successful **combines** elements of the two philosophies" [24].

2.8 Exercises

1. Jeffreys-Lindley's paradox

The **Jeffreys-Lindley's paradox** [40, 52] is an apparent disagreement between the Bayesian and Frequentist frameworks to a hypothesis testing situation.

In particular, assume that in a city 49,581 boys and 48,870 girls have been born in 20 years. Assume that the male births is distributed Binomial with probability θ . We want to test the null hypothesis H_0 . $\theta=0.5$ versus H_1 . $\theta\neq0.5$.

- •Show that the posterior model probability for the model under the null is approximately 0.95. Assume $\pi(H_0) = \pi(H_1) = 0.5$, and $\pi(\theta)$ equal to U(0,1) under H_1 .
- •Show that the *p*-value for this hypothesis test is equal to 0.0235 using the normal approximation, $Y \sim N(N \times \theta, N \times \theta \times (1-\theta))$.
- 2. We want to test H_0 . $\mu = \mu_0$ vs H_1 . $\mu \neq \mu_0$ given $y_i \stackrel{iid}{\sim} N(\mu, \sigma^2)$. Assume $\pi(H_0) = \pi(H_1) = 0.5$, and $\pi(\mu, \sigma) \propto 1/\sigma$ under the alternative hypothesis.

Show that

$$p(\mathbf{y}|\mathcal{M}_1) = \frac{\pi^{-N/2}}{2} \Gamma(N/2) 2^{N/2} \left(\frac{1}{\alpha_n \hat{\sigma}^2}\right)^{N/2} \left(\frac{N}{\alpha_n \hat{\sigma}^2}\right)^{-1/2} \frac{\Gamma(1/2) \Gamma(\alpha_n/2)}{\Gamma((\alpha_n+1)/2)}$$
 and $p(\mathbf{y}|\mathcal{M}_0) = (2\pi)^{-N/2} \left[\frac{2}{\Gamma(N/2)} \left(\frac{N}{2} \frac{\sum_{i=1}^{N} (y_i - \mu_0)^2}{N}\right)^{N/2}\right]^{-1}$. Then,

$$PO_{01} = \frac{p(\mathbf{y}|\mathcal{M}_0)}{p(\mathbf{y}|\mathcal{M}_1)}$$

$$= \frac{\Gamma((\alpha_n + 1)/2)}{\Gamma(1/2)\Gamma(\alpha_n/2)} (\alpha_n \hat{\sigma}^2/N)^{-1/2} \left[1 + \frac{(\mu_0 - \bar{y})^2}{\alpha_n \hat{\sigma}^2/N} \right]^{-\left(\frac{\alpha_n + 1}{2}\right)}$$

where
$$\alpha_n = N - 1$$
 and $\hat{\sigma}^2 = \frac{\sum_{i=1}^{N} (y_i - \bar{y})^2}{N-1}$.

Find the relationship between the posterior odds and the classical test statistic for the null hypothesis.

3. Math test continues

Using the setting of the **Example: Math test** in subsection 2.6.1, test H_0 . $\mu = \mu_0$ vs H_1 . $\mu \neq \mu_0$ where $\mu_0 = \{100, 100.5, 101, 101.5, 102\}$.

•What is the *p*-value for these hypothesis tests?

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•Find the posterior model probability of the null model for each μ_0 .

Objective and subjective Bayesian approaches

Cornerstone models: Conjugate families

We will introduce conjugate families in basic statistical models with examples, solving them analytically and computationally using R. We will have some mathematical, and computational exercises in R.

4.1 Motivation of conjugate families

Observing the three fundamental pieces of Bayesian analysis: the posterior distribution (parameter inference), the marginal likelihood (hypothesis testing), and the predictive distribution (prediction), equations 4.1, 4.2 and 4.3, respectively,

$$\pi(\boldsymbol{\theta}|\mathbf{y}) = \frac{p(\mathbf{y}|\boldsymbol{\theta}) \times \pi(\boldsymbol{\theta})}{p(\mathbf{y})},$$
(4.1)

$$p(\mathbf{y}) = \int_{\mathbf{\Theta}} p(\mathbf{y}|\boldsymbol{\theta})\pi(\boldsymbol{\theta})d\boldsymbol{\theta}, \tag{4.2}$$

and

$$p(\mathbf{Y}_0|\mathbf{y}) = \int_{\mathbf{\Theta}} p(\mathbf{Y}_0|\boldsymbol{\theta})\pi(\boldsymbol{\theta}|\mathbf{y})d\boldsymbol{\theta}, \tag{4.3}$$

we can understand that some of the initial limitations of the application of the Bayesian analysis were associated with the absence of algorithms to draw from non-standard posterior distributions (equation 4.1), and the lack of analytical solutions of the marginal likelihood (equation 4.2) and the predictive distribution (equation 4.3). Both issues requiring computational power.

Although there were algorithms to sample from non-standard posterior distributions since the second half of the last century [56, 38, 33], their particular application in the Bayesian framework emerged later [26, 76], maybe until the increasing computational power of desktop computers. However, it is also common practice nowadays to use models that have standard conditional posterior distributions to mitigate computational requirements. In addition, nice mathematical tricks plus computational algorithms [27, 17, 18] and approximations [77, 41] are used to obtain the marginal likelihood (prior predictive).

Despite these advances, there are two potentially conflicting desirable model specification features that we can see from equations 4.1, 4.2 and 4.3: (1) analytical solutions and (2) the posterior distribution in the same family as the prior distribution for a given likelihood. The latter is called *conjugate priors*, a family of priors that is closed under sampling [68].

These features are desirable as the former implies facility to perform hypothesis testing and predictive analysis, and the latter means invariance of the prior-to-posterior updating. Both features imply less computational burden.

We can easily achieve each of these features independently, for instance using improper priors for analytical tractability, and defining in a broad sense the family of prior distributions for prior conjugacy. However, these features are in conflict.

Fortunately, we can achieve these two nice characteristics if we assume that the data generating process is given by a distribution function in the exponential family. That is, given a random sample $\mathbf{y} = [y_1, y_2, \dots, y_N]^{\top}$, a probability density function $p(\mathbf{y}|\boldsymbol{\theta})$ belongs to the exponential family if it has the form

$$p(\mathbf{y}|\boldsymbol{\theta}) = \prod_{i=1}^{N} h(y_i) C(\boldsymbol{\theta}) \exp\left\{\eta(\boldsymbol{\theta})^{\top} \mathbf{T}(y_i)\right\}$$

$$= h(\mathbf{y}) C(\boldsymbol{\theta})^{N} \exp\left\{\eta(\boldsymbol{\theta})^{\top} \mathbf{T}(\mathbf{y})\right\}$$

$$= h(\mathbf{y}) \exp\left\{\eta(\boldsymbol{\theta})^{\top} \mathbf{T}(\mathbf{y}) - A(\boldsymbol{\theta})\right\},$$
(4.4)

where $h(\mathbf{y}) = \prod_{i=1}^N h(y_i)$ is a non-negative function, $\eta(\boldsymbol{\theta})$ is a known function of the parameters, $A(\boldsymbol{\theta}) = \log \left\{ \int_{\mathbf{Y}} h(\mathbf{y}) \exp \left\{ \eta(\boldsymbol{\theta})^\top \mathbf{T}(\mathbf{y}) \right\} d\mathbf{y} \right\} = -N \log(C(\boldsymbol{\theta}))$ is a normalization factor, and $\mathbf{T}(\mathbf{y}) = \sum_{i=1}^N \mathbf{T}(y_i)$ is the vector of sufficient statistics of the distribution (by the factorization theorem).

If the support of **y** is independent of θ , then the family is said to be *regular*, otherwise it is irregular. In addition, if we set $\eta = \eta(\theta)$, then the exponential family is said to be in the *canonical form*

$$p(\mathbf{y}|\boldsymbol{\theta}) = h(\mathbf{y})D(\boldsymbol{\eta})^N \exp\left\{\boldsymbol{\eta}^\top \mathbf{T}(\mathbf{y})\right\}$$
$$= h(\mathbf{y}) \exp\left\{\boldsymbol{\eta}^\top \mathbf{T}(\mathbf{y}) - B(\boldsymbol{\eta})\right\}.$$

A nice feature of this representation is that $\mathbb{E}[\mathbf{T}(\mathbf{y})|\boldsymbol{\eta}] = \nabla B(\boldsymbol{\eta})$ and $Var[\mathbf{T}(\mathbf{y})|\boldsymbol{\eta}] = \nabla^2 B(\boldsymbol{\eta})$.

4.1.1 Examples of exponential family distributions

1. Discrete distributions

Let's show that some of the most common distributions for random

variables that can take values on a finite or countably infinite set are part of the exponential family.

Poisson distribution

Given a random sample $\mathbf{y} = [y_1, y_2, \dots, y_N]^{\top}$ from a *Poisson distribution* let's show that $p(\mathbf{y}|\lambda)$ is in the exponential family.

$$p(\mathbf{y}|\lambda) = \prod_{i=1}^{N} \frac{\lambda^{y_i} \exp(-\lambda)}{y_i!}$$

$$= \frac{\lambda^{\sum_{i=1}^{N} y_i} \exp(-N\lambda)}{\prod_{i=1}^{N} y_i!}$$

$$= \frac{\exp(-N\lambda) \exp(\sum_{i=1}^{N} y_i \log(\lambda))}{\prod_{i=1}^{N} y_i!},$$

then $h(\mathbf{y}) = \left[\prod_{i=1}^N y_i!\right]^{-1}$, $\eta(\lambda) = \log(\lambda)$, $T(\mathbf{y}) = \sum_{i=1}^N y_i$ (sufficient statistic) and $C(\lambda) = \exp(-\lambda)$.

If we set $\eta = \log(\lambda)$, then

$$p(\mathbf{y}|\eta) = \frac{\exp(\eta \sum_{i=1}^{N} y_i - N \exp(\eta))}{\prod_{i=1}^{N} y_i!},$$

such that
$$B(\eta) = N \exp(\eta)$$
, then $\nabla(B(\eta)) = N \exp(\eta) = N\lambda = \mathbb{E}\left[\sum_{i=1}^N y_i \middle| \lambda\right]$, that is, $\mathbb{E}\left[\frac{\sum_{i=1}^N y_i}{N}\middle| \lambda\right] = \mathbb{E}[\bar{y}|\lambda] = \lambda$, and $\nabla^2(B(\eta)) = N \exp(\eta) = N\lambda = Var\left[\sum_{i=1}^N y_i\middle| \lambda\right] = N^2 \times Var\left[\bar{y}|\lambda\right]$, then $Var\left[\bar{y}|\lambda\right] = \frac{\lambda}{N}$.

Bernoulli distribution

Given a random sample $\mathbf{y} = [y_1, y_2, \dots, y_N]^{\top}$ from a *Bernoulli distribution* let's show that $p(\mathbf{y}|\theta)$ is in the exponential family.

$$p(\mathbf{y}|\theta) = \prod_{i=1}^{N} \theta^{y_i} (1 - \theta)^{1 - y_i}$$

= $\theta^{\sum_{i=1}^{N} y_i} (1 - \theta)^{N - \sum_{i=1}^{N} y_i}$
= $(1 - \theta)^N \exp\left\{\sum_{i=1}^{N} y_i \log\left(\frac{\theta}{1 - \theta}\right)\right\}$,

then
$$h(\mathbf{y}) = \mathbb{I}[y_i \in \{0, 1\}]$$
 (indicator function), $\eta(\theta) = \log\left(\frac{\theta}{1-\theta}\right)$, $T(\mathbf{y}) = \sum_{i=1}^{N} y_i$ and $C(\theta) = 1 - \theta$.

Write this distribution in the canonical form, and find the mean and variance of the sufficient statistic (Exercise 1).

Multinomial distribution

Given a random sample $\mathbf{y} = [\mathbf{y}_1, \mathbf{y}_2, \dots, \mathbf{y}_N]$ from a m-dimensional multinomial distribution, where $\mathbf{y}_i = [y_{i1}, \dots, y_{im}], \sum_{l=1}^m y_{il} = n, n$ independent trials each of which leads to a success for exactly one of m categories with probabilities $\boldsymbol{\theta} = [\theta_1, \theta_2, \dots, \theta_m], \sum_{l=1}^m \theta_l = 1$. Let's show that $p(\mathbf{y}|\boldsymbol{\theta})$ is in the exponential family.

$$p(\mathbf{y}|\boldsymbol{\theta}) = \prod_{i=1}^{N} \frac{n!}{\prod_{l=1}^{m} y_{il}!} \prod_{l=1}^{m} \theta_{l}^{y_{il}}$$

$$= \frac{(n!)^{N}}{\prod_{i=1}^{N} \prod_{l=1}^{m} y_{il}!} \exp\left\{\sum_{i=1}^{N} \sum_{l=1}^{m} y_{il} \log(\theta_{l})\right\}$$

$$= \frac{(n!)^{N}}{\prod_{i=1}^{N} \prod_{l=1}^{m} y_{il}!} \exp\left\{\left(N \times n - \sum_{i=1}^{N} \sum_{l=1}^{m-1} y_{il}\right) \log(\theta_{m}) + \sum_{i=1}^{N} \sum_{l=1}^{m-1} y_{il} \log(\theta_{l})\right\}$$

$$= \frac{(n!)^{N}}{\prod_{i=1}^{N} \prod_{l=1}^{m} y_{il}!} \theta_{m}^{N \times n} \exp\left\{\sum_{i=1}^{N} \sum_{l=1}^{m-1} y_{il} \log(\theta_{l}/\theta_{m})\right\},$$
then $h(\mathbf{y}) = \frac{(n!)^{N}}{\prod_{i=1}^{N} \prod_{l=1}^{m} y_{il}!}, \quad \eta(\boldsymbol{\theta}) = \left[\log\left(\frac{\theta_{1}}{\theta_{m}}\right) \dots \log\left(\frac{\theta_{m-1}}{\theta_{m}}\right)\right],$

$$T(\mathbf{y}) = \left[\sum_{i=1}^{N} y_{i1} \dots \sum_{i=1}^{N} y_{im-1}\right] \text{ and } C(\boldsymbol{\theta}) = \theta_{m}^{n}.$$

2. Continuous distributions

Let's show that some of the most common distributions for random variables that can take any value within a certain range or interval, often an infinite number of possible values, are part of the exponential family.

Normal distribution

Given a random sample $\mathbf{y} = [y_1, y_2, \dots, y_N]^{\top}$ from a normal distribution let's show that $p(\mathbf{y}|\mu, \sigma^2)$ is in the exponential family.

$$p(\mathbf{y}|\mu, \sigma^2) = \prod_{i=1}^{N} \frac{1}{2\pi\sigma^2} \exp\left\{-\frac{1}{2\sigma^2} (y_i - \mu)^2\right\}$$
$$= (2\pi)^{-N/2} (\sigma^2)^{-N/2} \exp\left\{-\frac{1}{2\sigma^2} \sum_{i=1}^{N} (y_i - \mu)^2\right\}$$
$$= (2\pi)^{-N/2} \exp\left\{-\frac{1}{2\sigma^2} \sum_{i=1}^{N} y_i^2 + \frac{\mu}{\sigma^2} \sum_{i=1}^{N} y_i - N\frac{\mu^2}{2\sigma^2} - \frac{N}{2} \log(\sigma^2)\right\},$$

then
$$h(\mathbf{y}) = (2\pi)^{-N/2}$$
, $\eta(\mu, \sigma^2) = \left[\frac{\mu}{\sigma^2} \frac{-1}{2\sigma^2}\right]$, $T(\mathbf{y}) = \left[\sum_{i=1}^{N} y_i \sum_{i=1}^{N} y_i^2\right]$ and $C(\mu, \sigma^2) = \exp\left\{-\frac{\mu^2}{2\sigma^2} - \frac{\log(\sigma^2)}{2}\right\}$.

Observe that

$$p(\mathbf{y}|\mu, \sigma^2) = (2\pi)^{-N/2} \exp\left\{\eta_1 \sum_{i=1}^N y_i + \eta_2 \sum_{i=1}^N y_i^2 - \frac{N}{2} \log(-2\eta_2) + \frac{N}{4} \frac{\eta_1^2}{\eta_2}\right\},\,$$

where
$$B(\eta) = \frac{N}{2} \log(-2\eta_2) - \frac{N}{4} \frac{\eta_1^2}{\eta_2}$$
. Then,

$$\nabla B(\boldsymbol{\eta}) = \begin{bmatrix} -\frac{N}{2} \frac{\eta_1}{\eta_2} \\ -\frac{N}{2} \frac{1}{\eta_2} + \frac{N}{4} \frac{\eta_1^2}{\eta_2^2} \end{bmatrix} = \begin{bmatrix} N \times \mu \\ N \times (\mu^2 + \sigma^2) \end{bmatrix} = \begin{bmatrix} \mathbb{E} \left[\sum_{i=1}^N y_i \big| \mu, \sigma^2 \right] \\ \mathbb{E} \left[\sum_{i=1}^N y_i^2 \big| \mu, \sigma^2 \right] \end{bmatrix}.$$

Multivariate normal distribution

Given $\mathbf{Y} = [\mathbf{y}_1 \ \mathbf{y}_2 \ \dots \ \mathbf{y}_p]$ a $N \times p$ matrix such that $\mathbf{y}_i \sim N_p(\boldsymbol{\mu}, \boldsymbol{\Sigma})$, $i = 1, 2, \dots, N$, that is, each *i*-th row of \mathbf{Y} follows a *multivariate* normal distribution. Then, assuming independence between rows, let's show that $p(\mathbf{y}_1, \mathbf{y}_2, \dots, \mathbf{y}_N | \boldsymbol{\mu}, \boldsymbol{\Sigma})$ is in the exponential family.

$$p(\mathbf{y}_{1}, \dots, \mathbf{y}_{N} | \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \prod_{i=1}^{N} (2\pi)^{-p/2} |\boldsymbol{\Sigma}|^{-1/2} \exp\left\{-\frac{1}{2} \left(\mathbf{y}_{i} - \boldsymbol{\mu}\right)^{\top} \boldsymbol{\Sigma}^{-1} \left(\mathbf{y}_{i} - \boldsymbol{\mu}\right)\right\}$$

$$= (2\pi)^{-pN/2} |\boldsymbol{\Sigma}|^{-N/2} \exp\left\{-\frac{1}{2} tr \left[\sum_{i=1}^{N} \left(\mathbf{y}_{i} - \boldsymbol{\mu}\right)^{\top} \boldsymbol{\Sigma}^{-1} \left(\mathbf{y}_{i} - \boldsymbol{\mu}\right)\right]\right\}$$

$$= (2\pi)^{-pN/2} |\boldsymbol{\Sigma}|^{-N/2} \exp\left\{-\frac{1}{2} tr \left[\left(\mathbf{S} + N \left(\boldsymbol{\mu} - \hat{\boldsymbol{\mu}}\right) \left(\boldsymbol{\mu} - \hat{\boldsymbol{\mu}}\right)^{\top}\right) \boldsymbol{\Sigma}^{-1}\right]\right\}$$

$$= (2\pi)^{-pN/2} \exp\left\{-\frac{1}{2} \left[\left(vec(\mathbf{S})^{\top} + Nvec\left(\hat{\boldsymbol{\mu}}\hat{\boldsymbol{\mu}}^{\top}\right)^{\top}\right) vec(\boldsymbol{\Sigma}^{-1})\right]$$

$$-2N\hat{\boldsymbol{\mu}}^{\top} vec\left(\boldsymbol{\mu}^{\top} \boldsymbol{\Sigma}^{-1}\right) + Ntr\left(\boldsymbol{\mu}\boldsymbol{\mu}^{\top} \boldsymbol{\Sigma}^{-1}\right) + N\log(|\boldsymbol{\Sigma}|)\right\},$$

where the second line uses the trace operator (tr), and its invariability under cyclic permutation is used in the third line. In addition, we add and subtract $\hat{\boldsymbol{\mu}} = \frac{1}{N} \sum_{i=1}^{N} \mathbf{y}_{i}$ in each parenthesis such that we get $\mathbf{S} = \sum_{i=1}^{N} (\mathbf{y}_{i} - \hat{\boldsymbol{\mu}}) (\mathbf{y}_{i} - \hat{\boldsymbol{\mu}})^{\top}$. We get the fourth line after collecting terms, and using some properties of the trace operator to introduce the vectorization operator (vec), that is, $tr(\mathbf{AB}) = vec(\mathbf{A}^{\top})^{\top} vec(\mathbf{B})$, and $vec(\mathbf{A} + \mathbf{B}) = vec(\mathbf{A}) + vec(\mathbf{B})$.

Then
$$h(\mathbf{y}) = (2\pi)^{-pN/2}$$
, $\eta(\boldsymbol{\mu}, \boldsymbol{\Sigma})^{\top} = \left[\left(vec\left(\boldsymbol{\Sigma}^{-1}\right) \right)^{\top} \left(vec\left(\boldsymbol{\mu}^{\top} \boldsymbol{\Sigma}^{-1}\right) \right)^{\top} \right]$, $T(\mathbf{y}) = \left[-\frac{1}{2} \left(vec\left(\mathbf{S}\right)^{\top} + Nvec\left(\hat{\boldsymbol{\mu}}\hat{\boldsymbol{\mu}}^{\top}\right)^{\top} \right) - N\hat{\boldsymbol{\mu}}^{\top} \right]^{\top}$ and $C(\boldsymbol{\mu}, \boldsymbol{\Sigma}) = \exp\left\{ -\frac{1}{2} \left(tr\left(\boldsymbol{\mu} \boldsymbol{\mu}^{\top} \boldsymbol{\Sigma}^{-1}\right) + \log(|\boldsymbol{\Sigma}|) \right) \right\}$.

4.2 Conjugate prior to exponential family

Theorem 4.2.1

The prior distribution $\pi(\boldsymbol{\theta}) \propto C(\boldsymbol{\theta})^{b_0} \exp \left\{ \eta(\boldsymbol{\theta})^{\top} \mathbf{a}_0 \right\}$ is conjugate to the exponential family (equation 4.4).

Proof

$$\pi(\boldsymbol{\theta}|\mathbf{y}) \propto C(\boldsymbol{\theta})^{b_0} \exp\left\{\eta(\boldsymbol{\theta})^{\top} \mathbf{a}_0\right\} \times h(\mathbf{y}) C(\boldsymbol{\theta})^N \exp\left\{\eta(\boldsymbol{\theta})^{\top} \mathbf{T}(\mathbf{y})\right\}$$

$$\propto C(\boldsymbol{\theta})^{N+b_0} \exp\left\{\eta(\boldsymbol{\theta})^{\top} (\mathbf{T}(\mathbf{y}) + \mathbf{a}_0)\right\}.$$

Observe that the posterior is in the exponential family, $\pi(\boldsymbol{\theta}|\mathbf{y}) \propto C(\boldsymbol{\theta})^{\beta_n} \exp\left\{\eta(\boldsymbol{\theta})^\top \alpha_n\right\}, \beta_n = N + b_0 \text{ and } \alpha_n = \mathbf{T}(\mathbf{y}) + \mathbf{a}_0.$

Remarks

We see comparing the prior and the likelihood that b_0 plays the role of a hypothetical sample size, and \mathbf{a}_0 plays the role of hypothetical sufficient statistics. This view helps the elicitation process.

We established the result in the *standard form* of the exponential family. We can also establish this result in the *canonical form* of the exponential family. Observe that given $\eta = \eta(\theta)$, another way to get a prior for η is to use the change of variable theorem given a bijective function.

In the setting where there is a regular conjugate prior, [23] show that we obtain a posterior expectation of the sufficient statistics that is a weighted average between the prior expectation and the likelihood estimate.

4.2.1 Examples: Theorem 4.2.1

1. Likelihood functions from discrete distributions

The Poisson-gamma model

Given a random sample $\mathbf{y} = [y_1, y_2, \dots, y_N]^{\top}$ from a Poisson distribution then a conjugate prior density for λ has the form

$$\pi(\lambda) \propto (\exp(-\lambda))^{b_0} \exp\{a_0 \log(\lambda)\}$$
$$= \exp(-\lambda b_0) \lambda^{a_0}$$
$$= \exp(-\lambda \beta_0) \lambda^{\alpha_0 - 1}.$$

This is the kernel of a gamma density in the *rate parametrization*, $G(\alpha_0, \beta_0)$, $\alpha_0 = a_0 + 1$ and $\beta_0 = b_0$. Then, a prior conjugate distribution for the Poisson likelihood is a gamma distribution.

Taking into account that $\sum_{i=1}^{N} y_i$ is a sufficient statistic for the Poisson distribution, then we can think about a_0 as the number of occurrences in b_0 experiments. Observe that

$$\pi(\lambda|\mathbf{y}) \propto \exp(-\lambda\beta_0)\lambda^{\alpha_0-1} \times \exp(-N\lambda)\lambda^{\sum_{i=1}^N y_i}$$
$$= \exp(-\lambda(N+\beta_0))\lambda^{\sum_{i=1}^N y_i + \alpha_0 - 1}.$$

As expected, this is the kernel of a gamma distribution, which means $\lambda | \mathbf{y} \sim G(\alpha_n, \beta_n)$, $\alpha_n = \sum_{i=1}^N y_i + \alpha_0$ and $\beta_n = N + \beta_0$.

Observe that α_0/β_0 is the prior mean, and α_0/β_0^2 is the prior variance. Then, $\alpha_0 \to 0$ and $\beta_0 \to 0$ imply a non-informative prior such that the posterior mean converges to the maximum likelihood estimator $\bar{y} = \frac{\sum_{i=1}^{N} y_i}{N}$,

¹Another parametrization of the gamma density is the scale parametrization where $\kappa_0 = 1/\beta_0$. See the health insurance example in Chapter 1.

$$\mathbb{E}\left[\lambda|\mathbf{y}\right] = \frac{\alpha_n}{\beta_n}$$

$$= \frac{\sum_{i=1}^{N} y_i + \alpha_0}{N + \beta_0}$$

$$= \frac{N\bar{y}}{N + \beta_0} + \frac{\alpha_0}{N + \beta_0}.$$

The posterior mean is a weighted average between sample and prior information. This is a general result from regular conjugate priors [23]. Observe that $\mathbb{E}[\lambda|\mathbf{y}] = \bar{y}, \lim N \to \infty$.

In addition, $\alpha_0 \to 0$ and $\beta_0 \to 0$ corresponds to $\pi(\lambda) \propto \frac{1}{\lambda}$, which is an improper prior. Improper priors may have bad consequences on Bayes factors (hypothesis testing), see bellow a discussion of this in the linear regression framework. In this example, we can get analytical solutions for the marginal likelihood and the predictive distribution (see the health insurance example and Exercise 3 in Chapter 1).

The Bernoulli-beta model

Given a random sample $\mathbf{y} = [y_1, y_2, \dots, y_N]^{\top}$ from a Bernoulli distribution then a conjugate prior density for θ has the form

$$\pi(\theta) \propto (1 - \theta)^{b_0} \exp\left\{a_0 \log\left(\frac{\theta}{1 - \theta}\right)\right\}$$
$$= (1 - \theta)^{b_0 - a_0} \theta^{a_0}$$
$$= \theta^{\alpha_0 - 1} (1 - \theta)^{\beta_0 - 1}.$$

This is the kernel of a beta density, $B(\alpha_0, \beta_0)$, $\alpha_0 = a_0 + 1$ and $\beta_0 = b_0 - a_0 + 1$. A prior conjugate distribution for the Bernoulli likelihood is a beta distribution. Given that b_0 is the hypothetical sample size, and a_0 is the hypothetical sufficient statistic, which is the number of successes, then $b_0 - a_0$ is the number of failures. This implies that α_0 is the number of prior successes plus one, and β_0 is the number of prior failures plus one. Given that the mode of a beta distributed random variable is $\frac{\alpha_0 - 1}{\alpha_0 + \beta_0 - 2} = \frac{a_0}{b_0}$, then we have the prior probability of success. Setting $\alpha_0 = 1$ and $\beta_0 = 1$, which implies a 0-1 uniform distribution, corresponds to a setting with 0 successes (and 0 failures) in 0 experiments.

Observe that

$$\pi(\theta|\mathbf{y}) \propto \theta^{\alpha_0 - 1} (1 - \theta)^{\beta_0 - 1} \times \theta^{\sum_{i=1}^N y_i} (1 - \theta)^{N - \sum_{i=1}^N y_i}$$
$$= \theta^{\alpha_0 + \sum_{i=1}^N y_i - 1} (1 - \theta)^{\beta_0 + N - \sum_{i=1}^N y_i - 1}.$$

The posterior distribution is beta, $\theta | \mathbf{y} \sim B(\alpha_n, \beta_n)$, $\alpha_n = \alpha_0 + \sum_{i=1}^N y_i$ and $\beta_n = \beta_0 + N - \sum_{i=1}^N y_i$, where the posterior mean $\mathbb{E}[\theta | \mathbf{y}] = \frac{\alpha_n}{\alpha_n + \beta_n} = \frac{\alpha_0 + N\bar{y}}{\alpha_0 + \beta_0 + N} = \frac{\alpha_0 + \beta_0}{\alpha_0 + \beta_0 + N} \frac{\alpha_0}{\alpha_0 + \beta_0} + \frac{N}{\alpha_0 + \beta_0 + N} \bar{y}$. The posterior mean is a weighted average between the prior mean and the maximum likelihood estimate.

El marginal likelihood in this setting is

$$p(\mathbf{y}) = \int_0^1 \frac{\theta^{\alpha_0 - 1} (1 - \theta)^{\beta_0 - 1}}{B(\alpha_0, \beta_0)} \times \theta^{\sum_{i=1}^N y_i} (1 - \theta)^{N - \sum_{i=1}^N y_i} d\theta$$
$$= \frac{B(\alpha_n, \beta_n)}{B(\alpha_0, \beta_0)},$$

where $B(\cdot, \cdot)$ is the beta function.

In addition, the predictive density is

$$\begin{split} p(Y_0|\mathbf{y}) &= \int_0^1 \theta^{y_0} (1-\theta)^{1-y_0} \times \frac{\theta^{\alpha_n-1} (1-\theta)^{\beta_n-1}}{B(\alpha_n,\beta_n)} d\theta \\ &= \frac{B(\alpha_n+y_0,\beta_n+1-y_0)}{B(\alpha_n,\beta_n)} \\ &= \frac{\Gamma(\alpha_n+\beta_n)\Gamma(\alpha_n+y_0)\Gamma(\beta_n+1-y_0)}{\Gamma(\alpha_n+\beta_n+1)\Gamma(\alpha_n)\Gamma(\beta_n)} \\ &= \left\{ \frac{\frac{\alpha_n}{\alpha_n+\beta_n}, \quad y_0 = 1}{\frac{\beta_n}{\alpha_n+\beta_n}, \quad y_0 = 0} \right\}. \end{split}$$

This is a Bernoulli distribution with probability of success equal to $\frac{\alpha_n}{\alpha_n + \beta_n}$.

The multinomial-Dirichlet model

Given a random sample $\mathbf{y} = [y_1, y_2, \dots, y_N]^{\top}$ from a multinomial distribution then a conjugate prior density for $\boldsymbol{\theta} = [\theta_1, \theta_2, \dots, \theta_m]$ has the form

$$\pi(\boldsymbol{\theta}) \propto \theta_m^{b_0} \exp\left\{\boldsymbol{\eta}(\boldsymbol{\theta})^\top \mathbf{a}_0\right\}$$
$$= \prod_{l=1}^{m-1} \theta_l^{a_{0l}} \theta_m^{b_0 - \sum_{l=1}^{m-1} a_{0l}}$$
$$= \prod_{l=1}^m \theta_l^{\alpha_{0l} - 1},$$

where
$$\boldsymbol{\eta}(\boldsymbol{\theta}) = \left[\log\left(\frac{\theta_1}{\theta_m}\right), \dots, \log\left(\frac{\theta_{m-1}}{\theta_m}\right)\right], \ \mathbf{a}_0 = \left[a_{01}, \dots, a_{am-1}\right]^\top,$$

$$\boldsymbol{\alpha}_0 = \left[\alpha_{01}, \alpha_{02}, \dots, \alpha_{0m}\right], \ \alpha_{0l} = a_{0l} + 1, \ l = 1, 2, \dots, m-1 \text{ and }$$

$$\alpha_{0m} = b_0 - \sum_{l=1}^{m-1} a_{0l} + 1.$$

This is the kernel of a Dirichlet distribution, that is, the prior distribution is $D(\alpha_0)$.

Observe that a_{0l} is the number of hypothetical number of times outcome l is observed over the hypothetical b_0 trials. Setting $\alpha_{0l} = 1$, that is a uniform distribution over the open standard simplex, implicitly we set $a_{0l} = 0$, which means that there are 0 occurrences of category l in $b_0 = 0$ experiments.

The posterior distribution of the multinomial-Dirichlet model is given by

$$\pi(\boldsymbol{\theta}|\mathbf{y}) \propto \prod_{l=1}^{m} \theta_{l}^{\alpha_{0l}-1} \times \prod_{l=1}^{m} \theta_{l}^{\sum_{i=1}^{N} y_{il}}$$
$$= \prod_{l=1}^{m} \theta_{l}^{\alpha_{0l} + \sum_{i=1}^{N} y_{il} - 1}.$$

This is the kernel of a Dirichlet distribution $D(\boldsymbol{\alpha}_n)$, $\boldsymbol{\alpha}_n = [\alpha_{n1}, \alpha_{n2}, \dots, \alpha_{nm}]$, $\alpha_{nl} = \alpha_{0l} + \sum_{i=1}^{N} y_{il}$, $l = 1, 2, \dots, m$. Observe that

$$\mathbb{E}[\theta_{j}|\mathbf{y}] = \frac{\alpha_{nj}}{\sum_{l=1}^{m} \left[\alpha_{0l} + \sum_{i=1}^{N} y_{il}\right]}$$

$$= \frac{\sum_{l=1}^{m} \alpha_{0l}}{\sum_{l=1}^{m} \left[\alpha_{0l} + \sum_{i=1}^{N} y_{il}\right]} \frac{\alpha_{0j}}{\sum_{l=1}^{m} \alpha_{0l}}$$

$$+ \frac{\sum_{l=1}^{m} \sum_{i=1}^{N} y_{il}}{\sum_{l=1}^{m} \sum_{i=1}^{N} y_{il}} \frac{\sum_{i=1}^{N} y_{ij}}{\sum_{l=1}^{m} \sum_{i=1}^{N} y_{il}}.$$

We have again that the posterior mean is a weighted average between the prior mean and the maximum likelihood estimate.

The marginal likelihood is

$$p(\mathbf{y}) = \int_{\mathbf{\Theta}} \frac{\prod_{l=1}^{m} \theta_{l}^{\alpha_{0l}-1}}{B(\boldsymbol{\alpha}_{0})} \times \prod_{i=1}^{N} \frac{n!}{\prod_{l=1}^{m} y_{il}} \prod_{l=1}^{m} \theta_{l}^{y_{il}} d\boldsymbol{\theta}$$

$$= \frac{N \times n!}{B(\boldsymbol{\alpha}_{0}) \prod_{i=1}^{N} \prod_{l=1}^{m} y_{il}!} \int_{\mathbf{\Theta}} \prod_{l=1}^{m} \theta_{l}^{\alpha_{0l} + \sum_{i=1}^{N} y_{il} - 1} d\boldsymbol{\theta}$$

$$= \frac{N \times n!}{B(\boldsymbol{\alpha}_{0}) \prod_{i=1}^{N} \prod_{l=1}^{m} y_{il}!} B(\boldsymbol{\alpha}_{n})$$

$$= \frac{N \times n! \Gamma(\sum_{l=1}^{m} \alpha_{0l})}{\Gamma(\sum_{l=1}^{m} \alpha_{0l} + N \times n)} \prod_{l=1}^{m} \frac{\Gamma(\alpha_{nl})}{\Gamma(\alpha_{0l}) \prod_{i=1}^{N} y_{il}!},$$

where
$$B(\boldsymbol{\alpha}) = \frac{\prod_{l=1}^{m} \Gamma(\alpha_l)}{\Gamma(\sum_{l=1}^{m} \alpha_l)}$$
.

Following similar steps we get the predictive density

$$p(Y_0|\mathbf{y}) = \frac{n!\Gamma\left(\sum_{l=1}^{m} \alpha_{nl}\right)}{\Gamma\left(\sum_{l=1}^{m} \alpha_{nl} + n\right)} \prod_{l=1}^{m} \frac{\Gamma\left(\alpha_{nl} + y_{0l}\right)}{\Gamma\left(\alpha_{nl}\right) y_{0l}!}.$$

This is a Dirichlet-multinomial distribution with parameters α_n .

Example: English premier league, Liverpool vs Manchester city

Let's see an example based on data from the English Premier league. In particular, we want to get the probability that in the following five matches Liverpool versus Manchester city, the former wins two games, and the latter three game. This is done based on the historical records of the last five matches where Liverpool was local between January 14th, 2018 and April tenth, 2022. There were two wins for Liverpool, two draws, and one win for Manchester city.²

We use two strategies to get the hyperparameters. First, we estimate the hyperparameters of the Dirichlet distribution using betting odds from bookmakers at 19:05 hours October sixth, 2022 (Colombia time). We got information from 24 bookmakers (see file DataOddsLIVvsMAN.csv),³ and transform these odds in probabilities using a simple standardization approach, then we use maximum

²https://www.11v11.com/teams/manchester-city/tab/opposingTeams/opposition/Liverpool/.

 $^{^3} https://www.oddsportal.com/soccer/england/premier-league/liverpool-manchester-city-WrqgEz5S/$

likelihood to estimate the hyperparameters. Second, we use empirical Bayes, that is, we estimate the hyperparameters optimizing the marginal likelihood.

$R\ code.\ Multinomial\text{-}Dirichlet\ model:\ Liverpool\ vs}\ Manchester\ city$

```
1 # Multinomial-Dirichlet example: Liverpool vs Manchester
2 Data <-read.csv("DataApplications/DataOddsLIVvsMAN.csv", sep</pre>
      = ",", header = TRUE)
3 # Change path
4 attach (Data)
5 library(dplyr)
6 Probs <- Data %>%
    mutate(pns1 = 1/home, pns2 = 1/draw, pns3 = 1/away)%>%
    mutate(SumInvOdds = pns1 + pns2 + pns3) %>%
    mutate(p1 = pns1/SumInvOdds, p2 = pns2/SumInvOdds, p3 =
      pns3/SumInvOdds) %>%
    select(p1, p2, p3)
11 # We get probabilities using simple standardization. There
      are more technical approaches to do this. See for
      instance Shin (1993) and Strumbelj (2014).
12 DirMLE <- sirt::dirichlet.mle(Probs)</pre>
13 # Use maximum likelihood to estimate parameters of the
14 # Dirichlet distribution
15 alpha0odds <- DirMLE$alpha
16 alpha0odds
        p1
                  p2
18 1599.122 1342.703 2483.129
20 y < -c(2, 2, 1)
21 # Historical records last five mathces
^{22} # Liverpool wins (2), draws (2) and Manchester
23 # city wins (1)
25 # Marginal likelihood
26 MarLik <- function(a0){
   n <- sum(y)
    Res1 <- sum(sapply(1:length(y),</pre>
    function(1){lgamma(a0[1]+y[1])-lgamma(a0[1])}))
    Res <- lgamma(sum(a0))-lgamma(sum(a0)+n)+Res1
    return(-Res)
32 }
33 EmpBay <- optim(alpha0odds, MarLik, method = "BFGS")
34 alpha0EB <- EmpBay$par
35 alpha0EB
36 p1
           p2
37 2362.622 2660.153 1279.510
38 # Bayes factor empirical Bayes vs betting odds.
39 # This is greather than 1 by construction
40 BF <- exp(-MarLik(alpha0EB))/exp(-MarLik(alpha0odds))
41 BF
42 2.085819
43 # Posterior distribution based on empirical Bayes
44 alphan <- alpha0EB + y
45 # Posterior parameters
46 S <- 100000
47 # Simulation draws from the Dirichlet distribution
48 thetas <- MCMCpack::rdirichlet(S, alphan)
49 colnames (thetas) <- c("Liverpool", "Draw", "Manchester")
```

$R\ code.\ Multinomial\text{-}Dirichlet\ model:\ Liverpool\ vs}\ Manchester\ city$

```
# Predictive distribution based on simulations
_{2} y0 <- c(2, 0, 3)
  # Liverpool two wins and Manchester city three wins in next
      five matches
  Pred <- apply(thetas, 1, function(p) {rmultinom(1, size =</pre>
      sum(y0), prob = p)
  ProYo <- sum(sapply(1:S,function(s){sum(Pred[,s]==y0)==3}))/</pre>
  ProY0
  0.0832
8 # Probability of y0
10 # Predictive distribution using analytical expression
  PredY0 <- function(y0){</pre>
    n <- sum(y0)
    Res1 <- sum(sapply(1:length(y), function(1){lgamma(alphan[</pre>
      1]+y0[1]) - lgamma(alphan[1])-lfactorial(y0[1])}))
    Res <- lfactorial(n) + lgamma(sum(alphan)) - lgamma(sum(
      alphan)+n) + Res1
    return(exp(Res))
16 }
17 PredYO(y0)
18 0.0833
```

We see that the Bayes factor gives evidence in favor of the hyperparameters based on empirical Bayes, this is by construction, as these hyperparameters maximize the marginal likelihood.

We observe that using the hyperparameters from empirical Bayes, the probability that in the next five games Liverpool wins two games and Manchester city wins three games is 8.33%. The result using the predictive distribution based on simulations is similar to the probability using the exact predictive.

2. Likelihood functions from continuous distributions

The normal-normal/inverse-gamma model

Given a random sample $\mathbf{y} = [y_1, y_2, \dots, y_N]^{\top}$ from a normal distribution, then the conjugate prior density has the form

$$\pi(\mu, \sigma^{2}) \propto \exp\left\{b_{0}\left(-\frac{\mu^{2}}{2\sigma^{2}} - \frac{\log \sigma^{2}}{2}\right)\right\} \exp\left\{a_{01}\frac{\mu}{\sigma^{2}} - a_{02}\frac{1}{\sigma^{2}}\right\}$$

$$= \exp\left\{b_{0}\left(-\frac{\mu^{2}}{2\sigma^{2}} - \frac{\log \sigma^{2}}{2}\right)\right\} \exp\left\{a_{01}\frac{\mu}{\sigma^{2}} - a_{02}\frac{1}{\sigma^{2}}\right\}$$

$$\times \exp\left\{-\frac{a_{01}^{2}}{2\sigma^{2}b_{0}}\right\} \exp\left\{\frac{a_{01}^{2}}{2\sigma^{2}b_{0}}\right\}$$

$$= \exp\left\{-\frac{b_{0}}{2\sigma^{2}}\left(\mu - \frac{a_{01}}{b_{0}}\right)^{2}\right\} \left(\frac{1}{\sigma^{2}}\right)^{\frac{b_{0}+1-1}{2}}$$

$$\times \exp\left\{\frac{1}{\sigma^{2}} - \frac{2b_{0}a_{02} + a_{01}^{2}}{2b_{0}}\right\}$$

$$= \underbrace{\left(\frac{1}{\sigma^{2}}\right)^{\frac{1}{2}} \exp\left\{-\frac{b_{0}}{2\sigma^{2}}\left(\mu - \frac{a_{01}}{b_{0}}\right)^{2}\right\}}_{1}$$

$$\times \underbrace{\left(\frac{1}{\sigma^{2}}\right)^{\frac{b_{0}-1}{2}} \exp\left\{-\frac{1}{\sigma^{2}} \frac{2b_{0}a_{02} - a_{01}^{2}}{2b_{0}}\right\}}_{2}.$$

The first part is the kernel of a normal density with mean $\mu_0 = a_{01}/\beta_0$ and variance σ^2/β_0 , $\beta_0 = b_0$ that is, $\mu|\sigma^2 \sim N(\mu_0, \sigma^2/\beta_0)$. The second part is the kernel of an inverse gamma density with shape parameter $\alpha_0/2 = \frac{\beta_0 - 3}{2}$, and scale parameter $\delta_0/2 = \frac{2\beta_0 a_{02} - a_{01}^2}{2\beta_0}$, $\sigma^2 \sim IG(\alpha_0/2, \delta_0/2)$. Observe that $b_0 = \beta_0$ is the hypothetical sample size, and a_{01} is the hypothetical sum of prior observations, then, it makes sense that a_{01}/β_0 and σ^2/β_0 are the prior mean and variance, respectively.

Therefore, the posterior distribution is also a normal-inverse gamma

distribution,

$$\pi(\mu, \sigma^{2}|\mathbf{y}) \propto \left(\frac{1}{\sigma^{2}}\right)^{1/2} \exp\left\{-\frac{\beta_{0}}{2\sigma^{2}}(\mu - \mu_{0})^{2}\right\} \left(\frac{1}{\sigma^{2}}\right)^{\alpha_{0}/2+1} \exp\left\{-\frac{\delta_{0}}{2\sigma^{2}}\right\}$$

$$\times (\sigma^{2})^{-N/2} \exp\left\{-\frac{1}{2\sigma^{2}}\sum_{i=1}^{N}(y_{i} - \mu)^{2}\right\}$$

$$= \left(\frac{1}{\sigma^{2}}\right)^{1/2} \exp\left\{-\frac{1}{2\sigma^{2}}\left(\beta_{0}(\mu - \mu_{0})^{2} + \sum_{i=1}^{N}(y_{i} - \bar{y})^{2} + N(\mu - \bar{y})^{2} + \delta_{0}\right)\right\}$$

$$\times \left(\frac{1}{\sigma^{2}}\right)^{\frac{\alpha_{0}+N}{2}+1} + \frac{(\beta_{0}\mu_{0} + N\bar{y})^{2}}{\beta_{0} + N} - \frac{(\beta_{0}\mu_{0} + N\bar{y})^{2}}{\beta_{0} + N}$$

$$= \left(\frac{1}{\sigma^{2}}\right)^{1/2} \exp\left\{-\frac{1}{2\sigma^{2}}\left((\beta_{0} + N)\left(\mu - \left(\frac{\beta_{0}\mu_{0} + N\bar{y}}{\beta_{0} + N}\right)\right)^{2}\right)\right\}$$

$$\times \left(\frac{1}{\sigma^{2}}\right)^{\frac{\alpha_{0}+N}{2}+1} \exp\left\{-\frac{1}{2\sigma^{2}}\left(\sum_{i=1}^{N}(y_{i} - \bar{y})^{2} + \delta_{0} + \frac{\beta_{0}N}{\beta_{0} + N}(\bar{y} - \mu_{0})^{2}\right)\right\}.$$

The first term is the kernel of a normal density, $\mu|\sigma^2, \mathbf{y} \sim N(\mu_n, \sigma_n^2)$, where $\mu_n = \frac{\beta_0 \mu_0 + N \bar{y}}{\beta_0 + N}$ and $\sigma_n^2 = \frac{\sigma^2}{\beta_n}$, $\beta_n = \beta_0 + N$. The second term is the kernel of an inverse gamma density, $\sigma^2|\mathbf{y} \sim IG(\alpha_n/2, \delta_n/2)$ where $\alpha_n = \alpha_0 + N$ and $\delta_n = \sum_{i=1}^N (y_i - \bar{y})^2 + \delta_0 + \frac{\beta_0 N}{\beta_0 + N} (\bar{y} - \mu_0)^2$. Observe that the posterior mean is a weighted average between prior and sample information. The weights depends on the sample sizes $(\beta_0 \text{ and } N)$.

The marginal posterior for σ^2 is inverse gamma with shape and scale parameters $\alpha_n/2$ and $\delta_n/2$, respectively. The marginal posterior of μ is

$$\pi(\mu|\mathbf{y}) \propto \int_0^\infty \left\{ \left(\frac{1}{\sigma^2}\right)^{\frac{\alpha_n+1}{2}+1} \exp\left\{-\frac{1}{2\sigma^2} (\beta_n(\mu-\mu_n)^2 + \delta_n)\right\} \right\} d\sigma^2$$

$$= \frac{\Gamma\left(\frac{\alpha_n+1}{2}\right)}{\left[\frac{\beta_n(\mu-\mu_n)^2+\delta_n}{2}\right]^{\frac{\alpha_n+1}{2}}}$$

$$\propto \left[\frac{\beta_n(\mu-\mu_n)^2+\delta_n}{2}\right]^{-\frac{\alpha_n+1}{2}} \left(\frac{\delta_n}{\delta_n}\right)^{-\frac{\alpha_n+1}{2}}$$

$$\propto \left[\frac{\alpha_n\beta_n(\mu-\mu_n)^2}{\alpha_n\delta_n} + 1\right]^{-\frac{\alpha_n+1}{2}},$$

where the second line due to having the kernel of an inverse gamma density with parameters $(\alpha_n + 1)/2$ and $-\frac{1}{2\sigma^2}(\beta_n(\mu - \mu_n)^2 + \delta_n)$.

This is the kernel of a Student's t distribution, $\mu|\mathbf{y} \sim t(\mu_n, \delta_n/\beta_n\alpha_n, \alpha_n)$, where $\mathbb{E}[\mu|\mathbf{y}] = \mu_n$ and $Var[\mu|\mathbf{y}] = \frac{\alpha_n}{\alpha_n-2} \left(\frac{\delta_n}{\beta_n\alpha_n}\right) = \frac{\delta_n}{(\alpha_n-2)\beta_n}$, $\alpha_n > 2$. Observe that the marginal posterior distribution for μ has heavier tails than the conditional posterior distribution due to incorporating uncertainty regarding σ^2 . The marginal likelihood is

$$\begin{split} p(\mathbf{y}) &= \int_{-\infty}^{\infty} \int_{0}^{\infty} \left\{ (2\pi\sigma^{2}/\beta_{0})^{-1/2} \exp\left\{ -\frac{1}{2\sigma^{2}/\beta_{0}} (\mu - \mu_{0})^{2} \right\} \frac{(\delta_{0}/2)^{\alpha_{0}/2}}{\Gamma(\alpha_{0}/2)} \left(\frac{1}{\sigma^{2}} \right)^{\alpha_{0}/2+1} \\ &\times \exp\left\{ -\frac{\delta_{0}}{2\sigma^{2}} \right\} (2\pi\sigma^{2})^{-N/2} \exp\left\{ -\frac{1}{2\sigma^{2}} \sum_{i=1}^{N} (y_{i} - \mu)^{2} \right\} \right\} d\sigma^{2} d\mu \\ &= \frac{(\delta_{0}/2)^{\alpha_{0}/2}}{\Gamma(\alpha_{0}/2)} (2\pi)^{-\left(\frac{N+1}{2}\right)} \beta_{0}^{1/2} \int_{-\infty}^{\infty} \int_{0}^{\infty} \left\{ \left(\frac{1}{\sigma^{2}} \right)^{\frac{\alpha_{0}+N+1}{2}+1} \right. \\ &\times \exp\left\{ -\frac{1}{2\sigma^{2}} (\beta_{0}(\mu - \mu_{0})^{2} + \sum_{i=1}^{N} (y_{i} - \mu)^{2} + \delta_{0}) \right\} \right\} d\sigma^{2} d\mu \\ &= \frac{(\delta_{0}/2)^{\alpha_{0}/2}}{\Gamma(\alpha_{0}/2)} (2\pi)^{-\left(\frac{N+1}{2}\right)} \beta_{0}^{1/2} \Gamma\left(\frac{N+1+\alpha_{0}}{2} \right) \\ &\times \int_{-\infty}^{\infty} \left[\frac{\beta_{0}(\mu - \mu_{0})^{2} + \sum_{i=1}^{N} (y_{i} - \mu)^{2} + \delta_{0}}{2} \right]^{-\frac{\alpha_{0}+N+1}{2}} d\mu \\ &= \frac{(\delta_{0}/2)^{\alpha_{0}/2}}{\Gamma(\alpha_{0}/2)} (2\pi)^{-\left(\frac{N+1}{2}\right)} \beta_{0}^{1/2} \Gamma\left(\frac{N+1+\alpha_{0}}{2} \right) \\ &\times \int_{-\infty}^{\infty} \left[\frac{\beta_{n}(\mu - \mu_{n})^{2} + \delta_{n}}{2} \right]^{-\frac{\alpha_{n}+1}{2}} d\mu \left(\frac{\delta_{n}/2}{\delta_{n}/2} \right)^{-\frac{\alpha_{n}+1}{2}} \\ &= \frac{(\delta_{0}/2)^{\alpha_{0}/2}}{\Gamma(\alpha_{0}/2)} (2\pi)^{-\left(\frac{N+1}{2}\right)} \beta_{0}^{1/2} \Gamma\left(\frac{\alpha_{n}+1}{2} \right) \left(\frac{\delta_{n}}{2} \right)^{-\frac{\alpha_{n}+1}{2}} \frac{\left(\frac{\delta_{n}\pi}{\beta_{n}} \right)^{1/2} \Gamma\left(\frac{\alpha_{n}}{2} \right)}{\Gamma\left(\frac{\alpha_{n}+1}{2}\right)} \\ &= \frac{\Gamma\left(\frac{\alpha_{n}}{2}\right)}{\Gamma\left(\frac{\alpha_{0}}{2}\right)} \frac{(\delta_{0}/2)^{\alpha_{0}/2}}{(\delta_{n}/2)^{\alpha_{n}/2}} \left(\frac{\beta_{0}}{\beta_{n}} \right)^{1/2} (\pi)^{-N/2}, \end{split}$$

where we take into account that $\int_{-\infty}^{\infty} \left[\frac{\beta_n (\mu - \mu_n)^2 + \delta_n}{2} \right]^{-\frac{\alpha_n + 1}{2}} d\mu \left(\frac{\delta_n / 2}{\delta_n / 2} \right)^{-\frac{\alpha_n + 1}{2}} = \int_{-\infty}^{\infty} \left[\frac{\beta_n \alpha_n (\mu - \mu_n)^2}{\delta_n \alpha_n} + 1 \right]^{-\frac{\alpha_n + 1}{2}} d\mu \left(\frac{\delta_n}{2} \right)^{-\frac{\alpha_n + 1}{2}}.$ The term in the integral is the kernel of a Student's t density, this means that the integral is equal to $\frac{\left(\frac{\delta_n \pi}{\beta_n}\right)^{1/2} \Gamma\left(\frac{\alpha_n}{2}\right)}{\Gamma\left(\frac{\alpha_n + 1}{2}\right)}.$

The predictive density is

$$\pi(Y_{0}|\mathbf{y}) \propto \int_{-\infty}^{\infty} \int_{0}^{\infty} \left\{ \left(\frac{1}{\sigma^{2}}\right)^{1/2} \exp\left\{-\frac{1}{2\sigma^{2}}(y_{0}-\mu)^{2}\right\} \left(\frac{1}{\sigma^{2}}\right)^{1/2} \exp\left\{-\frac{\beta_{n}}{2\sigma^{2}}(\mu-\mu_{n})^{2}\right\} \right\} \\
\times \left(\frac{1}{\sigma^{2}}\right)^{\alpha_{n}/2+1} \exp\left\{-\frac{\delta_{n}}{2\sigma^{2}}\right\} d\sigma^{2} d\mu \\
= \int_{-\infty}^{\infty} \int_{0}^{\infty} \left\{ \left(\frac{1}{\sigma^{2}}\right)^{\frac{\alpha_{n}+2}{2}+1} \exp\left\{-\frac{1}{2\sigma^{2}}((y_{0}-\mu)^{2}+\beta_{n}(\mu-\mu_{n})^{2}+\delta_{n})\right\} \right\} d\sigma^{2} d\mu \\
\propto \int_{-\infty}^{\infty} \left[\beta_{n}(\mu-\mu_{n})^{2}+(y_{0}-\mu)^{2}+\delta_{n}\right]^{-\left(\frac{\alpha_{n}}{2}+1\right)} d\mu \\
= \int_{-\infty}^{\infty} \left[(\beta_{n}+1) \left(\mu-\left(\frac{\beta_{n}\mu_{n}+y_{0}}{\beta_{n}+1}\right)\right)^{2}+\frac{\beta_{n}(y_{0}-\mu_{n})^{2}}{\beta_{n}+1}+\delta_{n} \right]^{-\left(\frac{\alpha_{n}}{2}+1\right)} d\mu \\
= \int_{-\infty}^{\infty} \left[1+\frac{(\beta_{n}+1)^{2} \left(\mu-\left(\frac{\beta_{n}\mu_{n}+y_{0}}{\beta_{n}+1}\right)\right)^{2}}{\beta_{n}(y_{0}-\mu_{n})^{2}+(\beta_{n}+1)\delta_{n}} \right]^{-\left(\frac{\alpha_{n}}{2}+1\right)} d\mu \\
\times \left(\frac{\beta_{n}(y_{0}-\mu_{n})^{2}+(\beta_{n}+1)\delta_{n}}{\beta_{n}+1}\right)^{-\left(\frac{\alpha_{n}}{2}+1\right)} \\
\propto \left(\frac{\beta_{n}(y_{0}-\mu_{n})^{2}+(\beta_{n}+1)\delta_{n}}{(\beta_{n}+1)^{2}(\alpha_{n}+1)}\right)^{\frac{1}{2}} \left(\frac{\beta_{n}(y_{0}-\mu_{n})^{2}+(\beta_{n}+1)\delta_{n}}{\beta_{n}+1}\right)^{-\left(\frac{\alpha_{n}}{2}+1\right)} \\
\propto (\beta_{n}(y_{0}-\mu_{n})^{2}+(\beta_{n}+1)\delta_{n})^{\left(\frac{\alpha_{n}+1}{2}\right)} \\
\propto \left[1+\frac{\beta_{n}\alpha_{n}}{(\beta_{n}+1)\delta_{n}\alpha_{n}}(y_{0}-\mu_{n})^{2}\right]^{-\left(\frac{\alpha_{n}+1}{2}\right)},$$

where we have that $\left[1 + \frac{(\beta_n+1)^2\left(\mu - \left(\frac{\beta_n\mu_n + y_0}{\beta_n+1}\right)\right)^2}{\beta_n(y_0 - \mu_n)^2 + (\beta_n+1)\delta_n}\right]^{-\left(\frac{\alpha_n}{2} + 1\right)}$ is the ker-

nel of a Student's t density with degrees of freedom α_n+1 and scale $\frac{\beta_n(y_0-\mu_n)^2+(\beta_n+1)\delta_n}{(\beta_n+1)^2(\alpha_n+1)}$.

The last expression is the kernel of a Student's t density, that is, $Y_0|\mathbf{y} \sim t\left(\mu_n, \frac{(\beta_n+1)\delta_n}{\beta_n\alpha_n}, \alpha_n\right).$

The multivariate normal-normal/inverse-Wishart model

We show in subsection 4.1 that the multivariate normal distribution is in the exponential family where

$$\begin{split} C(\boldsymbol{\mu}, \boldsymbol{\Sigma}) &= \exp\left\{-\frac{1}{2} \left(tr\left(\boldsymbol{\mu} \boldsymbol{\mu}^{\top} \boldsymbol{\Sigma}^{-1}\right) + \log(|\boldsymbol{\Sigma}|)\right)\right\}, \\ \eta(\boldsymbol{\mu}, \boldsymbol{\Sigma})^{\top} &= \left[\left(vec\left(\boldsymbol{\Sigma}^{-1}\right)\right)^{\top} \ \left(vec\left(\boldsymbol{\mu}^{\top} \boldsymbol{\Sigma}^{-1}\right)\right)^{\top}\right], \\ T(\mathbf{y}) &= \left[-\frac{1}{2} \left(vec\left(\mathbf{S}\right)^{\top} + Nvec\left(\hat{\boldsymbol{\mu}} \hat{\boldsymbol{\mu}}^{\top}\right)^{\top}\right) \ - N\hat{\boldsymbol{\mu}}^{\top}\right]^{\top} \end{split}$$

and

$$h(\mathbf{y}) = (2\pi)^{-pN/2}$$
.

Then, its conjugate prior distribution should have the form

$$\pi(\boldsymbol{\mu}, \boldsymbol{\Sigma}) \propto \exp\left\{-\frac{b_0}{2} \left(tr\left(\boldsymbol{\mu}\boldsymbol{\mu}^{\top}\boldsymbol{\Sigma}^{-1}\right) + \log(|\boldsymbol{\Sigma}|)\right)\right\}$$

$$\times \exp\left\{\mathbf{a}_{01}^{\top}vec\left(\boldsymbol{\Sigma}^{-1}\right) + \mathbf{a}_{02}^{\top}vec\left(\boldsymbol{\mu}^{\top}\boldsymbol{\Sigma}^{-1}\right)\right\}$$

$$= |\boldsymbol{\Sigma}|^{-b_0/2} \exp\left\{-\frac{b_0}{2} \left(tr\left(\boldsymbol{\mu}^{\top}\boldsymbol{\Sigma}^{-1}\boldsymbol{\mu}\right)\right) + tr\left(\mathbf{a}_{02}^{\top}\boldsymbol{\Sigma}^{-1}\boldsymbol{\mu}\right)\right\}$$

$$\times \exp\left\{\mathbf{a}_{01}^{\top}vec\left(\boldsymbol{\Sigma}^{-1}\right) + \frac{\mathbf{a}_{02}^{\top}\boldsymbol{\Sigma}^{-1}\mathbf{a}_{02}}{2b_0} - \frac{\mathbf{a}_{02}^{\top}\boldsymbol{\Sigma}^{-1}\mathbf{a}_{02}}{2b_0}\right\}$$

$$= |\boldsymbol{\Sigma}|^{-b_0/2} \exp\left\{-\frac{b_0}{2} \left(\boldsymbol{\mu} - \frac{\mathbf{a}_{02}}{b_0}\right)^{\top} \boldsymbol{\Sigma}^{-1} \left(\boldsymbol{\mu} - \frac{\mathbf{a}_{02}}{b_0}\right)\right\}$$

$$\times \exp\left\{-\frac{1}{2}tr\left(\left(\mathbf{A}_{01} - \frac{\mathbf{a}_{02}\mathbf{a}_{02}^{\top}}{b_0}\right) \boldsymbol{\Sigma}^{-1}\right)\right\}$$

$$= |\boldsymbol{\Sigma}|^{-1/2} \exp\left\{-\frac{b_0}{2} \left(\boldsymbol{\mu} - \frac{\mathbf{a}_{02}}{b_0}\right)^{\top} \boldsymbol{\Sigma}^{-1} \left(\boldsymbol{\mu} - \frac{\mathbf{a}_{02}}{b_0}\right)\right\}$$

$$\times |\boldsymbol{\Sigma}|^{-(\alpha_0 + p + 1)/2} \exp\left\{-\frac{1}{2}tr\left(\left(\mathbf{A}_{01} - \frac{\mathbf{a}_{02}\mathbf{a}_{02}^{\top}}{b_0}\right) \boldsymbol{\Sigma}^{-1}\right)\right\},$$

where b_0 is the hypothetical sample size, and \mathbf{a}_{01} and \mathbf{a}_{02} are p^2 and p dimensional vectors of prior sufficient statistics, where $\mathbf{a}_{01} = -\frac{1}{2}vec(\mathbf{A}_{01})$ such that \mathbf{A}_{01} is a $p \times p$ positive semi-definite matrix. Setting $b_0 = 1 + \alpha_0 + p + 1$, we have that the first part in the last expression is the kernel of a multivariate normal density with mean $\boldsymbol{\mu}_0 = \mathbf{a}_{02}/b_0$ and covariance $\frac{\boldsymbol{\Sigma}}{b_0}$, that is, $\boldsymbol{\mu}|\boldsymbol{\Sigma} \sim N_p\left(\boldsymbol{\mu}_0, \frac{\boldsymbol{\Sigma}}{\beta_0}\right)$, $b_0 = \beta_0$. It makes sense these hyperparameters because \mathbf{a}_{02} is the hypothetical sum of prior observations and b_0 is the hypothetical prior sample size. In addition, the second expression in the last line is the kernel of a inverse Wishart distribution with scale matrix $\boldsymbol{\Psi}_0 = \left(\mathbf{A}_{01} - \frac{\mathbf{a}_{02}\mathbf{a}_{02}}{b_0}\right)$ and α_0 degrees of freedom, that is, $\boldsymbol{\Sigma} \sim IW_p(\boldsymbol{\Psi}_0, \alpha_0)$. Observe that $\boldsymbol{\Psi}_0$ has the same structure as the first part of the sufficient statistics in $T(\mathbf{y})$, just that it should be understood as coming from prior hypothetical observations.

Therefore, the prior distribution in this setting is normal/inverse-Wishart, and given conjugacy, the posterior distribution is in the same family.

$$\pi(\boldsymbol{\mu}, \boldsymbol{\Sigma} | \mathbf{Y}) \propto (2\pi)^{-pN/2} |\boldsymbol{\Sigma}|^{-N/2} \exp\left\{-\frac{1}{2} tr \left[\left(\mathbf{S} + N \left(\boldsymbol{\mu} - \hat{\boldsymbol{\mu}}\right) \left(\boldsymbol{\mu} - \hat{\boldsymbol{\mu}}\right)^{\top}\right) \boldsymbol{\Sigma}^{-1}\right]\right\}$$

$$\times |\boldsymbol{\Sigma}|^{-1/2} \exp\left\{-\frac{\beta_0}{2} tr \left[\left(\boldsymbol{\mu} - \boldsymbol{\mu}_0\right) (\boldsymbol{\mu} - \boldsymbol{\mu}_0)^{\top} \boldsymbol{\Sigma}^{-1}\right]\right\} |\boldsymbol{\Sigma}|^{-(\alpha_0 + p + 1)/2}$$

$$\times \exp\left\{-\frac{1}{2} tr(\boldsymbol{\Psi}_0 \boldsymbol{\Sigma}^{-1})\right\}.$$

Taking into account that

$$N(\mu - \hat{\mu})(\mu - \hat{\mu})^{\top} + \beta_0(\mu - \mu_0)(\mu - \mu_0)^{\top} = (N + \beta_0)(\mu - \mu_n)(\mu - \mu_n)^{\top} + \frac{N\beta_0}{N + \beta_0}(\hat{\mu} - \mu_0)(\hat{\mu} - \mu_0)^{\top},$$

where $\mu_n=rac{N}{N+eta_0}\hat{m{\mu}}+rac{eta_0}{N+eta_0}m{\mu}_0$ is the posterior mean. We have

$$\pi(\boldsymbol{\mu}, \boldsymbol{\Sigma} | \mathbf{Y}) \propto |\boldsymbol{\Sigma}|^{-1/2} \exp\left\{-\frac{N+\beta_0}{2} tr\left[\left((\boldsymbol{\mu} - \boldsymbol{\mu}_n) (\boldsymbol{\mu} - \boldsymbol{\mu}_n)^{\top}\right) \boldsymbol{\Sigma}^{-1}\right]\right\} \times |\boldsymbol{\Sigma}|^{-(N+\alpha_0+p+1)/2} \times \exp\left\{-\frac{1}{2} tr\left[\left(\boldsymbol{\Psi}_0 + \mathbf{S} + \frac{N\beta_0}{N+\beta_0} (\hat{\boldsymbol{\mu}} - \boldsymbol{\mu}_0)(\hat{\boldsymbol{\mu}} - \boldsymbol{\mu}_0)^{\top}\right) \boldsymbol{\Sigma}^{-1}\right]\right\}.$$

Then,
$$\boldsymbol{\mu}|\boldsymbol{\Sigma}, \mathbf{Y} \sim N_p\left(\boldsymbol{\mu}_n, \frac{1}{\beta_n}\boldsymbol{\Sigma}\right)$$
, and $\boldsymbol{\Sigma}|\mathbf{Y} \sim IW\left(\boldsymbol{\Psi}_n, \alpha_n\right)$ where $\beta_n = N + \beta_0$, $\alpha_n = N + \alpha_0$ and $\boldsymbol{\Psi}_n = \boldsymbol{\Psi}_0 + \mathbf{S} + \frac{N\beta_0}{N + \beta_0}(\hat{\boldsymbol{\mu}} - \boldsymbol{\mu}_0)(\hat{\boldsymbol{\mu}} - \boldsymbol{\mu}_0)^{\top}$.

The marginal posterior of μ is given by $\int_{\mathcal{S}} \pi(\mu, \Sigma) d\Sigma$ where \mathcal{S} is the space of positive semi-definite matrices. Then,

$$\pi(\boldsymbol{\mu}|\mathbf{Y}) \propto \int_{\mathcal{S}} \left\{ |\mathbf{\Sigma}|^{-(\alpha_n + p + 2)/2} \right\} d\mathbf{\Sigma}$$

$$\exp \left\{ -\frac{1}{2} tr \left[\left(\beta_n \left(\boldsymbol{\mu} - \boldsymbol{\mu}_n \right) \left(\boldsymbol{\mu} - \boldsymbol{\mu}_n \right)^\top + \boldsymbol{\Psi}_n \right) \boldsymbol{\Sigma}^{-1} \right] \right\} d\mathbf{\Sigma}$$

$$\propto \left| \left(\beta_n \left(\boldsymbol{\mu} - \boldsymbol{\mu}_n \right) \left(\boldsymbol{\mu} - \boldsymbol{\mu}_n \right)^\top + \boldsymbol{\Psi}_n \right) \right|^{-(\alpha_n + 1)/2}$$

$$= \left[\left| \boldsymbol{\Psi}_n \right| \times \left| 1 + \beta_n \left(\boldsymbol{\mu} - \boldsymbol{\mu}_n \right)^\top \boldsymbol{\Psi}_n^{-1} \left(\boldsymbol{\mu} - \boldsymbol{\mu}_n \right) \right| \right]^{-(\alpha_n + 1)/2}$$

$$\propto \left(1 + \frac{1}{\alpha_n + 1 - p} \left(\boldsymbol{\mu} - \boldsymbol{\mu}_n \right)^\top \left(\frac{\boldsymbol{\Psi}_n}{(\alpha_n + 1 - p)\beta_n} \right)^{-1} (\boldsymbol{\mu} - \boldsymbol{\mu}_n) \right)^{-(\alpha_n + 1 - p + p)/2}$$

where the second line uses properties of the inverse Wishart distribution, and the third line uses a particular case of the Sylvester's determinant theorem.

We observe that the last line is the kernel of a multivariate t distribution, that is, $\boldsymbol{\mu}|\mathbf{Y} \sim t_p(v_n, \boldsymbol{\mu}_n, \boldsymbol{\Sigma}_n)$ where $v_n = \alpha_n + 1 - p$ and $\boldsymbol{\Sigma}_n = \frac{\Psi_n}{(\alpha_n + 1 - p)\beta_n}$.

The marginal likelihood is given by

$$p(\mathbf{Y}) = \frac{\Gamma_p \left(\frac{v_n}{2}\right)}{\Gamma_n \left(\frac{\alpha_0}{2}\right)} \frac{|\mathbf{\Psi}_0|^{\alpha_0/2}}{|\mathbf{\Psi}_n|^{\alpha_n/2}} \left(\frac{\beta_0}{\beta_n}\right)^{p/2} (2\pi)^{-Np/2},$$

where Γ_p is the multivariate gamma function (see Exercise 5).

The posterior predictive distribution is $\mathbf{Y}_0|\mathbf{Y} \sim t_p(v_n, \boldsymbol{\mu}_n, (\beta_n + 1)\boldsymbol{\Sigma}_n)$ (see Exercise 6).

Example: Tangency portfolio of US tech stocks

The tangency portfolio is the portfolio that maximizes the Sharpe ratio, where this is the excess of return of a portfolio standardized by its risk.

We want to find the shares **w** of a portfolio that maximizes the Sharpe ratio, where $\mu_{i,T+\kappa} = \mathbb{E}(R_{i,T+\kappa} - R_{f,T+\kappa} \mid \mathcal{I}_T)$, $R_{i,T+\kappa}$ and $R_{f,T+\kappa}$ are the returns of stock i and a risk-free asset. Observe that we have the expected value at period $T + \kappa$ of the excess return conditional on information up to $T(\mathcal{I}_T)$, and $\Sigma_{T+\kappa}$ is the covariance of the excess returns, which is a measure of risk. In particular,

$$\underset{\mathbf{w} \in \mathbb{R}^p}{\arg\max} \frac{\mathbf{w}^{\top} \boldsymbol{\mu}_{T+\kappa}}{\sqrt{\mathbf{w}^{\top} \boldsymbol{\Sigma}_{T+\kappa} \mathbf{w}}}; \quad \text{s.t.} \quad \mathbf{w}^{\top} \mathbf{1} = 1,$$

where the solution is

$$\mathbf{w}^* = rac{\mathbf{\Sigma}_{T+\kappa}^{-1} oldsymbol{\mu}_{T+\kappa}}{\mathbf{1}^{ op} \mathbf{\Sigma}_{T+\kappa}^{-1} oldsymbol{\mu}_{T+\kappa}}.$$

If we want to find the optimal portfolio for the next period under the assumption that the excess of returns follow a multivariate normal distribution, which is a common assumption in these applications, we can set $\kappa=1$, and use the predictive distribution of the excess of returns such that $\mu_{T+1}=\mu_n$ and $\Sigma_{T+1}=\frac{v_n}{v_n-2}(\beta_n+1)\Sigma_n$ given the previous predictive result.

We apply this framework to ten tech stocks of the US market between January first, 2021, and September ninth, 2022. In particular, we use information from Yahoo Finance for Apple (AAPL), Netflix (NFLX), Amazon (AMZN), Microsoft (MSFT), Google (GOOG), Meta (META), Tesla (TSLA), NVIDIA Corporation (NVDA), Intel (INTC), and PayPal (PYPL).

R code. Optimal tangency portfolio: Tech shares

```
1 library(quantmod)
2 library(xts)
3 library(ggplot2)
4 library(gridExtra)
5 # grid.arrange
6 graphics.off()
7 rm(list=ls())
8 # Data Range
9 sdate <- as.Date("2021-01-01")</pre>
10 edate <- as.Date("2022-09-30")</pre>
11 Date \leftarrow seq(sdate, edate, by = "day")
13 p <- length(tickers)</pre>
# AAPL: Apple, NFLX: Netflix, AMZN: Amazon,
# MSFT: Microsoft, GOOG: Google, META: Meta,
16 # TSLA: Tesla, NVDA: NVIDIA Corporation
17 # INTC: Intel, PYPL: PayPal
_{\rm 18} ss_stock <- getSymbols(tickers, from=sdate, to=edate, auto.
      assign = T)
19 ss_stock <- purrr::map(tickers,function(x) Ad(get(x)))</pre>
20 ss_stock <- as.data.frame(purrr::reduce(ss_stock, merge))</pre>
21 colnames(ss_stock) <- tickers</pre>
22 # This is to get stock prices
23 ss_rtn <- as.data.frame(apply(ss_stock, 2, function(x) {diff
      (log(x), 1)))
24 # Daily returns
25 t10yr <- getSymbols(Symbols = "DGS10", src = "FRED", from=
      sdate, to=edate, auto.assign = F)
_{\rm 26} # To get 10-Year US Treasury yield data from the
27 Federal Reserve Electronic Database (FRED)
28 t10yrd \leftarrow (1 + t10yr/100)^(1/365) -1
29 # Daily returns
30 t10yrd <- t10yrd[row.names(ss_rtn)]</pre>
31 Exc_rtn <- as.matrix(ss_rtn) - kronecker(t(rep(1, p)), as.</pre>
      matrix(t10yrd))
32 # Excesses of return
33 df <- as.data.frame(Exc_rtn)
34 df$Date <- as.Date(rownames(df))</pre>
35 # Get months
36 df $ Month <- months (df $ Date)
37 # Get years
38 df$Year <- format(df$Date, format="%y")
39 # Aggregate on months and year and get mean
40 Data <- sapply(1:p, function(i) {
   aggregate(df[, i] ~ Month + Year, df, mean)})
42 DataExcRtn <- matrix(0, length(Data[, 1] $Month), p)
43 for(i in 1:p){
    DataExcRtn[, i] <- as.numeric(Data[, i]$'df[, i]')</pre>
45 }
46 colnames (DataExcRtn) <- tickers
47 head(DataExcRtn)
```

R code. Optimal tangency portfolio: Tech shares

```
# Hyperparameters #
  N <- dim(DataExcRtn)[1]
  mu0 \leftarrow rep(0, p)
  beta0 <- 1
 Psi0 <- 100 * diag(p)
6 \text{ alpha0} \leftarrow p + 2
  # Posterior parameters #
  alphan <- N + alpha0
9 vn <- alphan + 1 - p
10 muhat <- colMeans(DataExcRtn)</pre>
nun \leftarrow N/(N + beta0) * muhat + beta0/(N + beta0) * mu0
12 S <- t(DataExcRtn - rep(1, N)%*%t(muhat))%*%(DataExcRtn -</pre>
       rep(1, N) %*%t(muhat))
13 Psin \leftarrow Psi0 + S + N*beta0/(N + beta0)*(muhat - mu0)%*%t(
      muhat - mu0)
14 betan <- N + beta0
15 Sigman <- Psin/((alphan + 1 - p)*betan)
16 Covarn <- (Sigman * (1 + betan)) * vn / (vn - 2)
17 Covari <- solve(Covarn)
18 OptShare <- t(Covari%*%mun/as.numeric((t(rep(1, p))%*%Covari</pre>
      %*%mun)))
19 colnames(OptShare) <- tickers</pre>
20 OptShare
    AAPL NFLX
                 AMZN
                          GOOG INTC META MSFT
      PYPL
  -0.019 0.248 0.102 -0.034 0.173 0.23 -0.022 -0.016 0.035
```

We find that the optimal tangency portfolio is composed by 24.8%, 10.2%, 17.3%, 23%, 3.5% and 30.1% weights of Netflix, Amazon, Intel, Meta, NVIDIA and PayPal, and -1.9%, -3.4%, -2.2% and -1.6% weights of Apple, Google, Microsoft and Tesla. A negative weight means being short in financial jargon, that is, borrowing a stock to sell it.

${\bf 4.3 \quad Linear \; regression: The \; conjugate \; normal-normal/inverse \\ \; gamma \; model }$

In this setting we analyze the conjugate normal-normal/inverse gamma model which is the workhorse in econometrics. In this model, the dependent variable

 y_i is related to a set of regressors $\mathbf{x}_i = (x_{i1}, x_{i2}, \dots, x_{iK})^{\top}$ in a linear way, that is, $y_i = \beta_1 x_{i1} + \beta_2 x_{i2} + \dots + \beta_K x_{iK} + \mu_i = \mathbf{x}_i^{\top} \beta + \mu_i$ where $\boldsymbol{\beta} = (\beta_1, \beta_2, \dots, \beta_K)^{\top}$ and $\mu_i \stackrel{iid}{\sim} N(0, \sigma^2)$ is an stochastic error such that $\mathbb{E}[\mu_i | \mathbf{x}_i] = 0$.

Defining
$$\mathbf{y} = \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_N \end{bmatrix}$$
, $\mathbf{X} = \begin{bmatrix} x_{11} & x_{12} & \dots & x_{1K} \\ x_{21} & x_{22} & \dots & x_{2K} \\ \vdots & \vdots & \vdots & \vdots \\ x_{N1} & x_{N2} & \dots & x_{NK} \end{bmatrix}$ and $\boldsymbol{\mu} = \begin{bmatrix} \mu_1 \\ \mu_2 \\ \vdots \\ \mu_N \end{bmatrix}$,

we can write the model in matrix form: $\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \mu$, where $\mu \sim N(\mathbf{0}, \sigma^2 \mathbf{I})$. This implies that $\mathbf{y} \sim N(\mathbf{X}\boldsymbol{\beta}, \sigma^2 \mathbf{I})$. Then, the likelihood function is

$$\begin{split} p(\mathbf{y}|\boldsymbol{\beta}, \sigma^2, \mathbf{X}) &= (2\pi\sigma^2)^{-\frac{N}{2}} \exp\left\{-\frac{1}{2\sigma^2}(\mathbf{y} - \mathbf{X}\boldsymbol{\beta})^\top (\mathbf{y} - \mathbf{X}\boldsymbol{\beta})\right\} \\ &\propto (\sigma^2)^{-\frac{N}{2}} \exp\left\{-\frac{1}{2\sigma^2}(\mathbf{y} - \mathbf{X}\boldsymbol{\beta})^\top (\mathbf{y} - \mathbf{X}\boldsymbol{\beta})\right\}. \end{split}$$

The conjugate priors for the parameters are

$$\beta | \sigma^2 \sim N(\beta_0, \sigma^2 \mathbf{B}_0),$$

 $\sigma^2 \sim IG(\alpha_0/2, \delta_0/2).$

Then, the posterior distribution is

$$\pi(\boldsymbol{\beta}, \sigma^{2}|\mathbf{y}, \mathbf{X}) \propto (\sigma^{2})^{-\frac{N}{2}} \exp\left\{-\frac{1}{2\sigma^{2}}(\mathbf{y} - \mathbf{X}\boldsymbol{\beta})^{\top}(\mathbf{y} - \mathbf{X}\boldsymbol{\beta})\right\}$$

$$\times (\sigma^{2})^{-\frac{K}{2}} \exp\left\{-\frac{1}{2\sigma^{2}}(\boldsymbol{\beta} - \boldsymbol{\beta}_{0})^{\top}\mathbf{B}_{0}^{-1}(\boldsymbol{\beta} - \boldsymbol{\beta}_{0})\right\}$$

$$\times \frac{(\delta_{0}/2)^{\alpha_{0}/2}}{\Gamma(\alpha_{0}/2)} \left(\frac{1}{\sigma^{2}}\right)^{\alpha_{0}/2+1} \exp\left\{-\frac{\delta_{0}}{2\sigma^{2}}\right\}$$

$$\propto (\sigma^{2})^{-\frac{K}{2}} \exp\left\{-\frac{1}{2\sigma^{2}}[\boldsymbol{\beta}^{\top}(\mathbf{B}_{0}^{-1} + \mathbf{X}^{\top}\mathbf{X})\boldsymbol{\beta} - 2\boldsymbol{\beta}^{\top}(\mathbf{B}_{0}^{-1}\boldsymbol{\beta}_{0} + \mathbf{X}^{\top}\mathbf{X}\hat{\boldsymbol{\beta}})]\right\}$$

$$\times \left(\frac{1}{\sigma^{2}}\right)^{(\alpha_{0}+N)/2+1} \exp\left\{-\frac{\delta_{0} + \mathbf{y}^{\top}\mathbf{y} + \boldsymbol{\beta}_{0}^{\top}\mathbf{B}_{0}^{-1}\boldsymbol{\beta}_{0}}{2\sigma^{2}}\right\},$$

where $\hat{\boldsymbol{\beta}} = (\mathbf{X}^{\top}\mathbf{X})^{-1}\mathbf{X}^{\top}\mathbf{y}$ is the maximum likelihood estimator. Adding and subtracting $\boldsymbol{\beta}_n^{\top}\mathbf{B}_n^{-1}\boldsymbol{\beta}_n$ to complete the square, where $\mathbf{B}_n = (\mathbf{B}_0^{-1} + \mathbf{X}^{\top}\mathbf{X})^{-1}$ and $\boldsymbol{\beta}_n = \mathbf{B}_n(\mathbf{B}_0^{-1}\boldsymbol{\beta}_0 + \mathbf{X}^{\top}\mathbf{X}\hat{\boldsymbol{\beta}})$,

$$\pi(\boldsymbol{\beta}, \sigma^{2} | \mathbf{y}, \mathbf{X}) \propto \underbrace{(\sigma^{2})^{-\frac{K}{2}} \exp\left\{-\frac{1}{2\sigma^{2}} (\boldsymbol{\beta} - \boldsymbol{\beta}_{n})^{\top} \mathbf{B}_{n}^{-1} (\boldsymbol{\beta} - \boldsymbol{\beta}_{n})\right\}}_{1} \times \underbrace{(\sigma^{2})^{-\left(\frac{\alpha_{n}}{2} + 1\right)} \exp\left\{-\frac{\delta_{n}}{2\sigma^{2}}\right\}}_{2}.$$

The first expression is the kernel of a normal density function, $\boldsymbol{\beta}|\sigma^2, \mathbf{y}, \mathbf{X} \sim N(\boldsymbol{\beta}_n, \sigma^2 \mathbf{B}_n)$. The second expression is the kernel of a inverse gamma density, $\sigma^2|\mathbf{y}, \mathbf{X} \sim IG(\alpha_n/2, \delta_n/2)$, where $\alpha_n = \alpha_0 + N$ and $\delta_n = \delta_0 + \mathbf{y}^{\mathsf{T}}\mathbf{y} + \boldsymbol{\beta}_0^{\mathsf{T}} \mathbf{B}_0^{-1} \boldsymbol{\beta}_0 - \boldsymbol{\beta}_n^{\mathsf{T}} \mathbf{B}_n^{-1} \boldsymbol{\beta}_n$.

Taking into account that

$$\begin{split} \boldsymbol{\beta}_n &= (\mathbf{B}_0^{-1} + \mathbf{X}^\top \mathbf{X})^{-1} (\mathbf{B}_0^{-1} \boldsymbol{\beta}_0 + \mathbf{X}^\top \mathbf{X} \hat{\boldsymbol{\beta}}) \\ &= (\mathbf{B}_0^{-1} + \mathbf{X}^\top \mathbf{X})^{-1} \mathbf{B}_0^{-1} \boldsymbol{\beta}_0 + (\mathbf{B}_0^{-1} + \mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top \mathbf{X} \hat{\boldsymbol{\beta}}, \end{split}$$

where $(\mathbf{B}_0^{-1} + \mathbf{X}^{\top}\mathbf{X})^{-1}\mathbf{B}_0^{-1} = \mathbf{I}_{\mathbf{K}} - (\mathbf{B}_0^{-1} + \mathbf{X}^{\top}\mathbf{X})^{-1}\mathbf{X}^{\top}\mathbf{X}$ [73]. Setting $\mathbf{W} = (\mathbf{B}_0^{-1} + \mathbf{X}^{\top}\mathbf{X})^{-1}\mathbf{X}^{\top}\mathbf{X}$ we have $\boldsymbol{\beta}_n = (\mathbf{I}_{\mathbf{K}} - \mathbf{W})\boldsymbol{\beta}_0 + \mathbf{W}\boldsymbol{\hat{\beta}}$, that is, the posterior mean of $\boldsymbol{\beta}$ is a weighted average between the sample and prior information, where the weights depend on the precision of each piece of information. Observe that when the prior covariance matrix is highly vague (non-informative), such that $\mathbf{B}_0^{-1} \to \mathbf{0}_{\mathbf{K}}$, we obtain $\mathbf{W} \to I_K$, such that $\boldsymbol{\beta}_n \to \hat{\boldsymbol{\beta}}$, that is, the posterior mean location parameter converges to the maximum likelihood estimator.

In addition, we know that the posterior conditional covariance matrix of the location parameters $\sigma^2(\mathbf{B}_0^{-1} + \mathbf{X}^\top \mathbf{X})^{-1} = \sigma^2(\mathbf{X}^\top \mathbf{X})^{-1} - \sigma^2((\mathbf{X}^\top \mathbf{X})^{-1}(\mathbf{B}_0 + (\mathbf{X}^\top \mathbf{X})^{-1})^{-1}(\mathbf{X}^\top \mathbf{X})^{-1})$ is positive semi-definite.⁴ Given that $\sigma^2(\mathbf{X}^\top \mathbf{X})^{-1}$ is the covariance matrix of the maximum likelihood estimator, we observe that prior information reduces estimation uncertainty.

Now, we calculate the posterior marginal distribution of β ,

$$\pi(\boldsymbol{\beta}|\mathbf{y}, \mathbf{X}) = \int_0^\infty \pi(\boldsymbol{\beta}, \sigma^2|\mathbf{y}, \mathbf{X}) d\sigma^2$$
$$= \int_0^\infty \left(\frac{1}{\sigma^2}\right)^{\frac{\alpha_n + K}{2} + 1} \exp\left\{-\frac{s}{2\sigma^2}\right\} d\sigma^2,$$

where $s = \delta_n + (\boldsymbol{\beta} - \boldsymbol{\beta}_n)^{\top} \mathbf{B}_n^{-1} (\boldsymbol{\beta} - \boldsymbol{\beta}_n)$. Then we can write

$$\begin{split} \pi(\boldsymbol{\beta}|\mathbf{y},\mathbf{X}) &= \int_0^\infty \left(\frac{1}{\sigma^2}\right)^{\frac{\alpha_n+K}{2}+1} \exp\left\{-\frac{s}{2\sigma^2}\right\} d\sigma^2 \\ &= \frac{\Gamma((\alpha_n+K)/2)}{(s/2)^{(\alpha_n+K)/2}} \int_0^\infty \frac{(s/2)^{(\alpha_n+K)/2}}{\Gamma((\alpha_n+K)/2)} (\sigma^2)^{-(\alpha_n+K)/2-1} \exp\left\{-\frac{s}{2\sigma^2}\right\} d\sigma^2. \end{split}$$

⁴A particular case of the Woodbury matrix identity.

The right term is the integral of the probability density function of an inverse gamma distribution with parameters $\nu = (\alpha_n + K)/2$ and $\tau = s/2$. Since we are integrating over the whole support of σ^2 , the integral is equal to 1, and therefore

$$\pi(\boldsymbol{\beta}|\mathbf{y}, \mathbf{X}) = \frac{\Gamma((\alpha_n + K)/2)}{(s/2)^{(\alpha_n + K)/2}}$$

$$\propto s^{-(\alpha_n + K)/2}$$

$$= [\delta_n + (\boldsymbol{\beta} - \boldsymbol{\beta}_n)^{\top} \mathbf{B}_n^{-1} (\boldsymbol{\beta} - \boldsymbol{\beta}_n)]^{-(\alpha_n + K)/2}$$

$$= \left[1 + \frac{(\boldsymbol{\beta} - \boldsymbol{\beta}_n)^{\top} \left(\frac{\delta_n}{\alpha_n} \mathbf{B}_n\right)^{-1} (\boldsymbol{\beta} - \boldsymbol{\beta}_n)}{\alpha_n}\right]^{-(\alpha_n + K)/2}$$

$$\propto \left[1 + \frac{(\boldsymbol{\beta} - \boldsymbol{\beta}_n)^{\top} \mathbf{H}_n^{-1} (\boldsymbol{\beta} - \boldsymbol{\beta}_n)}{\alpha_n}\right]^{-(\alpha_n + K)/2},$$

where $\mathbf{H}_n = \frac{\delta_n}{\alpha_n} \mathbf{B}_n$. This last expression is a multivariate t distribution for $\boldsymbol{\beta}$, $\boldsymbol{\beta} | \mathbf{y}, \mathbf{X} \sim t_K(\alpha_n, \boldsymbol{\beta}_n, \mathbf{H}_n)$.

Observe that as we have incorporated the uncertainty of the variance, the posterior for β changes from a normal to a t distribution, which has heavier tails, indicating more uncertainty.

The marginal likelihood of this model is

$$p(\mathbf{y}) = \int_0^\infty \int_{R^K} \pi(\boldsymbol{\beta}|\sigma^2, \mathbf{B}_0, \boldsymbol{\beta}_0) \pi(\sigma^2|\alpha_0/2, \delta_0/2) p(\mathbf{y}|\boldsymbol{\beta}, \sigma^2, \mathbf{X}) d\sigma^2 d\boldsymbol{\beta}.$$

Taking into account that $(\mathbf{y} - \mathbf{X}\boldsymbol{\beta})^{\top}(\mathbf{y} - \mathbf{X}\boldsymbol{\beta}) + (\boldsymbol{\beta} - \boldsymbol{\beta}_0)^{\top}\mathbf{B}_0^{-1}(\boldsymbol{\beta} - \boldsymbol{\beta}_0) = (\boldsymbol{\beta} - \boldsymbol{\beta}_n)^{\top}\mathbf{B}_n^{-1}(\boldsymbol{\beta} - \boldsymbol{\beta}_n) + m$, where $m = \mathbf{y}^{\top}\mathbf{y} + \boldsymbol{\beta}_0^{\top}\mathbf{B}_0^{-1}\boldsymbol{\beta}_0 - \boldsymbol{\beta}_n^{\top}\mathbf{B}_n^{-1}\boldsymbol{\beta}_n$, we have that

$$\begin{split} p(\mathbf{y}) &= \int_{0}^{\infty} \int_{R^{K}} \pi(\beta | \sigma^{2}) \pi(\sigma^{2}) p(\mathbf{y} | \beta, \sigma^{2}, \mathbf{X}) d\sigma^{2} d\beta \\ &= \int_{0}^{\infty} \pi(\sigma^{2}) \frac{1}{(2\pi\sigma^{2})^{N/2}} \exp\left\{-\frac{1}{2\sigma^{2}}m\right\} \frac{1}{(2\pi\sigma^{2})^{K/2} |\mathbf{B}_{0}|^{1/2}} \\ &\times \int_{R^{K}} \exp\left\{-\frac{1}{2\sigma^{2}} (\beta - \beta_{n})^{\top} \mathbf{B}_{n}^{-1} (\beta - \beta_{n})\right\} d\sigma^{2} d\beta \\ &= \int_{0}^{\infty} \pi(\sigma^{2}) \frac{1}{(2\pi\sigma^{2})^{N/2}} \exp\left\{-\frac{1}{2\sigma^{2}}m\right\} \frac{|\mathbf{B}_{n}|^{1/2}}{|\mathbf{B}_{0}|^{1/2}} d\sigma^{2} \\ &= \int_{0}^{\infty} \frac{(\delta_{0}/2)^{\alpha_{0}/2}}{\Gamma(\alpha_{0}/2)} \left(\frac{1}{\sigma^{2}}\right)^{\alpha_{0}/2+1} \exp\left\{\left(-\frac{\delta_{0}}{2\sigma^{2}}\right)\right\} \frac{1}{(2\pi\sigma^{2})^{N/2}} \exp\left\{-\frac{1}{2\sigma^{2}}m\right\} \frac{|\mathbf{B}_{n}|^{1/2}}{|\mathbf{B}_{0}|^{1/2}} d\sigma^{2} \\ &= \frac{1}{(2\pi)^{N/2}} \frac{(\delta_{0}/2)^{\alpha_{0}/2}}{\Gamma(\alpha_{0}/2)} \frac{|\mathbf{B}_{n}|^{1/2}}{|\mathbf{B}_{0}|^{1/2}} \int_{0}^{\infty} \left(\frac{1}{\sigma^{2}}\right)^{\frac{\alpha_{0}+N}{2}+1} \exp\left\{\left(-\frac{\delta_{0}+m}{2\sigma^{2}}\right)\right\} d\sigma^{2} \\ &= \frac{1}{\pi^{N/2}} \frac{\delta_{0}^{\alpha_{0}/2}}{\delta_{n}^{\alpha_{0}/2}} \frac{|\mathbf{B}_{n}|^{1/2}}{|\mathbf{B}_{0}|^{1/2}} \frac{\Gamma(\alpha_{n}/2)}{\Gamma(\alpha_{0}/2)}. \end{split}$$

We can show that $\delta_n = \delta_0 + \mathbf{y}^{\top} \mathbf{y} + \boldsymbol{\beta}_0^{\top} \mathbf{B}_0^{-1} \boldsymbol{\beta}_0 - \boldsymbol{\beta}_n^{\top} \mathbf{B}_n^{-1} \boldsymbol{\beta}_n = \delta_0 + (\mathbf{y} - \mathbf{X}\hat{\boldsymbol{\beta}})^{\top} (\mathbf{y} - \mathbf{X}\hat{\boldsymbol{\beta}}) + (\hat{\boldsymbol{\beta}} - \boldsymbol{\beta}_0)^{\top} ((\mathbf{X}^{\top} \mathbf{X})^{-1} + \mathbf{B}_0)^{-1} (\hat{\boldsymbol{\beta}} - \boldsymbol{\beta}_0)$ (see Exercise 7). Therefore, if we want to compare two models under this setting, the Bayes factor is

$$\begin{split} BF_{12} &= \frac{p(\mathbf{y}|\mathcal{M}_1)}{p(\mathbf{y}|\mathcal{M}_2)} \\ &= \frac{\frac{\delta_{10}^{\alpha_{10}/2}}{\delta_{1n}^{\alpha_{1n}/2}} \frac{|\mathbf{B}_{1n}|^{1/2}}{|\mathbf{B}_{10}|^{1/2}} \frac{\Gamma(\alpha_{1n}/2)}{\Gamma(\alpha_{10}/2)}}{\frac{\delta_{20}^{\alpha_{20}/2}}{\delta_{2n}^{\alpha_{20}/2}} \frac{|\mathbf{B}_{2n}|^{1/2}}{|\mathbf{B}_{20}|^{1/2}} \frac{\Gamma(\alpha_{2n}/2)}{\Gamma(\alpha_{20}/2)}}, \end{split}$$

Observe that *ceteris paribus*, the model having better fit, coherence be-

where subscripts 1 and 2 refer to each model, respectively.

tween sample and prior information regarding location parameters, higher prior to posterior precision and less parameters is favored by the Bayes factor. Observe that the Bayes factor rewards model fit as the sum of squared errors is in δ_n , the better fit (lower sum of squared errors), the better the Bayes factor. In addition, a weighted distance between sample and prior location parameters also appears in δ_n , the greater this distance, the worse is model support. The ratio of determinants between posterior and prior covariance matrices is also present, the higher this ratio, the better for the Bayes factor supporting a model due to information gains. To see the effect of model's parsimony, let's take the common situation in applications where $\mathbf{B}_{j0} = c\mathbf{I}_{K_j}$ then $|\mathbf{B}_{j0}| = c^{K_j}$ such that $\left(\frac{|\mathbf{B}_{20}|}{|\mathbf{B}_{10}|}\right)^{1/2} = \left(\frac{c^{K_2/2}}{c^{K_1/2}}\right)$, if $K_2/K_1 > 1$ and $c \to \infty$, the latter implying a non-informative prior, then $BF_{12} \to \infty$, this means infinite evidence supporting the parsimonious model no matter what sample information says. Comparing models having the same number of regressors $(K_1 = K_2)$ is not a safe ground as $|\mathbf{B}_0|$ depending on measure units of the regressors such that conclusions regarding model selection depending on this, which is not a nice property. This prevents against using non-informative priors when performing model selection in the Bayesian framework. Observe that this is not the case when $\alpha_0 \to 0$ and $\delta_0 \to 0$, which implies a non-informative prior for the variance parameter.⁵ We observe here that $\Gamma(\alpha_{j0})$ cancels out, $\alpha_{jn} \to N$ and $\delta_{jn} \to (\mathbf{y} - \mathbf{X}_j \hat{\boldsymbol{\beta}}_j)^\top (\mathbf{y} - \mathbf{X}_j \hat{\boldsymbol{\beta}}_j) + (\hat{\boldsymbol{\beta}}_j - \boldsymbol{\beta}_{j0})^\top ((\mathbf{X}_j^\top \mathbf{X}_j)^{-1} + \mathbf{B}_{j0})^{-1} (\hat{\boldsymbol{\beta}}_j - \boldsymbol{\beta}_{j0}),$ therefore there is not effect. This is due to σ^2 being a common parameter in both models. In general, we can use non-informative priors for common parameters to all models, but we cannot use non-informative priors for non-common parameters when performing model selection using the Bayes factor.

The posterior predictive is equal to

⁵[31] prevents against this common practice.

$$\pi(\mathbf{Y}_0|\mathbf{y}) = \int_0^\infty \int_{R^K} p(\mathbf{Y}_0|\boldsymbol{\beta}, \sigma^2, \mathbf{y}) \pi(\boldsymbol{\beta}|\sigma^2, \mathbf{y}) \pi(\sigma^2|\mathbf{y}) d\boldsymbol{\beta} d\sigma^2$$
$$= \int_0^\infty \int_{R^K} p(\mathbf{Y}_0|\boldsymbol{\beta}, \sigma^2) \pi(\boldsymbol{\beta}|\sigma^2, \mathbf{y}) \pi(\sigma^2|\mathbf{y}) d\boldsymbol{\beta} d\sigma^2,$$

where we take into account independence between \mathbf{Y}_0 and \mathbf{Y} . Given \mathbf{X}_0 , which is the $N_0 \times K$ matrix of regressors associated with \mathbf{Y}_0 , Then,

$$\begin{split} \pi(\mathbf{Y}_0|\mathbf{y}) &= \int_0^\infty \int_{R^K} \left\{ (2\pi\sigma^2)^{-\frac{N_0}{2}} \exp\left\{ -\frac{1}{2\sigma^2} (\mathbf{Y}_0 - \mathbf{X}_0 \boldsymbol{\beta})^\top (\mathbf{Y}_0 - \mathbf{X}_0 \boldsymbol{\beta})^\top \right\} \\ &\times (2\pi\sigma^2)^{-\frac{K}{2}} |\mathbf{B}_n|^{-1/2} \exp\left\{ -\frac{1}{2\sigma^2} (\boldsymbol{\beta} - \boldsymbol{\beta}_n)^\top \mathbf{B}_n^{-1} (\boldsymbol{\beta} - \boldsymbol{\beta}_n) \right\} \\ &\times \frac{(\delta_n/2)^{\alpha_n/2}}{\Gamma(\alpha_n/2)} \left(\frac{1}{\sigma^2} \right)^{\alpha_n/2+1} \exp\left\{ -\frac{\delta_n}{2\sigma^2} \right\} \right\} d\boldsymbol{\beta} d\sigma^2. \end{split}$$

Setting
$$\mathbf{M} = (\mathbf{X}_0^{\top} \mathbf{X}_0 + \mathbf{B}_n^{-1})$$
 and $\boldsymbol{\beta}_* = \mathbf{M}^{-1}(\mathbf{B}_n^{-1}\boldsymbol{\beta}_n + \mathbf{X}_0^{\top}\mathbf{Y}_0)$, we have $(\mathbf{Y}_0 - \mathbf{X}_0\boldsymbol{\beta})^{\top}(\mathbf{Y}_0 - \mathbf{X}_0\boldsymbol{\beta})^{\top} + (\boldsymbol{\beta} - \boldsymbol{\beta}_n)^{\top}\mathbf{B}_n^{-1}(\boldsymbol{\beta} - \boldsymbol{\beta}_n) = (\boldsymbol{\beta} - \boldsymbol{\beta}_*)^{\top}\mathbf{M}(\boldsymbol{\beta} - \boldsymbol{\beta}_*) + \boldsymbol{\beta}_n^{\top}\mathbf{B}_n^{-1}\boldsymbol{\beta}_n + \mathbf{Y}_0^{\top}\mathbf{Y}_0 - \boldsymbol{\beta}_*^{\top}\mathbf{M}\boldsymbol{\beta}_*$. Thus,

$$\pi(\mathbf{Y}_0|\mathbf{y}) \propto \int_0^\infty \left\{ \left(\frac{1}{\sigma^2}\right)^{-\frac{K+N_0+\alpha_n}{2}+1} \exp\left\{-\frac{1}{2\sigma^2}(\boldsymbol{\beta}_n^\top \mathbf{B}_n^{-1} \boldsymbol{\beta}_n + \mathbf{Y}_0^\top \mathbf{Y}_0 - \boldsymbol{\beta}_*^\top \mathbf{M} \boldsymbol{\beta}_* + \delta_n)\right\} \times \int_{R^K} \exp\left\{-\frac{1}{2\sigma^2}(\boldsymbol{\beta} - \boldsymbol{\beta}_*)^\top \mathbf{M} (\boldsymbol{\beta} - \boldsymbol{\beta}_*)\right\} d\boldsymbol{\beta} d\sigma^2,$$

where the term in the second integral is the kernel of a multivariate normal density with mean β_* and covariance matrix $\sigma^2 \mathbf{M}^{-1}$. Then,

$$\pi(\mathbf{Y}_0|\mathbf{y}) \propto \int_0^\infty \left(\frac{1}{\sigma^2}\right)^{\frac{N_0 + \alpha_n}{2} + 1} \exp\left\{-\frac{1}{2\sigma^2}(\boldsymbol{\beta}_n^\top \mathbf{B}_n^{-1} \boldsymbol{\beta}_n + \mathbf{Y}_0^\top \mathbf{Y}_0 - \boldsymbol{\beta}_*^\top \mathbf{M} \boldsymbol{\beta}_* + \delta_n)\right\} d\sigma^2,$$

which is the kernel of an inverse gamma density. Thus,

$$\pi(\mathbf{Y}_0|\mathbf{y}) \propto \left[\frac{\boldsymbol{\beta}_n^{\top} \mathbf{B}_n^{-1} \boldsymbol{\beta}_n + \mathbf{Y}_0^{\top} \mathbf{Y}_0 - \boldsymbol{\beta}_*^{\top} \mathbf{M} \boldsymbol{\beta}_* + \delta_n}{2}\right]^{-\frac{\alpha_n + N_0}{2}}.$$
Setting $\mathbf{C}^{-1} = \mathbf{I}_{N_0} + \mathbf{X}_0 \mathbf{B}_n \mathbf{X}_0^{\top}$ such that $\mathbf{C} = \mathbf{I}_{N_0} - \mathbf{X}_0 (\mathbf{B}_n^{-1} + \mathbf{X}_0 \mathbf{B}_n^{-1} + \mathbf{X}_0 \mathbf{B}_n^{-1})$

$$\begin{split} \mathbf{X}_{0}^{\top}\mathbf{X}_{0})^{-1}\mathbf{X}_{0}^{\top} &= \mathbf{I}_{N_{0}} - \mathbf{X}_{0}\mathbf{M}^{-1}\mathbf{X}_{0}^{\top},^{6} \text{ and } \boldsymbol{\beta}_{**} = \mathbf{C}^{-1}\mathbf{X}_{0}\mathbf{M}^{-1}\mathbf{B}_{n}^{-1}\boldsymbol{\beta}_{n}, \text{ then} \\ \boldsymbol{\beta}_{n}^{\top}\mathbf{B}_{n}^{-1}\boldsymbol{\beta}_{n} + \mathbf{Y}_{0}^{\top}\mathbf{Y}_{0} - \boldsymbol{\beta}_{*}^{\top}\mathbf{M}\boldsymbol{\beta}_{*} &= \boldsymbol{\beta}_{n}^{\top}\mathbf{B}_{n}^{-1}\boldsymbol{\beta}_{n} + \mathbf{Y}_{0}^{\top}\mathbf{Y}_{0} - (\boldsymbol{\beta}_{n}^{\top}\mathbf{B}_{n}^{-1} + \mathbf{Y}_{0}^{\top}\mathbf{X}_{0})\mathbf{M}^{-1}(\mathbf{B}_{n}^{-1}\boldsymbol{\beta}_{n} + \mathbf{X}_{0}^{\top}\mathbf{Y}_{0}) \\ &= \boldsymbol{\beta}_{n}^{\top}(\mathbf{B}_{n}^{-1} - \mathbf{B}_{n}^{-1}\mathbf{M}^{-1}\mathbf{B}_{n}^{-1})\boldsymbol{\beta}_{n} + \mathbf{Y}_{0}^{\top}\mathbf{C}\mathbf{Y}_{0} \\ &- 2\mathbf{Y}_{0}^{\top}\mathbf{C}\mathbf{C}^{-1}\mathbf{X}_{0}\mathbf{M}^{-1}\mathbf{B}_{n}^{-1}\boldsymbol{\beta}_{n} + \boldsymbol{\beta}_{**}^{\top}\mathbf{C}\boldsymbol{\beta}_{**} - \boldsymbol{\beta}_{**}^{\top}\mathbf{C}\boldsymbol{\beta}_{**} \\ &= \boldsymbol{\beta}_{n}^{\top}(\mathbf{B}_{n}^{-1} - \mathbf{B}_{n}^{-1}\mathbf{M}^{-1}\mathbf{B}_{n}^{-1})\boldsymbol{\beta}_{n} + (\mathbf{Y}_{0} - \boldsymbol{\beta}_{**})^{\top}\mathbf{C}(\mathbf{Y}_{0} - \boldsymbol{\beta}_{**}) \\ &- \boldsymbol{\beta}_{**}^{\top}\mathbf{C}\boldsymbol{\beta}_{**}, \end{split}$$

where $\boldsymbol{\beta}_n^{\top}(\mathbf{B}_n^{-1} - \mathbf{B}_n^{-1}\mathbf{M}^{-1}\mathbf{B}_n^{-1})\boldsymbol{\beta}_n = \boldsymbol{\beta}_{**}^{\top}\mathbf{C}\boldsymbol{\beta}_{**}$ and $\boldsymbol{\beta}_{**} = \mathbf{X}_0\boldsymbol{\beta}_n$ (see Exercise 8). Then,

$$\pi(\mathbf{Y}_0|\mathbf{y}) \propto \left[\frac{(\mathbf{Y}_0 - \mathbf{X}_0 \boldsymbol{\beta}_n)^{\top} \mathbf{C} (\mathbf{Y}_0 - \mathbf{X}_0 \boldsymbol{\beta}_n) + \delta_n}{2} \right]^{-\frac{\alpha_n + N_0}{2}}$$
$$\propto \left[\frac{(\mathbf{Y}_0 - \mathbf{X}_0 \boldsymbol{\beta}_n)^{\top} \left(\frac{\mathbf{C} \alpha_n}{\delta_n} \right) (\mathbf{Y}_0 - \mathbf{X}_0 \boldsymbol{\beta}_n)}{\alpha_n} + 1 \right]^{-\frac{\alpha_n + N_0}{2}}$$

The posterior predictive is a multivariate t distribution, $\mathbf{Y}_0|\mathbf{y} \sim t\left(\mathbf{X}_0\boldsymbol{\beta}_n, \frac{\delta_n(\mathbf{I}_{N_0} + \mathbf{X}_0\mathbf{B}_n\mathbf{X}_0^\top)}{\alpha_n}, \alpha_n\right)$ centered at $\mathbf{X}_0\boldsymbol{\beta}_n$.

Example: Demand of electricity

We study in this example the determinants of monthly demand of electricity by Colombian households. There is information of 2103 households, particularly, average price (USD/kWh), indicators of socioeconomic conditions of the neighborhood where the household is located (IndSocio1 is the lowest and IndSocio3 is the highest), an indicator if the household is located in a municipality that is above 1000 meters above the sea level, the number of rooms in the house, the number of members of the households, presence of children in the household (1 is yes), and monthly income (USD). The specification is

log(Electricity_i) =
$$\beta_1$$
 log(price_i) + β_2 IndSocio1_i + β_3 IndSocio2_i + β_4 Altitude_i
+ β_5 Nrooms_i + β_6 HouseholdMem_i + β_7 Children_i
+ β_8 log(Income_i) + β_9 + μ .

We use a non-informative vague prior setting such that $\alpha_0 = \delta_0 = 0.001$, $\boldsymbol{\beta}_0 = \mathbf{0}$ and $\mathbf{B}_0 = c_0 \mathbf{I}_k$, where $c_0 = 1000$ and k is the number of regressors.

The results from the R code (see below) is that the posterior mean of the own-price of electricity demand is -1.09, and the 95% symmetric credible interval is (-1.47, -0.71). Households in neighborhoods of low socioeconomic conditions and located in municipalities 1000 meters above the sea level consume less electricity, 32.7% and 19.7% on average, respectively. An additional

$$6$$
Using $(A + BDC)^{-1} = A^{-1} - A^{-1}B(D^{-1} + CA^{-1}B)^{-1}CA^{-1}$

room implies a 8.7% increase in electricity consumption, and another household member increases consumption in 5.9% on average. The income elasticity mean estimate is 0.074, which means that 10% increase of income increases 0.74% electricity demand.

We want to check the results of the Bayes factor comparing the previous specification (model 1) with other specification without considering the price of electricity (model 2), that is,

```
log(Electricity<sub>i</sub>) = \beta_1IndSocio1<sub>i</sub> + \beta_2IndSocio2<sub>i</sub> + \beta_3Altitude<sub>i</sub> + \beta_4Nrooms<sub>i</sub> + \beta_5HouseholdMem<sub>i</sub> + \beta_6Children<sub>i</sub> + \beta_7 log(Income<sub>i</sub>) + \beta_8 + \mu.
```

In particular, we check what happens as c_0 increases from 10^0 to 10^{20} . We see that when $c_0 = 1$, $BF_{12} = 8.68 \times 10^{+16}$, which means very strong evidence in favor of the model including the price of electricity. However, as c_0 increases, the Bayes factor decreases, which means evidence supporting the model 2, for instance, $BF_{12} = 3.11 \times 10^{-4}$ when $c_0 = 10^{20}$. This is an example of the problem of using non-informative priors to calculate the Bayes factor; there is very strong evidence to support the parsimonious model when $c_0 \to \infty$.

We can get the posterior predictive distribution of the monthly electricity demand of a household located in the lowest socioeconomic condition in a municipality located below 1000 meters above the sea level, 2 rooms, 3 members with children, a monthly income equal to USD 500, and an electricity price equal to USD/kWh 0.15. Figure 4.1 shows the histogram of the predictive posterior distribution, the highest posterior density credible interval at 95% is between kWh 44.4 and kWh 373.9, and the posterior mean is kWh 169.4.

R code. Demand of electricity, posterior predictive distribution

```
1 rm(list = ls())
2 set.seed(010101)
3 # Electricity demand
4 DataUt <- read.csv("DataApplications/Utilities.csv", sep = "
       ,", header = TRUE, fileEncoding = "latin1")
5 DataUtEst <- DataUt %>%
    filter(Electricity != 0)
7 attach(DataUtEst)
8 # Dependent variable: Monthly consumption (kWh) in log
9 Y <- log(Electricity)
10 # Regressors quantity including intercept
11 X <- cbind(LnPriceElect, IndSocio1, IndSocio2, Altitude,
Nrooms, HouseholdMem, Children, Lnincome, 1) _{\rm 12} # LnPriceElect: Price per kWh (USD) in log
13 # IndSocio1, IndSocio2, IndSocio3: Indicators socio-economic
        condition (1) is the lowest and (3) the highest
_{14} # Altitude: Indicator of household location (1 is more than
      1000 meters above sea level)
15 # Nrooms: Number of rooms in house
16 # HouseholdMem: Number of household members
17 # Children: Indicator por presence of children in household
       (1)
18 # Lnincome: Monthly income (USD) in log
19 k <- dim(X)[2]
20 N <- dim(X)[1]
21 # Hyperparameters
22 d0 <- 0.001/2
23 a0 <- 0.001/2
24 b0 <- rep(0, k)
25 B0 <- 1000*diag(k)
26 # Posterior parameters
27 bhat <- solve(t(X)%*%X)%*%t(X)%*%Y
28 Bn <- as.matrix(Matrix::forceSymmetric(solve(solve(B0) + t(X
      )%*\mbox{\ensuremath{\%}X))) # Force this matrix to be symmetric
29 bn \leftarrow Bn%*%(solve(B0)%*%b0 + t(X)%*%X%*%bhat)
30 dn <- as.numeric(d0 + t(Y)%*%Y+t(b0)%*%solve(B0)%*%b0-t(bn)%
      *%solve(Bn)%*%bn)
31 an <- a0 + N
32 Hn <- Bn*dn/an
33 # Posterior draws
_{\rm 34} S <- 10000 # Number of draws from posterior distributions
35 sig2 <- MCMCpack::rinvgamma(S,an/2,dn/2)</pre>
36 summary (coda::mcmc(sig2))
```

R code. Demand of electricity, posterior distribution

```
1 Iterations = 1:10000
2 Thinning interval = 1
3 Number of chains = 1
4 Sample size per chain = 10000
6 1. Empirical mean and standard deviation for each
7 variable, plus standard error of the mean:
                   SD
                            Naive SE Time-series SE
10 2.361e-01
                 7.617e-03 7.617e-05 7.617e-05
12 2. Quantiles for each variable:
13
14 2.5%
          25%
                50%
                       75% 97.5%
15 0.2217 0.2309 0.2360 0.2412 0.2513
17 Betas <- LaplacesDemon::rmvt(S, bn, Hn, an)
18 summary(coda::mcmc(Betas))
19 Iterations = 1:10000
20 Thinning interval = 1
21 Number of chains = 1
22 Sample size per chain = 10000
24 1. Empirical mean and standard deviation for each
25 variable, plus standard error of the mean:
                          SD
                                Naive SE Time-series SE
                Mean
28 LnPriceElect -1.09043 0.19459 0.0019459 0.0019459
               -0.32783 0.05294 0.0005294 0.0005294
29 IndSocio1
30 IndSocio2
               -0.05737 0.04557 0.0004557 0.0004557
31 Altitude
               -0.19780 0.02386 0.0002386 0.0002429
                0.08731 0.01094 0.0001094 0.0001119
32 Nrooms
33 HouseholdMem 0.05987 0.01334 0.0001334 0.0001334
                0.05696 0.03043 0.0003043 0.0003043
34 Children
                0.07447 0.01223 0.0001223 0.0001223
35 Lnincome
                2.52296 0.35077 0.0035077 0.0035077
38 2. Quantiles for each variable:
                    2.5%
                              25%
                                       50%
                                               75%
                                                       97.5%
41 LnPriceElect -1.472069 -1.22432 -1.08961 -0.95703 -0.71429
               42 IndSocio1
43 IndSocio2
               -0.244759 -0.21372 -0.19783 -0.18164 -0.15094
44 Altitude
45 Nrooms
                0.066432 0.07985 0.08709 0.09480 0.10864
46 HouseholdMem 0.033623 0.05089 0.05975
                                           0.06889 0.08596
47 Children
               -0.002259
                         0.03637
                                  0.05698
                                           0.07736
                                                    0.11681
                0.050536 0.06614
48 Lnincome
                                  0.07449
                                           0.08283
                                                    0.09852
                1.835507 2.28703 2.52165 2.76364 3.21199
49
50
```

R code. Demand of electricity, Bayes factor

```
1 # Log marginal function (multiply by -1 due to minimization)
2 LogMarLikLM <- function(X, c0){</pre>
    k \leftarrow dim(X)[2]
    N <- dim(X)[1]
    # Hyperparameters
    B0 \leftarrow c0*diag(k)
    b0 \leftarrow rep(0, k)
     # Posterior parameters
    bhat <- solve(t(X)%*%X)%*%t(X)%*%Y
    # Force this matrix to be symmetric
    Bn <- as.matrix(Matrix::forceSymmetric(solve(solve(B0) + t</pre>
      (X)%*%X)))
     bn <- Bn\%*\%(solve(B0)\%*\%b0 + t(X)\%*\%X\%*\%bhat)
    dn \leftarrow as.numeric(d0 + t(Y))%*%Y+t(b0)%*%solve(B0)%*%b0-t(bn)
      )%*%solve(Bn)%*%bn)
    an \leftarrow a0 + N
14
     # Log marginal likelihood
15
    logpy \leftarrow (N/2)*log(1/pi)+(a0/2)*log(d0)-(an/2)*log(dn) +
      0.5*log(det(Bn)/det(B0)) + lgamma(an/2)-lgamma(a0/2)
    return(-logpy)
18 }
19 cs <- c(10<sup>0</sup>, 10<sup>3</sup>, 10<sup>6</sup>, 10<sup>10</sup>, 10<sup>12</sup>, 10<sup>15</sup>, 10<sup>20</sup>)
20 # Observe -1 to recover the right sign
21 LogML <- sapply(cs, function(c) {-LogMarLikLM(c0=c, X = X)})</pre>
22 # Regressor without price
_{\rm 23} Xnew <- cbind(IndSocio1, IndSocio2, Altitude, Nrooms,
       HouseholdMem, Children, Lnincome, 1)
24 # Observe -1 to recover the right sign
25 LogMLnew <- sapply(cs, function(c) {-LogMarLikLM(c0=c,X =</pre>
       Xnew)})
26 # Bayes factor
27 BF <- exp(LogML - LogMLnew)
28 BF
29 8.687567e+16 1.006679e+05 3.108415e+03 3.108340e+01 3.108343
       e+00 9.829443e-02 3.108343e-04
30 # Empirical Bayes: Obtain c0 maximizing the log
31 marginal likelihood
32 c0 <- c0
33 EB <- optim(c0, fn = LogMarLikLM, method = "Brent", lower =
       0.0001, upper = 10^6, X = X)
34 EB$par
35 3.254822
36 EB$value
37 1404.108
38 EBnew <- optim(c0, fn = LogMarLikLM, method = "Brent", lower
        = 0.0001, upper = 10^6, X = Xnew)
39 EBnew $par
40 10.00597
41 EBnew$value
42 1422.199
43 # Change of order to take into account the -1 in the
       LogMarLikLM function
44 BFEM <- exp(EBnew$value - EB$value)
45 BFEM
46 71897938
```

$R\ code.\ Demand\ of\ electricity,\ predictive\ distribution$

```
Predictive distribution
  Xpred \leftarrow c(log(0.15), 1, 0, 0, 2, 3, 1, log(500), 1)
  Mean <- Xpred%*%bn
4 Hn <- dn*(1+t(Xpred)%*%Bn%*%Xpred)/an
5 ExpKwH <- exp(LaplacesDemon::rmvt(S, Mean, Hn, an))</pre>
6 summary (ExpKwH)
            24.06
  1st Qu.: 121.70
  Median : 169.37
          : 189.60
10 Mean
11 3rd Qu.: 234.19
12 Max.
          :1243.68
13 HDI <- HDInterval::hdi(ExpKwH, credMass = 0.95) # Highest
      posterior density credible interval
14 HDI
15 lower 44.40203
16 upper 373.86494
17 hist(ExpKwH, main = "Histogram: Monthly demand of
      electricity", xlab = "Monthly kWh", col = "blue", breaks
```

Histogram: Monthly demand of electricity



FIGURE 4.1
Histogram using the posterior predictive distribution of electricity demand

4.4 Multivariate linear regression: The conjugate normalnormal/inverse Wishart model

Let's study the multivariate regression setting where there are N-dimensional vectors \mathbf{y}_m , $m=1,2,\ldots,M$ such that $\mathbf{y}_m=\mathbf{X}\boldsymbol{\beta}_m+\mu_m$, \mathbf{X} is the set of common regressors, and μ_m is the N-dimensional vector of stochastic errors for each equation such that $\mathbf{U}=[\mu_1 \ \mu_2 \ \ldots \ \mu_M] \sim MN_{N,M}(\mathbf{0},\mathbf{I}_N,\boldsymbol{\Sigma})$, that is, a matrix variate normal distribution where $\boldsymbol{\Sigma}$ is the covariance matrix of each i-th row of $\mathbf{U}, i=1,2,\ldots,N$, and we are assuming independence between the rows. Then, $vec(\mathbf{U}) \sim N_{N\times M}(\mathbf{0},\boldsymbol{\Sigma}\otimes\mathbf{I_N})$.

This framework can be written in matrix form

$$\underbrace{\begin{bmatrix} y_{11} & y_{12} & \dots & y_{1M} \\ y_{21} & y_{22} & \dots & y_{2M} \\ \vdots & \vdots & \dots & \vdots \\ y_{N1} & y_{N2} & \dots & y_{NM} \end{bmatrix}}_{\mathbf{Y}} = \underbrace{\begin{bmatrix} x_{11} & x_{12} & \dots & x_{1K} \\ x_{21} & x_{22} & \dots & x_{2K} \\ \vdots & \vdots & \dots & \vdots \\ x_{N1} & x_{N2} & \dots & x_{NK} \end{bmatrix}}_{\mathbf{X}} \begin{bmatrix} \beta_{11} & \beta_{12} & \dots & \beta_{1M} \\ \beta_{21} & \beta_{22} & \dots & \beta_{2M} \\ \vdots & \vdots & \dots & \vdots \\ \beta_{K1} & \beta_{K2} & \dots & \beta_{KM} \end{bmatrix}}_{\mathbf{B}} + \underbrace{\begin{bmatrix} \mu_{11} & \mu_{12} & \dots & \mu_{1M} \\ \mu_{21} & \mu_{22} & \dots & \mu_{2M} \\ \vdots & \vdots & \dots & \vdots \\ \mu_{N1} & \mu_{N2} & \dots & \mu_{NM} \end{bmatrix}}_{\mathbf{H}}.$$

Therefore, $\mathbf{Y} \sim N_{N \times M}(\mathbf{XB}, \mathbf{\Sigma} \otimes \mathbf{I_N})^8$

$$\begin{split} p(\mathbf{Y}|\mathbf{B}, \mathbf{\Sigma}, \mathbf{X}) &\propto |\mathbf{\Sigma}|^{-N/2} \exp\left\{-\frac{1}{2} tr\left[(\mathbf{Y} - \mathbf{X}\mathbf{B})^{\top} (\mathbf{Y} - \mathbf{X}\mathbf{B}) \mathbf{\Sigma}^{-1}\right]\right\} \\ &= |\mathbf{\Sigma}|^{-N/2} \exp\left\{-\frac{1}{2} tr\left[\left(\mathbf{S} + (\mathbf{B} - \widehat{\mathbf{B}})^{\top} \mathbf{X}^{\top} \mathbf{X} (\mathbf{B} - \widehat{\mathbf{B}})\right) \mathbf{\Sigma}^{-1}\right]\right\}, \end{split}$$

where $\mathbf{S} = (\mathbf{Y} - \mathbf{X}\widehat{\mathbf{B}})^{\top}(\mathbf{Y} - \mathbf{X}\widehat{\mathbf{B}})$, $\widehat{\mathbf{B}} = (\mathbf{X}^{\top}\mathbf{X})^{-1}\mathbf{X}^{\top}\mathbf{Y}$ (see Exercise 9). The conjugate prior for this models is $\pi(\mathbf{B}, \mathbf{\Sigma}) = \pi(\mathbf{B}|\mathbf{\Sigma})\pi(\mathbf{\Sigma})$ where $\pi(\mathbf{B}|\mathbf{\Sigma}) \sim N_{K \times M}(\mathbf{B}_0, \mathbf{V}_0, \mathbf{\Sigma})$ and $\pi(\mathbf{\Sigma}) \sim IW(\mathbf{\Psi}_0, \alpha_0)$, that is,

 $^{^7}vec$ denotes the vectorization operation, and \otimes denotes the kronecker product.

⁸We can write down the former expression in a more familiar way using vectorization properties, $\underbrace{vec(Y)}_{\mathbf{y}} = \underbrace{(\mathbf{I}_M \otimes \mathbf{X})}_{\mathbf{Z}} \underbrace{vec(\mathbf{B})}_{\boldsymbol{\beta}} + \underbrace{vec(\mathbf{U})}_{\boldsymbol{\mu}}, \text{ where } \mathbf{y} \sim N_{N \times M}(\mathbf{Z}\boldsymbol{\beta}, \boldsymbol{\Sigma} \otimes \mathbf{I_N}).$

$$\pi(\mathbf{B}, \mathbf{\Sigma}) \propto |\mathbf{\Sigma}|^{-K/2} \exp\left\{-\frac{1}{2} tr \left[(\mathbf{B} - \mathbf{B}_0)^{\top} \mathbf{V}_0^{-1} (\mathbf{B} - \mathbf{B}_0) \mathbf{\Sigma}^{-1} \right] \right\}$$
$$\times |\mathbf{\Sigma}|^{-(\alpha_0 + M + 1)/2} \exp\left\{-\frac{1}{2} tr \left[\mathbf{\Psi}_0 \mathbf{\Sigma}^{-1} \right] \right\}.$$

The posterior distribution is given by

$$\begin{split} \pi(\mathbf{B}, \mathbf{\Sigma} | \mathbf{Y}, \mathbf{X}) &\propto p(\mathbf{Y} | \mathbf{B}, \mathbf{\Sigma}, \mathbf{X}) \pi(\mathbf{B} | \mathbf{\Sigma}) \pi(\mathbf{\Sigma}) \\ &\propto |\mathbf{\Sigma}|^{-\frac{N+K+\alpha_0+M+1}{2}} \\ &\times \exp\left\{-\frac{1}{2} tr\left[(\mathbf{\Psi_0} + \mathbf{S} + (\mathbf{B} - \mathbf{B_0})^{\top} \mathbf{V_0^{-1}} (\mathbf{B} - \mathbf{B_0}) \right.\right. \\ &\left. + (\mathbf{B} - \widehat{\mathbf{B}})^{\top} \mathbf{X}^{\top} \mathbf{X} (\mathbf{B} - \widehat{\mathbf{B}})) \mathbf{\Sigma}^{-1}\right]\right\}. \end{split}$$

Completing the squares on ${\bf B}$ and collecting the remaining terms in the bracket yields

$$\mathbf{\Psi}_0 + \mathbf{S} + (\mathbf{B} - \mathbf{B}_0)^{\top} \mathbf{V}_0^{-1} (\mathbf{B} - \mathbf{B}_0) + (\mathbf{B} - \widehat{\mathbf{B}})^{\top} \mathbf{X}^{\top} \mathbf{X} (\mathbf{B} - \widehat{\mathbf{B}}) = (\mathbf{B} - \mathbf{B}_n)^{\top} \mathbf{V}_n^{-1} (\mathbf{B} - \mathbf{B}_n) + \mathbf{\Psi}_n,$$
where

$$\mathbf{B}_{n} = (\mathbf{V}_{0}^{-1} + \mathbf{X}^{\top} \mathbf{X})^{-1} (\mathbf{V}_{0}^{-1} \mathbf{B}_{0} + \mathbf{X}^{\top} \mathbf{Y}) = (\mathbf{V}_{0}^{-1} + \mathbf{X}^{\top} \mathbf{X})^{-1} (\mathbf{V}_{0}^{-1} \mathbf{B}_{0} + \mathbf{X}^{\top} \mathbf{X} \widehat{\mathbf{B}}),$$

$$\mathbf{V}_{n} = (\mathbf{V}_{0}^{-1} + \mathbf{X}^{\top} \mathbf{X})^{-1},$$

Thus, the posterior distribution can be written as

$$\pi(\mathbf{B}, \mathbf{\Sigma}|\mathbf{Y}, \mathbf{X}) \propto |\mathbf{\Sigma}|^{-K/2} \exp\left\{-\frac{1}{2} tr \left[(\mathbf{B} - \mathbf{B}_n)^{\top} \mathbf{V}_n^{-1} (\mathbf{B} - \mathbf{B}_n) \mathbf{\Sigma}^{-1} \right] \right\} \times |\mathbf{\Sigma}|^{-\frac{N + \alpha_0 + M + 1}{2}} \exp\left\{-\frac{1}{2} tr \left[\mathbf{\Psi}_n \mathbf{\Sigma}^{-1} \right] \right\}.$$

That is $\pi(\mathbf{B}, \mathbf{\Sigma}|\mathbf{Y}, \mathbf{X}) = \pi(\mathbf{B}|\mathbf{\Sigma}, \mathbf{Y}, \mathbf{X})\pi(\mathbf{\Sigma}|\mathbf{Y}, \mathbf{X})$ where $\pi(\mathbf{B}|\mathbf{\Sigma}, \mathbf{Y}, \mathbf{X}) \sim N_{K\times M}(\mathbf{B}_n, \mathbf{V}_n, \mathbf{\Sigma})$ and $\pi(\mathbf{\Sigma}|\mathbf{Y}, \mathbf{X}) \sim IW(\mathbf{\Psi}_n, \alpha_n)$, $\alpha_n = N + \alpha_0$. Observe again that we can write down the posterior mean as a weighted average between prior and sample information such that $\mathbf{V}_0 \to \infty$ implies $\mathbf{B}_n \to \hat{\mathbf{B}}$, as we show in the univariate linear model.

The marginal posterior for \mathbf{B} is given by

$$\pi(\mathbf{B}|\mathbf{Y},\mathbf{X}) \propto \int_{\mathcal{S}} |\mathbf{\Sigma}|^{-(\alpha_n + K + M + 1)/2}$$

$$\times \exp\left\{-\frac{1}{2}tr\left\{\left[(\mathbf{B} - \mathbf{B}_n)^{\top}\mathbf{V}_n^{-1}(\mathbf{B} - \mathbf{B}_n) + \mathbf{\Psi}_n\right]\mathbf{\Sigma}^{-1}\right\}\right\} d\mathbf{\Sigma}$$

$$\propto |(\mathbf{B} - \mathbf{B}_n)^{\top}\mathbf{V}_n^{-1}(\mathbf{B} - \mathbf{B}_n) + \mathbf{\Psi}_n|^{-(K + \alpha_n)/2}$$

$$= \left[|\mathbf{\Psi}_n| \times |\mathbf{I}_K + \mathbf{V}_n^{-1}(\mathbf{B} - \mathbf{B}_n)\mathbf{\Psi}_n^{-1}(\mathbf{B} - \mathbf{B}_n)^{\top}|\right]^{-(\alpha_n + 1 - M + K + M - 1)/2}$$

$$\propto |\mathbf{I}_K + \mathbf{V}_n^{-1}(\mathbf{B} - \mathbf{B}_n)\mathbf{\Psi}_n^{-1}(\mathbf{B} - \mathbf{B}_n)^{\top}|^{-(\alpha_n + 1 - M + K + M - 1)/2}.$$

The second line uses the inverse Wishart distribution, the third line the Sylverter's theorem, and the last line is the kernel of a matrix t distribution, that is, $\mathbf{B}|\mathbf{Y}, \mathbf{X} \sim T_{K \times M}(\mathbf{B}_n, \mathbf{V}_n, \mathbf{\Psi}_n)$ with $\alpha_n + 1 - M$ degrees of freedom.

Observe that $vec(\mathbf{B})$ has mean $vec(\mathbf{B}_n)$ and variance $(\mathbf{V}_n \otimes \mathbf{\Psi}_n)/(\alpha_n - M - 1)$ based on its marginal distribution. On the other hand, the variance based on the conditional distribution is $\mathbf{V}_n \otimes \mathbf{\Sigma}$, where the mean of $\mathbf{\Sigma}$ is $\mathbf{\Psi}_n/(\alpha_n - M - 1)$.

The marginal likelihood is the following,

$$\begin{split} p(\mathbf{Y}) &= \int_{\mathcal{B}} \int_{\mathcal{S}} \left\{ (2\pi)^{-NM/2} |\mathbf{\Sigma}|^{-N/2} \exp\left\{ -\frac{1}{2} tr \left[\mathbf{S} + (\mathbf{B} - \hat{\mathbf{B}})^{\top} \mathbf{X}^{\top} \mathbf{X} (\mathbf{B} - \hat{\mathbf{B}}) \right] \mathbf{\Sigma}^{-1} \right\} \\ &\times (2\pi)^{-KM/2} |\mathbf{V}_0|^{-M/2} |\mathbf{\Sigma}|^{-K/2} \exp\left\{ -\frac{1}{2} tr \left[(\mathbf{B} - \mathbf{B}_0)^{\top} \mathbf{V}_0^{-1} (\mathbf{B} - \mathbf{B}_0) \mathbf{\Sigma}^{-1} \right] \right\} \\ &\times \frac{|\Psi_0|^{\alpha_0/2}}{2^{\alpha_0M/2} \Gamma_M(\alpha_0/2)} |\mathbf{\Sigma}|^{-(\alpha_0+M+1)/2} \exp\left\{ -\frac{1}{2} tr \left[\Psi_0 \mathbf{\Sigma}^{-1} \right] \right\} \right\} d\mathbf{\Sigma} d\mathbf{B} \\ &= (2\pi)^{-M(N+K)/2} |\mathbf{V}_0|^{-M/2} \frac{|\Psi_0|^{\alpha_0/2}}{2^{\alpha_0M/2} \Gamma_M(\alpha_0/2)} \\ &\times \int_{\mathcal{B}} \int_{\mathcal{S}} \left\{ |\mathbf{\Sigma}|^{-(\alpha_0+N+K+M+1)/2} \exp\left\{ -\frac{1}{2} tr \left[\mathbf{S} + (\mathbf{B} - \hat{\mathbf{B}})^{\top} \mathbf{X}^{\top} \mathbf{X} (\mathbf{B} - \hat{\mathbf{B}}) + (\mathbf{B} - \mathbf{B}_0)^{\top} \mathbf{V}_0^{-1} (\mathbf{B} - \mathbf{B}_0) + \Psi_0 \right] \mathbf{\Sigma}^{-1} \right\} \right\} d\mathbf{\Sigma} d\mathbf{B} \\ &= (2\pi)^{-M(N+K)/2} |\mathbf{V}_0|^{-M/2} \frac{|\Psi_0|^{\alpha_0/2}}{2^{\alpha_0M/2} \Gamma_M(\alpha_0/2)} 2^{M(\alpha_n+K)/2} \Gamma_M((\alpha_n+K)/2) \\ &\times \int_{\mathcal{B}} \left| \mathbf{S} + (\mathbf{B} - \hat{\mathbf{B}})^{\top} \mathbf{X}^{\top} \mathbf{X} (\mathbf{B} - \hat{\mathbf{B}}) + (\mathbf{B} - \mathbf{B}_0)^{\top} \mathbf{V}_0^{-1} (\mathbf{B} - \mathbf{B}_0) + \Psi_0 \right]^{-(\alpha_n+K)/2} d\mathbf{B} \\ &= (2\pi)^{-M(N+K)/2} |\mathbf{V}_0|^{-M/2} \frac{|\Psi_0|^{\alpha_0/2}}{2^{\alpha_0M/2} \Gamma_M(\alpha_0/2)} 2^{M(\alpha_n+K)/2} \Gamma_M((\alpha_n+K)/2) \\ &\times \int_{\mathcal{B}} \left| (\mathbf{B} - \hat{\mathbf{B}}_n)^{\top} \mathbf{V}_n^{-1} (\mathbf{B} - \hat{\mathbf{B}}_n) + \Psi_n \right|^{-(\alpha_n+K)/2} d\mathbf{B} \\ &= (2\pi)^{-M(N+K)/2} |\mathbf{V}_0|^{-M/2} \frac{|\Psi_0|^{\alpha_0/2}}{2^{\alpha_0M/2} \Gamma_M(\alpha_0/2)} 2^{M(\alpha_n+K)/2} \Gamma_M((\alpha_n+K)/2) \\ &\times \int_{\mathcal{B}} \left[|\Psi_n| \times |\mathbf{I}_K + \mathbf{V}_n^{-1} (\mathbf{B} - \hat{\mathbf{B}}_n) \Psi_n^{-1} (\mathbf{B} - \hat{\mathbf{B}}_n)^{\top} \right]^{-(\alpha_n+K)/2} d\mathbf{B} \\ &= |\Psi_n|^{-(\alpha_n+K)/2} (2\pi)^{-M(N+K)/2} |\mathbf{V}_0|^{-M/2} \frac{|\Psi_0|^{\alpha_0/2} 2^{M(\alpha_n+K)/2} \Gamma_M((\alpha_n+K)/2)}{2^{\alpha_0M/2} \Gamma_M(\alpha_0/2)} \\ &\times \int_{\mathbf{B}} \left| \mathbf{I}_K + \mathbf{V}_n^{-1} (\mathbf{B} - \hat{\mathbf{B}}_n) \Psi_n^{-1} (\mathbf{B} - \hat{\mathbf{B}}_n)^{\top} \right]^{-(\alpha_n+K)/2} d\mathbf{B} \\ &= |\Psi_n|^{-(\alpha_n+K)/2} (2\pi)^{-M(N+K)/2} |\mathbf{V}_0|^{-M/2} \frac{|\Psi_0|^{\alpha_0/2} 2^{M(\alpha_n+K)/2} \Gamma_M((\alpha_0/2))}{2^{\alpha_0M/2} \Gamma_M(\alpha_0/2)} \\ &\times \pi^{MK/2} \frac{\Gamma_M((\alpha_n+1-M+K+M-1)/2)}{\Gamma_M((\alpha_n+1-M+K+M-1)/2)} |\Psi_n|^{K/2} |\mathbf{V}_n|^{M/2} \\ &= \frac{|\mathbf{V}_n|^{M/2} |\Psi_0|^{\alpha_0/2} \Gamma_M(\alpha_0/2)}{\Gamma_M(\alpha_0/2)} \pi^{-MN/2}. \end{aligned}$$

The third equality follows from having the kernel of a inverse Wishart distribution, the fifth from the Silvester's theorem, and the seventh from having the kernel of a matrix t distribution.

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Observe that this last expression is the multivariate case of the marginal likelihood of the univariate regression model. Taking into account that

$$\begin{split} (\mathbf{A} + \mathbf{B})^{-1} &= \mathbf{A}^{-1} - (\mathbf{A}^{-1} + \mathbf{B}^{-1})^{-1} \mathbf{A}^{-1} \\ &= \mathbf{B}^{-1} - (\mathbf{A}^{-1} + \mathbf{B}^{-1})^{-1} \mathbf{B}^{-1} \\ &= \mathbf{A}^{-1} (\mathbf{A}^{-1} + \mathbf{B}^{-1}) \mathbf{B}^{-1}, \end{split}$$

we can show that $\Psi_n = \Psi_0 + \mathbf{S} + (\hat{\mathbf{B}} - \mathbf{B}_0)^\top \mathbf{V}_n (\hat{\mathbf{B}} - \mathbf{B}_0)$ (see Exercise 7). Therefore, the marginal likelihood rewards fit (smaller sum of squares, \mathbf{S}), similarity between prior and sample information regarding location parameters, and information gains in variability from \mathbf{V}_0 to \mathbf{V}_n .

Given a matrix of regressors \mathbf{X}_0 for N_0 unobserved units, the predictive density of \mathbf{Y}_0 given \mathbf{Y} , $\pi(\mathbf{Y}_0|\mathbf{Y})$ is a matrix t distribution $T_{N_0,M}(\alpha_n - M + 1, \mathbf{X}_0\mathbf{B}_n, \mathbf{I}_{N_0} + \mathbf{X}_0\mathbf{V}_n\mathbf{X}_0^\top, \mathbf{\Psi}_n)$ (see Exercise 6). Observe that the prediction is centered at $\mathbf{X}_0\mathbf{B}_n$, and the covariance matrix of $vec(\mathbf{Y}_0)$ is $\frac{(\mathbf{I}_{N_0} + \mathbf{X}_0\mathbf{V}_n\mathbf{X}_0^\top)\otimes \mathbf{\Psi}_n}{\alpha_n - M - 1}$.

4.5 Summary

We introduce the conjugate family models for discrete and continuous data. These models are the basic Bayesian framework to due its mathematical tractability as we get closed-form expressions for the posterior distributions, the marginal likelihood, and the predictive distribution. We also present the Bayesian linear univariate and multivariate regression frameworks under conjugate families. This is the cornerstone to perform regression analysis in the Bayesian setting.

4.6 Exercises

- 1. Write in the canonical form the distribution of the Bernoulli example, and find the mean and variance of the sufficient statistic.
- 2. Given a random sample $\mathbf{y} = [y_1, y_2, \dots, y_N]^{\top}$ from N binomial experiments each having known size n_i and same unknown probability θ . Show that $p(\mathbf{y}|\theta)$ is in the exponential family, and find the posterior distribution, the marginal likelihood and the predictive distribution of the binomial-beta model assuming the number of trials is known.
- 3. Given a random sample $\mathbf{y} = [y_1, y_2, \dots, y_N]^{\top}$ from a *exponential distribution*. Show that $p(\mathbf{y}|\lambda)$ is in the exponential family, and find

the posterior distribution, marginal likelihood and predictive distribution of the exponential-gamma model.

- 4. Given $\mathbf{y} \sim N_N(\boldsymbol{\mu}, \boldsymbol{\Sigma})$, that is, a multivariate normal distribution show that $p(\mathbf{y}|\boldsymbol{\mu}, \boldsymbol{\Sigma})$ is in the exponential family.
- 5. Find the marginal likelihood in the normal/inverse-Wishart model.
- 6. Find the posterior predictive distribution in the normal/inverse-Wishart model, and show that $\mathbf{Y}_0|\mathbf{Y} \sim T_{N_0,M}(\alpha_n M + 1, \mathbf{X}_0 \mathbf{B}_n, \mathbf{I}_{N_0} + \mathbf{X}_0 \mathbf{V}_n \mathbf{X}_0^\top, \mathbf{\Psi}_n)$.
- 7. Show that $\delta_n = \delta_0 + (\mathbf{y} \mathbf{X}\hat{\boldsymbol{\beta}})^{\top}(\mathbf{y} \mathbf{X}\hat{\boldsymbol{\beta}}) + (\hat{\boldsymbol{\beta}} \boldsymbol{\beta}_0)^{\top}((\mathbf{X}^{\top}\mathbf{X})^{-1} + \mathbf{B}_0)^{-1}(\hat{\boldsymbol{\beta}} \boldsymbol{\beta}_0)$ in the linear regression model, and that $\boldsymbol{\Psi}_n = \boldsymbol{\Psi}_0 + \mathbf{S} + (\hat{\mathbf{B}} \mathbf{B}_0)^{\top}\mathbf{V}_n(\hat{\mathbf{B}} \mathbf{B}_0)$ in the linear multivariate regression model.
- 8. Show that in the linear regression model $\boldsymbol{\beta}_n^{\top}(\mathbf{B}_n^{-1} \mathbf{B}_n^{-1}\mathbf{M}^{-1}\mathbf{B}_n^{-1})\boldsymbol{\beta}_n = \boldsymbol{\beta}_{**}^{\top}\mathbf{C}\boldsymbol{\beta}_{**}$ and $\boldsymbol{\beta}_{**} = \mathbf{X}_0\boldsymbol{\beta}_n$.
- 9. Show that $(\mathbf{Y} \mathbf{X}\mathbf{B})^{\top}(\mathbf{Y} \mathbf{X}\mathbf{B}) = \mathbf{S} + (\mathbf{B} \widehat{\mathbf{B}})^{\top}\mathbf{X}^{\top}\mathbf{X}(\mathbf{B} \widehat{\mathbf{B}})$ where $\mathbf{S} = (\mathbf{Y} \mathbf{X}\widehat{\mathbf{B}})^{\top}(\mathbf{Y} \mathbf{X}\widehat{\mathbf{B}}), \hat{\mathbf{B}} = (\mathbf{X}^{\top}\mathbf{X})^{-1}\mathbf{X}^{\top}\mathbf{Y}$ in the multivariate regression model.

10. What is the probability that the Sun will rise tomorrow?

This is the most famous Richard Price's example developed in the Appendix of the Bayes' theorem paper [5]. Here, we implicitly use Laplace's Rule of Succession to solve this question. In particular, if we were a priori uncertain about the probability the Sun will rise on a specified day, we can assume a prior uniform distribution over (0,1), that is, a beta (1,1) distribution. Then, what is the probability that the Sun will rise?

- 11. Using information from Public Policy Polling in September 27th-28th for the 2016 presidential five-way race in USA, there are 411, 373 and 149 sampled people supporting Hillary Clinton, Donald Trump and other, respectively.
 - •Find the posterior probability of the percentage difference of people supporting Hillary versus Trump according to this data using a non-informative prior, that is, $\alpha_0 = [1 \ 1 \ 1]$ in the multinomial-Dirichlet model. What is the probability of having more supports of Hillary vs Trump?
 - What is the probability that sampling one hundred independent individuals 44, 40 and 16 support Hillary, Trump and other, respectively?

12. Math test example continues

You have a random sample of math scores of size N=50 from a normal distribution, $Y_i \sim \mathcal{N}(\mu, \sigma)$. The sample mean and variance are

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equal to 102 and 10, respectively. Using the normal-normal/inverse-gamma model where $\mu_0 = 100$, $\beta_0 = 1$, $\alpha_0 = \delta_0 = 0.001$

- •Get a 95% confidence and credible interval for μ .
- •What is the posterior probability that $\mu > 103$?

13. Demand of electricity example continues

Set c_0 such that maximizes the marginal likelihood in the specifications with and without electricity price in the example of demand of electricity (empirical Bayes). Then, calculate the Bayes factor, and conclude if there is evidence supporting the inclusion of the price of electricity in the demand equation.

14. Utility demand

Use the file Utilities.csv to estimate a multivariate linear regression model where $\mathbf{Y}_i = [\log(\text{electricity}_i) \log(\text{water}_i) \log(\text{gas}_i)]$ as function of $\log(\text{electricity price}_i)$, $\log(\text{water price}_i)$, $\log(\text{gas price}_i)$, $\operatorname{IndSocio1}_i$, $\operatorname{IndSocio2}_i$, $\operatorname{Altitude}_i$, Nrooms_i , $\operatorname{HouseholdMem}_i$, $\operatorname{Children}_i$, and $\log(\operatorname{Income}_i)$. Set a non-informative prior framework, $\mathbf{B}_0 = [0]_{11\times 3}$, $\mathbf{V}_0 = 1000\mathbf{I}_{11}$, $\mathbf{\Psi}_0 = 1000\mathbf{I}_3$ and $\alpha_0 = 3$, where we have K = 11 (regressors plus intercept) and M = 3 (equations) in this exercise.

- (a) Find the posterior mean estimates and the highest posterior density intervals at 95% of $\bf B$ and $\bf \Sigma$. Use the marginal distribution and the conditional distribution to obtain the posterior estimates of $\bf B$, and compare the results.
- (b) Find the Bayes factor comparing the baseline model in this exercise with the same specification but using the income in dollars. Now, calculate the Bayes factor using the income in thousand dollars. Is there any difference?
- (c) Find the predictive distribution for the monthly demand of electricity, water and gas in the baseline specification of a household located in the lowest socioeconomic condition in a municipality located below 1000 meters above the sea level, 2 rooms, 3 members with children, a monthly income equal to USD 500, an electricity price equal to USD/kWh 0.15, a water price equal to USD/M³ 0.70, and a gas price equal to USD/M³ 0.75.

Simulation methods

5.1	The inverse transform method
5.2	Method of composition
5.3	Accept and reject algorithm
5.4	Importance sampling
5.5.1	Markov chain Monte Carlo methods Some theory Gibbs sampler Metropolis-Hastings
5.6	Sequential Monte Carlo
5.7	Hamiltonian Monte Carlo

Convergence diagnostics

Part II Regression models: A GUIded tour

We describe how to perform Bayesian inference in some of the most common univariate models: normal-inverse gamma, logit, probit, multinomial probit and logit, ordered probit, negative binomial, tobit, quantile regression, and Bayesian bootstrap in linear models. We show the posterior distributions of the parameters and some applications. In addition, we demonstrate how to perform inference in various models using three levels of programming skills: our graphical user interface (GUI), packages from **R**, and direct programming of the posterior distributions. The first requires no programming skills, the second requires an intermediate level, and the third demands more advanced skills. We also include mathematical and computational exercises

6.1 The Gaussian linear model

The Gaussian linear model specifies $\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \mu$ such that $\mu \sim N(\mathbf{0}, \sigma^2 \mathbf{I_N})$ is an stochastic error, \mathbf{X} is a $N \times K$ matrix of regressors, $\boldsymbol{\beta}$ is a K-dimensional vector of location coefficients, σ^2 is the variance of the model (scale parameter), \mathbf{y} is a N-dimensional vector of a dependent variable, and N is the sample size. We describe this model using the conjugate family in Section 4.3, that is, $\pi(\boldsymbol{\beta}, \sigma^2) = \pi(\boldsymbol{\beta} | \sigma^2) \times \pi(\sigma^2)$, and this allowed to get the posterior marginal distribution for $\boldsymbol{\beta}$ and σ^2 .

We assume independent prior in this section, that is, $\pi(\boldsymbol{\beta}, \sigma^2) = \pi(\boldsymbol{\beta}) \times \pi(\sigma^2)$, where $\boldsymbol{\beta} \sim N(\boldsymbol{\beta}_0, \mathbf{B}_0)$ and $\sigma^2 \sim IG(\alpha_0/2, \delta_0/2)$, $\alpha_0/2$ and $\delta_0/2$ are the shape and rate parameters. This setting allows getting the posterior conditional distributions, that is, $\pi(\boldsymbol{\beta}|\sigma^2, \mathbf{y}, \mathbf{X})$ and $\pi(\sigma^2|\boldsymbol{\beta}, \mathbf{y}, \mathbf{X})$, which in turn allows to use the Gibbs sampler algorithm to perform posterior inference of $\boldsymbol{\beta}$ and σ^2 .

The likelihood function in this model is

$$p(\mathbf{y}|\boldsymbol{\beta}, \sigma^2, \mathbf{X}) = (2\pi\sigma^2)^{-\frac{N}{2}} \exp\left\{-\frac{1}{2\sigma^2}(\mathbf{y} - \mathbf{X}\boldsymbol{\beta})^\top (\mathbf{y} - \mathbf{X}\boldsymbol{\beta})\right\}.$$

Then, the conditional posterior distributions are

$$\boldsymbol{\beta}|\sigma^2, \mathbf{y}, \mathbf{X} \sim N(\boldsymbol{\beta}_n, \sigma^2 \mathbf{B}_n),$$

and

$$\sigma^2 | \boldsymbol{\beta}, \mathbf{y}, \mathbf{X} \sim IG(\alpha_n/2, \delta_n/2),$$

where
$$\mathbf{B}_n = (\mathbf{B}_0^{-1} + \sigma^{-2}\mathbf{X}^{\top}\mathbf{X})^{-1}$$
, $\boldsymbol{\beta}_n = \mathbf{B}_n(\mathbf{B}_0^{-1}\boldsymbol{\beta}_0 + \sigma^{-2}\mathbf{X}^{\top}\mathbf{y})$, $\alpha_n = \alpha_0 + N$ and $\delta_n = \delta_0 + (\mathbf{y} - \mathbf{X}\boldsymbol{\beta})^{\top}(\mathbf{y} - \mathbf{X}\boldsymbol{\beta})$ (see Exercise 1 in this chapter).¹

Example: The market value of soccer players in Europe

Let's analyze the determinants of the market value of soccer players in Europe. In particular, we use the dataset 1ValueFootballPlayers.csv which is in folder **DataApp** in our github repository https://github.com/besmarter/BSTApp. This dataset was used by [72] to finding the determinants of high performance soccer players in the five most important national leagues in Europe.

The specification of the model is

$$log(Value_i) = \beta_1 + \beta_2 Perf_i + \beta_3 Age_i + \beta_4 Age_i^2 + \beta_5 Nat Team_i + \beta_6 Goals_i + \beta_7 Exp_i + \beta_8 Exp_i^2 + \mu_i,$$

where *Value* is the market value in Euros (2017), *Perf* is a measure of performance, *Age* is the players' age in years, *NatTem* is an indicator variable that takes the value of 1 if the player has been on the national team, *Goals* is the number of goals scored by the player during his career, and *Exp* is his experience in years.

We assume that the dependent variable distributes normal, then we use a normal-inverse gamma model using vague conjugate priors where $\mathbf{B}_0 = 1000\mathbf{I}_8$, $\beta_0 = \mathbf{0}_8$, $\alpha_0 = 0.001$ and $\delta_0 = 0.001$. We perform a Gibbs sampler with 5,000 MCMC iterations plus a burn-in equal to 5,000, and a thinning parameter equal to 1.

We can see in the results of the algorithms that we get similar results using the three approaches: GUI, package and our function. For instance, the value of a top soccer player in Europe increases 134% (exp(0.85) -1)) on average when he has played in the national team, the credible interval at 95% is (86%, 197%).

¹This model can be extended to consider heterosked asticity such that $y_i \sim N(\mathbf{x}_i^\top \beta, \sigma^2/\tau_i)$, where $\tau_i \sim G(v/2, v/2)$. See exercise 2 for details.

R code. The value of soccer players, using R packages

```
1 rm(list = ls())
2 set.seed(010101)
3 ####################### Linear regression: Value of
      4 Data <- read.csv("DataApplications/1ValueFootballPlayers.csv
      ", sep = ",", header = TRUE, fileEncoding = "latin1")
5 attach(Data)
6 y <- log(Value)
7 # Value: Market value in Euros (2017) of soccer players
8 # Regressors quantity including intercept
9 X <- cbind(1, Perf, Age, Age2, NatTeam, Goals, Exp, Exp2)
10 # Perf: Performance. Perf2: Performance squared. Age: Age;
      Age: Age squared.
11 # NatTeam: Indicator of national team. Goals: Scored goals.
      Goals2: Scored goals squared
12 # Exp: Years of experience. Exp2: Years of experience
      squared. Assists: Number of assists
13 k <- dim(X)[2]
14 N <- dim(X)[1]
15 # Hyperparameters
16 d0 <- 0.001/2
17 a0 <- 0.001/2
18 b0 <- rep(0, k)
19 c0 <- 1000
20 B0 <- c0*diag(k)
21 B0i <- solve(B0)
22 # MCMC parameters
23 mcmc <- 5000
24 burnin <- 5000
25 tot <- mcmc + burnin
26 thin <- 1
27 # Posterior distributions using packages: MCMCpack sets the
      model in terms of the precision matrix
28 posterior <- MCMCpack::MCMCregress(y~X-1, b0=b0, B0 = B0i,</pre>
      c0 = a0, d0 = d0, burnin = burnin, mcmc = mcmc, thin =
      thin)
29 summary(coda::mcmc(posterior))
30 Iterations = 1:5000
31 Thinning interval = 1
32 Number of chains = 1
33 Sample size per chain = 5000
34 1. Empirical mean and standard deviation for each variable,
35 plus standard error of the mean:
36 Mean
            SD Naive SE Time-series SE
37 X
            3.695499 2.228060 3.151e-02
                                              3.151e-02
                                              6.079e-05
38 XPerf
            0.035445 0.004299 6.079e-05
39 XAge
            0.778410 0.181362 2.565e-03
                                              2.565e-03
            -0.016617 0.003380 4.781e-05
40 XAge2
                                              4.781e - 05
41 XNatTeam 0.850362 0.116861 1.653e-03
                                              1.689e-03
42 XGoals
            0.009097 0.001603 2.266e-05
                                              2.266e-05
43 XExp
            0.206208 0.062713 8.869e-04
                                              8.428e-04
44 XExp2
            -0.006992 0.002718 3.844e-05
                                              3.719e-05
45 sigma2
            0.969590 0.076091 1.076e-03
                                              1.076e-03
```

R. code. The value of soccer players, programming our Gibbs sampler

```
1 # Posterior distributions programming the Gibbs sampling
2 # Auxiliary parameters
3 XtX <- t(X) %*%X
4 bhat <- solve(XtX)%*%t(X)%*%y
5 an <- a0 + N
6 # Gibbs sampling functions
7 PostSig2 <- function(Beta){</pre>
8 dn <- d0 + t(y - X%*%Beta)%*%(y - X%*%Beta)
    sig2 \leftarrow invgamma::rinvgamma(1, shape = an/2, rate = dn/2)
    return(sig2)
11 }
12 PostBeta <- function(sig2){</pre>
Bn <- solve(B0i + sig2^(-1)*XtX)</pre>
    bn <- Bn%*%(B0i%*%b0 + sig2^(-1)*XtX%*%bhat)
    Beta <- MASS::mvrnorm(1, bn, Bn)
16
    return (Beta)
17 }
18 PostBetas <- matrix(0, mcmc+burnin, k)
19 PostSigma2 <- rep(0, mcmc+burnin)</pre>
20 Beta <- rep(0, k)</pre>
21 for(s in 1:tot){
   sig2 <- PostSig2(Beta = Beta)
    PostSigma2[s] <- sig2
23
    Beta <- PostBeta(sig2 = sig2)
    PostBetas[s,] <- Beta
25
26 }
27 keep <- seq((burnin+1), tot, thin)</pre>
28 PosteriorBetas <- PostBetas[keep,]</pre>
29 colnames(PosteriorBetas) <- c("Intercept", "Perf", "Age", "</pre>
      Age2", "NatTeam", "Goals", "Exp", "Exp2")
30 summary(coda::mcmc(PosteriorBetas))
31 Iterations = 1:5000
32 Thinning interval = 1
33 Number of chains = 1
34 Sample size per chain = 5000
35 1. Empirical mean and standard deviation for each variable,
36 plus standard error of the mean:
              SD Naive SE Time-series SE
38 Intercept 3.663230 2.194363 3.103e-02
                                                 3.103e-02
             0.035361 0.004315 6.102e-05
                                                 6.102e-05
39 Perf
40 Age
             0.780374 0.178530 2.525e-03
                                                 2.525e-03
             -0.016641 0.003332 4.713e-05
                                                 4.713e-05
41 Age2
42 NatTeam
             0.850094 0.119093 1.684e-03
                                                 1.684e-03
             0.009164 0.001605 2.270e-05
43 Goals
                                                 2.270e-05
44 Exp
             0.205965 0.062985 8.907e-04
                                                  8.596e-04
                                                  3.701e-05
             -0.007006 0.002731 3.862e-05
45 Exp2
46 PosteriorSigma2 <- PostSigma2[keep]
47 summary(coda::mcmc(PosteriorSigma2))
48 Iterations = 1:5000
49 Thinning interval = 1
50 Number of chains = 1
51 Sample size per chain = 5000
52 1. Empirical mean and standard deviation for each variable,
53 plus standard error of the mean:
                              Naive SE Time-series SE
54 Mean
                    SD
55 0.973309
                  0.077316
                                                   0.001116
                                   0.001093
```

6.2 The logit model

In the logit model the dependent variable is binary, $Y_i = \{1, 0\}$, then it follows a Bernoulli distribution, $Y_i \overset{ind}{\sim} B(\pi_i)$, that is, $p(Y_i = 1) = \pi_i$, such that $\pi_i = \frac{\exp\{\mathbf{x}_i^{\mathsf{T}}\boldsymbol{\beta}\}}{1+\exp\{\mathbf{x}_i^{\mathsf{T}}\boldsymbol{\beta}\}}$.

The likelihood function of the logit model is

$$p(\mathbf{y}|\boldsymbol{\beta}, \mathbf{X}) = \prod_{i=1}^{N} \pi_i^{y_i} (1 - \pi_i)^{1 - y_i}$$
$$= \prod_{i=1}^{N} \left(\frac{\exp\left\{\mathbf{x}_i^{\top} \boldsymbol{\beta}\right\}}{1 + \exp\left\{\mathbf{x}_i^{\top} \boldsymbol{\beta}\right\}} \right)^{y_i} \left(\frac{1}{1 + \exp\left\{\mathbf{x}_i^{\top} \boldsymbol{\beta}\right\}} \right)^{1 - y_i}.$$

We can specify a Normal distribution as prior, $\beta \sim N(\beta_0, \mathbf{B}_0)$. Then, the posterior distribution is

$$\pi(\boldsymbol{\beta}|\mathbf{y}, \mathbf{X}) \propto \prod_{i=1}^{N} \left(\frac{\exp\left\{\mathbf{x}_{i}^{\top} \boldsymbol{\beta}\right\}}{1 + \exp\left\{\mathbf{x}_{i}^{\top} \boldsymbol{\beta}\right\}} \right)^{y_{i}} \left(\frac{1}{1 + \exp\left\{\mathbf{x}_{i}^{\top} \boldsymbol{\beta}\right\}} \right)^{1 - y_{i}} \times \exp\left\{ -\frac{1}{2} (\boldsymbol{\beta} - \boldsymbol{\beta}_{0})^{\top} \mathbf{B}_{\mathbf{0}}^{-1} (\boldsymbol{\beta} - \boldsymbol{\beta}_{\mathbf{0}}) \right\}.$$

The logit model does not have a standard posterior distribution. Then, a random walk Metropolis–Hastings algorithm can be used to obtain draws from the posterior distribution. A potential proposal is a multivariate Normal centered at the current value, with covariance matrix $\tau^2(\mathbf{B}_0^{-1}+\widehat{\boldsymbol{\Sigma}}^{-1})^{-1}$, where $\tau>0$ is a tuning parameter and $\widehat{\boldsymbol{\Sigma}}$ is the sample covariance matrix from the maximum likelihood estimation [53].²

Observe that $\log(p(\mathbf{y}|\boldsymbol{\beta}, \mathbf{X})) = \sum_{i=1}^{N} y_i \mathbf{x}_i^{\top} \boldsymbol{\beta} - \log(1 + \exp(\mathbf{x}_i^{\top} \boldsymbol{\beta}))$. We can use this expression when calculating the acceptance parameter in the computational implementation of the Metropolis-Hastings algorithm. In particular, the acceptance parameter is

$$\alpha = \min \left\{ 1, \exp(\log(p(\mathbf{y}|\boldsymbol{\beta}^c, \mathbf{X})) + \log(\pi(\boldsymbol{\beta}^c)) - (\log(p(\mathbf{y}|\boldsymbol{\beta}^{(s-1)}, \mathbf{X})) + \log(\pi(\boldsymbol{\beta}^{(s-1)}))) \right\}$$

where β^c and $\beta^{(s-1)}$ are the draws from the proposal distribution and the previous iteration of the Markov chain, respectively.³

Example: Simulation exercise

 $^{^2}$ Tuning parameters should be set in a way such that one obtains reasonable diagnostic criteria and acceptation rates.

³Formulating the acceptance rate using log helps to mitigate computational problems.

Let's do a simulation exercise to check the performance of the algorithm.

Set $\boldsymbol{\beta} = \begin{bmatrix} 0.5 & 0.8 & -1.2 \end{bmatrix}^{\top}$, $x_{ik} \sim N(0,1)$, k = 2,3 and $i = 1,2,\ldots,10000$. We set as hyperparameters $\boldsymbol{\beta}_0 = \begin{bmatrix} 0 & 0 & 0 \end{bmatrix}^{\top}$ and $\mathbf{B}_0 = 1000\mathbf{I}_3$. The tune parameter for the Metropolis-Hastings algorithm is equal to 1. We obtain an acceptance rate of 0.46, and the diagnostics suggest that the posterior chains behave well. We also see that the posterior results using MCMClogit from MCMCpack and our Metropolis-Hastings algorithm are very similar. In general, the 95% credible intervals encompass the population values, and the mean and median are very close to these values.

The logit model

$R.\ code.\ The\ value\ of\ soccer\ players,\ using\ R$ packages

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```
################# Logit: Simulation
      ############################
2 # Simulate data
3 rm(list = ls())
4 set.seed(010101)
5 N <- 10000 # Sample size
_{6} B <- c(0.5, 0.8, -1.2) # Population location parameters
7 x2 <- rnorm(N) # Regressor
8 x3 <- rnorm(N) # Regressor</pre>
9 X <- cbind(1, x2, x3) # Regressors
10 XB <- X%*%B
11 PY \leftarrow exp(XB)/(1 + exp(XB)) # Probability of Y = 1
_{12} Y <- rbinom(N, 1, PY) # Draw Y's
13 table(Y) # Frequency
# write.csv(cbind(Y, x2, x3), file = "DataSimulations/")
      LogitSim.csv") # Export data
_{15} # MCMC parameters
16 iter <- 5000; burnin <- 1000; thin <- 5; tune <- 1
17 # Hyperparameters
18 K <- dim(X)[2]
19 b0 <- rep(0, K)
20 cO <- 1000
21 BO <- c0*diag(K)
22 B0i <- solve(B0)
23 # Posterior distributions using packages: MCMCpack sets the
      model in terms of the precision matrix
24 RegLog <- MCMCpack::MCMClogit(Y~X-1, mcmc = iter, burnin =
      burnin, thin = thin, b0 = b0, B0 = B0i, tune = tune)
25 summary(RegLog)
26 Iterations = 1001:5996
27 Thinning interval = 5
28 Number of chains = 1
29 Sample size per chain = 1000
30 1. Empirical mean and standard deviation for each variable,
31 plus standard error of the mean:
32 Mean
            SD Naive SE Time-series SE
       0.4896 0.02550 0.0008064
                                        0.001246
33 X
34 Xx2 0.8330 0.02730 0.0008632
                                        0.001406
35 Xx3 -1.2104 0.03049 0.0009643
                                        0.001536
36 2. Quantiles for each variable:
        2.5%
                  25%
                          50%
                                  75%
                                         97.5%
       0.4424
               0.4728
                       0.4894
                                0.5072
                                        0.5405
38 X
39 Xx2 0.7787
               0.8159
                       0.8327
                                0.8505
                                        0.8852
40 Xx3 -1.2758 -1.2296 -1.2088 -1.1902 -1.1513
```

$R.\ code.\ The\ value\ of\ soccer\ players,\ programming\ our\ M-H\ algorithm$

```
1 # Posterior distributions programming the Metropolis-
      Hastings algorithm
2 MHfunc \leftarrow function(y, X, b0 = rep(0, dim(X)[2] + 1), B0 =
      1000*diag(dim(X)[2] + 1), tau = 1, iter = 6000, burnin =
       1000, thin = 5){
    Xm <- cbind(1, X) # Regressors</pre>
    K <- dim(Xm)[2] # Number of location parameters</pre>
    BETAS <- matrix(0, iter + burnin, K) # Space for posterior
       chains
    Reg <- glm(y ~ Xm - 1, family = binomial(link = "logit"))</pre>
      # Maximum likelihood estimation
    BETA <- Reg$coefficients # Maximum likelihood parameter
      estimates
    tot <- iter + burnin # Total iterations M-H algorithm
    COV <- vcov(Reg) # Maximum likelihood covariance matrix
9
    COVt <- tau^2*solve(solve(BO) + solve(COV)) # Covariance
      matrix for the proposal distribution
    Accep <- rep(0, tot) # Space for calculating the
      acceptance rate
    # Create progress bar in case that you want to see
      iterations progress
    pb <- winProgressBar(title = "progress bar", min = 0,</pre>
13
    max = tot, width = 300)
14
    for(it in 1:tot){
      BETAc \leftarrow BETA + MASS::mvrnorm(n = 1, mu = rep(0, K),
16
      Sigma = COVt) # Candidate location parameter
      likecand <- sum((Xm%*%BETAc) * Y - apply(Xm%*%BETAc, 1,
17
      function(x) log(1 + exp(x))) # Log likelihood for the
      candidate
      likepast <- sum((Xm%*%BETA) * Y - apply((Xm%*%BETA), 1,
      function(x) log(1 + exp(x)))) # Log likelihood for the
      actual draw
      priorcand <- (-1/2)*crossprod((BETAc - b0), solve(B0))%*</pre>
      %(BETAc - b0) # Log prior for candidate
      priorpast <- (-1/2)*crossprod((BETA - b0), solve(B0))%*%</pre>
      (BETA - b0) # Log prior for actual draw
      alpha <- min(1, exp(likecand + priorcand - likepast -
      priorpast)) #Probability of selecting candidate
      u \leftarrow runif(1) # Decision rule for selecting candidate
      if(u < alpha){</pre>
23
        BETA <- BETAc # Changing reference for candidate if
24
      selected
         Accep[it] <- 1 # Indicator if the candidate is
      accepted
26
      BETAS[it, ] <- BETA # Saving draws
27
      setWinProgressBar(pb, it, title=paste( round(it/tot*100,
28
       0),
      "% done"))
29
    }
30
31
    close(pb)
    keep <- seq(burnin, tot, thin)</pre>
32
    return(list(Bs = BETAS[keep[-1], ], AceptRate = mean(Accep
       [keep[-1]])))
34 }
```

The logit model

R. code. The value of soccer players, programming our M-H algorithm

```
Posterior <- MHfunc(y = Y, X = cbind(x2, x3), iter = iter,</pre>
      burnin = burnin, thin = thin) # Running our M-H function
       changing some default parameters.
paste("Acceptance rate equal to", round(Posterior$AceptRate,
       2), sep = " ")
3 "Acceptance rate equal to 0.46"
4 PostPar <- coda::mcmc(Posterior$Bs)</pre>
5 # Names
6 colnames(PostPar) <- c("Cte", "x1", "x2")
7 # Summary posterior draws
8 summary(PostPar)
9 Iterations = 1:1000
10 Thinning interval = 1
11 Number of chains = 1
12 Sample size per chain = 1000
13 1. Empirical mean and standard deviation for each variable,
14 plus standard error of the mean:
15 Mean
          SD Naive SE Time-series SE
16 Cte 0.4893 0.02427 0.0007674
                                       0.001223
17 x 1
      0.8309 0.02699 0.0008536
                                       0.001440
18 x2 -1.2107 0.02943 0.0009308
                                       0.001423
19 2. Quantiles for each variable:
      2.5%
               25%
                       50%
                                75%
                                      97.5%
20
21 Cte 0.4431 0.4721 0.4899 0.5059 0.5344
      0.7817 0.8123 0.8305 0.8505 0.8833
22 x 1
23 x2 -1.2665 -1.2309 -1.2107 -1.1911 -1.1538
_{24} # Trace and density plots
25 plot(PostPar)
26 # Autocorrelation plots
27 coda::autocorr.plot(PostPar)
28 # Convergence diagnostics
29 coda::geweke.diag(PostPar)
30 Fraction in 1st window = 0.1
31 Fraction in 2nd window = 0.5
32 Cte
         x 1
                x2
33 -0.975 -3.112 1.326
34 coda::raftery.diag(PostPar,q=0.5,r=0.05,s = 0.95)
35 Quantile (q) = 0.5
36 Accuracy (r) = +/- 0.05
_{37} Probability (s) = 0.95
38 Burn-in Total Lower bound Dependence
39 (M)
                               factor (I)
           (N) (Nmin)
40 Cte 6
               731 385
                                   1.90
41 x1 6
                                   1.83
               703
                    385
42 x2 6
               725
                    385
                                   1.88
43 coda::heidel.diag(PostPar)
44 Stationarity start p-value
45 test
               iteration
_{
m 46} Cte passed
                    1
                              0.4436
47 x1 passed
                    101
                              0.3470
                              0.0872
48 x2 passed
                    1
49 Halfwidth Mean
                  Halfwidth
50 test
51 Cte passed
                 0.489 0.00240
52 x1 passed
                 0.832 0.00268
                -1.211 0.00279
53 x2 passed
```

6.3 The probit model

The probit model also has as dependent variable a binary outcome. In this case, there is a latent variable $(y_i^*, \text{ unobserved})$ that defines the structure of the estimation problem. In particular,

$$Y_i = \begin{cases} 0, \ Y_i^* \le 0 \\ 1, \ Y_i^* > 0 \end{cases},$$

such that $Y_i^* = \mathbf{x}_i^{\top} \boldsymbol{\beta} + \mu_i$, $\mu_i \stackrel{i.i.d.}{\sim} N(0,1)$. This implies $P(Y_i = 1) = \pi_i = 0$ $\Phi(\mathbf{x}_{i}^{\top}\boldsymbol{\beta}).$

[1] implemented data augmentation [75] to apply a Gibbs sampling algorithm in this model. Augmenting this model with Y_i^* , we can have the likelihood contribution from observation $i, p(y_i|y_i^*) = \mathbb{1}_{y_i=0} \mathbb{1}_{y_i^* \leq 0} + \mathbb{1}_{y_i=1} \mathbb{1}_{y_i^* > 0}$, where $\mathbb{1}_A$ is an indicator function that takes the value of 1 when condition A is satisfied.

The posterior distribution is $\pi(\boldsymbol{\beta}, \boldsymbol{y^*}|\boldsymbol{y}, \boldsymbol{X}) \propto \prod_{i=1}^{N} \left[\mathbbm{1}_{y_i=0} \mathbbm{1}_{y_i^* \leq 0} + \mathbbm{1}_{y_i=1} \mathbbm{1}_{y_i^* > 0}\right] \times$ $N_N(\boldsymbol{y}^*|\boldsymbol{X}\boldsymbol{\beta},\boldsymbol{I}_n)\times N_K(\boldsymbol{\beta}|\boldsymbol{\beta}_0,\boldsymbol{B}_0)$ when taking a Gaussian distribution as prior $\boldsymbol{\beta} \sim N_k(\boldsymbol{\beta}_0, \boldsymbol{B}_0)$. This implies

$$y_i^*|\boldsymbol{\beta}, \boldsymbol{y}, \boldsymbol{X} \sim \left\{ \begin{aligned} TN_{(-\infty,0]}(\mathbf{x}_i^{\top} \boldsymbol{\beta}, \mathbf{1}) &, \ \mathbf{y_i} = \mathbf{0} \\ TN_{(0,\infty)}(\mathbf{x}_i^{\top} \boldsymbol{\beta}, \mathbf{1}) &, \ \mathbf{y_i} = \mathbf{1} \end{aligned} \right\},$$

$$\boldsymbol{\beta}|\boldsymbol{y}^*, \boldsymbol{X} \sim N(\boldsymbol{\beta}_n, \boldsymbol{B}_n),$$

where
$$B_n = (B_0^{-1} + X^{\top}X)^{-1}$$
, and $\beta_n = B_n(B_0^{-1}\beta_0 + X^{\top}y^*)$.

Example: Determinants of hospitalization

We use the dataset named **2HealthMed.csv**, which is in folder **DataApp** in our github repository (https://github.com/besmarter/BSTApp and was used by [63]. Our dependent variable is a binary indicator with a value equal to 1 if an individual was hospitalized in 2007, and 0 otherwise.

The specification of the model is

$$Hosp_{i} = \beta_{1} + \beta_{2}SHI_{i} + \beta_{3}Female_{i} + \beta_{4}Age_{i} + \beta_{5}Age_{i}^{2} + \beta_{6}Est2_{i} + \beta_{7}Est3_{i} + \beta_{8}Fair_{i} + \beta_{9}Good_{i} + \beta_{10}Excellent_{i},$$

where SHI is a binary variable equal to 1 if the individual is in a subsidized

⁴The variance in this model is set to 1 due to identification restrictions. Observe that $P(Y_i = 1 | \mathbf{x_i}) = \mathbf{P}(\mathbf{Y_i^*} > \mathbf{0} | \mathbf{x_i}) = \mathbf{P}(\mathbf{x_i^\top} \boldsymbol{\beta} + \mu_i > \mathbf{0} | \mathbf{x_i}) = \mathbf{P}(\mu_i > -\mathbf{x_i^\top} \boldsymbol{\beta} | \mathbf{x_i}) = \mathbf{P}(\mathbf{c} \times \mu_i > -\mathbf{c} \times \mathbf{x_i^\top} \boldsymbol{\beta} | \mathbf{x_i}) \ \forall c > 0$. Multiplying for a positive constant does not affect the probability of $Y_i = 1$. 5TN denotes a truncated normal density.

health care program and 0 otherwise, Female is an indicator of gender, Age in years, Est2 and Est3 are indicators of socioeconomic status, the reference is Est1, which is the lowest, and self perception of health status where bad is the reference.

Let's set $\beta_0 = \mathbf{0}_{10}$, $\mathbf{B}_0 = \mathbf{I}_{10}$, iterations, burn-in and thinning parameters equal to 10000, 1000 and 1, respectively. We use the package *bayesm* to perform this exercise. We asked to program a Gibbs sampler algorithm in the exercises.

We find evidence that gender and self-perceived health status affect the probability of hospitalization. Women have a higher probability of being hospitalized than men, and a better perception of health status decreases this probability.

R. code. Determinants of hospitalization

```
nydata <- read.csv("DataApplications/2HealthMed.csv", header</pre>
        = T, sep = ",")
2 attach (mydata)
3 str(mydata)
4 K <- 10 # Number of regressors
5 b0 <- rep(0, K) # Prio mean
6 BOi <- diag(K) # Prior precision (inverse of covariance)
7 Prior <- list(betabar = b0, A = B0i) # Prior list</pre>
8 y <- Hosp # Dependent variables</pre>
9 X <- cbind(1, SHI, Female, Age, Age2, Est2, Est3, Fair, Good
       , Excellent) # Regressors
10 Data \leftarrow list(y = y, X = X) # Data list
11 Mcmc <- list(R = 10000, keep = 1, nprint = 0) # MCMC
       parameters
12 RegProb <- bayesm::rbprobitGibbs(Data = Data, Prior = Prior,</pre>
       Mcmc = Mcmc) # Inference using bayesm package
13 PostPar <- coda::mcmc(RegProb$betadraw) # Posterior draws
14 colnames(PostPar) <- c("Cte", "SHI", "Female", "Age", "Age2", "Est2", "Est3", "Fair", "Good", "Excellent") # Names
15 summary(PostPar) # Posterior summary
16 Iterations = 1:10000
17 Thinning interval = 1
18 Number of chains = 1
19 Sample size per chain = 10000
20 2. Quantiles for each variable:
           2.5%
                                     50%
                        25%
                                                 75%
                                                           97.5%
21
22 Cte
              -1.22e+00 -1.03e+00 -9.43e-01 -8.50e-01 -0.671744
              -1.24e-01 -4.63e-02 -6.30e-03 3.26e-02 0.104703
23 SHI
24 Female
              2.80e-02 9.65e-02
                                   1.28e-01
                                               1.60e-01
                                                          0.223123
25 Age
             -7.55e-03 -2.50e-03
                                    1.25e-04
                                                2.80e-03
                                                          0.007646
             -4.98e-05 9.05e-06 4.02e-05 7.07e-05
-1.89e-01 -1.23e-01 -8.84e-02 -5.32e-02
                                                          0.000128
26 Age2
27 Est2
                                                           0.012714
             -2.13e-01 -1.03e-01 -4.73e-02 1.01e-02 0.109527
28 Est3
29 Fair
             -7.09e-01 -5.69e-01 -4.93e-01 -4.16e-01 -0.269494
             -1.42e+00 -1.28e+00 -1.20e+00 -1.12e+00 -0.982533
30 Good
31 Excellent -1.33e+00 -1.15e+00 -1.06e+00 -9.74e-01 -0.795881
```

6.4 The multinomial probit model

The multinomial probit model is used to model the choice of the l-th alternative over a set L mutually exclusive options. We observe

$$y_{il} = \begin{cases} 1, & y_{il}^* \ge \max\{\boldsymbol{y}_i^*\} \\ 0, & \text{otherwise} \end{cases},$$

such that $\boldsymbol{y}_i^* = \boldsymbol{X}_i \boldsymbol{\delta} + \boldsymbol{\mu}_i$, $\boldsymbol{\mu}_i \overset{i.i.d.}{\sim} N(\boldsymbol{0}, \boldsymbol{\Sigma})$, \boldsymbol{y}_i^* is an unobserved latent L dimensional vector, $\boldsymbol{X}_i = \begin{bmatrix} (1 \ \boldsymbol{c}_i^\top) \otimes \boldsymbol{I}_L \ \boldsymbol{A}_i \end{bmatrix}$ is an $L \times j$ matrix of regressors for each alternative, $l = 1, 2, \ldots, L, j = L \times (1 + dim\{\boldsymbol{c}_i\}) + a, \boldsymbol{c}_i$ is a vector of the individuals' specific characteristics, \boldsymbol{A}_i is an $L \times a$ matrix of alternative-varying regressors, a is the number of alternative-varying regressors, and $\boldsymbol{\delta}$ is a j dimensional vector of parameters.

We take into account simultaneously the alternative-varying regressors (alternative attributes) and alternative-invariant regressors (individual characteristics). 6 \boldsymbol{y}_i^* can be stacked up into a multiple regression with correlated stochastic errors, $\boldsymbol{y}^* = \boldsymbol{X}\boldsymbol{\delta} + \boldsymbol{\mu}$, where $\boldsymbol{y}^* = [\boldsymbol{y}_1^{*\top}, \boldsymbol{y}_2^{*\top}, \dots, \boldsymbol{y}_N^{*\top}], \boldsymbol{X} = [\boldsymbol{X}_1^{\top}, \boldsymbol{X}_2^{\top}, \dots, \boldsymbol{X}_N^{\top}]^{\top}$, and $\boldsymbol{\mu} = [\boldsymbol{\mu}_1^{\top}, \boldsymbol{\mu}_2^{\top}, \dots, \boldsymbol{\mu}_N^{\top}]^{\top}$. Following the practice of expressing \boldsymbol{y}_{il}^* relative to \boldsymbol{y}_{iL}^* by letting $\boldsymbol{w}_i = \boldsymbol{v}_{il}^{\top}$.

Following the practice of expressing y_{il}^* relative to y_{iL}^* by letting $\boldsymbol{w}_i = [w_{i1}, w_{i2}, \dots, w_{iL-1}]^\top$, $w_{il} = y_{il}^* - y_{iL}^*$, we can write $\boldsymbol{w}_i = \boldsymbol{R}_i \boldsymbol{\beta} + \boldsymbol{\epsilon}_i$, $\boldsymbol{\epsilon}_i \sim N(\mathbf{0}, \boldsymbol{\Omega})$, where $\boldsymbol{R}_i = [(1 \ \boldsymbol{c}_i^\top) \otimes \boldsymbol{I}_{L-1} \ \boldsymbol{\Delta} \boldsymbol{A}_i]$ is an $L-1 \times k$ matrix where $\boldsymbol{\Delta} \boldsymbol{A}_i = \boldsymbol{A}_{li} - \boldsymbol{A}_{Li}, l = 1, 2, \dots, L-1$, that is, the last row of \boldsymbol{A}_i is subtracted from each row of \boldsymbol{A}_i , and $\boldsymbol{\beta}$ is a k dimensional vector, $k = (L-1) \times (1 + dim\{\boldsymbol{c}_i\}) + a$.

Observe that $\boldsymbol{\beta}$ contains the same last a elements as $\boldsymbol{\delta}$, that is, alternative specific attributes coefficients, but the first $(L-1)\times(1+\dim\{\boldsymbol{c}_i\})$ -th elements are $\delta_{jl}-\delta_{jL},\ j=1+\dim\{\boldsymbol{c}_i\},\ l=1,2,\ldots,L-1$, that is, the difference between the coefficients of each qualitative response and the L-th alternative for the individuals' characteristics. This makes it difficult to interpret the multinomial probit coefficients.

Note that in multinomial models, for each alternative specific attribute, it is only required to estimate one coefficient for all alternatives, whereas for individuals' characteristics (non-alternative specific regressors), it is necessary to estimate L-1 coefficients (the coefficient of the base alternative is set equal to 0).

The likelihood function in this model is $p(\boldsymbol{\beta}, \boldsymbol{\Omega} | \boldsymbol{y}, \boldsymbol{R}) = \prod_{i=1}^{N} \prod_{l=1}^{L} p_{il}^{y_{il}}$ where $p_{il} = p(y_{il}^* \geq \max(\boldsymbol{y}_i^*))$.

We assume independent priors, $\beta \sim N(\beta_0, B_0)$ and $\Omega^{-1} \sim W(\alpha_0, \Sigma_0)$. We can employ Gibbs sampling in this model because this is a standard

⁶Note that this model is not identified if Σ is unrestricted. The likelihood function is the same if a scalar random variable is added to each of the L latent regressions.

⁷W denotes the Wishart density.

Bayesian linear regression model when data augmentation in \boldsymbol{w} is used. The posterior conditional distributions are

$$\boldsymbol{\beta}|\boldsymbol{\Omega},\boldsymbol{w}\sim N(\boldsymbol{\beta}_n,\boldsymbol{B}_n),$$

$$\boldsymbol{\Omega}^{-1}|\boldsymbol{\beta},\boldsymbol{w}\sim W(\alpha_n,\boldsymbol{\Sigma}_n),$$
where $\boldsymbol{B}_n=(\boldsymbol{B}_0^{-1}+\boldsymbol{X}^{*\top}\boldsymbol{X}^*)^{-1},\;\boldsymbol{\beta}_n=\boldsymbol{B}_n(\boldsymbol{B}_0^{-1}\boldsymbol{\beta}_0+\boldsymbol{X}^{*\top}\boldsymbol{w}^*),\;\boldsymbol{\Omega}^{-1}=$

$$\boldsymbol{C}^{\top}\boldsymbol{C},\;\boldsymbol{X}_i^{*\top}=\boldsymbol{C}^{\top}\boldsymbol{R}_i,\;\boldsymbol{w}_i^*=\boldsymbol{C}^{\top}\boldsymbol{w}_i,\;\boldsymbol{X}^*=\begin{bmatrix}\boldsymbol{X}_1^*\\\boldsymbol{X}_2^*\\\vdots\\\boldsymbol{X}_N^*\end{bmatrix},\;\alpha_n=\alpha_0+N,\;\boldsymbol{\Sigma}_n=$$

 $(\boldsymbol{\Sigma}_0 + \sum_{i=1}^N (\boldsymbol{w}_i - \boldsymbol{R}_i \boldsymbol{\beta})^{\top} (\boldsymbol{w}_i - \boldsymbol{R}_i \boldsymbol{\beta}))^{-1}.$

We can collapse the multinomial vector \mathbf{y}_i into the indicator variable $d_i = \sum_{l=1}^{L-1} l \times \mathbb{1}_{\max(\mathbf{w}_l) = \mathbf{w}_{il}}$. Then the distribution of $\mathbf{w}_i | \boldsymbol{\beta}, \boldsymbol{\Omega}^{-1}, d_i$ is an L-1 dimensional Gaussian distribution truncated over the appropriate cone in \mathcal{R}^{L-1} . [54] propose drawing from the univariate conditional distributions $\mathbf{w}_{il} | \mathbf{w}_{i,-l}, \boldsymbol{\beta}, \boldsymbol{\Omega}^{-1}, d_i \sim TN_{I_{il}}(m_{il}, \tau_{il}^2)$, where

$$I_{il} = \begin{cases} w_{il} > \max(\boldsymbol{w}_{i,-l}, 0), & d_i = l \\ w_{il} < \max(\boldsymbol{w}_{i,-l}, 0), & d_i \neq l \end{cases},$$

and permuting the columns and rows of Ω^{-1} so that the *l*-th column and row is the last,

$$oldsymbol{\Omega}^{-1} = egin{bmatrix} oldsymbol{\Omega}_{-l,-l} & oldsymbol{\omega}_{-l,l} \ oldsymbol{\omega}_{l,-1} & \omega_{l,l} \end{bmatrix}^{-1} = egin{bmatrix} oldsymbol{\Omega}_{-l,-l}^{-1} + au_{ll}^{-2} oldsymbol{f} oldsymbol{f}^ op & -oldsymbol{f} au_{ll}^{-2} \ - au_{ll}^{-2} oldsymbol{f}^ op & au_{ll}^{-2} \end{bmatrix}$$

where $\boldsymbol{f} = \boldsymbol{\Omega}_{-l,-l}^{-1} \boldsymbol{\omega}_{-l,l}$, $\tau_{ll}^2 = \omega_{ll} - \boldsymbol{\omega}_{l,-l} \boldsymbol{\Omega}_{-l,-1}^{-1} \boldsymbol{\omega}_{-l,l}$, $m_{il} = \boldsymbol{r}_{il}^{\top} \boldsymbol{\beta} + \boldsymbol{f}^{\top} (\boldsymbol{w}_{i,-l} - \boldsymbol{R}_{i,-l} \boldsymbol{\beta})$, $\boldsymbol{w}_{i,-l}$ is an L-2 dimensional vector of all components of \boldsymbol{w}_i excluding w_{il} , \boldsymbol{r}_{il} is the l-th row of \boldsymbol{R}_i , $l = 1, 2, \ldots, L-1$.

The identified parameters are obtained by normalizing with respect to one of the diagonal elements $\frac{1}{\omega_{1,1}^{0.5}}\boldsymbol{\beta}$ and $\frac{1}{\omega_{1,1}}\boldsymbol{\Omega}.^9$

Example: Choice of fishing mode

Let's do the famous example of *choice of fishing mode* from [14].

6.5 Summary

We present ...

⁸Observe that the identification issue in this model is due to scaling w_{il} by a positive constant does not change the value of d_i .

⁹Our GUI is based on the *bayesm* package that takes into account this identification restriction to display the outcomes of the posterior chains.

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6.6 Exercises

1. Get the posterior conditional distributions of the Gaussian linear model assuming independent priors $\pi(\boldsymbol{\beta}, \sigma^2) = \pi(\boldsymbol{\beta}) \times \pi(\sigma^2)$, where $\boldsymbol{\beta} \sim N(\boldsymbol{\beta}_0, \mathbf{B}_0)$ and $\sigma^2 \sim IG(\alpha_0/2, \delta_0/2)$.

- 2. Show that the posterior conditional distributions of the Gaussian linear model with heteroskedasticity assuming independent priors $\pi(\boldsymbol{\beta}, \sigma^2, \tau) = \pi(\boldsymbol{\beta}) \times \pi(\sigma^2) \times \prod_{i=1}^N \pi(\tau_i)$, where $\boldsymbol{\beta} \sim N(\boldsymbol{\beta}_0, \mathbf{B}_0)$, $\sigma^2 \sim IG(\alpha_0/2, \delta_0/2)$ and $\tau_i \sim G(v/2, v/2)$ are $\boldsymbol{\beta}|\sigma^2, \tau, \mathbf{y}, \mathbf{X} \sim N(\boldsymbol{\beta}_n, \mathbf{B}_n)$, $\sigma^2|\boldsymbol{\beta}, \tau, \mathbf{y}, \mathbf{X} \sim IG(\alpha_n, \delta_n)$ and $\tau_i|\boldsymbol{\beta}, \sigma^2, \mathbf{y}, \mathbf{X} \sim G(v_{1n}, v_{2in})$, where $\tau = [\tau_1, \dots, \tau_n]^\top$, $\mathbf{B}_n = (\mathbf{B}_0^{-1} + \sigma^{-2}\mathbf{X}^\top \mathbf{\Psi} \mathbf{X})^{-1}$, $\boldsymbol{\beta}_n = \mathbf{B}_n(\mathbf{B}_0^{-1} \boldsymbol{\beta}_0 + \sigma^{-2}\mathbf{X}^\top \mathbf{\Psi} \mathbf{y})$, $\alpha_n = \alpha_0 + N$, $\delta_n = \delta_0 + (\mathbf{y} \mathbf{X} \boldsymbol{\beta})^\top \mathbf{\Psi} (\mathbf{y} \mathbf{X} \boldsymbol{\beta})$, $v_{1n} = v + 1$, $v_{2in} = v + \sigma^{-2}(y_i \mathbf{x}_i^\top \boldsymbol{\beta})^2$, and $\mathbf{\Psi} = \text{diagonal } \{\tau_i\}$.
- 3. The market value of soccer players in Europe continues

Use the setting of the previous exercise to perform inference using a Gibbs sampling algorithm of the the market value of soccer players in Europe setting v=5 and same other hyperparameters as the homoscedastic case. Is there any meaningful difference for the coefficient associated with the national team compared to the application in the homoscedastic case?

4. Example: Determinants of hospitalization continues

Program a Gibbs sampling algorithm in the application of determinants of hospitalization.

Multivariate models

Time series models

Panel data models

Bayesian model average

- 10.1 Calculating the marginal likelihood
- 10.1.1 Savage-Dickey density ratio
- 10.1.2 Gelfand-Dey method
- 10.1.3 Chib's methods

Part III

Advanced methods: Theory, applications and programming

Hierarchical models

11.1 Direchlet processes

Causal inference

Machine learning

- 13.1 Cross validation and Bayes factors
- 13.2 Regularization
- 13.3 Bayesian additive regression trees
- 13.4 Gaussian processes

 $Spatial\ econometric\ models$

Further topics

15.1	Approximate	Bayesian	computation

- 15.2 Variational Bayes
- 15.3 Integrated nested Laplace approximations
- 15.4 Bayesian exponential tilted empirical likelihood

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