TDDE07 - Lab 2

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1. Linear and polynomial regression

1a)

Before looking at the data, we came to the conclusion that the intercept should be about -10 degrees, with a cyclic temperature throughout the year and a maximum at 7 months. The variance was adjusted in order for the prior regression curve to be in accordance with our prior beliefs. Draws were made from the prior and plotted along with the temperature readings for comparison, as can be seen in Figure 1.

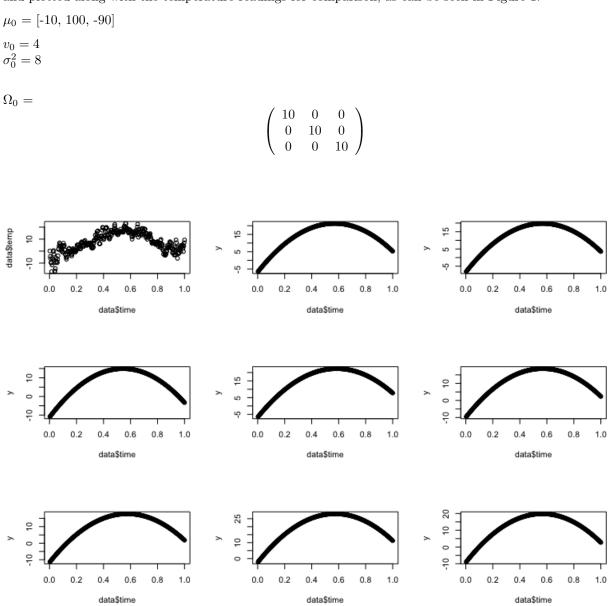


Figure 1: Plotting the chosen hyperparameters

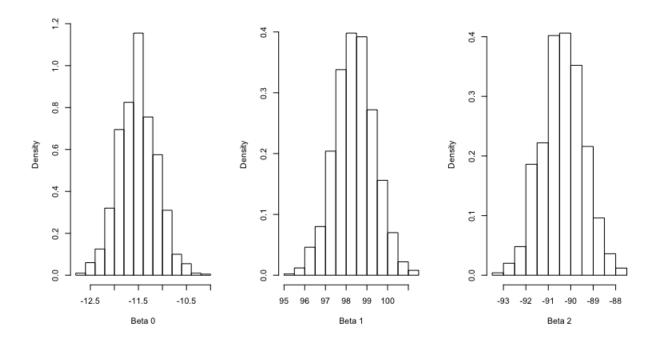
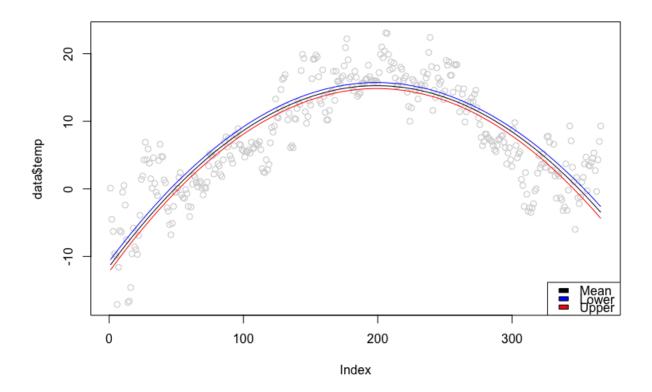


Figure 2: Marginal posterior of the beta parameters

The figure below shows the plotted data along with the curve for the posterior median of the regression function. Furthermore, a 95 % credible interval has been calculated. The interval bands doesn't contain most of the data points due to the high credibility.



 $Figure \ 3: \ Posterior \ credible \ intervals$

1c)

The highest expected temperature was calculated using the derivative of the time function, with respect to time.

$$\frac{\partial f}{\partial t} = \beta_1 + 2\beta_2 t = 0 \iff t = -\frac{\beta_1}{2\beta_2}$$
$$0 \le t \le 1 \Rightarrow 0 \le \bar{x} \le 366 \iff \bar{x} = 366 * t$$

The histogram of \bar{x} can be seen in Figure 4.

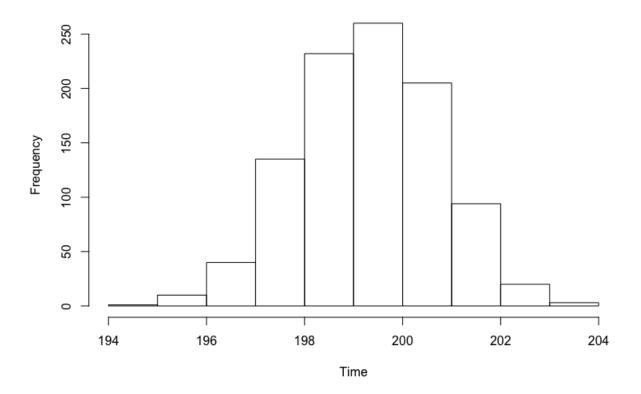


Figure 4: \bar{x}

	Min.	1st Qu.	Median.	Mean	3rd Qu.	Max.
1	194.54	198.30	199.27	199.29	200.26	203.37

Table 1: Summary of \bar{x}

This is in line with the output from which.max(y_med) = 200, where y_med is the posterior mean from a).

1d)

Using the posterior parameters from 1b) as the first three betas, the new beta parameters would be set to zero due to suspicion that the introduced variables might not be needed. Regarding the covariance matrix, we want low variance (high bias) to avoid overfitting, giving high diagonal values in Ω_0 for the new variables, with the rest set to zero since we can't say anything about the covariance between the new variables.

β_0 .	β_1	β_2	β_3	β_4	β_5	β_6	β_7
-11.53113	98.43113	-90.38911	0	0	0	0	0

Table 2: μ_0

$\Omega_0 =$									
	/ 376	183.5	122.5	0	0	0	0	0 \	
	183.5	132.5	92.5	0	0	0	0	0	
	122.5	92.5	83.7	0	0	0	0	0	
	0	0	0	100	0	0	0	0	
	0	0	0	0	100	0	0	0	
	376 183.5 122.5 0 0	0	0	0	0	100	0	0	

2. Posterior approximation for classification with logistic regression

2b)

Numerical values for beta:

β_0 .	β_1	β_2	β_3	β_4	β_5	β_6	β_7
0.62672884	0.01979113	0.18021897	0.16756670	-0.14459669	-0.08206561	-1.35913317	-0.02468351

Table 3: β

Numerical values for the hessian:

Hessian =

$\left(\begin{array}{c} 2.2660225\\ 0.0033388 \end{array}\right)$	0.00000		-1.1791e - 02 -3.1254e - 05	$\begin{array}{c} 0.04578072 \\ 0.00014149 \end{array}$	-3.0293e - 02 -3.5885e - 05		-0.098023 -0.000144
-0.065451		6.2181e - 03		0.00189628	-3.2404e - 06		0.0017527
$\begin{bmatrix} -0.011791 \\ 0.0457807 \end{bmatrix}$		-3.5582e - 04			-1.3408e - 04 -3.2993e - 04	0.00==000	0.0005437 0.0005120
-0.030293			-1.4249e - 02 -1.3408e - 04		-3.2993e - 04 7.1846e - 04	0.00520825 0.0051841	0.0003120 0.0010952
-0.188748	3 5.0668e - 04	-6.1345e - 03	-1.4689e - 03	0.00320825	5.1841e - 03	0.1512621	0.0067688
$\setminus -0.098023$	9 -1.4442e - 04	1.7527e - 03	5.4371e - 04	0.00051201	1.0952e - 03	0.0067688	0.0199722

Posterior distribution of NSmallChild parameter

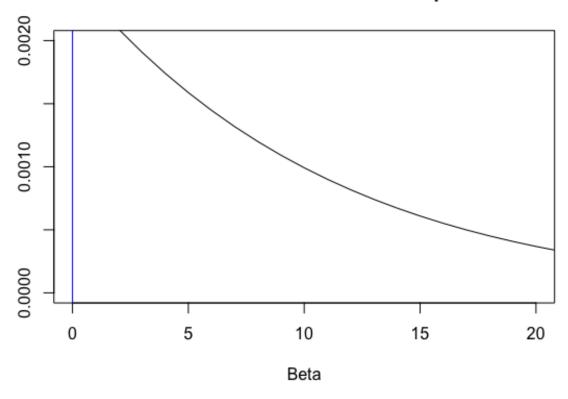


Figure 5: Plotting the chosen hyperparameters

According to the figure, this parameter is very close to zero and thus, this variable is not of importance.

2c)

Given input specified in Table 3, the probability was calculated by collecting sample draws from the beta posterior distribution and counting the occurrences where $\mathtt{Work} = 1$ and dividing it with the number of draws. This generated a probability p = 22.3%.

husbandInc.	educYears	$\exp Y ears$	expYears2	age	nSmallChild	nBigChild
10	8	10	1	40	1	1

Table 4: μ_0

1 - Code

```
library(mvtnorm)
library(invgamma)
data = read.table("TempLinkoping.txt", header=TRUE)
####### A #######
I \leftarrow diag(3)
mu < -c(-10, 100, -90)
omega <- 10 * I
v <- 4
s2 <- 8
x \leftarrow seq(0, 10, by=0.01)
# Inverse chisquare from lab1
draw sigma <- function(v, s2) {</pre>
  X_draw <- rchisq(1, v)</pre>
  return (v * s2 / X_draw)
# Given temperature regression function
temp <- function(beta, time) {</pre>
  return (beta[1] + beta[2] * time + beta[3] * (time ^ 2)) + rnorm(0,1)
par(mfrow=c(3,3))
plot(data$time, data$temp)
for (i in 1:8) {
  sigma_sq = draw_sigma(v, s2)
  beta = rmvnorm(1, mu, sigma_sq*solve(omega))
  y = temp(beta[1,], data$time)
  plot(data$time, y)
dev.off()
####### B #######
n_draws <- 1000
ones <- c(rep(1, length(data$time)))</pre>
x2 <- data$time^2</pre>
X <- cbind(ones, data$time, x2)</pre>
beta_hat <- solve(t(X)%*%X)%*%t(X)%*%data$temp
omega_n \leftarrow t(X) \%*\% X + omega
mu_n <- solve(omega_n) %*% (t(X) %*% X %*% beta_hat + omega %*% mu)</pre>
v_n <- v + length(data$time)</pre>
s2_n <- (v*s2 + (t(data\$temp)%*%data\$temp + t(mu)%*%omega%*%mu - t(mu_n)%*%omega_n%*%mu_n))/v_n
sigma_post = draw_sigma(v_n, s2_n)
```

```
beta_post = rmvnorm(n_draws, mu_n, solve(omega_n)*sigma_post[1])
Y = matrix(nrow=n_draws, ncol=366)
for (i in 1:n_draws) {
 Y[i,] = temp(beta_post[i,], data$time)
y_med = c()
y_up = c()
y_low = c()
for (i in 1:366) {
  y_med = c(y_med, median(Y[,i]))
  y_low = c(y_low, quantile(Y[,i], 0.025))
 y_{up} = c(y_{up}, quantile(Y[,i], 0.975))
par(mfrow=c(1,3))
hist(beta_post[,1], freq = FALSE, xlab = "Beta 0", main='')
hist(beta_post[,2], freq = FALSE, xlab = "Beta 1", main='')
hist(beta_post[,3], freq = FALSE, xlab = "Beta 2", main='')
dev.off()
plot(data$temp, type='p', col='lightgray')
lines(y_med, type='l')
lines(y_low, type='l', col='red')
lines(y_up, type='l', col='blue')
legend("bottomright",
       legend = c("Mean", "Lower", "Upper"),
       fill = c("black", "blue", "red"))
####### C #######
x_max = -366 * beta_post[,2]/(2 * beta_post[,3])
hist(x_max, main='', xlab='Time')
dev.off()
####### D #######
# Prior
\# Mu_0 = [...mu, 0, 0, 0, 0] due to suspicion that the introduced variables might not be needed
# Omega_0, want low variance (high bias to avoid overfitting) => High diagonal values in omega_Ofor the
new_mu = c(mu_n, rep(0, 5))
new_omega <- 100 * diag(8)
new_omega[1:3,1:3] = omega_n
```

2 - Code

```
womenData = read.table("WomenWork.txt", header=TRUE)
####### A #######
# Added a zero in the model formula so that R doesn't add an extra intercept
# A . in the model formula means to add all other variables in the dataset as features
glmModel <- glm(Work ~ 0 + ., data = womenData, family = binomial)</pre>
####### B #######
# Param setup
tau <- 10
nParams <- 8
mu <- as.vector(rep(0, nParams))</pre>
X <- as.matrix(womenData[,2:9])</pre>
y <- as.matrix(womenData[,1])</pre>
initVal <- as.vector(rep(0,nParams))</pre>
sigma <- tau^2 * diag(nParams)</pre>
initBetas <- rep(0,nParams)</pre>
# Calculate the Log Posterior Logistic
logPost <- function(beta,y,X,mu,Sigma) {</pre>
  # Calculate predictions by multiplying data with betas
  pred <- X%*%beta
  # Log likelihood function, derived by taking the log of the likelihood function prod(e^pred*y/(1+e^pr
  logLik <- sum(y * pred - log(1 + exp(pred)))</pre>
  logPrior <- dmvnorm(beta, matrix(0,nParams,1), Sigma, log=TRUE);</pre>
  return (logPrior + logLik)
}
OptimResults<-optim(initBetas,logPost,gr=NULL,y,X,mu,sigma,method=c("BFGS"),control=list(fnscale=-1),he
postCov <- -solve(OptimResults$hessian)</pre>
st_div <- sqrt(diag(postCov))</pre>
betaMode <- OptimResults$par</pre>
nSmallChild <- postDraws[,7]</pre>
postDist \leftarrow dnorm(x = seq(0,10,0.01), mean = betaMode[7], sd = st_div[7])
# plot posterior distribution NSmallChild parameter
cred_int <- quantile(postDraws, c(0.025, 0.975))</pre>
plot(postDist,
     xlim = c(0,100),
     ylim = c(0,0.002),
     type = '1')
abline(v = cred_int[1], col='red')
```

```
abline(v = cred_int[2], col='blue')
######## C #########
y \leftarrow c(constant = 1,
       husbandInc = 10,
       educYears = 8,
       expYears = 10,
       expYears2 = 1,
       age = 40,
       nSmallChild = 1,
       nBigChild = 1)
n_draws <- 1000
predDist <- function(n, beta, sigma, y) {</pre>
 y_draws = c()
  for (i in 1:n) {
    #beta draw
    betaDraw = as.vector(rmvnorm(1, mean = beta, sigma = sigma))
    #probability
    p <- exp(y%*%betaDraw)/(1+exp(y%*%betaDraw))</pre>
    #prediction of y
   y_{draw} = rbinom(1, 1, p)
    y_draws = c(y_draws, y_draw)
 }
 return (y_draws)
draws = predDist(n_draws, betaMode, postCov, y)
workProb <- sum(draws == 1)</pre>
```