## Bayesian Learning

Lab 2

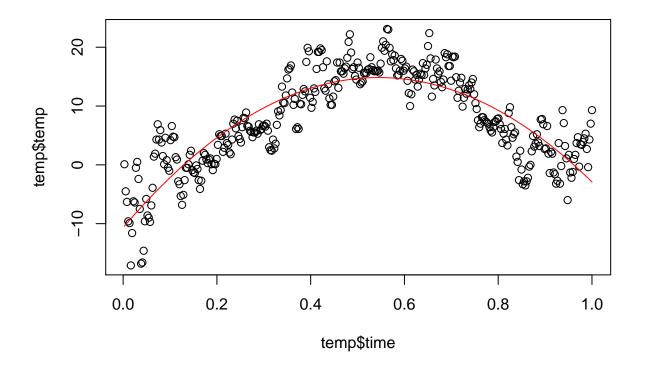
Emil K Svensson and Rasmus Holm 2017-05-02

## Question 1

```
temp <- read.table("../data/TempLinkoping2016.txt", header=T)
mod <- lm(temp ~ time + I(time^2), data=temp)

idx <- order(temp$time)
x <- temp$time[idx]
y <- fitted(mod)[idx]

plot(temp$time, temp$temp)
lines(x, y, col='red', type='l')</pre>
```



Prior

$$\begin{split} \sigma^2 &\sim \text{Inv} - \chi^2(\nu_0, \sigma_0^2) \\ \beta | \sigma^2 &\sim N(\mu_0, \sigma^2 \Omega_0^{-1}) \end{split}$$

Likelihood

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

Posterior

$$\sigma^2 | \mathbf{y} \sim \text{Inv} - \chi^2(\nu_n, \sigma_n^2)$$
  
 $\beta | \sigma^2, \mathbf{y} \sim \text{N}(\mu_n, \sigma^2 \Omega_n^{-1})$ 

where

$$\mu_n = (\mathbf{X}^{\mathsf{T}} \mathbf{X} + \Omega_0)^{-1} (\mathbf{X}^{\mathsf{T}} \mathbf{X} \hat{\beta} + \Omega_0 \mu_0)$$

$$\Omega_n = \mathbf{X}^{\mathsf{T}} \mathbf{X} + \Omega_0$$

$$\nu_n = \nu_0 + n$$

$$\nu_n \sigma_n^2 = \nu_0 \sigma_0^2 + (\mathbf{y}^{\mathsf{T}} \mathbf{y} + \mu_0^{\mathsf{T}} \Omega_0 \mu_0 - \mu_n^{\mathsf{T}} \Omega_n \mu_n)$$

**a**)

```
mu0 <- c(0, 0, 0)
omega0 <- diag(3) * 0.05
nu0 <- 1
sigmasq0 <- 20
hyperparams <- list(mu=mu0, omega=omega0, nu=nu0, sigmasq=sigmasq0)</pre>
```

b)

```
library(geoR)
library(MASS)

time <- data.frame(rep(1,nrow(temp)), temp$time, temp$time^2)

mtime <- as.matrix(time)

mtemp <- matrix(temp$temp, ncol = 1)

prior_estimate <- function(data, params) {
    sigmasq <- rinvchisq( n = 1, df = params$nu, scale = params$sigmasq)
    betacoef <- mvrnorm(n = 1, mu = params$mu, Sigma = sigmasq * solve(params$omega) )

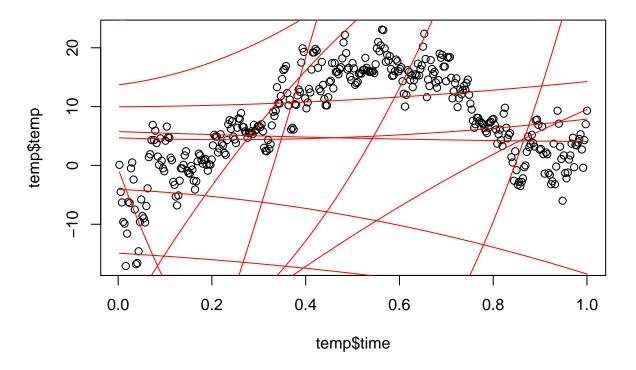
    data %*% betacoef
}</pre>
```

```
plot(temp$time, temp$temp)

x <- sort(temp$time)

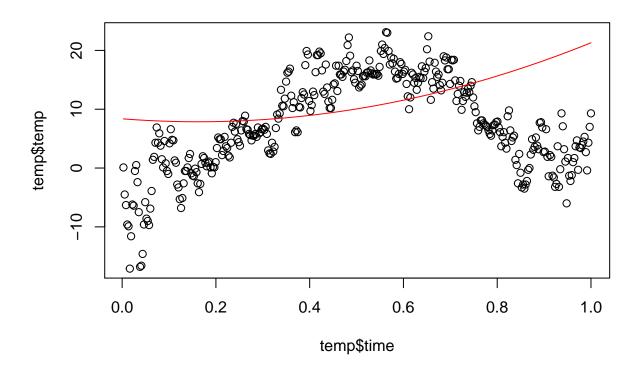
set.seed(12345)

for (i in 1:20){
    y <- prior_estimate(mtime, hyperparams)[order(temp$time)]
    lines(x, y, col='red', type='l')
}</pre>
```



```
set.seed(12345)

x <- sort(temp$time)
y <- rowMeans(sapply(1:1000, function(x) prior_estimate(mtime, hyperparams)[order(temp$time)]))
plot(temp$time, temp$temp)
lines(x, y, col='red', type='l')</pre>
```



**c**)

```
posterior_param_sample <- function(X, y, hyperparams){
    XX <- t(X) %*% X

    betahat <- solve(XX) %*% t(X) %*% y

mun <- solve(XX + hyperparams$omega) %*%
        (XX %*% betahat + hyperparams$omega %*% hyperparams$mu)

omegan <- XX + hyperparams$omega

nun <- hyperparams$nu + nrow(X)

nunsigmasqn <- hyperparams$nu * hyperparams$sigmasq +
        (t(y) %*% y +
        t(hyperparams$mu) %*% hyperparams$omega %*% hyperparams$mu -
        t(mun) %*% omegan %*% mun )

sigmasqn <- nunsigmasqn / nun

sigmasq <- rinvchisq(n = 1, df=nun, scale=sigmasqn)
    beta <- mvrnorm(n = 1, mu = mun, Sigma = as.numeric(sigmasq) * solve(omegan))</pre>
```

```
list(beta = beta, sigmasq = sigmasq)
}

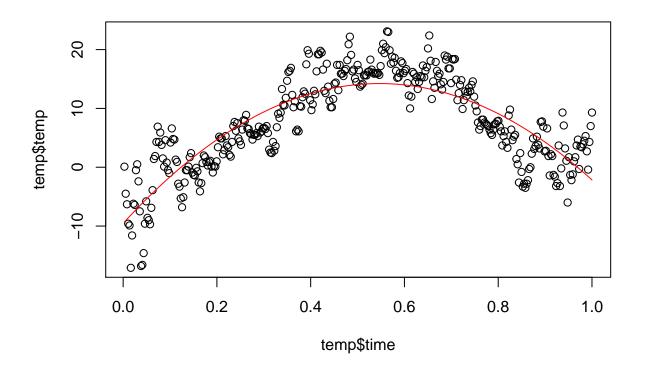
posterior_estimate <- function(X, y, hyperparams){
    sample <- posterior_param_sample(X, y, hyperparams)
    X %*% sample$beta
}

plot(temp$time, temp$temp)

set.seed(12345)

idx <- order(temp$time)
x <- temp$time[idx]
y <- posterior_estimate(mtime, mtemp, hyperparams)[idx]

lines(x, y, col='red', type='l')</pre>
```



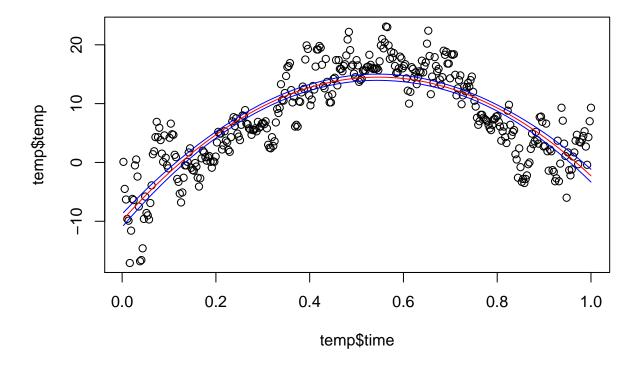
```
set.seed(12345)

ests <- sapply(1:1000, FUN = function(x) posterior_estimate(mtime, mtemp, hyperparams))
cred_interval <- apply(ests, MARGIN = 1, quantile, probs = c(0.05, 0.95))

idx <- order(temp$time)
x <- temp$time[idx]
y1 <- rowMeans(ests)[idx]</pre>
```

```
y2 <- cred_interval[1,][idx]
y3 <- cred_interval[2,][idx]

plot(temp$time, temp$temp)
lines(x, y1, col='red', type='l')
lines(x, y2, col='blue', type='l')
lines(x, y3, col='blue', type='l')</pre>
```



d)

```
set.seed(12345)
betas <- sapply(1:1000, FUN = function(x) posterior_param_sample(mtime, mtemp, hyperparams)$beta)
hot <- mean(-betas[2,] / (2 * betas[3,]))
hot * 366 # July 27, 2016 (Wed)
## [1] 200.0608</pre>
```

**e**)

Set  $\mu_0$  to zeros and a high  $\Omega_0$  that expresses a high degree of certainty in our prior.

## Question 2

```
women <- read.table("../data/WomenWorks.txt", header = TRUE)</pre>
a)
glmModel <- glm(Work ~ 0 + ., data = women, family = binomial)</pre>
summary(glmModel)
##
## Call:
## glm(formula = Work ~ 0 + ., family = binomial, data = women)
## Deviance Residuals:
      Min
                10
                     Median
                                   30
                                           Max
## -2.1662 -0.9299 0.4391
                               0.9494
                                        2.0582
## Coefficients:
##
              Estimate Std. Error z value Pr(>|z|)
## Constant
                          1.52307
                                    0.423 0.672274
              0.64430
## HusbandInc -0.01977
                           0.01590 -1.243 0.213752
## EducYears 0.17988
                           0.07914
                                    2.273 0.023024 *
## ExpYears
               0.16751
                           0.06600
                                    2.538 0.011144 *
## ExpYears2
               -0.14436
                           0.23585
                                    -0.612 0.540489
              -0.08234
                           0.02699 -3.050 0.002285 **
## Age
## NSmallChild -1.36250
                           0.38996 -3.494 0.000476 ***
## NBigChild
             -0.02543
                           0.14172 -0.179 0.857592
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
      Null deviance: 277.26 on 200 degrees of freedom
##
## Residual deviance: 222.73 on 192 degrees of freedom
## AIC: 238.73
## Number of Fisher Scoring iterations: 4
b)
library(mvtnorm)
logprior <- function(beta, mean, sigma){</pre>
    dmvnorm(beta, mean = mean, sigma = sigma, log = TRUE)
}
loglikelihood <- function(beta, X, Y){</pre>
   linear_prediction <- t(X) %*% beta</pre>
   probabilities <- (Y * linear_prediction) - log(1 + exp(linear_prediction))</pre>
```

```
loglike <- sum(probabilities)</pre>
    ## if (abs(loglike) == Inf)
           loglike = -20000
    loglike
}
logposterior <- function(beta, X, Y, mean, sigma){</pre>
   loglikelihood(beta, X, Y) + logprior(beta, mean, sigma)
}
tau <- 10
mu < -rep(0,8)
sigma \leftarrow tau^2 * diag(8)
womenX <- as.matrix(women[,2:ncol(women)])</pre>
womenY <- as.matrix(women[,1])</pre>
optpost <- optim(par = matrix(rep(0, 8), ncol = 1),
                  fn = logposterior, method = "BFGS", hessian = TRUE,
                  X = t(womenX), Y = womenY,
                  mean = mu, sigma = sigma,
                  control=list(fnscale=-1))
posterior_beta_sample <- function(n, mu, sigma){</pre>
    rmvnorm(n, mean = mu, sigma = sigma)
}
mu <- optpost$par</pre>
sigma <- -solve(optpost$hessian)</pre>
set.seed(12345)
beta_samples <- posterior_beta_sample(n=1000, mu=mu, sigma=sigma)
cred_intervals <- apply(beta_samples, 2, quantile, prob=c(0.025, 0.975))</pre>
colnames(cred_intervals) <- colnames(women)[-1]</pre>
```

The 95% credibility intervals below show that the intercept, husband income, transformed years in work experience, and the number of children older than 6 years are statistical insignificant, i.e. 0 (no impact) is contained in the intervals. This indicates that it may be better to remove those variables from the model.

## cred\_intervals

**c**)

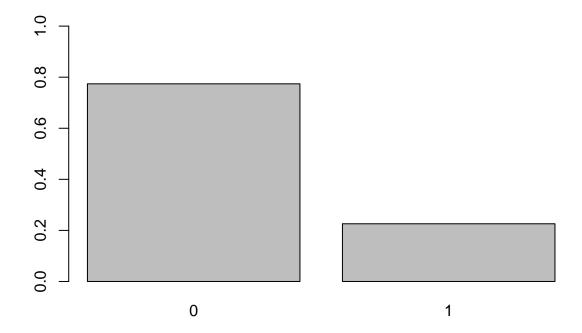
```
Constant HusbandInc EducYears
                                          ExpYears ExpYears2
                                                                      Age
## 2.5% -2.313434 -0.05091541 0.03095675 0.04723847 -0.6143409 -0.13476231
## 97.5% 3.533276 0.00780642 0.32739377 0.29656573 0.2943841 -0.03095853
        NSmallChild NBigChild
##
## 2.5%
         -2.1296887 -0.3023168
## 97.5% -0.5759195 0.2692414
```

8

```
posterior_predictive_sample <- function(X, beta){
    linear_prediction <- t(X) %*% beta
    probability <- exp(linear_prediction) / (1 + exp(linear_prediction))
    rbinom(n=1, size=1, prob=probability)
}

x <- matrix(c(1, 10, 8, 10, (10 / 10)^2, 40, 1, 1), ncol=1)
prediction_samples <- apply(beta_samples, 1, function(beta) posterior_predictive_sample(x, beta))

counts <- table(prediction_samples)
barplot(counts / sum(counts), ylim=c(0, 1))</pre>
```



The barplot above indicates that it is much more probable that given the data the woman is not working under our model.