## Bayesian Learning

Lab 2

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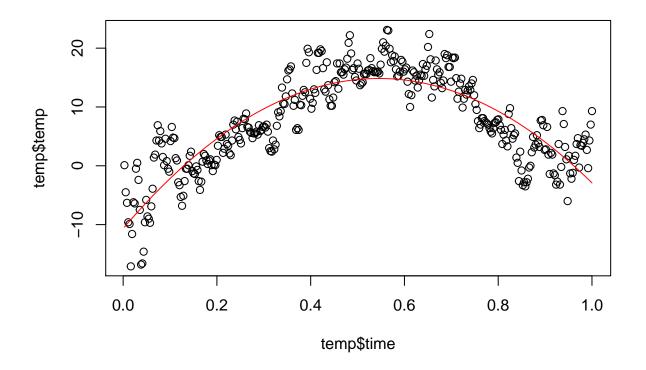
## Question 1

```
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) )

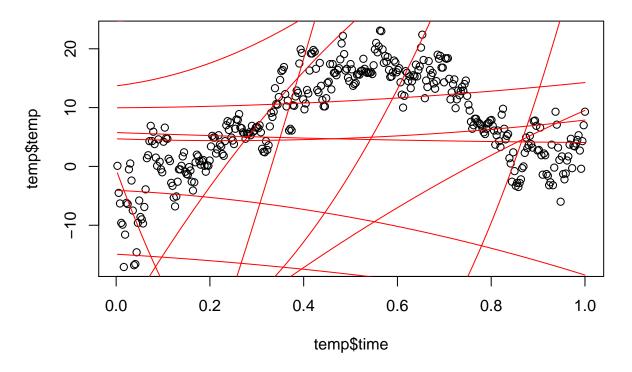
    data %*% betacoef
}</pre>
```

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

x <- sort(temp$time)

set.seed(12345)

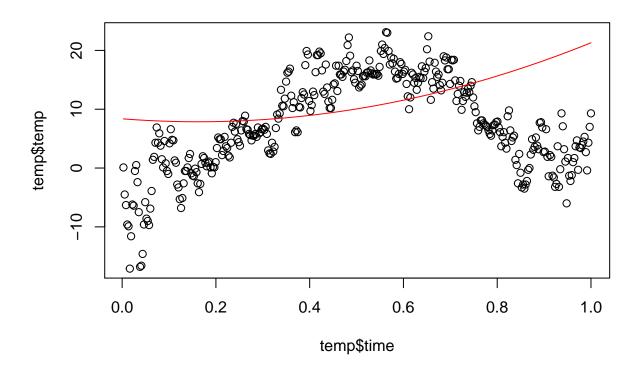
for (i in 1:20){
    y <- prior_estimate(mtime, hyperparams)[order(temp$time)]
    lines(x, y, col='red', type='l')
}</pre>
```



```
set.seed(12345)

x <- sort(temp$time)
y <- rowMeans(sapply(1:1000, function(x) prior_estimate(mtime, hyperparams)[order(temp$time)]))

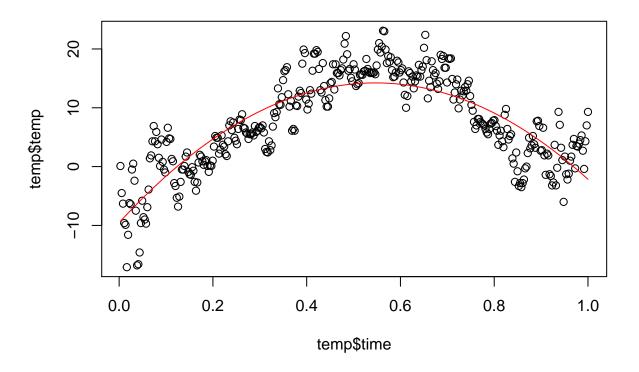
plot(temp$time, temp$temp)
lines(x, y, col='red', type='l')</pre>
```



**c**)

```
posterior_param_sample <- function(X, y, hyperparams){</pre>
    XX \leftarrow t(X) %% X
    betahat <- solve(XX) %*% t(X) %*% y
    mun <- solve(XX + hyperparams$omega) %*%</pre>
        (XX %*% betahat + hyperparams$omega %*% hyperparams$mu)
    omegan <- XX + hyperparams$omega
    nun <- hyperparams$nu + nrow(X)</pre>
    \verb|nunsigmasqn <- hyperparams$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)</pre>
    beta <- mvrnorm(n = 1, mu = mun, Sigma = as.numeric(sigmasq) * solve(omegan))
    list(beta = beta, sigmasq = sigmasq)
}
posterior_estimate <- function(X, y, hyperparams){</pre>
    sample <- posterior_param_sample(X, y, hyperparams)</pre>
    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>
```



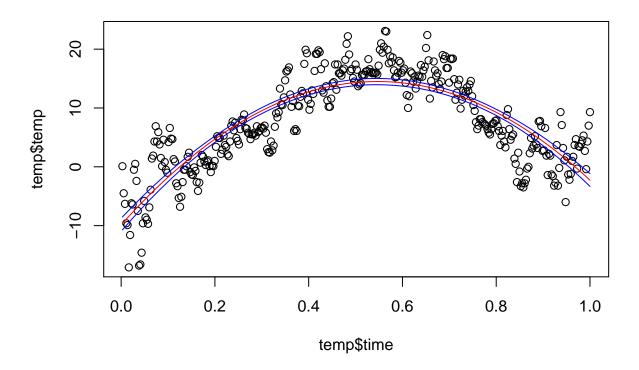
```
set.seed(12345)

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]
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')</pre>
```

```
lines(x, y3, col='blue', type='l')
```



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 \leftarrow rep(0,8)
sigma <- tau^2 * diag(8)
womenX <- as.matrix(women[,2:ncol(women)])</pre>
womenY <- as.matrix(women[,1])</pre>
optpost <- optim(par = matrix(rep(0.1,8), ncol = 1),</pre>
                  fn = logposterior, method = "BFGS", hessian = TRUE,
                  X = t(womenX), Y = womenY,
                  mean = mu, sigma = sigma,
                  control=list(fnscale=-1))
posterior_sample <- function(mu, sigma){</pre>
    rmvnorm(1, mean = mu, sigma = sigma)
}
mu <- optpost$par</pre>
sigma <- -solve(optpost$hessian)</pre>
samples <- sapply(1:1000, FUN = function(x) posterior_sample(mu=mu, sigma=sigma))</pre>
cred_intervals <- apply(samples, 1, quantile, prob=c(0.025, 0.975))</pre>
cred_intervals
##
                            [,2]
                                       [,3]
                                                   [,4]
                                                               [,5]
                                                                            [,6]
               [,1]
## 2.5% -2.357327 -0.05077317 0.0395291 0.04207453 -0.5765345 -0.13383125
## 97.5% 3.320715 0.01122911 0.3340529 0.28791992 0.3215241 -0.02841438
               [,7]
## 2.5% -2.097121 -0.3019207
## 97.5% -0.621485 0.2865123
c)
```