## Lab 7

#### PSTAT 115, Fall 2019

Nov. 20, 2019

# **Objectives**

• Bayesian simple linear regression using stan

## Simple linear regression model

The simple linear regression model is:

$$y_i = \beta x_i + \alpha + \epsilon_i$$
  $i = 1, \dots, n$ 

 $\epsilon_i \sim \text{normal}(0, \sigma^2)$  and  $\epsilon_i$ 's are independent.

This model can be written as

$$y_i - \beta x_i - \alpha \sim \text{normal}(0, \sigma^2)$$

which means

$$y_i \sim \text{normal}(\beta x_i + \alpha, \sigma^2)$$

. And we will use this last expression of the model to code simple linear regression with stan.

Now the parameters that we have are  $\beta$ ,  $\alpha$  and  $\sigma$ . In frequentist view, we find the likelihood and compute the MLE then mission's complete. But in Bayesian context, we can specify some priors, find the posterior distribution and estimate those parameters with posterior means/medians/modes etc. Of course, the posterior distribution might be complicated and we need to turn to MCMC for help. That's when we would like to use stan to tackle problems.

We code the model in a stan file as follows. Note: in stan, if you don't specifiy the prior, it will use the improper priors for the unbounded parameters.

```
data {
  int<lower=0> N;
  vector[N] x;
  vector[N] y;
}

parameters {
  real alpha;
  real beta;
  real<lower=0> sigma;
}

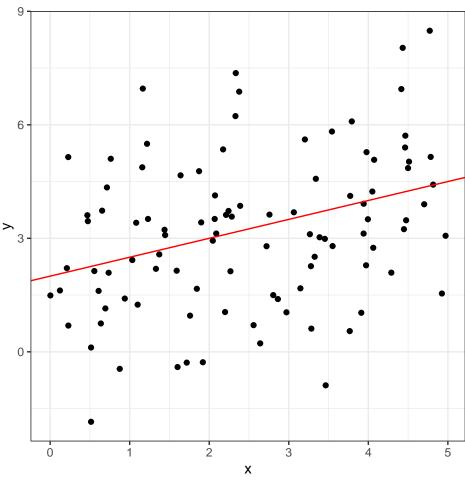
model {
  y ~ normal(alpha + x*beta, sigma);
}
```

## Stan Example

#### Fitting Stan model

Important Code for HW

```
library(rstan)
set.seed(123)
x <- runif(100, 0,5)
a <- 2
b <- 0.5
y <- b*x+a+rnorm(100, mean = 0, sd = 2)
n <- length(y)
ggplot(data = data.frame(x=x,y=y))+geom_point(aes(x=x,y=y))+geom_abline(slope = b, intercept = a, col =</pre>
```



Now let's look at the estimates of posterior means.

mean(beta\_samples)

## [1] 0.4615147

mean(alpha\_samples)

## [1] 1.989233

mean(sigma\_samples)

## [1] 1.960732

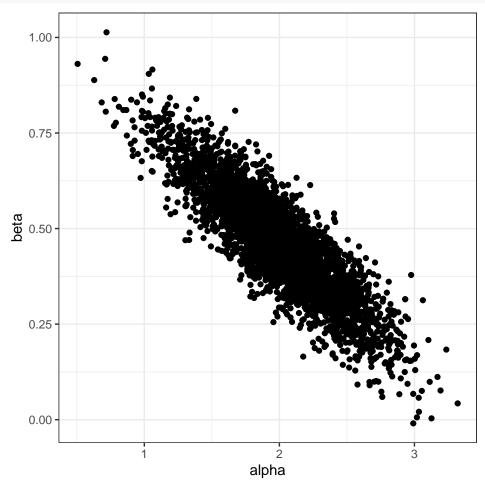
Let's say if we are interested in the predict value when x=5, that is  $\hat{y}=\alpha+5\beta$ . Then we can get samples of this predict value. The sample mean will give us a good estimate of the posterior mean of this predict value at x=5.

```
y_hat <- alpha_samples + 5*beta_samples
mean(y_hat)</pre>
```

#### ## [1] 4.296807

We can also explore the posterior relationship between  $\alpha$  and  $\beta$ . What is the relationship between these two parameters a posteriori? Can you explain intuitively why this relationship exists?

ggplot(tibble(alpha=alpha\_samples, beta=beta\_samples)) + geom\_point(aes(x=alpha, y=beta)) + theme\_bw()



#### Making 60% and 90% confidence bands plot for y

Important Code for HW

```
xgrid \leftarrow seq(0, 5, by=0.1)
xgrid
## [1] 0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0 1.1 1.2 1.3 1.4 1.5 1.6 1.7 1.8
## [20] 1.9 2.0 2.1 2.2 2.3 2.4 2.5 2.6 2.7 2.8 2.9 3.0 3.1 3.2 3.3 3.4 3.5 3.6 3.7
## [39] 3.8 3.9 4.0 4.1 4.2 4.3 4.4 4.5 4.6 4.7 4.8 4.9 5.0
compute curve <- function(sample) {</pre>
  alpha <- sample[1]</pre>
  beta <- sample[2]</pre>
  y_values <- alpha + beta*xgrid
res <- apply(cbind(alpha_samples, beta_samples), 1, compute_curve)
#each col of res is for a set of alpha, beta values
\#each row of res is different y values computed from different sets of alpha beta for a fixed x values.
quantiles <- apply(res, 1, function(x) quantile(x, c(0.05, 0.20, 0.80, 0.95)))
posterior_mean <- rowMeans(res)</pre>
posterior_mean <- apply(res, 1, median)</pre>
tibble(x=xgrid,
q05=quantiles[1,],
q20=quantiles[2,],
q80=quantiles[3,],
q95=quantiles[4,],
mean=posterior_mean) %>%
ggplot() +
geom_ribbon(aes(x=xgrid, ymin=q05, ymax=q95), alpha=0.2) +
geom_ribbon(aes(x=xgrid, ymin=q20, ymax=q80), alpha=0.5) +
geom_line(aes(x=xgrid, y=posterior_mean), size=1) +
theme_bw()
```

