#### Linear regression using Stan

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### But what does linear regression mean? (1)

$$y \sim \mathcal{D}(\boldsymbol{\theta})$$

- ightharpoonup where heta indicates a vector of parameters;
- In standard **linear** regression models, we want to specify a functional from for  $\mathbb{E}[y|X=x]$  **independently** of the the (conditional) variance  $\mathbb{V}(y|X=x)$ ;
- There are only a few choices of D that allow to do so;
- ► The most popular one is the Normal distribution. Other options include the Student's t and the Double exponential distribution;
- The former is by far the most widely adopted;
- ▶ Therefore,  $\theta = [\mu, \sigma]$  and:

$$y \sim N(\mu, \sigma)$$

### But what does linear regression mean? (2)

For i = 1, ..., N independent observations, we typically write:

$$y_i \sim N(\mu_i, \sigma)$$
  
 $\mu_i = \mathbb{E}[y|X = x] = \alpha + \sum_{p=1}^P \beta_p x_{p,i}$ 

or, equivalently:

$$y_i = \mu_i + \varepsilon_i$$

$$\mu_i = \mathbb{E}[y|X = x] = \alpha + \sum_{p=1}^{P} \beta_p x_{p,i}$$

$$\varepsilon_i \sim \mathsf{N}(0, \sigma^2)$$

This is the classic homoscedastic linear regression model.

#### How many parameters?

- ▶ The classical linear regression model has P + 2 parameters:
  - $1. \sigma$
  - $2. \alpha$
  - 3. all the  $\beta_p$
- ▶ We therefore need to set a prior on each of these parameters;
- ➤ To understand what kind of priors we need, it is important to understand what each of these parameters governs in the distribution on y<sub>i</sub>;
- In particular,
  - 1.  $\sigma = \sqrt{\mathbb{V}(y)}$  governs the **spread** of y, marginally;
  - 2.  $\alpha = \mathbb{E}[y|X=0]$  governs the **location** of y, marginally;
  - 3.  $\beta_p = \beta_p \approx \mathbb{E}[y|X_p = x_p + 1] \mathbb{E}[y|X_p = x_p]$  describes the average change in y when we shift  $x_p$  by one unit.

### Calibrating the prior(s) (1)

- If we have any knowledge about y<sub>i</sub>, we can use this information to set reasonable prior distributions on each of these parameters;
- Going back to Thomas' example on yield, we might know what the expected yield of some crop might be, and how volatile this average value can be;
- In other words, we might have **prior** information on both  $\alpha$  and  $\sigma$ ;
- Then, in case P=1 and  $x_i$  represents, for example, soil moisture, then the literature (or common sense/agronomic expertise) can provide upper and lower bounds for the prior on  $\beta$ ;
- ▶ The same line of reasoning also apply in case of P > 1.

### Calibrating the prior(s) (2)

- There are however cases in which such details are not or only partially known;
- Then how do we set reasonable priors?
- In the next slides we are going to explore a simulation-based calibration approach to solve this puzzle;
- ► The idea is to:
  - 1. choose a prior for  $\sigma$ ,  $\alpha$  and  $\beta_p$  and draw potential parameter values from these distribution;
  - 2. plug these parameter values into  $N(\alpha + \sum_{p=1}^{P} \beta_p x_{p,i}, \sigma)$  and simulate potential values of  $y_i$ .

#### Load data and libraries

```
library(truncnorm)
library(rstan)

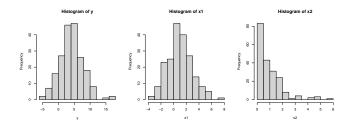
setwd("your working directory")

load("linear_regression_data.RData")
source("aux_functions.R")

set.seed(123)
```

### Visualize the data (1)

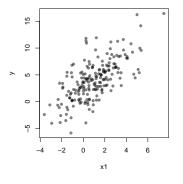
```
par(mfrow=c(1,3), pty="s")
hist(y)
hist(x1)
hist(x2)
```

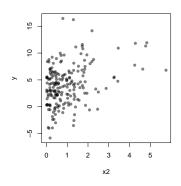


```
par(mfrow=c(1,1), pty="m")
```

### Visualize the data (2)

```
par(mfrow=c(1,2), pty="s")
plot(x1, y, pch = 16, col = rgb(0,0,0,.5))
plot(x2, y, pch = 16, col = rgb(0,0,0,.5))
```

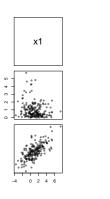




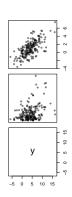
```
par(mfrow=c(1,1), pty="m")
```

### Visualize the data (3)

```
par(pty="s")
pairs(cbind(x1,x2,y), pch = 16, col = rgb(0,0,0,.5))
```







```
par(pty="m")
```

### Begin sampling from the prior: $\sigma$ (1)

- We begin by studying how  $\sigma$  may impact  $\mathbf{y} = [y_1, \dots, y_N]$ ;
- Since the  $\sigma$  can only take up positive values, the **prior** distribution must satisfy this restriction;
- ► There are several options: Exponential distribution, Gamma distribution, Truncated Normal distribution and others;
- ► Here we use the Truncated Normal:

$$\sigma \sim N_+(0,\lambda)$$

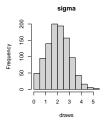
ightharpoonup Calibrating the prior for  $\sigma$  consist in choosing a sensible value for  $\lambda$ .

#### Begin sampling from the prior: $\sigma$ (2)

▶ For example, for  $\lambda = 2$ , this is how 1000 draws from  $N_+(0,2)$  look like:

```
std_dev <- rtruncnorm(1000, 0, Inf, 2)

par(pty="s")
hist(std_dev, main = "sigma", xlab = "draws")</pre>
```



```
par(pty="s")
```

### Begin sampling from the prior: $\sigma$ (3)

- ▶ We can now begin to study what  $N_+(0,2)$  means in terms of potential values of  $\mathbf{y}$ :
  - 1. Sample one parameter value from  $N_+(0,\lambda)$ , and get  $\sigma_v^s$ ;
  - 2. Sample N = 200 observations from  $N(0, \sigma_y^s)$  and get  $\tilde{\mathbf{y}}^s$ ;
  - 3. Repeat the process S = 1000 times.

```
y_sim <- matrix(nrow=1000, ncol=200)
for (s in 1:1000) {
    std_dev <- rtruncnorm(1, 0, Inf, 2)
    y_sim[s, ] <- rnorm(200, 0, std_dev)
}</pre>
```

- ▶ The  $N \times S$  matrix  $y_{sim} = [y^1, ..., y^S]$  approximates the so-called **Prior Predictive Distribution** (PrPD) of y;
- ▶ We can then use visualization to study the properties of the PrPD and check whether our assumption makes any sense in face of the observed data, **y**.

## Visualizing the PrPD (1)

## [1] 1000

- 1. For each draw  $\tilde{\mathbf{y}}^s$  calculate  $max^s = \max(\tilde{\mathbf{y}}^s)$ ;
- 2. For each draw  $\tilde{\mathbf{y}}^s$  calculate  $min^s = min(\tilde{\mathbf{y}}^s)$ ;
- 3. For each draw  $\tilde{\mathbf{y}}^s$  calculate  $sd^s = sd(\tilde{\mathbf{y}}^s)$

```
max_y <- apply(y_sim, 1, max)
min_y <- apply(y_sim, 1, min)
sd_y <- apply(y_sim, 1, sd)</pre>
```

These are obviously  $S \times 1$  vectors:

```
length(max_y)

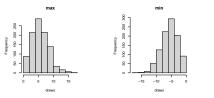
## [1] 1000
length(min_y)

## [1] 1000
length(sd_y)
```

### Visualizing the PrPD (2)

Plot these quantities:

```
par(mfrow=c(1,3), pty="s")
hist(max_y, main = "max", xlab = "draws")
hist(min_y, main = "min", xlab = "draws")
hist(sd_y, main = "sd", xlab = "draws")
```





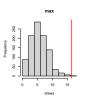
```
par(mfrow=c(1,1), pty="m")
```

Each histogram represents the extent of maximum/minumim values, and standard deviations that can be obtained setting  $\lambda=2$ .

### Visualizing the PrPD (3)

We can check consistency between the summary statistics of the PrPD and our observed data points by overlaying min(y), max(y) and sd(y) to the corresponding histogram:

```
par(mfrow=c(1,3), pty="s")
hist(max_y, main = "max", xlab = "draws")
abline(v=max(y),col="red", lwd=2)
hist(min_y, main = "min", xlab = "draws")
abline(v=min(y),col="blue", lwd=2)
hist(sd_y, main = "sd", xlab = "draws")
abline(v=sd(y),col="green", lwd=2)
```







```
par(mfrow=c(1,1), pty="m")
```

### Visualizing the PrPD (3)

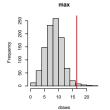
- The standard deviation (SD) of the PrPD is smaller than the observed SD;
- Also the maximum value does not quite match;
- ▶ We can try changing *lambda* to, say, 3:

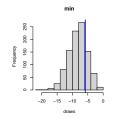
```
y_sim <- matrix(nrow=1000, ncol=200)
for (s in 1:1000) {
    std_dev <- rtruncnorm(1, 0, Inf, 3)
    y_sim[s, ] <- rnorm(200, 0, std_dev)
}

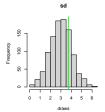
max_y <- apply(y_sim, 1, max)
min_y <- apply(y_sim, 1, min)
sd_y <- apply(y_sim, 1, sd)</pre>
```

### Visualizing the PrPD (4)

```
par(mfrow=c(1,3), pty="s")
hist(max_y, main = "max", xlab = "draws")
abline(v=max(y),col="red", lwd=2)
hist(min_y, main = "min", xlab = "draws")
abline(v=min(y),col="blue", lwd=2)
hist(sd_y, main = "sd", xlab = "draws")
abline(v=sd(y),col="green", lwd=2)
```







```
par(mfrow=c(1,1), pty="m")
```

▶ SD is now well calibrated, but to have better max and min we will need to tweak  $\alpha$  and  $\beta$ .

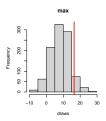
### Calibrating $\alpha$ (1)

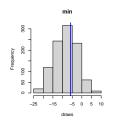
- ▶ Recall that  $\alpha = \mathbb{E}[y|X=0]$ ;
- ▶ Therefore, the range of plausible value of  $\alpha$  should reflect the expected range of average values for y;
- Since y includes both positive and negative values, using a **Normal** prior for  $\alpha$  should do;
- ▶ We begin with  $\alpha \sim N(0,5)$ ;
- Notice that this prior implies that the range of  $\mathbb{E}[y|X=0]$  is roughly [-15,+15] and values between [-5,+5] are more likely.

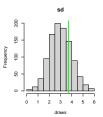
```
y_sim <- matrix(nrow=1000, ncol=200)
for (s in 1:1000) {
   std_dev <- rtruncnorm(1, 0, Inf, 3)
   mu <- rnorm(1, 0, 5)
   y_sim[s, ] <- rnorm(200, mu, std_dev)
}</pre>
```

### Calibrating $\alpha$ (2)

```
par(mfrow=c(1,3), pty="s")
hist(max_y, main = "max", xlab = "draws")
abline(v=max(y),col="red", lwd=2)
hist(min_y, main = "min", xlab = "draws")
abline(v=min(y),col="blue", lwd=2)
hist(sd_y, main = "sd", xlab = "draws")
abline(v=sd(y),col="green", lwd=2)
```





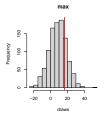


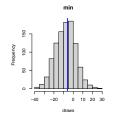
```
par(mfrow=c(1,1), pty="m")
```

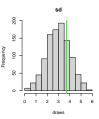
#### Calibrating $\alpha$ (3)

 $\triangleright$  Setting  $\alpha$  to N(0, 10), instead leads to:

```
par(mfrow=c(1,3), pty="s")
hist(max_y, main = "max", xlab = "draws")
abline(v=max(y),col="red", lwd=2)
hist(min_y, main = "min", xlab = "draws")
abline(v=min(y),col="blue", lwd=2)
hist(sd_y, main = "sd", xlab = "draws")
abline(v=sd(y),col="green", lwd=2)
```







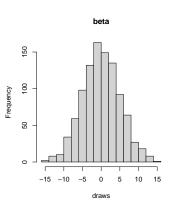
```
par(mfrow=c(1,1), pty="m")
```

### Calibrating $\alpha$ (4)

- Observe that, although these priors might seem at first quite informative, they are not!
- ▶ In fact, although the most likely values under the priors set so far are close to those observed in the dataset, the range of plausible *y* extends quite beyond that range;
- Prior distributions of this nature are typically referred to as weakly informative;
- There is a very large chunk of the literature suggesting that weakly informative priors should be preferred to completely uninformative ones;
- There are several reasons for this, but one very important one in probabilistic programming is that they usually trigger computational issues;
- Examples of uninformative priors that should not be used include the historically very popular Uniform $(-\infty, +\infty)$ .

### Calibrating $\beta_1$ and $\beta_2$ (1)

▶ Recall that  $\beta_p \approx \mathbb{E}[y|X_p = x_p + 1] - \mathbb{E}[y|X_p = x_p]$ . Since  $\bar{\mathbf{y}} = 4.07$ , setting  $\beta_p \sim \mathsf{N}(0,5)$  implies that shifting  $x_p$  by one unit would change the expected value of y by:



### Calibrating $\beta_1$ and $\beta_2$ (2)

- Notably, by using a normal prior centered at zero, we are implicitly assuming that the most likely change in \( \mathbb{E}[y|X\_p = x\_p] \) before seeing the data is small;
- This is a conservative choice and is typically adopted in regression problems;
- ▶ However, notice also that the N(0,5) prior allows the 'slopes' to potentially extend up to roughly [-15, +15] and beyond;
- It is the likelihood-prior interaction that will cause a change in the **posterior** distribution of  $\beta_p$ .

### Calibrating $\beta_1$ and $\beta_2$ (3)

- To simulate the impact of
  - 1.  $\sigma \sim N_{+}(0,3)$
  - 2.  $\alpha \sim N(0, 10)$
  - 3.  $\beta_1 = \beta_2 \sim N(0,5)$

on the PrPD of y, I have prepared a small Stan script;

- When we need to calibrate more than one prior, wiring down the whole data generating process in Stan is a convenient way to avoid hardcoding the entire scheme in R;
- I will go through this program very briefly, just so you have an understanding of how it works.

### Stan program for prior calibration (1)

We first load the script using the R function stan\_model():

```
rstan_options(auto_write = TRUE)
lin_reg_prior <- stan_model("linear_regression_prior.stan")</pre>
```

▶ Before using using the lin\_reg\_prior object to simulate the data, let us have a look at what this code looks like!

### Stan program for prior calibration (2)

- The input to any Stan program is a data list;
- ► This list contains all the (hyper)parameter values, the data and any additional variable necessary to run the code contained in the script

```
dat_list <- list(</pre>
  N = length(y),
  x1 = x1
  x2 = x2
  v = v.
  mu_b1 = 0,
  mu_b2 = 0,
  sigma_b1 = 5,
  sigma_b2 = 5,
  mu_alpha = 0,
  sigma alpha = 10.
  lambda_sigma = 3
```

### Stan program for prior calibration (3)

- N indicates the sample size;
- $\blacktriangleright$  x1 and x2 are  $N \times 1$  vector for the two covariates;
- $\triangleright$  y indicates the  $N \times 1$  vector **y**;
- ▶ mu\_b1 and mu\_b1 indicate the mean of the priors on  $\beta_1$  and  $\beta_2$ , respectively;
- ▶ sigma\_b1 and sigma\_b1 indicate the standard deviation of the priors on  $\beta_1$  and  $\beta_2$ , respectively;
- mu\_alpha indicates the mean of the prior on α;
- ▶ sigma\_alpha indicates the standard deviation of the prior on  $\alpha$ ;
- lambda\_sigma indicates the standard deviation of the prior on σ.

#### Stan program for prior calibration (4)

ightharpoonup We can now simulate  $\tilde{\mathbf{y}}^s$ !

 $\blacktriangleright$  and extract the  $N \times S$  matrix of simulated outcome values:

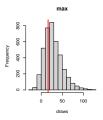
```
prior_sample_fit <- extract(prior_sample)
y_sim <- prior_sample_fit$y_pred</pre>
```

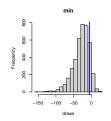
We can next repeat the visual assessment introduced in the earlier slides. We begin by calculating our summary statistics:

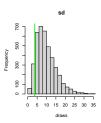
```
max_y <- apply(y_sim, 1, max)
min_y <- apply(y_sim, 1, min)
sd_y <- apply(y_sim, 1, sd)</pre>
```

#### Calibrated PrPD (1)

```
par(mfrow=c(1,3), pty="s")
hist(max_y, main = "max", xlab = "draws")
abline(v=max(y),col="red", lwd=2)
hist(min_y, main = "min", xlab = "draws")
abline(v=min(y),col="blue", lwd=2)
hist(sd_y, main = "sd", xlab = "draws")
abline(v=sd(y),col="green", lwd=2)
```







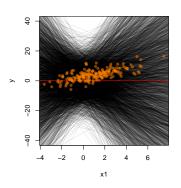
```
par(mfrow=c(1,1), pty="m")
```

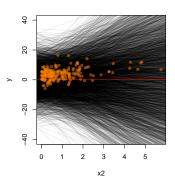
#### Calibrated PrPD (2)

- We can also visualize the bundle of regression lines implied by  $\beta_1 = \beta_2 \sim N(0,5)$ ;
- To do this we can use the

```
par(mfrow=c(1,2))
plot_prior_lines(x1,
                 prior_sample_fit$alpha,
                 prior_sample_fit$beta1,
                 prior_sample_fit$y_pred,
                 vlim = c(-40, 40), xlab = "x1")
plot_prior_lines(x2,
                 prior_sample_fit$alpha,
                 prior_sample_fit$beta2,
                 prior_sample_fit$y_pred,
                 vlim = c(-40, 40), xlab = "x2")
par(mfrow=c(1,1))
```

# Calibrated PrPD (3)





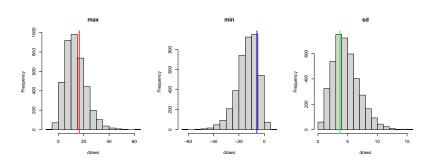
### Calibrated PrPD (4)

- Things to observe:
  - This model is general. Perhaps too general... It implies very extreme modelling assumption (i.e., potentially nonsensical regression lines!);
  - 2. Conditional on the two covariates  $x_1$  and  $x_2$ , the prior on the intercept maybe also too wide (look at how wide the bundle of regression lies is around zero!);
- Let us see what changes by adjusting both distributions!

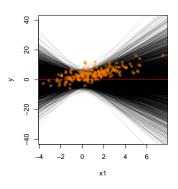
#### Calibrated PrPD (5)

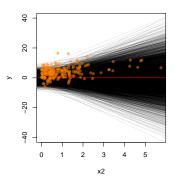
Adjusting  $\beta_1$  and  $\beta_2$  to N(0,2),  $\alpha$  to N(0,2.5) and re-running the simulation yields:

```
dat_list$sigma_b1 <- 2
dat_list$sigma_b2 <- 2
dat_list$sigma_alpha <- 2.5</pre>
```



# Calibrated PrPD (6)





### Default priors (1)

- This calibration exercise demonstrates that, when knowledge on y and  $x_p$  is limited, prior calibration can be tedious;
- Moreover, when the number of regressors increases and/or the model includes non-linearities and interactions, choosing the most suitable distribution for  $\beta_p$  becomes very hard;
- ▶ Also, new sample information could yield rather different (and bad!) PrPD if the distribution of the new y,  $x_1$  and  $x_2$  changes too much;
- A commonly adopted solution to this issue consists of standardizing our variables;
- ► These transformation remove scale-dependence and allow to defined one-fits-all priors on the parameters of interest;
- Let's see how!

# Default priors (2)

Standardizing some numeric variable z means:

$$\dot{z} = \frac{z - \bar{z}}{\operatorname{sd}(z)}$$

- ► Therefore, any ż will have mean zero and standard deviation 1 by construction;
- Applying this straightforward transformation to our data simplifies enormously the calibration procedure discussed above;
- ▶ To begin with, by standardizing y, we force its mean to zero, so  $\alpha = 0$  by construction;
- lacktriangle We therefore do not need to calibrate the prior for lpha anymore;
- Moreover, since  $\mathbb{V}(\dot{y}) = 1$ , setting  $\sigma \sim \mathsf{N}_+(0,1)$  already provides a good default for the prior on the standard deviation!

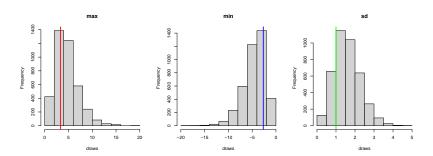
### Default priors (3)

- Finally, since  $\mathbb{V}(\dot{y}) = 1$  and the range of  $\dot{y}$  is artificially constraint to roughly [-3, +3] (three SD of  $\dot{y}$ ), giving both  $\beta_1$  and  $\beta_2$  a N(0,1) prior seems a sensible choice;
- ► To see what these new values yield in terms of PrPD, let us redefine out data list accordingly:

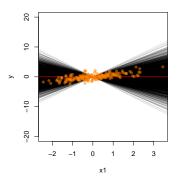
```
v std <- as.vector(scale(v))</pre>
x1_std <- as.vector(scale(x1))</pre>
x2_std <- as.vector(scale(x2))</pre>
dat_list <- list(</pre>
  N = length(y_std),
  x1 = x1_std, x2 = x2_std,
  y = y_std,
 mu b1 = 0.
 mu_b2 = 0,
  sigma_b1 = 1, sigma_b2 = 1,
  mu_alpha = 0, sigma_alpha = 0,
  lambda_sigma = 1
```

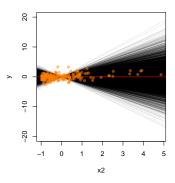
## Default priors (4)

# Default priors (5)



# Default priors (6)





#### Estimation

Once all the priors have been setup, we can use Stan to sample from the implied joint posterior distribution:

$$f(\sigma, \beta_1, \beta_2 | \mathbf{y}, X_1 = x_1, X_2 = x_2)$$

▶ To do this, we need to load the second program:

```
lin_reg <- stan_model("linear_regression.stan")</pre>
```

► Fitting the model to the data now only requesires a slighly different sampling statement:

### Exploring the results (1)

To quickly visualize the results, we can call the function summary as we would do with the standard 1m output:

```
params <- c("beta1", "beta2", "sigma")</pre>
summary(fit, pars = params)$summary[,1:3]
##
             mean
                       se mean
                                       sd
## beta1 0.7648427 0.0007385151 0.04126889
## beta2 0.4857550 0.0007458866 0.04067350
## sigma 0.5618872 0.0004903962 0.02896625
params <- c("beta1", "beta2", "sigma")</pre>
summary(fit, pars = params)$summary[,4:8]
             2.5% 25% 50% 75%
##
                                                    97.5%
## beta1 0.6822056 0.7378445 0.7646251 0.7917823 0.8481729
```

```
## beta1 0.6822056 0.7378445 0.7646251 0.7917823 0.8481729
## beta2 0.4045790 0.4586613 0.4854272 0.5127030 0.5666763
## sigma 0.5080042 0.5416115 0.5608089 0.5806676 0.6213943
```

# Exploring the results (2)

- Because of standardization, we need to transform all the coefficients back to the original scale of the data;
- ▶ Otherwise,  $\hat{\beta}_p \approx \mathbb{E}[y|X_p = x_p + \operatorname{sd}(y)] \mathbb{E}[y|X_p = x_p]$ ;
- To do this, it suffices to rescale the standardized coefficients  $\hat{\beta}_p$  by the standard deviation of  $x_p$  and y:

$$\beta_{p} = \hat{\beta}_{p} \times \frac{\mathsf{sd}(y)}{\mathsf{sd}(x_{p})}$$

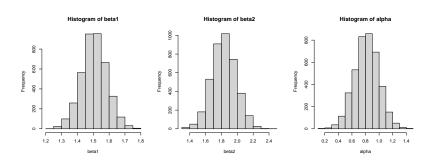
▶ Moreover, we can retrieve the intercept through:

$$\alpha = \bar{\mathbf{y}} - \sum_{p=1}^{P} \beta_p \bar{\mathbf{x}}_p$$

#### Exploring the results (3)

```
beta1 <- sample_fit$beta1 * (sd(y)/sd(x1))
beta2 <- sample_fit$beta2 * (sd(y)/sd(x2))
alpha <- mean(y)-(beta1*mean(x1))-(beta2*mean(x2))

par(mfrow=c(1,3), pty="s")
hist(beta1)
hist(beta2)
hist(alpha)</pre>
```

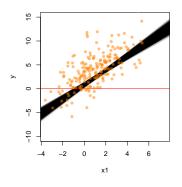


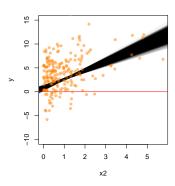
#### par(mfrow=c(1,1), pty="m")

# Exploring the results (4)

```
summary(beta1)
##
    Min. 1st Qu. Median Mean 3rd Qu. Max.
##
    1.223 1.453 1.506
                         1.506
                                1.559
                                       1.789
summary(beta2)
##
    Min. 1st Qu. Median Mean 3rd Qu. Max.
##
    1.303 1.729 1.830
                         1.831
                                       2.423
                                1.933
summary(alpha)
##
     Min. 1st Qu. Median Mean 3rd Qu.
                                     Max.
##
   0.1201
          0.6977 0.8171 0.8154 0.9368 1.4617
```

# Exploring the results (5)





## What if we used lm() instead?

```
summary(lm(y \sim x1 + x2))
##
## Call:
## lm(formula = y \sim x1 + x2)
##
## Residuals:
##
      Min 10 Median 30
                                   Max
## -5.6352 -1.4044 0.1549 1.3290 5.4982
##
## Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 0.81122 0.23174 3.50 0.000574 ***
## x1
        1.50803 0.07953 18.96 < 2e-16 ***
           1.83350 0.15226 12.04 < 2e-16 ***
## x2
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.082 on 197 degrees of freedom
## Multiple R-squared: 0.689, Adjusted R-squared: 0.6859
## F-statistic: 218.3 on 2 and 197 DF, p-value: < 2.2e-16
```

#### Leveraging the posterior

- So far, it might seems that all the extra effort to setup a probabilistic program to estimate a linear regression might be worthless in fact of a simple lm() call;
- However, having samples from the joint posterior unlocks several features that only belong in the Bayesian realm;
- First of all, it is worth stressing that having  $f(\beta_p|\mathbf{y},X=x)$  is different from having just  $\mathbb{E}[\beta_p]$  and its standard error;
- If the parameters in  $f(\sigma, \beta_1, \beta_2 | \mathbf{y}, X_1 = x_1, X_2 = x_2)$  are uncorrelated, then extracting any component from the  $S \times (P+2)$  matrix  $[\sigma, \alpha, \beta_1, \beta_2]$ , provides S samples from the marginal posterior of that parameter;
- These are the quantities that we are going to exploit.

## Posterior probabilities (1)

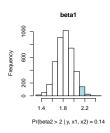
- Consider the marginal posterior of  $\beta_2$ , which we store in the vector beta2;
- We can use this object to calculate probabilities. For example, what is the probability that  $\beta_2 > 2$ ?
- ► This can be easily calculated as the average count of posterior parameter values that are greater than 2:

```
thr <- 2
prob <- mean(beta2>thr); prob
```

```
## [1] 0.13825
```

#### Posterior probabilities (2)

We can also visualize this probability using the histogram above:



```
par(pty="m")
```

#### Credible Intervals

- We can summarize the marginal posterior of our parameters via intervals;
- These are called Credible Intervals (CrI);
- They are naturally interpreted as the as the proportion of parameter values that fall between (i.e., what is the range of most likely values) an upper and lower value;
- Therefore, CrI are simply computed as the  $\gamma$  and  $1 \gamma$  quantiles of  $\beta_2$ . For  $\gamma = 0.025$  we have the 95% CrI:

```
quantile(beta2, c(.025, .975))
```

```
## 2.5% 97.5%
## 1.525164 2.136232
```

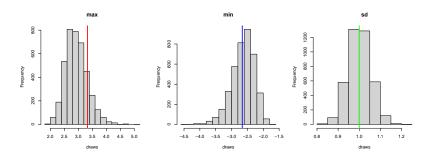
# Model checking (internal consistency - 1)

- Another useful tool in the Bayesian regression toolbox is the sp-called Posterior Predictive Distribution (PoPD);
- ▶ Much like we used the PrPD to simulate *y* under different prior configurations, we use the PoPD to simulate *y* under the estimated posterior;
- ► The idea is to check the consistency between the simulated *y* and the observed counterpart;
- This automatically computed in the Stan program above and can be easily accessed by extracting y\_pred from the fitted model:

```
y_pred <- sample_fit$y_pred</pre>
```

#### Model checking (internal consistency - 2)

```
par(mfrow=c(1,3), pty="s")
hist(apply(y_pred, 1, max), main = "max", xlab = "draws")
abline(v=max(y_std),col="red", lwd=2)
hist(apply(y_pred, 1, min), main = "min", xlab = "draws")
abline(v=min(y_std),col="blue", lwd=2)
hist(apply(y_pred, 1, sd), main = "sd", xlab = "draws")
abline(v=sd(y_std),col="green", lwd=2)
```



```
par(mfrow=c(1,1), pty="m")
```

# Model checking (internal consistency - 3)

```
plot_prior_dens(y_pred, y_std, main = "PoPD")
```

