

- Dependent variable
- Response variable
- Outcome

- Independent variable
- Predictor variable
- Explanatory variable
- Covariate



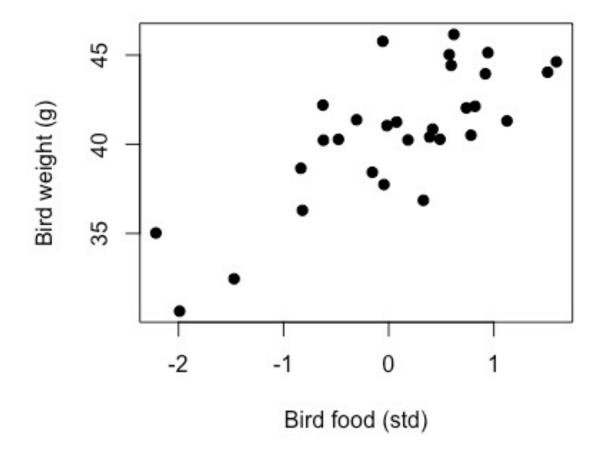
```
#set seed for reproducing results
set.seed(1)
#number of data points
N <- 30</pre>
```

```
#set seed for reproducing results
set.seed(1)
#number of data points
N <- 30
#simulate predictors (standardized)
food_std <- rnorm(N, 0, 1)</pre>
```

```
#set seed for reproducing results
set.seed(1)
#number of data points
N <- 30
#simulate predictors (standardized)
food_std <- rnorm(N, 0, 1)
#generating intercept and slope values
alpha <- 40
beta <- 3
#simulate linear predictor
mu <- alpha + beta * food_std</pre>
```

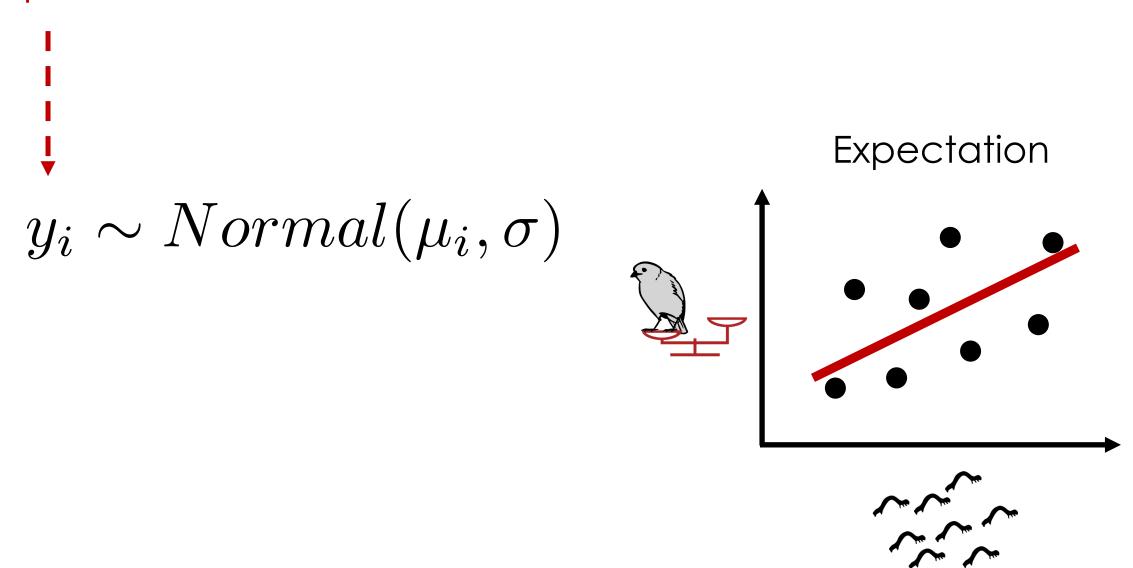
```
#set seed for reproducing results
set.seed(1)
#number of data points
N < -30
#simulate predictors (standardized)
food_std <- rnorm(N, 0, 1)
#generating intercept and slope values
alpha <- 40
beta <- 3
#simulate linear predictor
mu <- alpha + beta * food_std</pre>
#process error (wrt relationship
between food and bird weight)
sigma <- 3
#simulate y values
bird_weight <- rnorm(N, mu, sigma)</pre>
```

```
#set seed for reproducing results
set.seed(1)
#number of data points
N < -30
#simulate predictors (standardized)
food_std <- rnorm(N, 0, 1)</pre>
#generating intercept and slope values
alpha <- 40
beta <- 3
#simulate linear predictor
mu <- alpha + beta * food_std</pre>
#process error (wrt relationship
between food and bird weight)
sigma <- 3
#simulate y values
bird_weight <- rnorm(N, mu, sigma)</pre>
#plot
plot(food_std, bird_weight,
     pch = 19,
     xlab = 'Bird food (std)',
     ylab = 'Bird weight (g)')
```



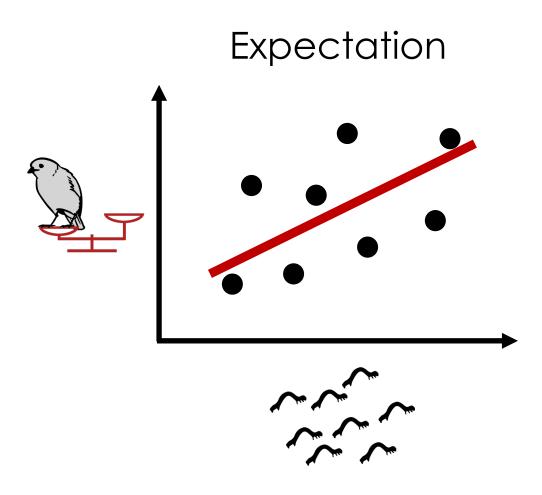
Expectation

Response



Response Expected value



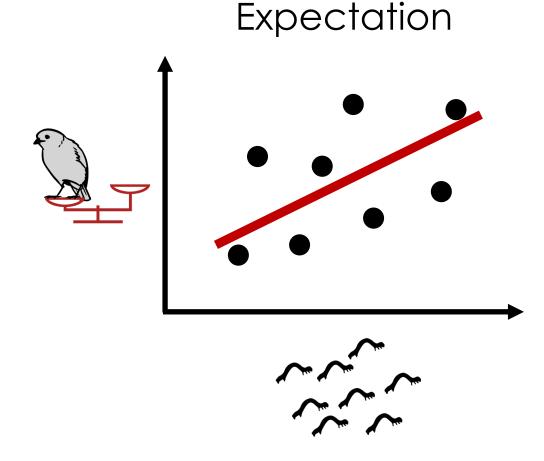


Response Expected value Expectation $y_i \sim Normal(\mu_i, \sigma)$ $\mu_i = \alpha + \beta \times x_i$ Intercept Slope

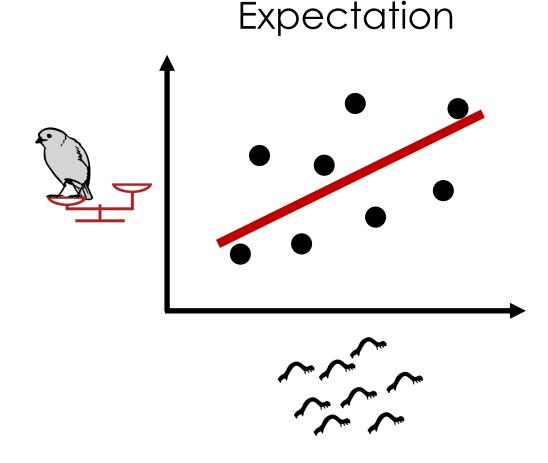
Predictor

Response Expected value Residual Error Expectation $y_i \sim Normal(\mu_i, \sigma)$ $\mu_i = \alpha + \beta \times x_i$ Intercept Slope Predictor

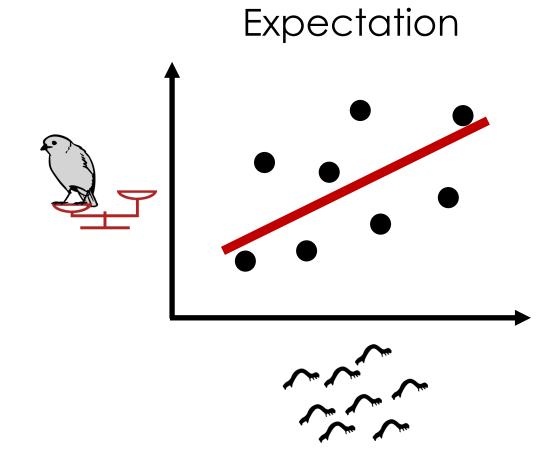
$$y_i \sim Normal(\mu_i, \sigma)$$
$$\mu_i = \alpha + \beta \times x_i$$

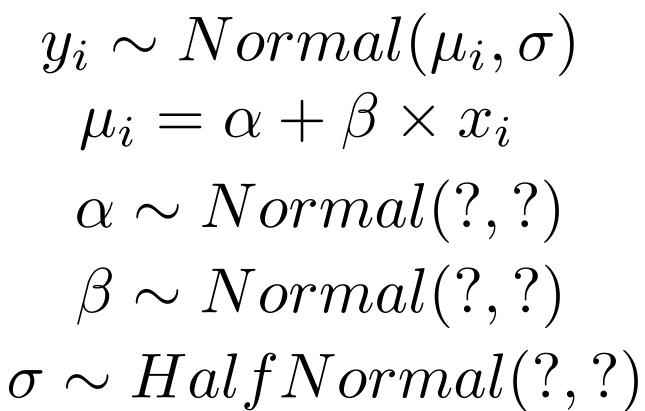


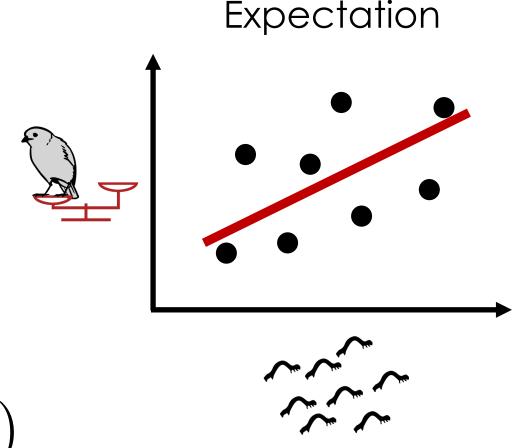
 $y_{i} \sim Normal(\mu_{i}, \sigma)$ $\mu_{i} = \alpha + \beta \times x_{i}$ $\alpha \sim Normal(?,?)$

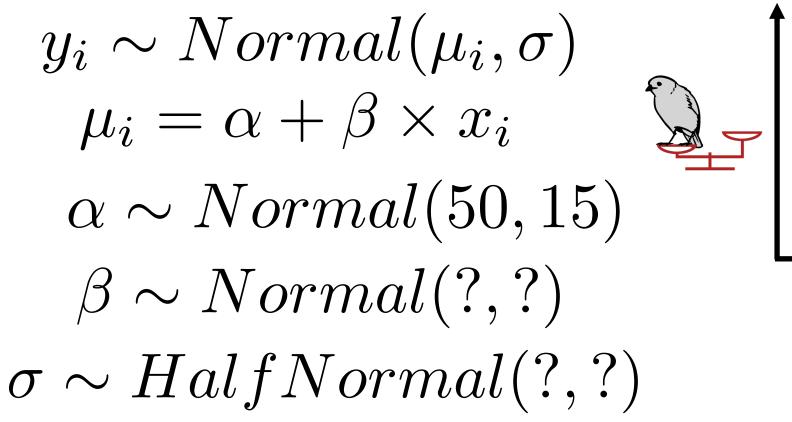


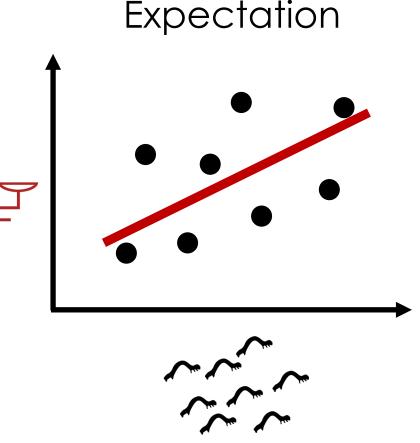
 $y_{i} \sim Normal(\mu_{i}, \sigma)$ $\mu_{i} = \alpha + \beta \times x_{i}$ $\alpha \sim Normal(?, ?)$ $\beta \sim Normal(?, ?)$

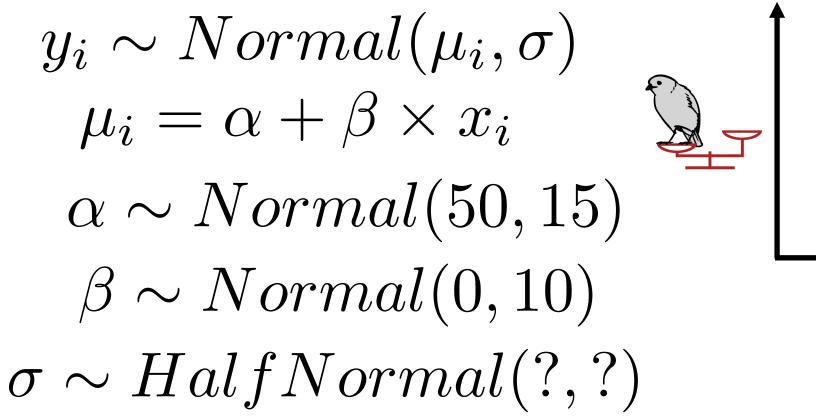


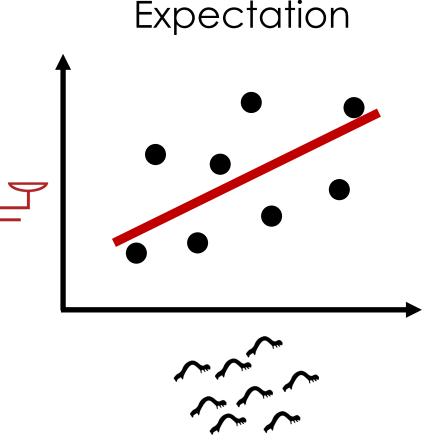


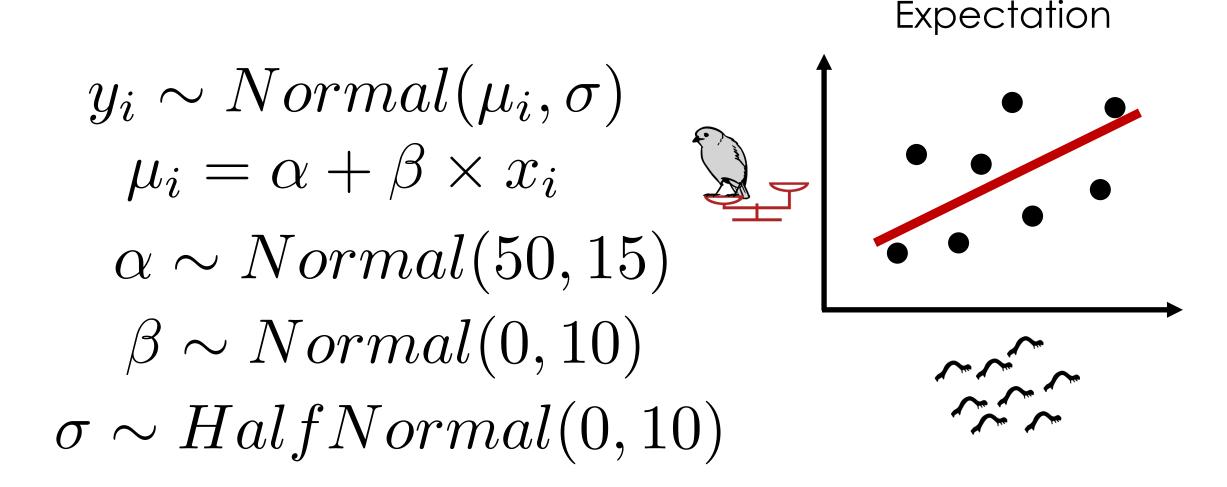












$$y_{i} \sim Normal(\mu_{i}, \sigma) - -- Pr(y, x | \alpha, \beta, \sigma)$$

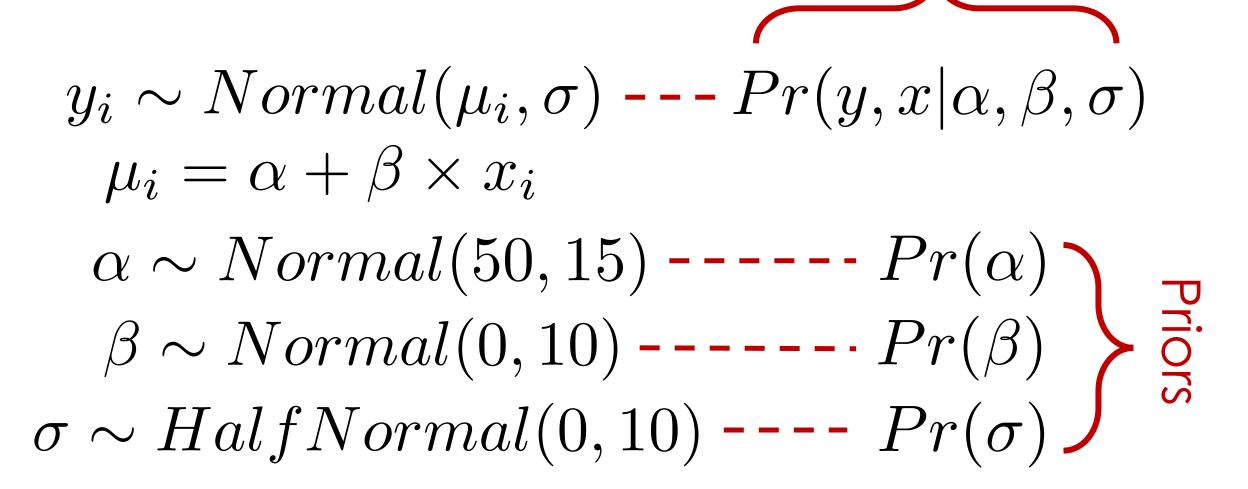
$$\mu_{i} = \alpha + \beta \times x_{i}$$

$$\alpha \sim Normal(50, 15) - -- Pr(\alpha)$$

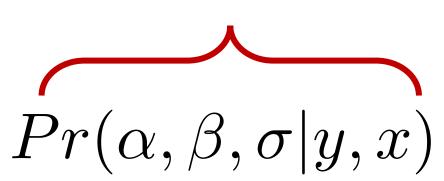
$$\beta \sim Normal(0, 10) - -- Pr(\beta)$$

$$\sigma \sim HalfNormal(0, 10) - -- Pr(\sigma)$$

Likelihood



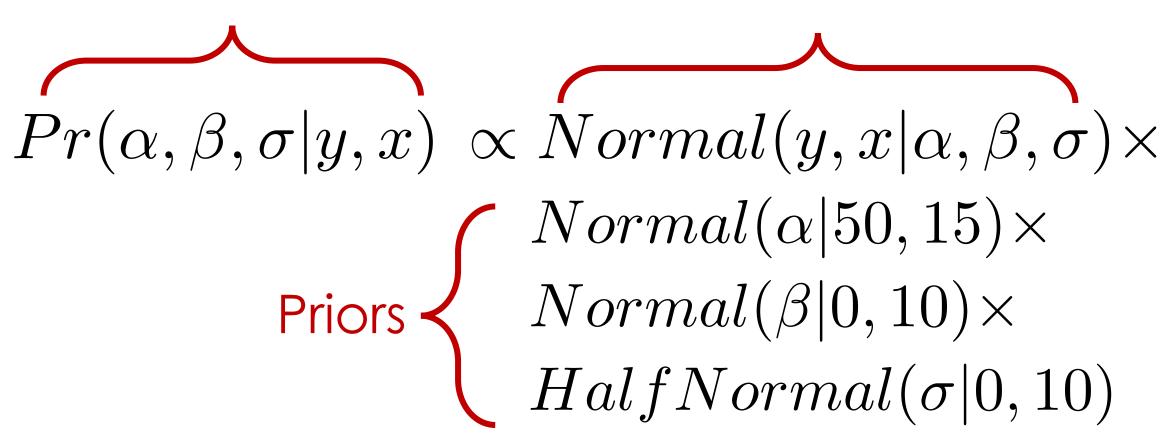
Posterior



Posterior Likelihood $Pr(\alpha,\beta,\sigma|y,x) \propto Normal(y,x|\alpha,\beta,\sigma) \times$

Posterior

Likelihood



MCMC to the rescue!





in R Studio

$$y_{i} \sim Normal(\mu_{i}, \sigma)$$

$$\mu_{i} = \alpha + \beta \times x_{i}$$

$$\alpha \sim Normal(50, 15)$$

$$\beta \sim Normal(0, 10)$$

$$\sigma \sim HalfNormal(0, 10)$$

Fitting the model

Diagnostic	Description	What you're looking for
\hat{R}	How well the chains 'agree' about the posterior density	<= 1.01

n.eff

Number of effective samples from posterior

posterior

Sampling anomaly

0 divergences

> 400 for

4 chains

Simulated data

looks like

observed data

Posterior predictive check

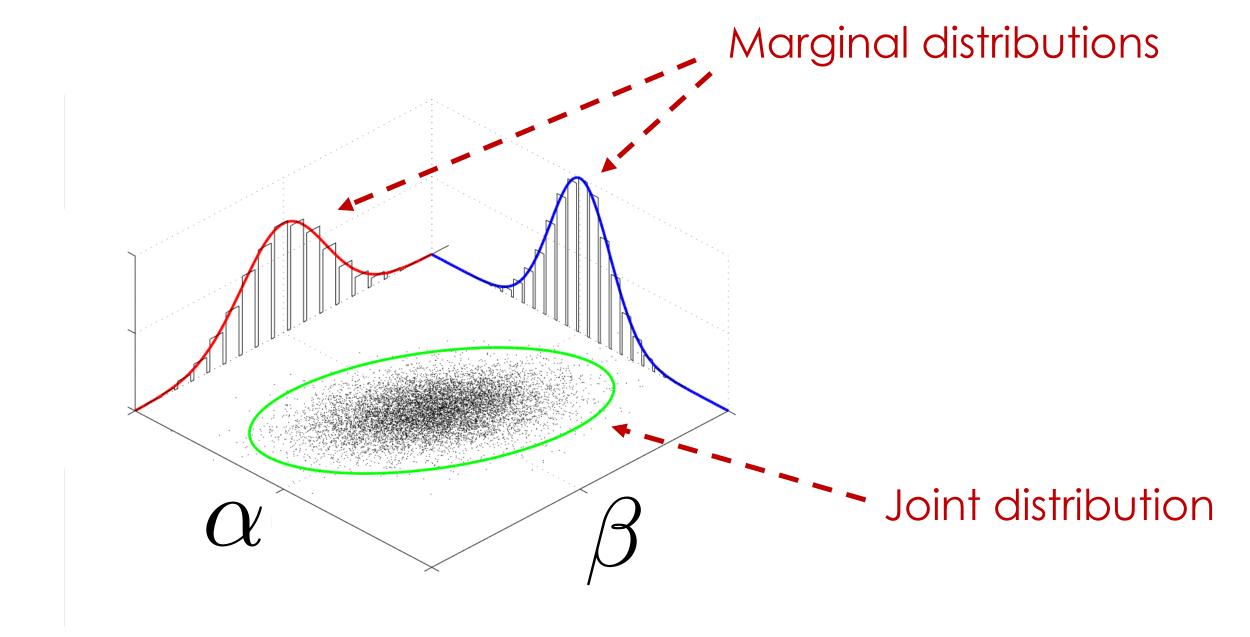
Realizations of response variable generated from

Divergences

(Hamiltonian Monte Carlo specific)

Assessing the model

Summarizing the model output



Plot results