## Linear Models

Lecture 02

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#### **Linear Models Basics**

Pretty much everything we a going to see in this course will fall under the umbrella of either linear or generalized linear models.

In previous classes most of your time has likely been spent with the *iid* case,

$$y_i = \beta_0 + \beta_1 x_{i1} + \dots + \beta_p x_{ip} + \epsilon_i$$
$$\epsilon_i \stackrel{iid}{\sim} N(0, \sigma^2)$$

these models can also be expressed as,

$$y_i \stackrel{\text{iid}}{\sim} N(\beta_0 + \beta_1 x_{i1} + \dots + \beta_p x_{ip}, \sigma^2)$$

#### Some notes on notation

- Observed values and scalars will usually be lower case letters, e.g.  $x_i, y_i, z_{ij}.$
- Parameters will usually be greek symbols, e.g.  $\mu$ ,  $\sigma$ ,  $\rho$ .
- Vectors and matrices will be shown in bold, e.g.  $\mu, X, \Sigma$ .
- Elements of a matrix (or vector) will be referenced with {}s, e.g.  ${\{Y\}}_i, {\{\Sigma\}}_{ij}$
- Random variables will be indicated by  $\sim$ , e.g.  $x \sim Norm(0, 1), z \sim Gamma(1, 1)$
- Matrix / vector transposes will be indicated with ', e.g. A', (1 B)'

#### Linear model - matrix notation

We can also express a linear model using matrix notation as follows,

$$Y = X \beta + \epsilon$$

$$n \times 1 = n \times p p \times 1 = n \times 1$$

$$\epsilon_{n \times 1} \sim N(0, \sigma^2 1_n)$$

$$n \times 1 = n \times 1$$

or alternatively as,

$$\mathbf{Y}_{n\times 1} \sim N \left( \mathbf{X}_{n\times p} \mathbf{\beta}_{p\times 1}, \sigma^2 \mathbb{1}_n \right)$$

#### Multivariate Normal Distribution - Review

For an n-dimension multivate normal distribution with covariance  $\Sigma$  (positive semidefinite) can be written as

$$\mathbf{Y}_{n\times 1} \sim N(\boldsymbol{\mu}, \boldsymbol{\Sigma}_{n\times n})$$

where 
$$\left\{\boldsymbol{\Sigma}\right\}_{ij}=\rho_{ij}\sigma_{i}\sigma_{j}$$

$$\begin{pmatrix} y_1 \\ y_2 \\ \vdots \\ y_n \end{pmatrix} \sim N \begin{pmatrix} \begin{pmatrix} \mu_1 \\ \mu_2 \\ \vdots \\ \mu_n \end{pmatrix}, \begin{pmatrix} \rho_{11}\sigma_1\sigma_1 & \rho_{12}\sigma_1\sigma_2 & \cdots & \rho_{1n}\sigma_1\sigma_n \\ \rho_{21}\sigma_2\sigma_1 & \rho_{22}\sigma_2\sigma_2 & \cdots & \rho_{2n}\sigma_2\sigma_n \\ \vdots & \vdots & \ddots & \vdots \\ \rho_{n1}\sigma_n\sigma_1 & \rho_{n2}\sigma_n\sigma_2 & \cdots & \rho_{nn}\sigma_n\sigma_n \end{pmatrix}$$

### **Multivariate Normal Distribution - Density**

For the n dimensional multivate normal given on the last slide, its density is given by

$$f(\boldsymbol{Y}|\boldsymbol{\mu},\boldsymbol{\Sigma}) = (2\pi)^{-n/2} \det(\boldsymbol{\Sigma})^{-1/2} \exp\left(-\frac{1}{2}(\boldsymbol{Y}-\boldsymbol{\mu})' \boldsymbol{\Sigma}^{-1} (\boldsymbol{Y}-\boldsymbol{\mu})\right)$$

and its log density is given by

$$\log f(\mathbf{Y}|\boldsymbol{\mu}, \boldsymbol{\Sigma}) = -\frac{n}{2}\log 2\pi - \frac{1}{2}\log \det(\boldsymbol{\Sigma}) - \frac{1}{2}(\mathbf{Y}-\boldsymbol{\mu})' \boldsymbol{\Sigma}^{-1} (\mathbf{Y}-\boldsymbol{\mu})$$

#### Some useful matrix identities

The following come from the Matrix Cookbook Chapters 1 & 2.

$$(AB)' = B'A'$$

$$(A + B)' = A' + B'$$

$$(A')^{-1} = (A^{-1})'$$

$$(ABC ...)^{-1} = ... C^{-1}B^{-1}A^{-1}$$

$$\det(AB) = \det(A)$$

## Maximum Likelihood - $\beta$ (iid)

## Maximum Likelihood - $\sigma^2$ (iid)

## A Quick Example

### Parameters -> Synthetic Data

Lets generate some simulated data where the underlying model is known and see how various regression procedures function.

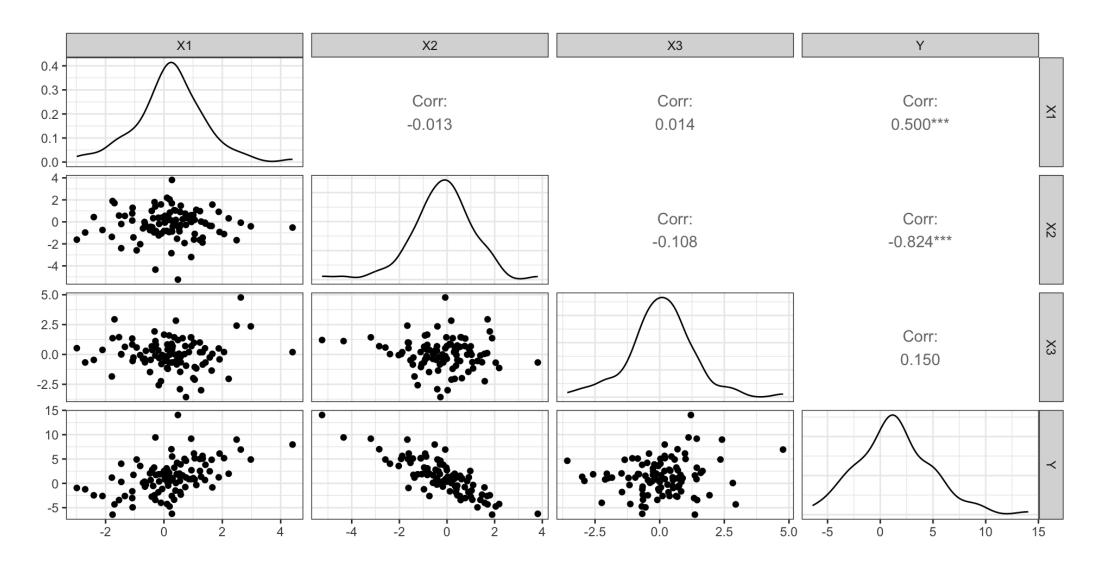
$$Y_i = \beta_0 + \beta_1 X_{1i} + \beta_2 X_{2i} + \beta_3 X_{3i} + \epsilon_i$$
  
 $\epsilon_i \sim N(0, 1)$   
 $\beta_0 = 0.7, \ \beta_1 = 1.5, \ \beta_2 = -2.2, \ \beta_3 =$ 

#### **Model Matrix**

```
1 X = model.matrix(~X1+X2+X3, d)
 2 as tibble(X)
# A tibble: 100 \times 4
   `(Intercept)`
                     X1
                              X2
                                      X3
          <dbl> <dbl> <dbl>
                                   <dbl>
                                 -1.46
 1
                 0.557 0.897
 2
                 0.758
                        0.375
                                 -0.945
 3
                 0.273
                        3.81
                                 -0.675
                        -0.0745
                                 0.514
                 1.41
 4
              1 1.01
                      0.623
 5
                                 -1.99
                 0.942
                        -0.00618 0.700
 6
              1 1.66
                       1.57
                              0.0478
              1 - 1.09 0.766
                                1.33
 8
 9
               1 - 0.296 \quad 1.40
                                 -0.0914
10
               1 - 0.0604 \quad 0.396
                                 -0.0527
# i 90 more rows
```

## Pairs plot

```
1 GGally::ggpairs(d, progress = FALSE)
```



## Least squares fit

Let  $\hat{Y}$  be our estimate for Y based on our estimate of  $oldsymbol{eta}$ ,

$$\hat{\mathbf{Y}} = \hat{\beta_0} + \hat{\beta_1} \, \mathbf{X}_1 + \hat{\beta_2} \, \mathbf{X}_2 + \hat{\beta_3} \, \mathbf{X}_3 = \mathbf{X} \, \hat{\boldsymbol{\beta}}$$

The least squares estimate,  $\hat{m{\beta}_{1s}}$ , is given by

$$\hat{\boldsymbol{\beta}}_{ls} = \underset{\boldsymbol{\beta}}{\operatorname{argmin}} \sum_{i=1}^{n} (Y_i - \boldsymbol{X}_{i\cdot}\boldsymbol{\beta})^2$$

Previously we showed that,

$$\hat{\boldsymbol{\beta}}_{1s} = (\boldsymbol{X}'\boldsymbol{X})^{-1}\boldsymbol{X}'\boldsymbol{Y}$$

#### Beta estimate

```
1 (beta_hat = solve(t(X) %*% X, t(X)) %*% d$Y)
                  [,1]
(Intercept) 0.5522298
X1
         1.4708769
X2
           -2.1761159
Х3
            0.1535830
 1 1 = lm(Y \sim X1 + X2 + X3, data=d)
 2 l$coefficients
(Intercept)
                    X1
                                X2
                                            X3
  0.5522298 1.4708769 -2.1761159 0.1535830
```

# Bayesian regression model

## **Basics of Bayes**

We will be fitting the same model as described above, we just need to provide some additional information in the form of a prior for our model parameters (the  $\beta$ s and  $\sigma^2$ ).

$$f(\theta|\mathbf{x}) = \frac{f(\mathbf{x}|\theta) \pi(\theta)}{\int f(\mathbf{x}|\theta) d\theta}$$
$$\propto f(\mathbf{x}|\theta) \pi(\theta)$$

#### brms

We will be using a package called brms for most of our Bayesian model fitting

- it has a convenient model specification syntax
- mostly sensible prior defaults
- supports most of the model types we will be exploring
- uses Stan behind the scenes

## brms + linear regression

```
1 b = brms::brm(Y \sim X1 + X2 + X3, data=d, chains = 2, silent = 2)
Running /opt/homebrew/Cellar/r/4.3.1/lib/R/bin/R CMD SHLIB foo.c
using C compiler: 'Apple clang version 14.0.3 (clang-1403.0.22.14.1)'
using SDK: ''
clang -I"/opt/homebrew/Cellar/r/4.3.1/lib/R/include" -DNDEBUG
I"/Users/rundel/Library/R/arm64/4.3/library/Rcpp/include/" -
I"/Users/rundel/Library/R/arm64/4.3/library/RcppEigen/include/" -
I"/Users/rundel/Library/R/arm64/4.3/library/RcppEigen/include/unsupported" -
I"/Users/rundel/Library/R/arm64/4.3/library/BH/include" -
I"/Users/rundel/Library/R/arm64/4.3/library/StanHeaders/include/src/" -
I"/Users/rundel/Library/R/arm64/4.3/library/StanHeaders/include/" -
I"/Users/rundel/Library/R/arm64/4.3/library/RcppParallel/include/" -
I"/Users/rundel/Library/R/arm64/4.3/library/rstan/include" -DEIGEN NO DEBUG -DBOOST DISABLE ASSERTS
-DBOOST PENDING INTEGER LOG2 HPP -DSTAN THREADS -DBOOST NO AUTO PTR -include
'/Users/rundel/Library/R/arm64/4.3/library/StanHeaders/include/stan/math/prim/fun/Eigen.hpp' -
D REENTRANT -DRCPP PARALLEL USE TBB=1 -I/opt/homebrew/include
                                                                  -fPIC -I/opt/homebrew/include -c
foo.c -o foo.o
In file included from <built-in>:1:
In file included from
/Users/rundel/Library/R/arm64/4.3/library/StanHeaders/include/stan/math/prim/fun/Eigen.hpp:22:
In file included from /Users/rundel/Library/R/arm64/4.3/library/RcppEigen/include/Eigen/Dense:1:
In file included from /Users/rundel/Library/R/arm64/4.3/library/RcppEigen/include/Eigen/Core:88:
/IIana / mandal / Tibana / D / ama ( / / / 2 / libana / Dama i ana / inalada / Eisan / ana / Gana / util / Masana h. (20.1)
```

#### Model results

```
1 b
 Family: gaussian
  Links: mu = identity; sigma = identity
Formula: Y \sim X1 + X2 + X3
  Data: d (Number of observations: 100)
  Draws: 2 chains, each with iter = 2000; warmup = 1000; thin = 1;
        total post-warmup draws = 2000
Population-Level Effects:
         Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk ESS Tail ESS
                                        0.75 1.00
                                                      2908
             0.55
                       0.10
                               0.36
                                                               1697
Intercept
                                                      2361
                                                               1437
X1
             1.47
                       0.09 1.29 1.64 1.00
X2
            -2.18
                       0.07 \quad -2.32 \quad -2.02 \quad 1.00
                                                      2366
                                                              1528
             0.15
                       0.08
                               -0.01
                                     0.30 1.00
Х3
                                                      2681
                                                               1479
```

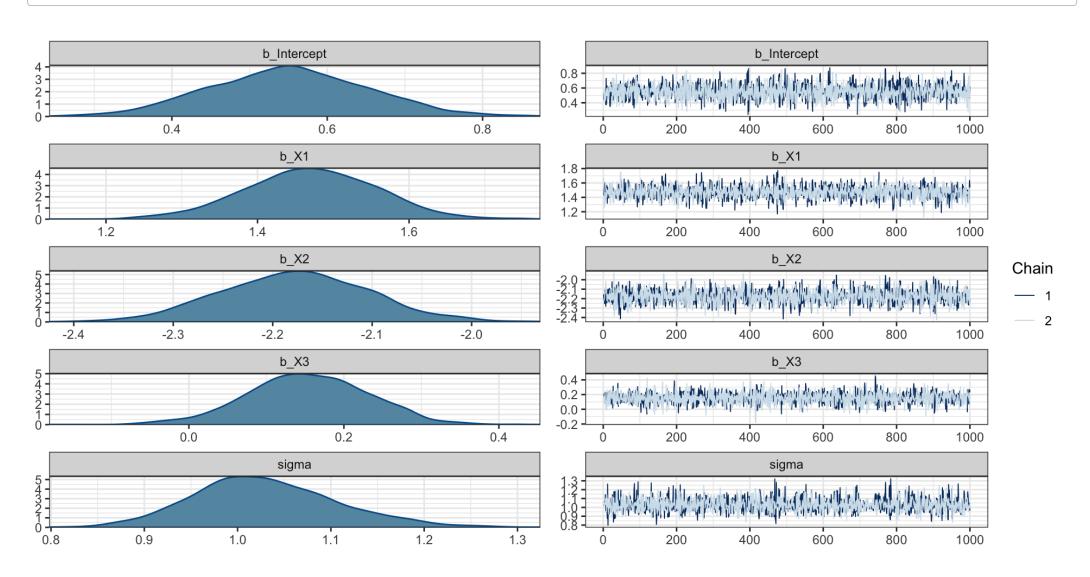
#### Family Specific Parameters:

```
Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS sigma 1.03 0.08 0.89 1.19 1.00 2691 1361
```

Draws were sampled using sampling(NUTS). For each parameter, Bulk\_ESS and Tail\_ESS are effective sample size measures, and Rhat is the potential scale reduction factor on split chains (at convergence, Rhat = 1).

### Model visual summary

1 plot(b)



### What about the priors?

```
1 brms::prior summary(b)
                           class coef group resp dpar nlpar lb ub
                 prior
                                                                         source
                (flat)
                               b
                                                                        default
                (flat)
                                   X1
                                                                    (vectorized)
                (flat)
                                   X2
                                                                   (vectorized)
                (flat)
                                   Х3
                                                                    (vectorized)
                                                                        default
student t(3, 1.1, 3.1) Intercept
  student t(3, 0, 3.1)
                           sigma
                                                                        default
```

#### tidybayes

```
(post = b |>
     tidybayes::gather draws(b Intercept, b X1, b X2, b X3, sigma)
 3)
# A tibble: 10,000 × 5
# Groups: .variable [5]
   .chain .iteration .draw .variable .value
   <int> <int> <int> <chr> <dbl>
       1
                  1
                       1 b Intercept 0.440
 1
                       2 b Intercept 0.524
 2
       1
 3
       1
                        3 b Intercept 0.544
                       4 b Intercept 0.502
                  4
 4
       1
                        5 b Intercept 0.567
 5
       1
                       6 b Intercept 0.569
 6
       1
                  6
 7
                       7 b Intercept 0.373
       1
                       8 b Intercept 0.731
                  8
 8
       1
 9
       1
                        9 b Intercept 0.379
       1
                       10 b Intercept 0.514
10
                 10
# i 9,990 more rows
```

#### tidybayes - posterior summaries

```
1 (post_sum = post |>
2    group_by(.variable, .chain) |>
3    summarize(
4    post_mean = mean(.value),
5    post_median = median(.value),
6    .groups = "drop"
7    )
8  )

# A tibble: 10 × 4
   .variable   .chain post mean post median
```

```
<chr>
        <int> <dbl>
                              <dbl>
               1 0.556
1 b Intercept
                             0.555
2 b Intercept
               2 0.546
                         0.546
               1 1.47
                             1.47
3 b X1
               2 1.47
                             1.47
4 b X1
                   -2.18
                             -2.18
5 b X2
                   -2.18
                             -2.18
6 b X2
7 b X3
                   0.157
                            0.157
8 b X3
                 0.151
                         0.149
9 sigma
                   1.03
                             1.02
10 sigma
                    1.03
                             1.02
```

#### tidybayes + ggplot - traceplot

```
post |>
ggplot(aes(x=.iteration, y=.value, color=as.character(.chain))) +
geom_line(alpha=0.33) +
facet_wrap(~.variable, scale="free_y") +
labs(color="Chain")
```

## Tidy Bayes + ggplot - Density plot

```
post |>
ggplot(aes(x=.value, fill=as.character(.chain))) +
geom_density(alpha=0.5) +
facet_wrap(~.variable, scale="free_x") +
labs(fill="Chain")
```

## **Comparing Approaches**

```
(pt est = post sum |>
     filter(.chain == 1) |>
     ungroup() |>
     mutate(
     truth = c(beta, sigma),
 6
       ols = c(l$coefficients,
                 sd(l$residuals))
     ) |>
     select(
       .variable, truth,
10
11
       ols, post mean
12
13 )
```

## **Comparing Approaches**

```
post |>
filter(.chain == 1) |>
ggplot(aes(x=.value)) +
geom_density(alpha=0.5, fill="lightblue") +
facet_wrap(~.variable, scale="free_x") +
geom_vline(
data = pt_est |> tidyr::pivot_longer(cols = truth:post_mean, names_t
aes(xintercept = value, color=pt_est),
alpha = 0.5, linewidth=1.5
```

## **Comparing Approaches**

