

# Bayes approach

Saneesh

2022-12-02

## Generalized linear models

```
library(brms)
```

```
## Loading required package: Rcpp
```

```
## Loading 'brms' package (version 2.18.0). Useful instructions  
## can be found by typing help('brms'). A more detailed introduction  
## to the package is available through vignette('brms_overview').
```

```
##
```

```
## Attaching package: 'brms'
```

```
## The following object is masked from 'package:stats':
```

```
##
```

```
##      ar
```

```
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.3.2 --
```

```
## v ggplot2 3.4.0      v purrr  0.3.5
```

```
## v tibble  3.1.8      v dplyr  1.0.10
```

```
## v tidyr   1.2.1      v stringr 1.4.1
```

```
## v readr   2.1.3      v forcats 0.5.2
```

```
## -- Conflicts ----- tidyverse_conflicts() --
```

```
## x dplyr::filter() masks stats::filter()
```

```
## x dplyr::lag()    masks stats::lag()
```

```
library(report)
```

```
cardata <- brm(vs ~ mpg * drat,
```

```
              data = mtcars,
```

```
              family = bernoulli())
```

```
## Compiling Stan program...
```

```
## Start sampling
```

```

##
## SAMPLING FOR MODEL '63230c89284e82c150c4ad8a7dc4eb7c' NOW (CHAIN 1).
## Chain 1:
## Chain 1: Gradient evaluation took 0 seconds
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Chain 1: Adjust your expectations accordingly!
## Chain 1:
## Chain 1:
## Chain 1: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 1: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 1: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 1: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 1: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 1: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 1: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 1: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 1: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 1: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 1: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 1: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 1:
## Chain 1: Elapsed Time: 0.366 seconds (Warm-up)
## Chain 1:                0.257 seconds (Sampling)
## Chain 1:                0.623 seconds (Total)
## Chain 1:
##
## SAMPLING FOR MODEL '63230c89284e82c150c4ad8a7dc4eb7c' NOW (CHAIN 2).
## Chain 2:
## Chain 2: Gradient evaluation took 0 seconds
## Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Chain 2: Adjust your expectations accordingly!
## Chain 2:
## Chain 2:
## Chain 2: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 2: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 2: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 2: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 2: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 2: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 2: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 2: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 2: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 2: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 2: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 2: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 2:
## Chain 2: Elapsed Time: 0.317 seconds (Warm-up)
## Chain 2:                0.244 seconds (Sampling)
## Chain 2:                0.561 seconds (Total)
## Chain 2:
##
## SAMPLING FOR MODEL '63230c89284e82c150c4ad8a7dc4eb7c' NOW (CHAIN 3).
## Chain 3:
## Chain 3: Gradient evaluation took 0 seconds

```

```

## Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Chain 3: Adjust your expectations accordingly!
## Chain 3:
## Chain 3:
## Chain 3: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 3: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 3: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 3: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 3: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 3: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 3: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 3: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 3: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 3: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 3: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 3: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 3:
## Chain 3: Elapsed Time: 0.311 seconds (Warm-up)
## Chain 3:                0.271 seconds (Sampling)
## Chain 3:                0.582 seconds (Total)
## Chain 3:
##
## SAMPLING FOR MODEL '63230c89284e82c150c4ad8a7dc4eb7c' NOW (CHAIN 4).
## Chain 4:
## Chain 4: Gradient evaluation took 0 seconds
## Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Chain 4: Adjust your expectations accordingly!
## Chain 4:
## Chain 4:
## Chain 4: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 4: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 4: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 4: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 4: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 4: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 4: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 4: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 4: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 4: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 4: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 4: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 4:
## Chain 4: Elapsed Time: 0.309 seconds (Warm-up)
## Chain 4:                0.278 seconds (Sampling)
## Chain 4:                0.587 seconds (Total)
## Chain 4:

```

```
summary(cardata)
```

```

## Family: bernoulli
## Links: mu = logit
## Formula: vs ~ mpg * drat
## Data: mtcars (Number of observations: 32)
## Draws: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;

```

```

##           total post-warmup draws = 4000
##
## Population-Level Effects:
##           Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## Intercept    -36.54     23.37   -87.11    6.24 1.00     751     1139
## mpg           1.99      1.20    -0.22    4.53 1.00     758     1154
## drat          5.94      5.78    -5.17   17.64 1.00     800     1151
## mpg:drat     -0.34      0.28    -0.91    0.21 1.00     754     1120
##
## 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).

report(cardata)

## Warning: Response residuals not available to calculate mean square error. (R)MSE
##   is probably not reliable.

## Start sampling

##
## SAMPLING FOR MODEL '63230c89284e82c150c4ad8a7dc4eb7c' NOW (CHAIN 1).
## Chain 1:
## Chain 1: Gradient evaluation took 0 seconds
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Chain 1: Adjust your expectations accordingly!
## Chain 1:
## Chain 1:
## Chain 1: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 1: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 1: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 1: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 1: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 1: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 1: Iteration:  1001 / 2000 [ 50%] (Sampling)
## Chain 1: Iteration:  1200 / 2000 [ 60%] (Sampling)
## Chain 1: Iteration:  1400 / 2000 [ 70%] (Sampling)
## Chain 1: Iteration:  1600 / 2000 [ 80%] (Sampling)
## Chain 1: Iteration:  1800 / 2000 [ 90%] (Sampling)
## Chain 1: Iteration:  2000 / 2000 [100%] (Sampling)
## Chain 1:
## Chain 1: Elapsed Time: 0.063 seconds (Warm-up)
## Chain 1:                0.069 seconds (Sampling)
## Chain 1:                0.132 seconds (Total)
## Chain 1:
##
## SAMPLING FOR MODEL '63230c89284e82c150c4ad8a7dc4eb7c' NOW (CHAIN 2).
## Chain 2:
## Chain 2: Gradient evaluation took 0 seconds
## Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Chain 2: Adjust your expectations accordingly!
## Chain 2:
## Chain 2:

```

```

## Chain 2: Iteration:    1 / 2000 [ 0%] (Warmup)
## Chain 2: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 2: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 2: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 2: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 2: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 2: Iteration:  1001 / 2000 [ 50%] (Sampling)
## Chain 2: Iteration:  1200 / 2000 [ 60%] (Sampling)
## Chain 2: Iteration:  1400 / 2000 [ 70%] (Sampling)
## Chain 2: Iteration:  1600 / 2000 [ 80%] (Sampling)
## Chain 2: Iteration:  1800 / 2000 [ 90%] (Sampling)
## Chain 2: Iteration:  2000 / 2000 [100%] (Sampling)
## Chain 2:
## Chain 2: Elapsed Time: 0.072 seconds (Warm-up)
## Chain 2:                0.072 seconds (Sampling)
## Chain 2:                0.144 seconds (Total)
## Chain 2:
##
## SAMPLING FOR MODEL '63230c89284e82c150c4ad8a7dc4eb7c' NOW (CHAIN 3).
## Chain 3:
## Chain 3: Gradient evaluation took 0 seconds
## Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Chain 3: Adjust your expectations accordingly!
## Chain 3:
## Chain 3:
## Chain 3: Iteration:    1 / 2000 [ 0%] (Warmup)
## Chain 3: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 3: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 3: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 3: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 3: Iteration:  1000 / 2000 [ 50%] (Warmup)
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## Chain 3: Iteration:  1400 / 2000 [ 70%] (Sampling)
## Chain 3: Iteration:  1600 / 2000 [ 80%] (Sampling)
## Chain 3: Iteration:  1800 / 2000 [ 90%] (Sampling)
## Chain 3: Iteration:  2000 / 2000 [100%] (Sampling)
## Chain 3:
## Chain 3: Elapsed Time: 0.061 seconds (Warm-up)
## Chain 3:                0.065 seconds (Sampling)
## Chain 3:                0.126 seconds (Total)
## Chain 3:
##
## SAMPLING FOR MODEL '63230c89284e82c150c4ad8a7dc4eb7c' NOW (CHAIN 4).
## Chain 4:
## Chain 4: Gradient evaluation took 0 seconds
## Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Chain 4: Adjust your expectations accordingly!
## Chain 4:
## Chain 4:
## Chain 4: Iteration:    1 / 2000 [ 0%] (Warmup)
## Chain 4: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 4: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 4: Iteration:   600 / 2000 [ 30%] (Warmup)

```

```

## Chain 4: Iteration: 800 / 2000 [ 40%] (Warmup)
## Chain 4: Iteration: 1000 / 2000 [ 50%] (Warmup)
## Chain 4: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 4: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 4: Iteration: 1400 / 2000 [ 70%] (Sampling)
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## Chain 4: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 4: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 4:
## Chain 4: Elapsed Time: 0.062 seconds (Warm-up)
## Chain 4: 0.069 seconds (Sampling)
## Chain 4: 0.131 seconds (Total)
## Chain 4:

## Warning: Response residuals not available to calculate mean square error. (R)MSE
## is probably not reliable.

## We fitted a Bayesian logistic model (estimated using MCMC sampling with 4
## chains of 2000 iterations and a warmup of 1000) to predict vs with mpg
## (formula: vs ~ mpg * drat). Priors over parameters were set as student_t
## (location = 0.00, scale = 2.50) distributions. The model's explanatory power is
## substantial (R2 = 0.51, 95% CI [0.40, 0.57]). Within this model:
##
## - The effect of b Intercept (Median = -35.37, 95% CI [-87.11, 6.24]) has a
## 95.17% probability of being negative (< 0), 95.15% of being significant (<
## -0.09), and 94.92% of being large (< -0.54). The estimation successfully
## converged (Rhat = 1.001) but the indices are unreliable (ESS = 798)
## - The effect of b mpg (Median = 1.95, 95% CI [-0.22, 4.53]) has a 96.20%
## probability of being positive (> 0), 95.40% of being significant (> 0.09), and
## 88.52% of being large (> 0.54). The estimation successfully converged (Rhat =
## 1.001) but the indices are unreliable (ESS = 753)
## - The effect of b drat (Median = 5.84, 95% CI [-5.17, 17.64]) has a 84.50%
## probability of being positive (> 0), 84.03% of being significant (> 0.09), and
## 82.15% of being large (> 0.54). The estimation successfully converged (Rhat =
## 1.002) but the indices are unreliable (ESS = 756)
## - The interaction effect of drat on b mpg (Median = -0.34, 95% CI [-0.91,
## 0.21]) has a 88.55% probability of being negative (< 0), 81.08% of being
## significant (< -0.09), and 23.08% of being large (< -0.54). The estimation
## successfully converged (Rhat = 1.001) but the indices are unreliable (ESS =
## 750)
##
## Following the Sequential Effect eXistence and sIgnificance Testing (SEXIT)
## framework, we report the median of the posterior distribution and its 95% CI
## (Highest Density Interval), along the probability of direction (pd), the
## probability of significance and the probability of being large. The thresholds
## beyond which the effect is considered as significant (i.e., non-negligible) and
## large are |0.09| and |0.54|. Convergence and stability of the Bayesian sampling
## has been assessed using R-hat, which should be below 1.01 (Vehtari et al.,
## 2019), and Effective Sample Size (ESS), which should be greater than 1000
## (Burkner, 2017). We fitted a Bayesian logistic model (estimated using MCMC
## sampling with 4 chains of 2000 iterations and a warmup of 1000) to predict vs
## with drat (formula: vs ~ mpg * drat). Priors over parameters were set as
## uniform (location = , scale = ) distributions. The model's explanatory power is
## substantial (R2 = 0.51, 95% CI [0.40, 0.57]). Within this model:

```

```

##
## - The effect of b Intercept (Median = -35.37, 95% CI [-87.11, 6.24]) has a
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## (Highest Density Interval), along the probability of direction (pd), the
## probability of significance and the probability of being large. The thresholds
## beyond which the effect is considered as significant (i.e., non-negligible) and
## large are |0.09| and |0.54|. Convergence and stability of the Bayesian sampling
## has been assessed using R-hat, which should be below 1.01 (Vehtari et al.,
## 2019), and Effective Sample Size (ESS), which should be greater than 1000
## (Burkner, 2017). We fitted a Bayesian logistic model (estimated using MCMC
## sampling with 4 chains of 2000 iterations and a warmup of 1000) to predict vs
## with mpg (formula: vs ~ mpg * drat). Priors over parameters were set as uniform
## (location = , scale = ) distributions. The model's explanatory power is
## substantial (R2 = 0.51, 95% CI [0.40, 0.57]). Within this model:
##
## - The effect of b Intercept (Median = -35.37, 95% CI [-87.11, 6.24]) has a
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## successfully converged (Rhat = 1.001) but the indices are unreliable (ESS =
## 750)
##
## Following the Sequential Effect eXistence and sIgnificance Testing (SEXIT)
## framework, we report the median of the posterior distribution and its 95% CI
## (Highest Density Interval), along the probability of direction (pd), the

```

```
## probability of significance and the probability of being large. The thresholds
## beyond which the effect is considered as significant (i.e., non-negligible) and
## large are |0.09| and |0.54|. Convergence and stability of the Bayesian sampling
## has been assessed using R-hat, which should be below 1.01 (Vehtari et al.,
## 2019), and Effective Sample Size (ESS), which should be greater than 1000
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## with drat (formula: vs ~ mpg * drat). Priors over parameters were set as
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## 1.001) but the indices are unreliable (ESS = 753)
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## 0.21]) has a 88.55% probability of being negative (< 0), 81.08% of being
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## successfully converged (Rhat = 1.001) but the indices are unreliable (ESS =
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## Following the Sequential Effect eXistence and sIgnificance Testing (SEXIT)
## framework, we report the median of the posterior distribution and its 95% CI
## (Highest Density Interval), along the probability of direction (pd), the
## probability of significance and the probability of being large. The thresholds
## beyond which the effect is considered as significant (i.e., non-negligible) and
## large are |0.09| and |0.54|. Convergence and stability of the Bayesian sampling
## has been assessed using R-hat, which should be below 1.01 (Vehtari et al.,
## 2019), and Effective Sample Size (ESS), which should be greater than 1000
## (Burkner, 2017).
```

Mixed model with random slopes.

### Question

>Is there a general relationship between petal and sepal width? and how it differs by species?

```
library(brms)
library(tidyverse)
library(report)

ggplot(iris, aes(Petal.Width, Sepal.Width, colour= Species))+
  geom_point()
```



