Bayesian Estimation of Recruitment Trends in Alberta (BERTA) Tutorial

Christopher L. Cahill

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Goals

- Understand general structure of BERTA model used in Cahill et al. (2021).
- Gain an overview of available modeling options, including tweaking priors and MCMC run parameters.
- Become familiar with browser(), get_fit(), future_pwalk().
- Learn how .R and .stan scripts are working together to subset data, fit a Bayesian stock reduction analysis model to those data, and then save the model fit with a unique file name identifier.
- Practice debugging.

Packages

Let's load the packages we will use:

```
library(tidyverse)
## -- Attaching packages ------ tidyverse 1.3.1 --
## v ggplot2 3.3.5
                    v purrr
                            0.3.4
## v tibble 3.1.3
                   v dplyr
                            1.0.7
## v tidyr
         1.1.3
                   v stringr 1.4.0
                    v forcats 0.5.1
## v readr
           2.0.1
## -- Conflicts ----- tidyverse conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                  masks stats::lag()
library(rstan)
## Warning: package 'rstan' was built under R version 4.1.2
## Loading required package: StanHeaders
## rstan (Version 2.21.3, GitRev: 2e1f913d3ca3)
```

```
## For execution on a local, multicore CPU with excess RAM we recommend calling
## options(mc.cores = parallel::detectCores()).
## To avoid recompilation of unchanged Stan programs, we recommend calling
## rstan_options(auto_write = TRUE)

## Do not specify '-march=native' in 'LOCAL_CPPFLAGS' or a Makevars file

##
## Attaching package: 'rstan'

## The following object is masked from 'package:tidyr':
##
## extract

library(furrr)

## Loading required package: future

library(future)
library(ggplot2)
```

Data

We will work with the Fall Walleye Index Netting (FWIN) dataset used in Cahill et al. 2021, which included all lakes with ≥ 3 FWIN surveys during 2000-2018:

```
data <- readRDS(here::here("data/BERTA-wide-0-25.rds"))
glimpse(data)</pre>
```

```
## Rows: 236
## Columns: 46
## Groups: name [55]
                <int> 3526, 3526, 3526, 3526, 3916, 3916, 3916, 3969, 3969, 3969, ~
## $ WBID
## $ year
                <dbl> 5, 6, 13, 19, 6, 11, 14, 9, 12, 15, 17, 7, 12, 17, 4, 11, 1~
                <chr> "milk river ridge reservoir", "milk river ridge reservoir",~
## $ name
## $ nnet
                <int> 18, 20, 11, 12, 8, 12, 10, 11, 12, 12, 12, 18, 15, 15, 4, 6~
## $ n
                <int> 201, 283, 232, 158, 189, 117, 132, 357, 171, 186, 201, 373,~
## $ effort
                <dbl> 18.0, 20.0, 11.0, 12.0, 8.0, 6.0, 5.0, 11.0, 6.0, 6.0, 6.0,~
## $ X_TTM_c
                <dbl> 676088.4, 676088.4, 676088.4, 676088.4, 652638.4, 652638.4, 
                <dbl> 5469124, 5469124, 5469124, 5469124, 6050150, 6050150, 60501~
## $ Y_TTM_c
## $ p_aged
                <dbl> 1.0000000, 1.0000000, 1.0000000, 0.9430380, 1.0000000, 0.98~
                <dbl> 12.22278, 12.22278, 12.22278, 12.22278, 13.93477, 13.93477,~
## $ omega
## $ linf
                <dbl> 56.72357, 56.72357, 56.72357, 56.72357, 51.38603, 51.38603,~
## $ vbk
                <dbl> 0.2154797, 0.2154797, 0.2154797, 0.2154797, 0.2711781, 0.27~
## $ a50
                <dbl> 7, 7, 7, 7, 4, 4, 4, 5, 5, 5, 5, 5, 5, 5, 5, 4, 4, 4, 4, 5, 5,~
                <dbl> 3.409773, 3.409773, 3.409773, 3.409773, 3.100920, 3.100920,~
## $ beta_wl
## $ X_long
                <dbl> -112.5735, -112.5735, -112.5735, -112.5735, -112.6363, -112~
## $ Y_lat
                <dbl> 49.36904, 49.36904, 49.36904, 49.36904, 54.59751, 54.59751,~
                <dbl> 1355.0, 1355.0, 1355.0, 1355.0, 527.1, 527.1, 527.1, 970.7,~
## $ Area_Ha
                <int> 1605, 1605, 1605, 1605, 1293, 1293, 1293, 1293, 1293, 1293, ~
## $ DD5
```

```
## $ Max_Depth <dbl> NA, NA, NA, NA, 27.5, 27.5, 27.5, 27.4, 27.4, 27.4, 27.4, 1~
## $ Mean_Depth <dbl> NA, NA, NA, NA, 14.3, 14.3, 14.3, 9.2, 9.2, 9.2, 9.2, 6.9, ~
## $ '1'
              <dbl> 23, 21, 5, 2, 0, 4, 4, 2, 10, 3, 23, 24, 5, 6, 1, 18, 0, 0,~
## $ '2'
              <dbl> 13, 67, 43, 6, 0, 4, 7, 24, 4, 14, 41, 11, 5, 10, 7, 14, 0,~
## $ '3'
              <dbl> 18, 55, 13, 8, 5, 17, 44, 53, 5, 12, 14, 16, 5, 4, 15, 13, ~
## $ '4'
              <dbl> 30, 46, 10, 11, 3, 16, 13, 78, 4, 8, 17, 50, 11, 2, 7, 5, 1~
## $ '5'
              <dbl> 27, 35, 17, 13, 41, 3, 8, 15, 12, 0, 8, 55, 5, 37, 17, 9, 1~
## $ '6'
              <dbl> 26, 20, 16, 30, 80, 1, 23, 6, 26, 11, 7, 38, 5, 18, 0, 7, 7~
## $ '7'
              <dbl> 16, 11, 18, 40, 45, 2, 11, 8, 37, 1, 4, 27, 6, 15, 0, 5, 8,~
## $ '8'
              <dbl> 23, 10, 14, 11, 9, 1, 3, 4, 4, 12, 4, 39, 4, 5, 2, 1, 6, 45~
## $ '9'
              <dbl> 16, 7, 20, 8, 2, 1, 1, 14, 1, 23, 0, 7, 8, 12, 0, 4, 4, 85,~
## $ '10'
              <dbl> 6, 1, 14, 3, 2, 18, 1, 26, 3, 38, 7, 10, 17, 7, 3, 4, 3, 3,~
## $ '11'
              <dbl> 0, 1, 3, 6, 0, 30, 1, 38, 6, 9, 16, 23, 6, 1, 0, 5, 3, 1, 0~
## $ '12'
              <dbl> 0, 0, 0, 3, 0, 12, 0, 45, 2, 3, 25, 39, 1, 3, 0, 2, 1, 0, 0~
## $ '13'
              <dbl> 0, 0, 1, 2, 1, 0, 5, 3, 14, 2, 5, 3, 9, 0, 2, 0, 0, 0, 0~
## $ '14'
              <dbl> 0, 1, 1, 0, 1, 1, 6, 18, 7, 5, 3, 1, 0, 4, 0, 1, 1, 1, 0, 0~
## $ '15'
              <dbl> 0, 0, 0, 1, 0, 0, 2, 10, 5, 4, 1, 0, 1, 5, 0, 0, 0, 0, 0, 0
## $ '16'
              <dbl> 0, 0, 0, 1, 0, 0, 2, 2, 3, 8, 2, 0, 1, 4, 0, 0, 0, 0, 0, 0, ~
## $ '17'
              <dbl> 0, 0, 0, 0, 0, 0, 0, 3, 3, 12, 3, 0, 7, 3, 0, 0, 0, 0, 0~
## $ '18'
              <dbl> 0, 0, 0, 0, 0, 0, 0, 1, 2, 10, 3, 0, 1, 4, 0, 0, 0, 0, 0~
## $ '19'
              <dbl> 0, 0, 0, 0, 0, 0, 0, 0, 2, 0, 1, 0, 3, 0, 0, 0, 0, 0, 0, 0, ~
## $ '20'
              <dbl> 0, 0, 0, 0, 0, 0, 0, 0, 1, 2, 4, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0
## $ '21'
              <dbl> 0, 0, 0, 0, 0, 0, 0, 0, 0, 3, 0, 0, 0, 1, 0, 0, 0, 0, 0, ~
## $ '22'
              <dbl> 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 2, 0, 0, 0, 0, 0, ~
## $ '23'
              ## $ '24'
              ## $ '25'
              ## $ lake
              <dbl> 1, 1, 1, 1, 2, 2, 2, 3, 3, 3, 4, 4, 4, 5, 5, 5, 5, 6, 6,~
```

```
stocking <- readRDS(here::here("data/stocking_matrix_ha.rds"))
glimpse(stocking)</pre>
```

```
## num [1:106, 1:39] 0 0 0 0 0 0 0 0 0 0 0 0 ...
## - attr(*, "dimnames")=List of 2
## ..$ : chr [1:106] "berry creek reservoir" "jensen reservoir" "milk river ridge reservoir" "travers
## ..$ : NULL
```