

Skagit Creel Analysis

Thomas Buehrens, Kale Bentley, Andrew Fowler & Amy Edwards

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Purpose

The purpose of this document is to record the steps and code necessary to reproduce the Skagit steelhead fishery creel analysis for 2021.

Requirements

All analyses require R software ([link](#)) (v3.4.3) for data retrieval, data processing, and summarizing model results, and Stan software ([link](#)) for Hamiltonian Monte Carlo (HMC) simulation. For Stan to work, rtools must also be installed: ([link](#)).

Functions

We also need a couple of helper functions which we will load from the functions folder, which we will load using `sapply`

```
wd_functions<-"functions"  
sapply(FUN = source, paste(wd_functions, list.files(wd_functions), sep="/"))
```

Packages

In addition to `purrr`, we also need a few packages that are not included with the base installation of R, so we begin by installing them (if necessary) and then loading them.

```
#####  
# Load packages, install and load if not already  
#####  
using("tidyverse",  
      "rstan",  
      "RColorBrewer",  
      "readxl",  
      "readr",  
      #"ggplot2",  
      "tinytex",  
      "here",  
      "plyr",  
      "lubridate",  
      "devtools",  
      "xlsx",  
      "cowplot",  
      "ggpubr",  
      "chron",  
      "suncalc",  
      "shinytan",  
      "loo",  
      "data.table",  
      "RColorBrewer",  
      "reshape2",  
      "MASS",  
      "timeDate"  
)
```

User inputs

```
#####  
# Specify relative working directories for sub-folders  
#####  
wd_LUTs <-"lookup tables"      # Location of look-up tables (maybe could be merged with data files??)  
wd_data  <-"data"              # Location where data files are stored  
wd_source_files<-"source files" # Location of source file (code working "behind the scenes")
```

```

wd_models <- "models"           # Location of model files
wd_outputs <- "results"         # Location of saved output (summary figures/tables and model results)

#####
# Specify names of .csv data files
#####
effort_file_name<- "03_Effort_dat - improved_tiein_entry - 2019_Skagit_creel_JSH_2-17-19.csv"
interview_file_name<- "03_Interview-dat_hand-corr-times_2-28-2019.csv"

#####
# Denote data of interest (used to filter data below)
#####
# Specify filter type(s) to extract data by (Enter "Y" or "N")
by.Year<- "N" # If "Y", will filter by full calendar year(s) (Jan. 1 - Dec. 31)
by.YearGroup<- "N" # If "Y", will filter by a "Year Group", which go from May 1st Yr1 - April 30 Yr2
by.Season<- "N" # If "Y", will filter by "season", which is either Summer (May 1st - Oct 31st) or
by.StreamName<- "Y" # If "Y", will filter by stream name
by.Date<- "N" # If "Y", will filter by a date range

# Specify date ranges for "Year Groups" and "Seasons"
YearBegin<- 121 # day of year a "YearGroup" begins (FYI - 121 = May 1st in a non-leap year)
summerBegin<-121
summerEnd<- 304 # FYI - 304 = Oct. 31st (in a non-leap year)
winterBegin<-305
winterEnd<- 120

# Specify filter unit(s)
YearGroup.of.Interest<- c("2017-2018")
Season.of.Interest<- c("Winter")
Year.of.Interest<- c("2017")
StreamName.of.Interest<-c("Skagit")
Begin.Date<- c("2016-05-01") #Format must be "yyyy-mm-dd"
End.Date<- c("2017-03-31") #Format must be "yyyy-mm-dd"

#####
# Denote catch group of interest (species_origin_fate)
#####
catch.group.of.interest<-c("SH_W_R")

#####
# Identify dates when fishery was closed by section
#####
total.closed.dates<-0 # Total number of dates that at least one section of the river was closed (i.e.

# NOTE: if "total.closed.dates" >0, use the following format to enter closure dates and section, where
# the first column is the list of individual dates (by row) the fishery was closed date
# the number of additional columns equals the number of "final" sections based on "final.effort.s
# the enter the following values below each section:
# Enter 1 if the section was open and enter 0 if the section was closed

# Date , Section-1, Section-2
closed.Dates.Sections<-c("2019-02-11", 0, 0 )

```

Data Preparation

```
# Load LUTs
source(paste0(wd_source_files, "/Load_LUTs.R"))

# Load creel data and format
source(paste0(wd_source_files, "/Import_Skagit_Creel_Data_and_Format.R"))

# Extract data of interest and format
## add code that shows options for filtering data by date/year/season/location
source(paste0(wd_source_files, "/05_Extract_Data_of_Interest_and_Calculate_Fields_2019-04-08.R"))

#Run source summary file
## add code that shows options for "catch groups"
source(paste0(wd_source_files, "/06_Summarize_Effort_and_Catch_Data_for_TimeSeries_Model_2019-04-23.R"))

##KB note: I will work on updating the code in the "05" and "06" file at some point soon; also, creatin
```

Run Analysis

```
# Denote whether you want to run a new model or load "saved" results from a previous model run
model_source<-c("load_saved") #enter either "run_new" or "load_saved"

# Assign a "Model_Run" number (if model_source == run_new, results will be saved to a new sub-folder; i
Model_Run<-4 #Enter numeric number (NOTE: be careful not to over-write previous models runs by enteri

# Denote which creel model you want to run
creel_models[,1:3] #model summary table
model_number<-c(4)

# Specify time period to stratify data by - day vs. week
model_period<-c("day") #enter "day" or "week"

# Specify parameter values for model priors
value_cauchyDF_sigma_eps_C = 1 # the hyperhyper scale (degrees of freedom) parameter in the hyperprior
value_cauchyDF_sigma_eps_E = 1 # the hyperhyper scale (degrees of freedom) parameter in the hyperprior
value_cauchyDF_sigma_r_E = 1 # the hyperhyper scale (degrees of freedom) parameter in the hyperprior
value_cauchyDF_sigma_r_C = 1 # the hyperhyper scale (degrees of freedom) parameter in the hyperprior
value_normal_sigma_omega_C_0 = 1 #the SD hyperparameter in the prior distribution omega_C_0; normal
value_normal_sigma_omega_E_0 = 3 # the SD hyperparameter in the prior distribution omega_E_0; normal
value_lognormal_sigma_b = 1 # the SD hyperparameter in the prior distribution b; default = 1
value_normal_sigma_B1 = 5 # the SD hyperparameter in the prior distribution B1; default = 5
value_normal_mu_mu_C = log(0.02) # the mean hyperparameter in the prior distribution mu_C; median (log
value_normal_sigma_mu_C = 1.5 # the SD hyperparameter in the prior distribution mu_C; normal sd (log-
value_normal_mu_mu_E = log(15) # the mean hyperparameter in the prior distribution mu_E; median effort
value_normal_sigma_mu_E = 2 # the SD hyperparameter in the prior distribution mu_E; normal sd (log-s
value_betashape_phi_E_scaled = 1 # the rate (alpha) and shape (beta) hyperparameters in phi_E_scaled;
value_betashape_phi_C_scaled = 1 # the rate (alpha) and shape (beta) hyperparameters in phi_C_scaled;
value_cauchyDF_sigma_mu_C = 1 # the hyperhyper SD parameter in the hyperprior distribution sigma
value_cauchyDF_sigma_mu_E = 1 # the hyperhyper SD parameter in the hyperprior distribution sigma
```

```

# Specific Stan model arguments
n_chain<-4          # set the number of Markov chains. The default is 4.
n_iter<-200         # set the number of iterations for each chain (including warmup). The default is 2
n_cores<-4          # set the number of cores to use when executing the chains in parallel. The default
n_warmup<-100       # set the length of warm-up (aka burn-in) iterations per chain. The default is n
n_thin<-1           # set the thinning rate (aka, the period for saving samples). The default is 1, wh
adapt_delta<-0.80    # set adapt delta, which is the target average proposal acceptance probability dur
max_treepdepth<-10  # set the max tree depth; default is 8; NOTE: this sets the max depth of tree used

# Create sub-folders for output (if they don't already exist)
source(paste0(wd_source_files, "/Create_output_subfolder.R"), print.eval = TRUE)

# Run source code to prepare data for model
source(paste0(wd_source_files, "/Prepare_Data_For_Model.R "))

# Run source code to generate creel estimates
source(paste0(wd_source_files, "/RunNew_or_LoadSaved_Creel_Model.R"))

# Generate summaries of model inputs and outputs
if(model_source == "run_new"){ source(paste0(wd_source_files, "/Summarize_Model_Inputs_and_Outputs.R

```

Summarize Results

```

#convergence diagnostics
launch_diagnostics<-c("No") #Enter "Yes" to launch ShinyShin diagnostics
if(launch_diagnostics=="Yes"){launch_shinystan(output$res_stan)}

# generate plots and tables of creel estimates
source(paste0(wd_source_files, "/Generate_Summaries_of_Creel_Estimates.R"))

# KB note: update so table/plots of results are shown in PDF document

```

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In order to reproduce this pdf or html page you need to have a LaTeX application installed. Running this snippet of code will automatically install tinytex on your machine so you can render pdfs and html:

```

#tinytex::install_tinytex()

```