### Assignment 6

#### anonymous

#### 1 General information

#### Reporting accuracy

For posterior statistics of interest, only report digits for which the Monte Carlo standard error (MCSE) is zero.

Example: If you estimate  $E(\mu) = 1.234$  with  $MCSE(E(\mu)) = 0.01$ , you should report  $E(\mu) = 1.2$ .

See lecture video 4.1, the chapter notes, and a case study for more information

This is the template for assignment 6. You can download the qmd-file or copy the code from this rendered document after clicking on </> Code in the top right corner.

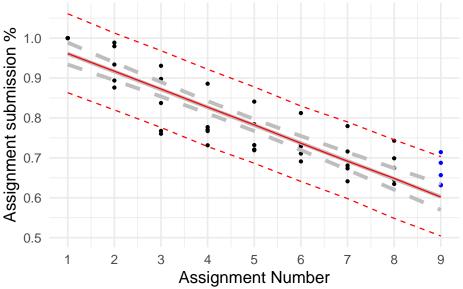
Please replace the instructions in this template by your own text, explaining what you are doing in each exercise.

## 2 Stan warm-up: linear model of BDA retention with Stan (2 points)

Write your answers/code here!

```
assignment <- rep(1:8, 5)
# These are in some sense our test data: the proportion of students handing in the last a
# sorted by year.
# Usually, we would not want to split our data like that and instead
# use e.g. Leave-One-Out Cross-Validation (LOO-CV, see e.g. http://mc-stan.org/loo/index.
# to evaluate model performance.
propstudents9 = c(121/176, 153/242, 218/332, 190/301, 175/245)
# The total number of assignments
no_assignments = 9
# The assignment numbers for which we want to generate predictions
x_predictions = 1:no_assignments
# (Cmd)Stan(R) expects the data to be passed in the below format:
model_data = list(N=length(assignment),
                 x=assignment,
                 y=propstudents,
                 no_predictions=no_assignments,
                 x_predictions=x_predictions)
# This reads the file at the specified path and tries to compile it.
# If it fails, an error is thrown.
retention_model = cmdstan_model("./assignment6_linear_model.stan")
# This "out <- capture.output(...)" construction suppresses output from cmdstanr
# See also https://github.com/stan-dev/cmdstanr/issues/646
out <- capture.output(</pre>
    # Sampling from the model happens here:
   fit <- retention_model$sample(data=model_data, refresh=0, show_messages=FALSE)</pre>
# This extracts the draws from the sampling result as a data.frame.
draws_df = fit$draws(format="draws_df")
# This does some data/draws wrangling to compute the 5, 50 and 95 percentiles of
# the mean at the specified covariate values (x_predictions).
# It can be instructive to play around with each of the data processing steps
# to find out what each step does, e.g. by removing parts from the back like "|> gather(
# and printing the resulting data.frame.
mu_quantiles_df = draws_df |>
      select(starts_with(c("mu_pred"))) |>
      apply(2, quantile, c(0.05, 0.5, 0.95)) |>
      t() |> data.frame(x=x_predictions) |> gather(pct,y,-x)
# Same as above, but for the predictiions.
y_quantiles_df = draws_df |>
      select(starts_with(c("y_pred"))) |>
      apply(2, quantile, c(0.05, 0.5, 0.95)) |>
      t() |> data.frame(x=x_predictions) |> gather(pct,y,-x)
```

```
# Plotting happens here:
ggplot() +
  # scatter plot of the training data:
  geom_point(aes(x, y), data = data.frame(x=assignment, y=propstudents), size = 1) +
  # scatter plot of the test data:
  geom_point(aes(x, y), data = data.frame(x=no_assignments, y=propstudents9), size = 1, or example of the size = 1.
  # you have to tell us what this plots:
  geom_line(aes(x,y,linetype=pct), data=mu_quantiles_df, color = 'grey', linewidth=1.5) +
  # you have to tell us what this plots:
  geom_line(aes(x,y,linetype=pct), data=y_quantiles_df, color = 'red') +
  # adding xticks for each assignment:
  scale_x_continuous(breaks=1:no_assignments) +
  # adding labels to the plot:
 labs(y = "Assignment submission %", x= "Assignment Number") +
  # specifying that line types repeat:
  scale_linetype_manual(values=c(2,1,2)) +
  # remove the legend for the linetypes:
  guides(linetype = "none")
```



Write your answers/code here!

# 3 Generalized linear model: Bioassay with Stan (4 points)

Write your answers/code here!

```
data("bioassay")
```

Write your answers/code here!

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