

# Assignment 8

## LOO-CV model comparison

anonymous

### 1 General information

This is the template for [assignment 8](#). 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 A hierarchical model for chicken weight time series

#### 2.1 Exploratory data analysis

#### 2.2 (a)

```
# Useful functions: ggplot, aes(x=...), geom_histogram
```

#### 2.3 (b)

```
# Useful functions: ggplot, aes(x=...,y=...,group=...,color=...), geom_line
```

#### 2.4 Linear regression

#### 2.5 (c)

In `brms`, a regression can be specified as below, see also [below \(#m\)](#) or [the last template](#). Fill in the appropriate variables, data, and likelihood family. Specify the priors, then run the model (by removing `#! eval: false` below).

```
priors <- c(
  prior(normal(0, <value>), coef = "Time"),
  prior(normal(0, <value>), coef = "Diet2"),
  prior(normal(0, <value>), coef = "Diet3"),
  prior(normal(0, <value>), coef = "Diet4")
)

f1 <- brms::brm(
```

```

# This specifies the formula
<OUTCOME> ~ 1 + <PREDICTOR> + <PREDICTOR>,
# This specifies the dataset
data = <data>,
# This specifies the observation model family
family = <observation_family>,
# This passes the priors specified above to brms
prior = priors,
# This causes brms to cache the results
file = "additional_files/assignment8/f1"
)

```

## 2.6 (d)

```

# Useful functions: brms::pp_check

```

## 2.7 (e)

```

# Useful functions: brms::pp_check(..., type = ..., group=...)

```

## 2.8 Log-normal linear regression

### 2.9 (f)

```

log_priors <- c(
  prior(normal(0, log(3)), coef = "Time"),
  prior(normal(0, log(5)), coef = "Diet2"),
  prior(normal(0, log(5)), coef = "Diet3"),
  prior(normal(0, log(5)), coef = "Diet4")
)

```

## 2.10 Hierarchical log-normal linear regression

### 2.11 (g)

### 2.12 (h)

## 2.13 Model comparison using the ELPD

### 2.14 (i)

```

# Useful functions: loo, loo_compare

```

## 2.15 (j)

```
# Useful functions: plot(loo(...), label_points = TRUE)
```

## 2.16 (k)

## 2.17 Model comparison using the RMSE

## 2.18 (l)