

MTHM017 - Advanced Topics in Statistics, Assignment

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Declaration of AI Assistance: I have used OpenAI's ChatGPT tool in creating this report.

AI-supported/AI-integrated use is permitted in this assessment. I acknowledge the following uses of GenAI tools in this assessment:

1. I have used GenAI tools to check and debug my code.
2. I have used GenAI tools to proofread and correct grammar or spelling errors.
3. I have used GenAI tools to give me feedback on a draft.

I declare that I have referenced use of GenAI outputs within my assessment in line with the University referencing guidelines.

A. Bayesian Inference

1.

[6 marks] Read in the data, then for each person produce a histogram of that given person's reaction times. The range of the x axis should be the same on each histogram. Visually compare the reaction time distributions of schizophrenic and non-schizophrenic individuals. What differences/similarities can you observe? Reference the histograms of specific individuals to support your conclusions.

I will begin by making the dataset more usable and clear.

```
# Renaming first column
names(rtimes)[1] <- "PatientType"
# Classifying the patient type, 1-11 non-schiz, 12-17 schiz
rtimes$PatientType <- c(rep("non-schizophrenic", 11), rep("schizophrenic", 6))
rtimes1 <- rtimes
head(rtimes1)
```

##		PatientType	T1	T2	T3	T4	T5	T6	T7	T8	T9	T10	T11	T12	T13	T14	T15
## 1	non-schizophrenic	312	272	350	286	268	328	298	356	292	308	296	372	396	402	280	
## 2	non-schizophrenic	354	346	384	342	302	312	322	376	306	402	320	298	308	414	304	
## 3	non-schizophrenic	256	284	320	274	324	268	370	430	314	312	362	256	342	388	302	
## 4	non-schizophrenic	260	294	306	292	264	290	272	268	344	362	330	280	354	320	334	
## 5	non-schizophrenic	204	272	250	260	314	308	246	236	208	268	272	264	308	236	238	
## 6	non-schizophrenic	590	312	286	310	778	364	318	316	316	298	344	262	274	330	312	
##		T16	T17	T18	T19	T20	T21	T22	T23	T24	T25	T26	T27	T28	T29	T30	
## 1		330	254	282	350	328	332	308	292	258	340	242	306	328	294	272	
## 2		422	388	422	426	338	332	426	478	372	392	374	430	388	354	368	
## 3		366	298	396	274	226	328	274	258	220	236	272	322	284	274	356	
## 4		276	418	288	338	350	350	324	286	322	280	256	218	256	220	356	
## 5		350	272	252	252	236	306	238	350	206	260	280	274	318	268	210	
## 6		310	376	326	346	334	282	292	282	300	290	302	300	306	294	444	

Creating Histograms

We want to extract the data from these columns `rtimes[, 2:31]`, for all 17 patients. We will plot 17 histograms, displaying the distribution of the 30 reaction times.

In order to keep a consistent range across the 17 histograms, we need to find the absolute minimum and maximum values in this dataset. To do this, we extract all the data, 17 rows * 30 trials.

```
DataValues <- unlist(rtimes1[, 2:31])
range(DataValues)
```

```
## [1] 204 1714
```

```
# We want to store these values
```

```
x_min <- 204
x_max <- 1714
```

Below, we can observe two sets of histograms, the first containing the eleven non-schizophrenic patients, and the seconds containing the 6 schizophrenic patients.

We reshape to data frame to long format, as having the reaction time data in one column makes much easier to code the histogram. To create a separate plot for each patient, we use `facet_wrap()` to group data by PatientID.

```
# Reshaping to long format
rtimes_long <- rtimes1 %>%
  pivot_longer(cols = starts_with("T"), # Columns T1 to T30
               names_to = "Trial",
               values_to = "ReactionTime") %>%
  mutate(PatientID = rep(1:17, each = 30)) %>% # Add patientID
  select(PatientID, PatientType, ReactionTime)

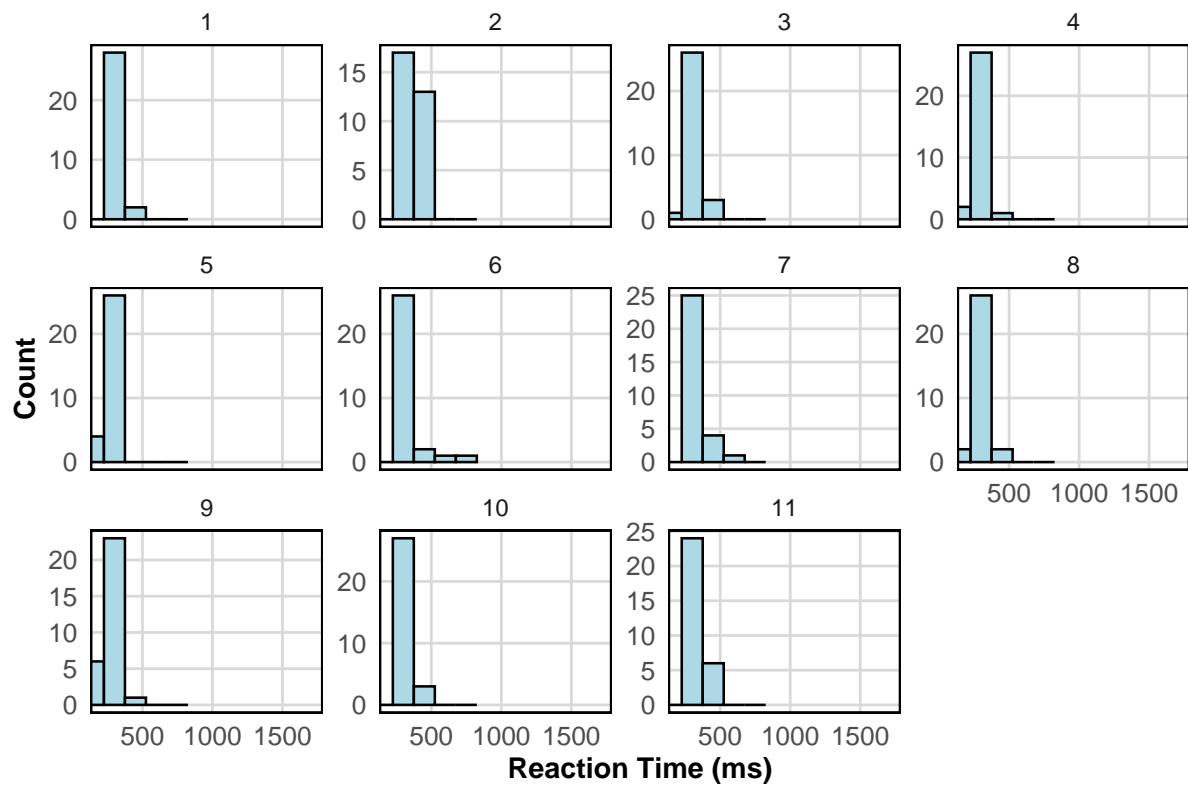
# Splitting into two datasets
non_schizo <- rtimes_long %>% filter(PatientType == "non-schizophrenic")
schizo <- rtimes_long %>% filter(PatientType == "schizophrenic")

# Plotting non-schizophrenic histograms
hist1 <- ggplot(non_schizo, aes(x = ReactionTime)) +
  geom_histogram(binwidth = 150, fill = "lightblue", color = "black") +
  facet_wrap(~ PatientID, ncol = 4, scales = "free_y") + # 11 plots, 4 columns
  coord_cartesian(xlim = c(x_min, x_max)) + # Fixed x-axis
  labs(title = "Figure 1: Non-Schizophrenic Reaction Times",
       x = "Reaction Time (ms)",
       y = "Count") +
  custom_theme

# Plotting schizophrenic histograms
hist2 <- ggplot(schizo, aes(x = ReactionTime)) +
  geom_histogram(binwidth = 150, fill = "salmon", color = "black") +
  facet_wrap(~ PatientID, ncol = 3, scales = "free_y") + # 6 plots, 3 columns
  coord_cartesian(xlim = c(x_min, x_max)) + # Fixed x-axis
  labs(title = "Figure 2: Schizophrenic Reaction Times",
       x = "Reaction Time (ms)",
       y = "Count") +
  custom_theme

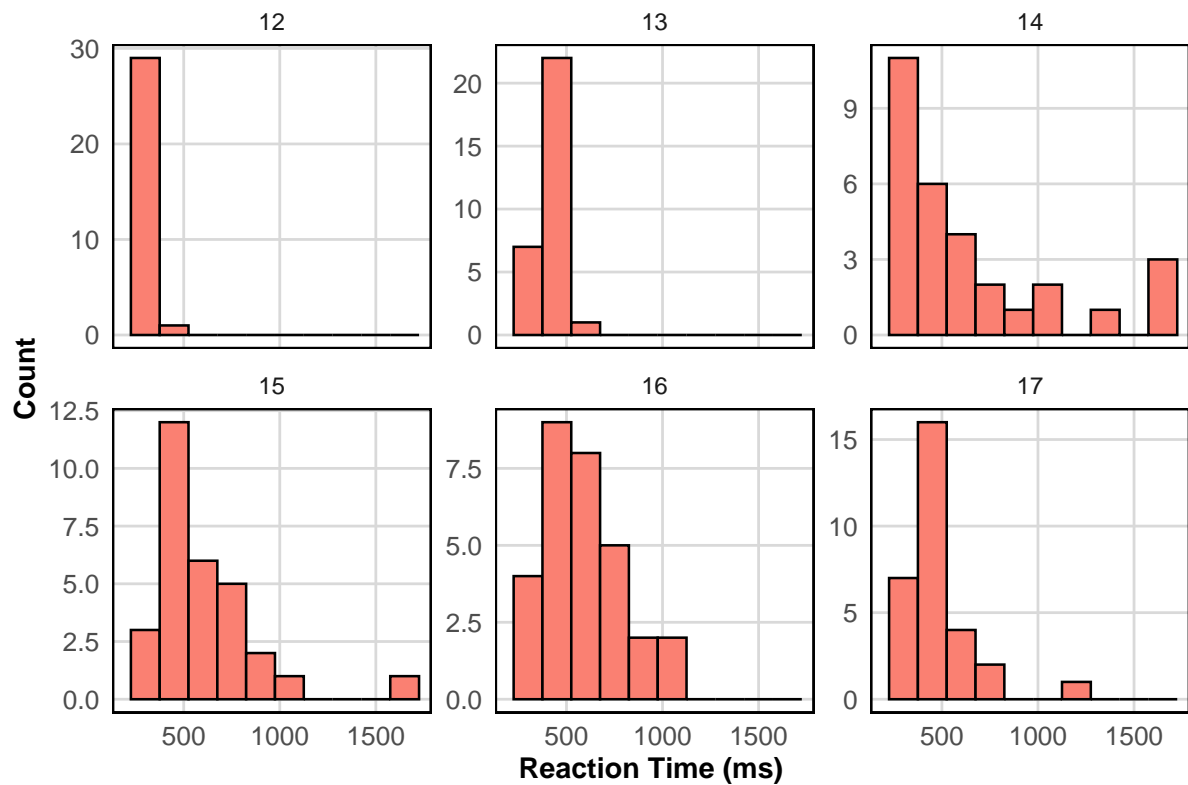
print(hist1)
```

Figure 1: Non-Schizophrenic Reaction Times



```
print(hist2)
```

Figure 2: Schizophrenic Reaction Times



In Figure 1 we can see that Non-schizophrenic patients such as patient 1 have a narrower and concentrated histogram, with reaction times completely clustered around the second bin (150-300ms). For example, every Non-schizophrenic patient's results show that at least two thirds of their reaction times were faster than 300ms.

In contrast, the Schizophrenic patients histograms in Figure 2 exhibit a much wider spread of reaction times, with some patients having reactions times exceeding 1500ms. This indicates greater variability, for instance, patient 14 has a wide range of reaction times with many being greater than 1000ms, whilst also having a significant count below 500ms. This suggests that Schizophrenic patients have a mixture of normal and delayed responses, and this aligns with the theory of attention deficits and motor retardation in schizophrenics. This is generally observed by their inconsistent distributions and right skew, compared to the tighter uniform distributions of non-schizophrenics

Nonetheless, both groups share similarities in their distributions. They both have peaks within the first two bins 0-500ms, as you can see when comparing patient 8 (non) to 13 (schizophrenic). However, only schizophrenics have reaction time values in the higher bands (greater than 800ms)

Overall, these histograms support the hypothesis that schizophrenics experience attention deficits and motor reflex retardation, as seen by the wider spread of values and more irregular histograms for schizophrenic individuals.

2. [5 marks] The above model uses the logarithm of measured reaction times. Explain why taking the

logarithm is necessary here (referencing the relevant output), then perform the transformation yourself. For each person compute the standard deviation of the log transformed reaction times of that individual.

-Calculate standard deviations: For each of the 17 people, compute the standard deviation of their 30 log-transformed reaction times.

Log-transforming the reaction times is necessary for the model above for two main reasons. Firstly, there is skewness in the data as seen in the Schizophrenic reaction times histograms, for example, Patient 14 and 15 are right-skewed distributions with extreme values (over 1500ms), whilst non-schizophrenic's have a tighter distribution. Following this, schizophrenics don't seem to follow a normal distribution, thus, log transforming reduces the skewness and helps stabilise the variance from the effect of outliers/skew. This enables the data to fit the models normal distributions assumption better. Finally, the log-transformation scales the data, making the mean and variance more interpretable. For example, the original range of 204ms to 1714ms, becomes 5.31 to 7.44 (3 s.f.) when log-transformed. Put simply, handling smaller numbers is easier.

```
log_rtimes_long <- rtimes_long %>%  
  mutate(LogReactionTime = log(ReactionTime))  
range(log_rtimes_long$LogReactionTime) # checking the range also lets us know if any observ
```

```
## [1] 5.318120 7.446585
```

Now we compute the standard deviation of each patients 30 log-transformed reaction times. This is useful as it measures the within personal variability on the log scale.

```
# Compute standard deviation of log-transformed reaction times for each person
sd_log <- log_rtimes_long %>%
  group_by(PatientID) %>%
  summarise(SD_Log_RT = sd(LogReactionTime))

# Print results
head(sd_log)
```

```
## # A tibble: 6 x 2
##   PatientID SD_Log_RT
##   <int>     <dbl>
## 1         1     0.127
## 2         2     0.129
## 3         3     0.169
## 4         4     0.150
## 5         5     0.145
## 6         6     0.224
```

3. [5 marks] List the parameters of the model and assign non-informative uniform prior distributions to each parameter, paying attention to the values these parameters are allowed take.

The model

Non-Schizophrenic Reaction Times: For the responses of the i th non-schizophrenic person ($i = 1, 2, \dots, 11$)

$$y_{ij} \sim N(\alpha_i, \sigma_y^2), \quad i = 1, 2, \dots, 11, \quad j = 1, 2, \dots, 30$$

Schizophrenic Reaction Times:

For the responses of the i th schizophrenic individual ($i = 12, 13, \dots, 17$), with probability $(1 - \lambda)$ there is no delay, and the response is normally distributed with mean α_i and variance σ_y^2 ; and with probability λ the response is delayed so that the observations have mean $\alpha_i + \tau$ and variance σ_y^2 .

$$y_{ij} \sim N(\alpha_i + \tau z_{ij}, \sigma_y^2)$$

$$z_{ij} \sim \text{Bernoulli}(\lambda), \quad i = 12, 13, \dots, 17, \quad j = 1, 2, \dots, 30$$

Distributions of α :

For non-schizophrenic individuals we assume that α_i follows a normal distribution with mean μ and variance σ_α^2

$$\alpha_i \sim N(\mu, \sigma_\alpha^2), \quad i = 1, 2, \dots, 11$$

For schizophrenics:

$$\alpha_i \sim N(\mu + \beta, \sigma_\alpha^2), \quad i = 12, 13, \dots, 17$$

The parameters of the model, and their non-informative uniform prior distributions:

for Non-specific Individuals:

- α_i (for $i = 1$ to 17): Person-specific means on the log scale.

```
alpha[i] ~ dunif(-10, 10)
```

- μ : Mean of α_i for non-schizophrenics.

```
mu ~ dunif(-10, 10)
```

- $\tau_y = 1/\sigma_y^2$: Precision of log-reaction times.

```
tau_y ~ dunif(0, 10)
```

- $\tau_\alpha = 1/\sigma_\alpha^2$: Precision of α_i .

```
tau_alpha ~ dunif(0, 10)
```

for Schizophrenic Individuals:

- β : Additional variable in α_i mean for schizophrenics.

```
beta ~ dunif(-10, 10)
```

- τ : Delay parameter for schizophrenics ($\tau > 0$).

```
tau ~ dunif(0, 10)
```

- λ : Probability of delay ($0 \leq \lambda \leq 1$).

```
lambda ~ dunif(0, 1)
```

4.