

Cervical Dystonia

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```
#Loading Libraries
```

```
library(haven)
library(data.table)
library(tidyverse)
library(skimr)
library(gridExtra)
library(gtsummary)
library(expss)
```

```
#Loading in Data
```

```
cerv <- read_dta("cdystonia.dta")
```

```
glimpse(cerv)
```

```
## Observations: 631
```

```
## Variables: 7
```

```
## $ week    <dbl> 0, 2, 4, 8, 12, 16, 0, 2, 4, 8, 12, 16, 0, 2, 4, 8, 12, 16, ...
## $ site    <dbl+lbl> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, ...
## $ id      <dbl> 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 3, 3, 3, 3, 3, 3, 4, 4, ...
## $ treat   <dbl+lbl> 2, 2, 2, 2, 2, 2, 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 3, ...
## $ age     <dbl> 65, 65, 65, 65, 65, 65, 70, 70, 70, 70, 70, 70, 64, 64, 64, ...
## $ sex     <dbl+lbl> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, ...
## $ twstrs  <dbl> 32, 30, 24, 37, 39, 36, 60, 26, 27, 41, 65, 67, 44, 20, 23, ...
```

```
head(cerv)
```

```
## # A tibble: 6 x 7
```

```
##   week      site    id    treat    age      sex twstrs
##   <dbl> <dbl+lbl> <dbl> <dbl+lbl> <dbl> <dbl+lbl> <dbl>
## 1     0     1 [1]     1 2 [5000U]    65     1 [F]    32
## 2     2     1 [1]     1 2 [5000U]    65     1 [F]    30
## 3     4     1 [1]     1 2 [5000U]    65     1 [F]    24
## 4     8     1 [1]     1 2 [5000U]    65     1 [F]    37
## 5    12     1 [1]     1 2 [5000U]    65     1 [F]    39
## 6    16     1 [1]     1 2 [5000U]    65     1 [F]    36
```

```
#Mutating the ID variable to capture the Site Location
```

```
cerv_dyst <- cerv %>%
  mutate(id = paste0(site, id)) %>%
  select(-site)
```

```
glimpse(cerv_dyst)
```

```
## Observations: 631
## Variables: 6
## $ week    <dbl> 0, 2, 4, 8, 12, 16, 0, 2, 4, 8, 12, 16, 0, 2, 4, 8, 12, 16, ...
## $ id      <chr> "11", "11", "11", "11", "11", "11", "12", "12", "12", "12", ...
## $ treat   <dbl+lbl> 2, 2, 2, 2, 2, 2, 1, 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 3, ...
## $ age     <dbl> 65, 65, 65, 65, 65, 65, 70, 70, 70, 70, 70, 70, 64, 64, 64, ...
## $ sex     <dbl+lbl> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, ...
## $ twstrs  <dbl> 32, 30, 24, 37, 39, 36, 60, 26, 27, 41, 65, 67, 44, 20, 23, ...
```

```
#Mutating the sex variable to numeric, Female = 1
cerv_dyst <- cerv_dyst %>%
  mutate(sex = as.numeric(sex)) %>%
  apply_labels(treat = "Treatment", sex = "Sex")
```

Exploratory Analysis

```
#Table recoding
table_cerv <- cerv_dyst %>%
  mutate(
    sex = if_else(sex == 1, "Female", "Male"),
    treat = if_else(treat == 1, "10000U", if_else(
      treat == 2, "5000U", "Placebo")),
    week = paste0("Week ", week),
    week = factor(week, levels = c("Week 0",
                                   "Week 2",
                                   "Week 4",
                                   "Week 8",
                                   "Week 12",
                                   "Week 16")))

#Summary Table by Gender, Week, Treatment
table_cerv %>%
  select(-id, -week, -treat) %>%
  as.data.frame() %>%
  tbl_summary(by = sex) %>%
  add_p() %>%
  bold_p()
```

Characteristic	Female, N = 395 ¹	Male, N = 236 ¹	p-value ²
Age [years]	54 (47, 65)	57 (43, 66)	0.9
TWSTRS-total score	44 (34, 52)	40 (31, 50)	0.012

¹Statistics presented: median (IQR)

²Statistical tests performed: Wilcoxon rank-sum test

```
table_cerv %>%
  select(-id, -sex, -week, -age) %>%
  as.data.frame() %>%
  tbl_summary(by = treat) %>%
  add_p() %>%
  bold_p()
```

Characteristic	10000U, N = 213 ¹	5000U, N = 211 ¹	Placebo, N = 207 ¹	p-value ²
TWSTRS-total score	44 (33, 51)	43 (32, 51)	43 (33, 50)	>0.9

¹Statistics presented: median (IQR)

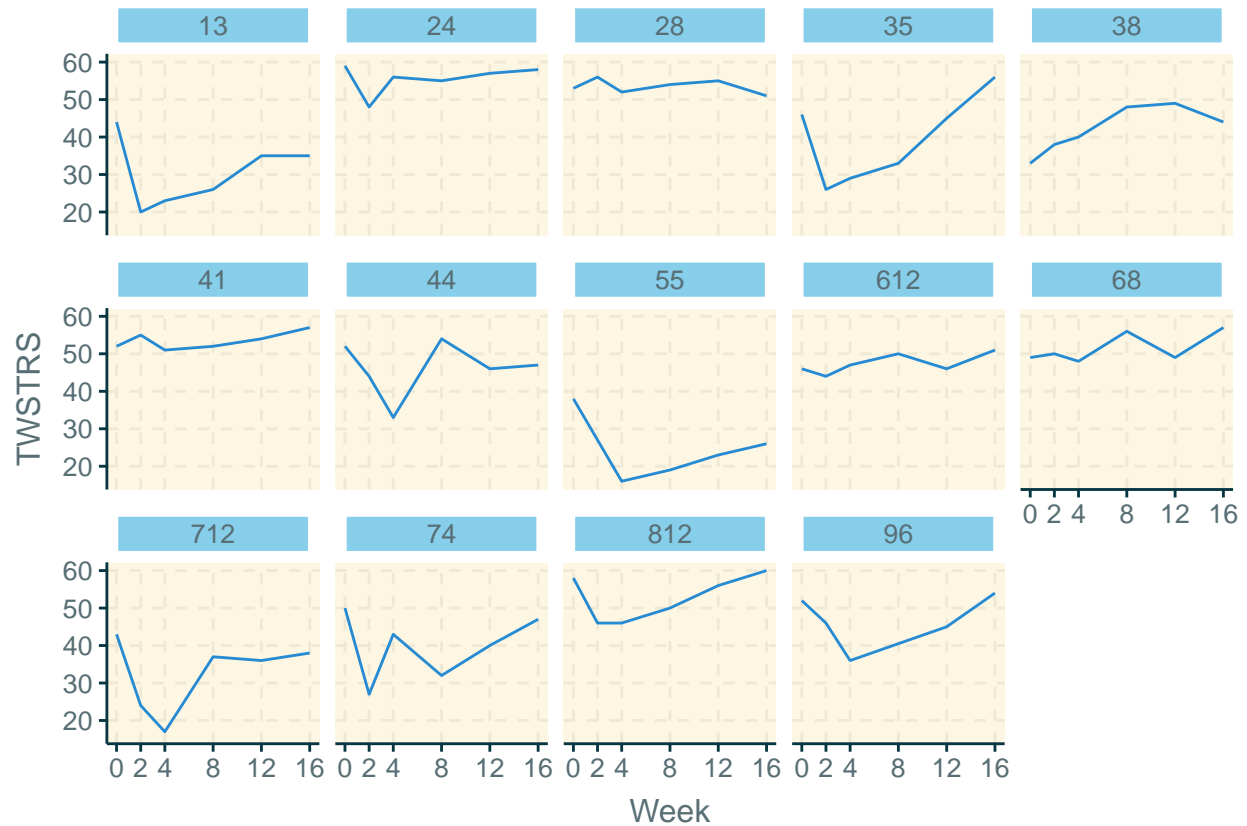
²Statistical tests performed: Kruskal-Wallis test

```
#Histogram of Age and Rating Scales
age_hist <- ggplot(data = cerv_dyst, aes(x = age)) +
  geom_histogram(col = "white", fill = "seagreen3")

twstrs_hist <- ggplot(data = cerv_dyst, aes(x = twstrs)) +
  geom_histogram(col = "white", fill = "seagreen3") +
  labs(x = "Toronto Western Spasmodic Torticollis Rating Scale (TWSTRS)")

set.seed(123)
id_sample <- sample(cerv_dyst$id, 15)

# Plots of 15 random sample individuals
ggplot(data = cerv_dyst %>% filter(id %in% id_sample),
  aes(x = week, y = twstrs)) +
  geom_line() +
  facet_wrap(~id, ncol = 5) +
  theme(strip.background = element_rect(colour="white",
    fill="skyblue",
    size=4,
    linetype="solid")) +
  scale_x_continuous(breaks = c(0, 2, 4, 8, 12, 16)) +
  labs(x = "Week",
    y = "TWSTRS")
```



```
# Every individual's change through the weeks with average
ggplot(data = cerv_dyst, aes(x = week, y = twstrs)) +
  geom_line(alpha=0.3, aes(group = id)) +
  geom_smooth(col = "seagreen4", size = 2) +
  scale_x_continuous(breaks = c(0, 2, 4, 8, 12, 16)) +
  labs(x = "Week",
       y = "TWSTRS")
```

```
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
```

