

Laryngectomy Analysis

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Introduction

This document explores the relationship between hospital stay in days and the number of complications among patients in the `analysis` dataset from the `Open.Visualization.Academy` package.

Methods

The data for this analysis come from the `analysis` dataset in the `Open.Visualization.Academy` R package.

Results

Scatterplot with linear trend (no confidence bounds)

```
ggplot(analysis, aes(x = complications_count, y = hospital_stay_days)) +  
  geom_point() +  
  geom_smooth(method = 'lm', se = FALSE) +  
  labs(  
    x = 'Complication count',  
    y = 'Hospital stay (days)'  
  )  
`geom_smooth()` using formula = 'y ~ x'
```

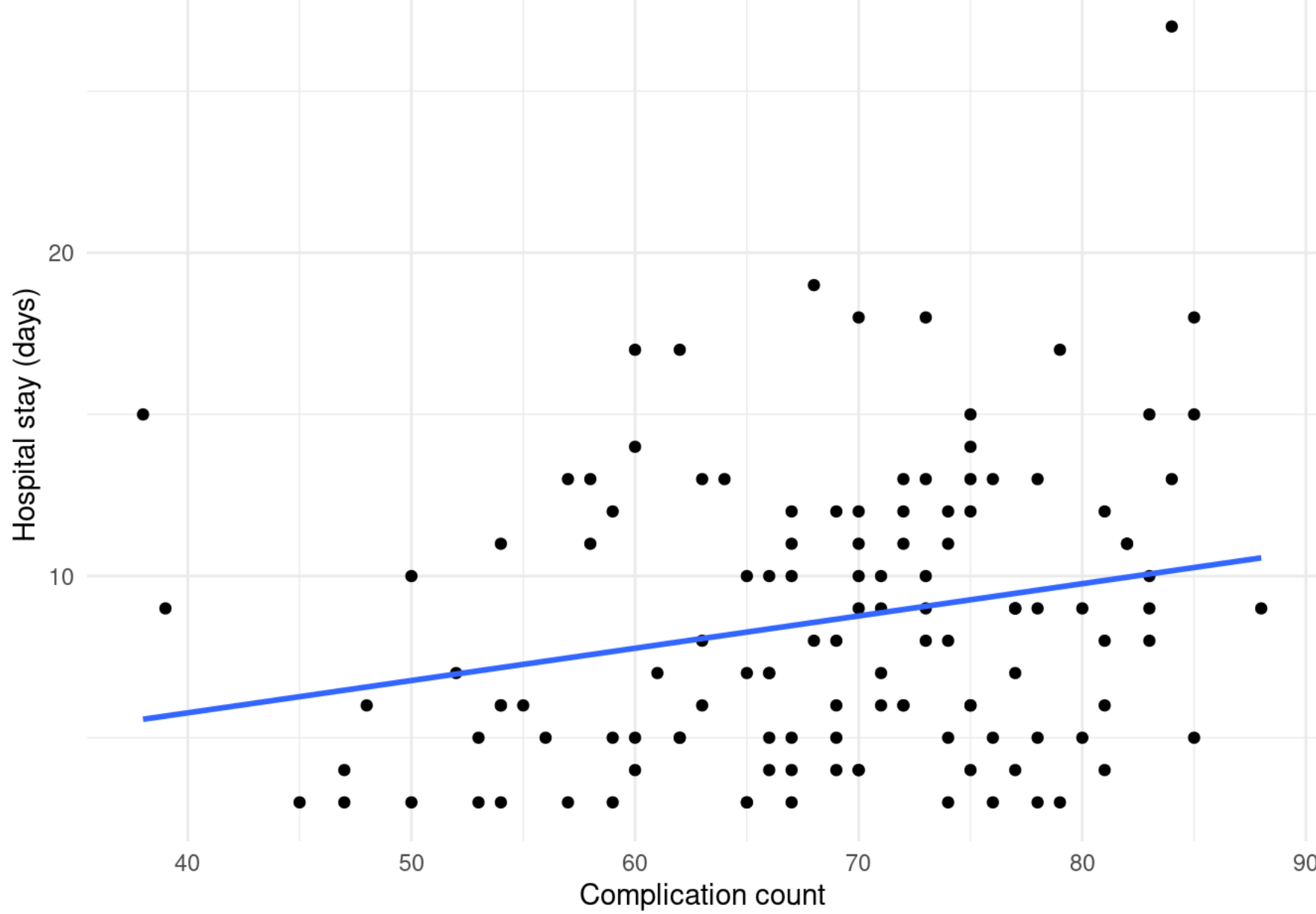


Figure 1: Scatterplot of hospital stay (days) versus complication count with linear trend.

Scatterplot with transparency and linear trend

```
ggplot(analysis, aes(x = complications_count, y = hospital_stay_days)) +  
  geom_point(alpha = 0.3) +  
  geom_smooth(method = 'lm', se = FALSE) +  
  labs(  
    x = 'Complication count',  
    y = 'Hospital stay (days)'  
  )  
`geom_smooth()` using formula = 'y ~ x'
```

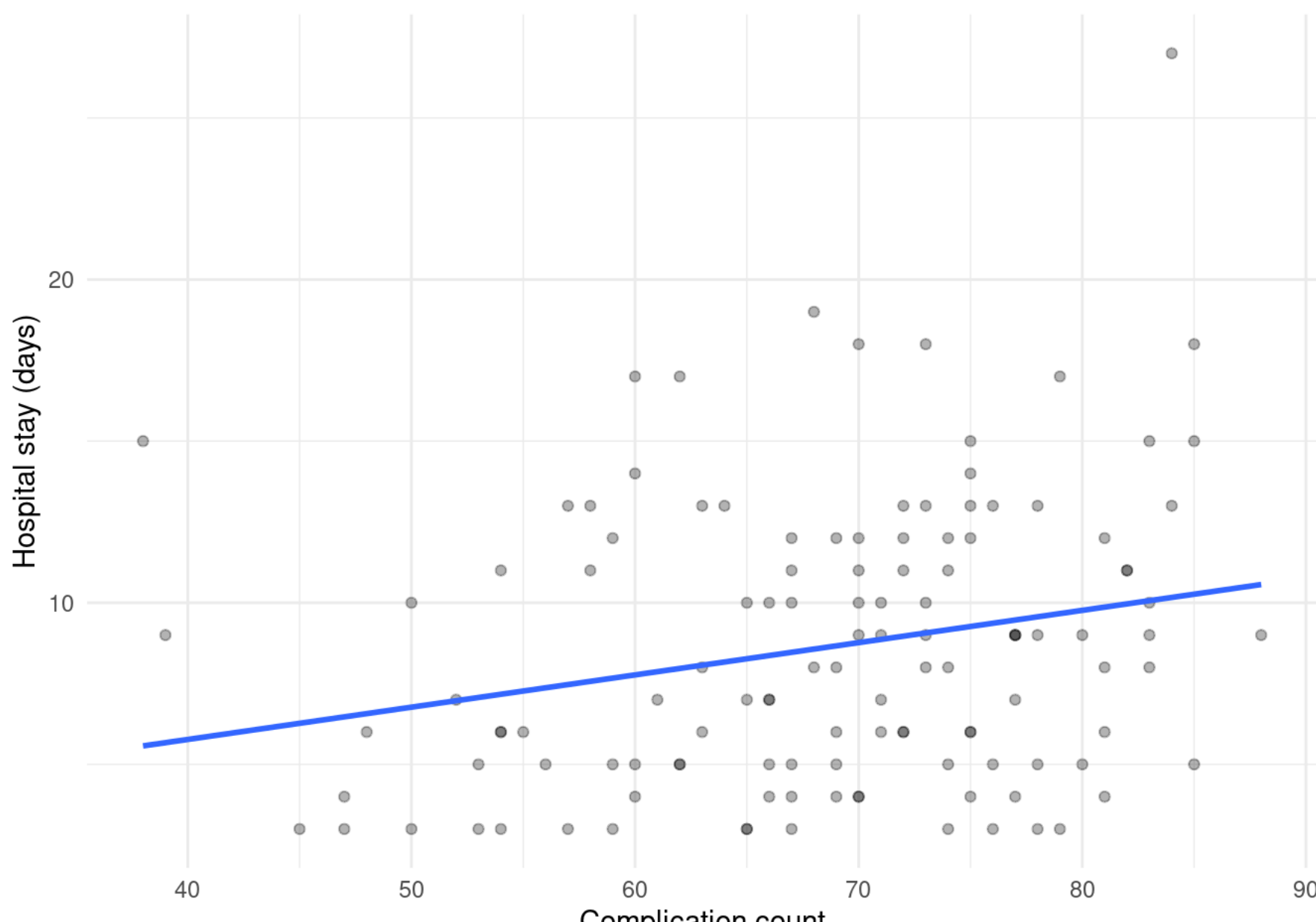


Figure 2: Scatterplot of hospital stay versus complication count using point transparency to reduce overplotting.

Sunflower-like plot for overlapping points

```
library(dplyr)  
library(tidyr)  
  
# Calculate counts at each (x, y) position  
counted_data <- analysis |>  
  group_by(complications_count, hospital_stay_days) |>  
  summarise(n = n(), .groups = 'drop')  
  
# Create sunflower-like plot with radiating lines  
create_sunflower <- function(data) {  
  data |>  
  filter(n > 1) |>  
  rowwise() |>  
  mutate(  
    angles = list(seq(0, 2 * pi, length.out = n + 1)[1:n]),  
    x_end = list(complications_count + 0.03 * cos(angles)),  
    y_end = list(hospital_stay_days + 0.03 * sin(angles))  
  ) |>  
  unnest(cols = c(angles, x_end, y_end))  
}  
  
sunflower_segments <- create_sunflower(counted_data)  
  
ggplot() +  
  geom_segment(  
    data = sunflower_segments,  
    aes(x = complications_count, y = hospital_stay_days,  
        xend = x_end, yend = y_end),  
    alpha = 0.7  
  ) +  
  geom_point(  
    data = counted_data,  
    aes(x = complications_count, y = hospital_stay_days),  
    size = 1  
  ) +  
  labs(  
    x = 'Complication count',  
    y = 'Hospital stay (days)'  
  )
```

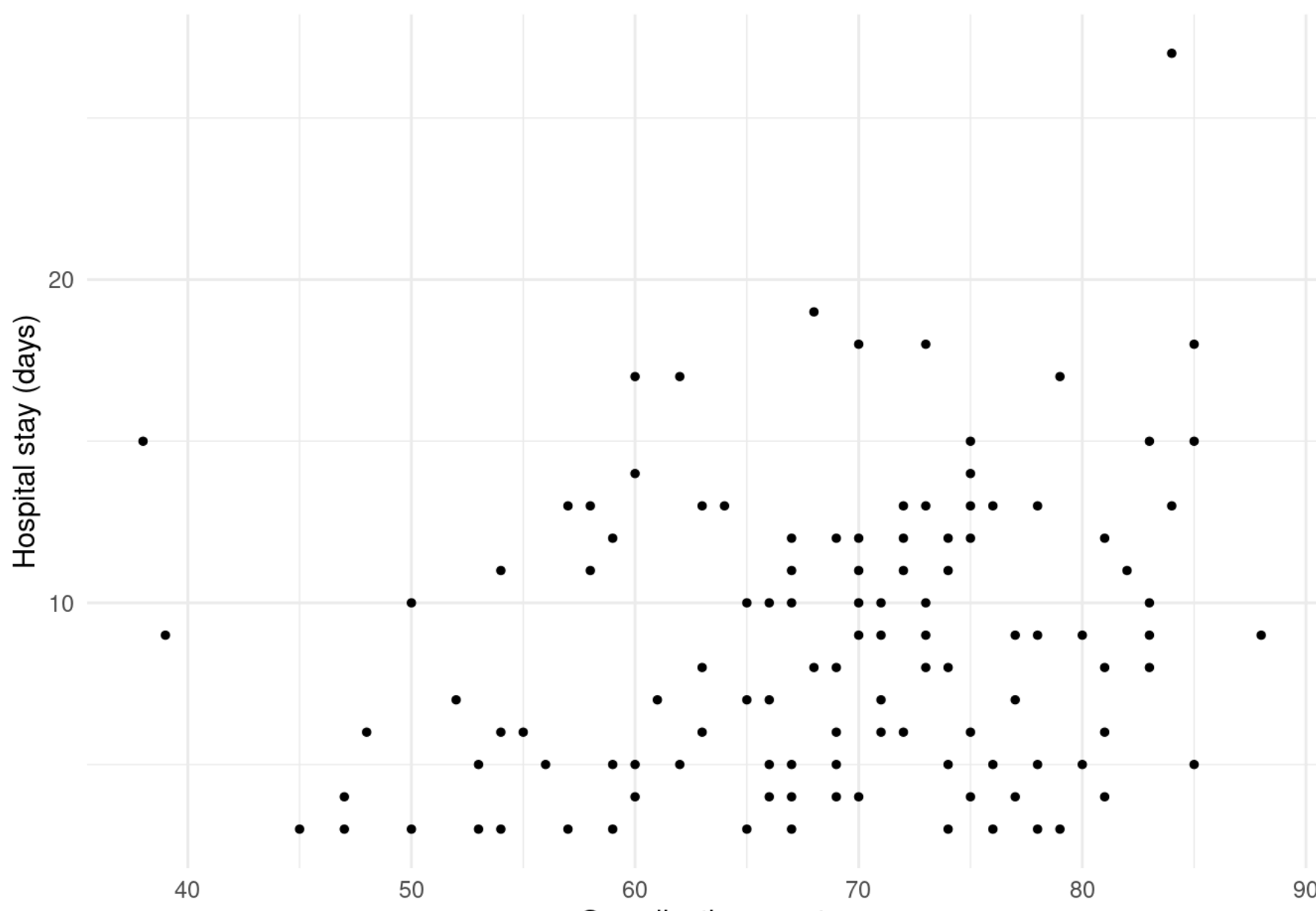


Figure 3: Sunflower plot showing overlapping combinations of complication count and hospital stay.

Hospital stay vs. complication count as a categorical variable

Side-by-side boxplot

```
ggplot(analysis,  
  aes(x = factor(complications_count), y = hospital_stay_days)) +  
  geom_boxplot() +  
  labs(  
    x = 'Complication count (categorical)',  
    y = 'Hospital stay (days)'  
  )
```

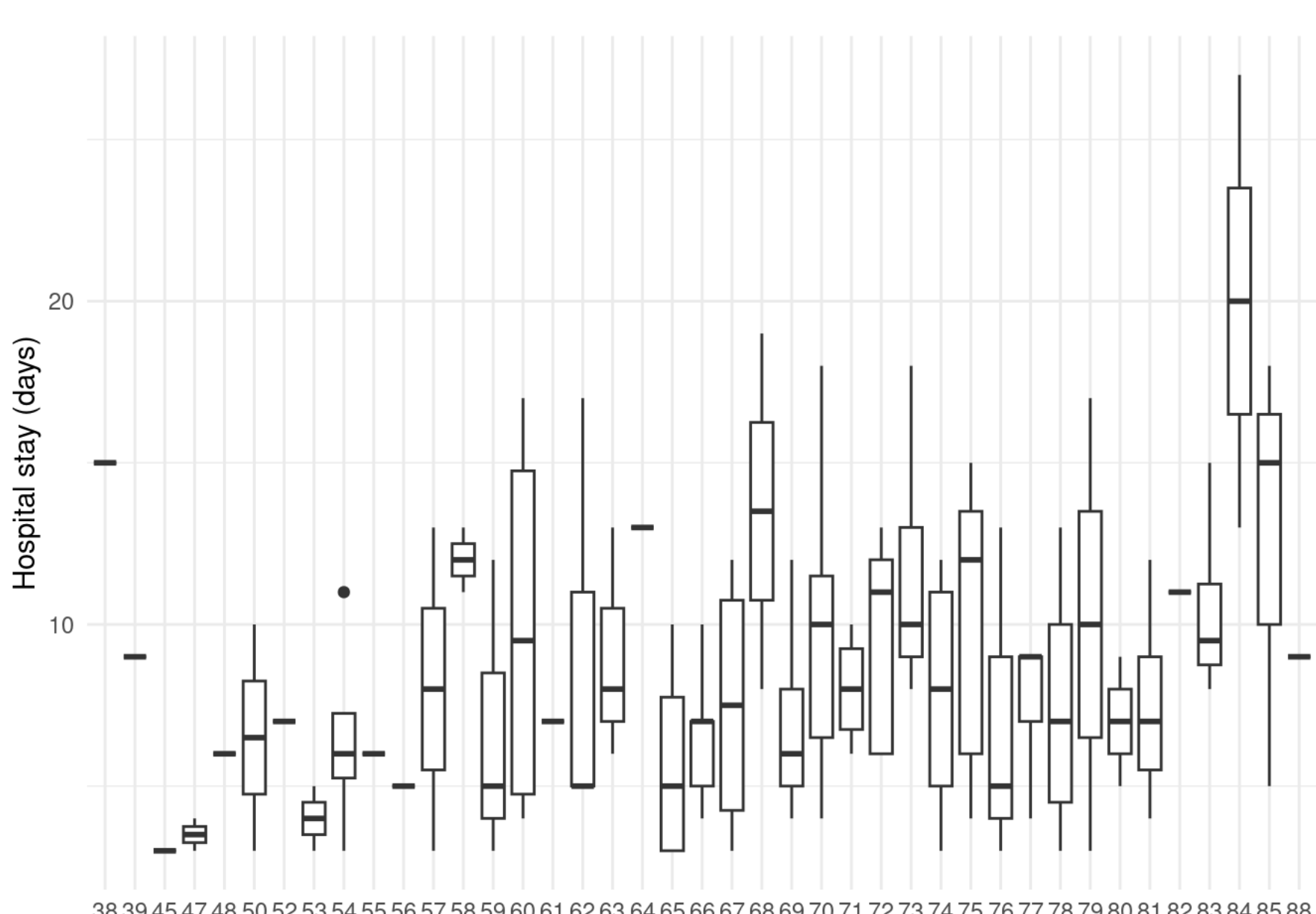


Figure 4: Hospital stay (days) by complication count: side-by-side boxplots.

Violin plot

```
ggplot(analysis,  
  aes(x = factor(complications_count), y = hospital_stay_days)) +  
  geom_violin(trim = FALSE) +  
  labs(  
    x = 'Complication count (categorical)',  
    y = 'Hospital stay (days)'  
  )
```

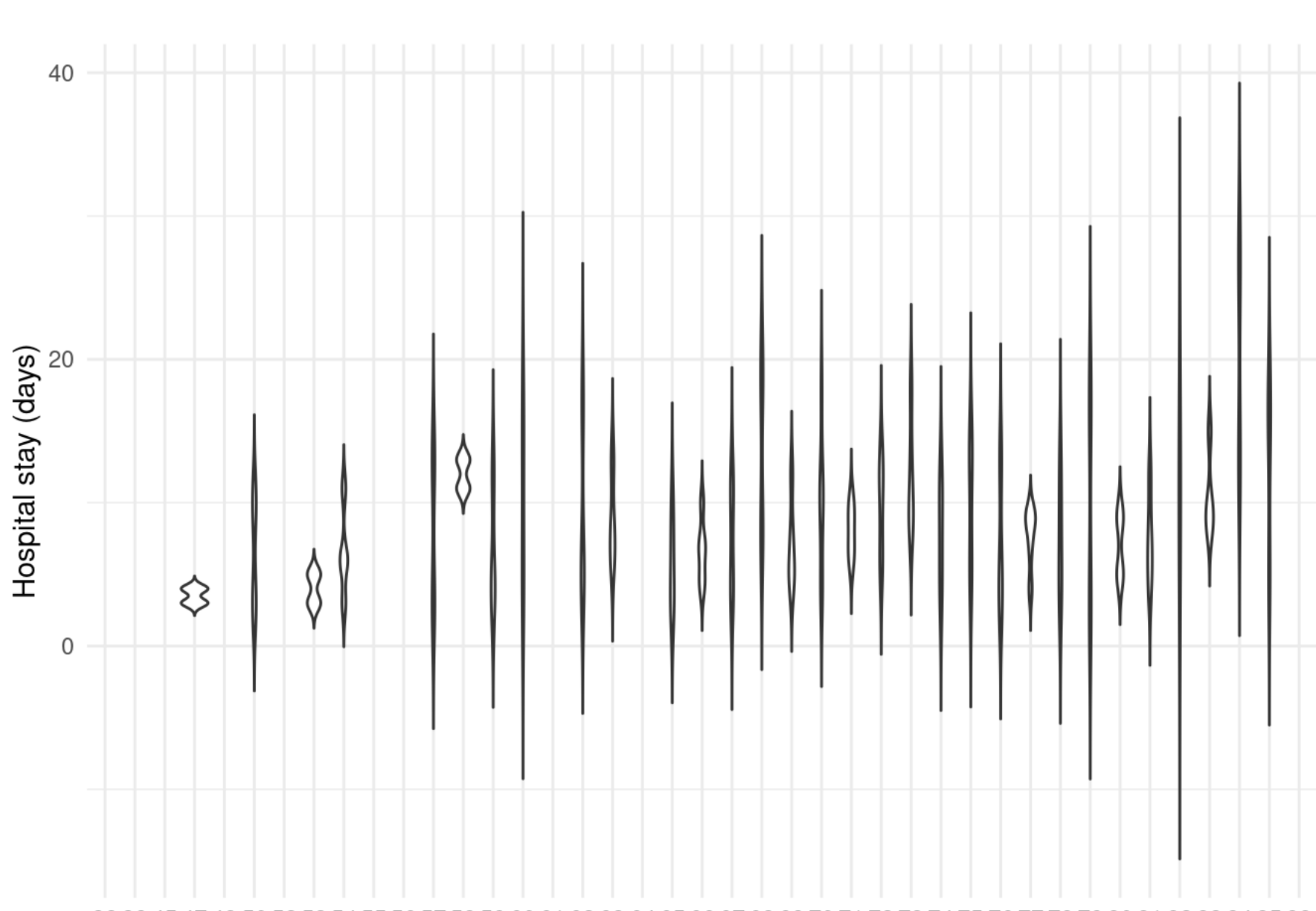


Figure 5: Hospital stay (days) by complication count: violin plots.

Discussion

These visualizations show how hospital stay in days changes with the number of complications. The scatterplots (Figures [Figure 1](#) and [Figure 2](#)) and sunflower plot (Figure [Figure 3](#)) provide a detailed view of individual patient-level relationships. The boxplot and violin plot (Figures [Figure 4](#) and [Figure 5](#)) summarize the distribution of hospital stays for each complication count.