

# Regression to the mean modeling

Mean NRS pain rating of 6.2 at 0.5 correlation

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## Generate 2x2 covariance matrix

Generate a covariance matrix using an SD of 1.2, 1.7, and 2.2, and correlation of 0.5.

```
cor <- matrix(c(1, 0.5, 0.5, 1), ncol = 2)
std_1 <- c(1.2, 1.2)
std_2 <- c(1.7, 1.7)
std_3 <- c(2.2, 2.2)
cov_1 <- cor2cov(cor.mat = cor,
                 sd = std_1)
cov_1

##      [,1] [,2]
## [1,] 1.44 0.72
## [2,] 0.72 1.44

cov_2 <- cor2cov(cor.mat = cor,
                 sd = std_2)
cov_2

##      [,1] [,2]
## [1,] 2.890 1.445
## [2,] 1.445 2.890

cov_3 <- cor2cov(cor.mat = cor,
                 sd = std_3)
cov_3

##      [,1] [,2]
## [1,] 4.84 2.42
## [2,] 2.42 4.84
```

---

Mean = 6.2, SD = 1.2, cor = 0.5

## Generate and summarise data

### Base data

```
# Set the random seed for reproducibility
set.seed(2019)

# Generate the data
six_1.base <- as.data.frame(mvrnorm(n = 150, mu = c(6.2, 6.2), Sigma = cov_1))

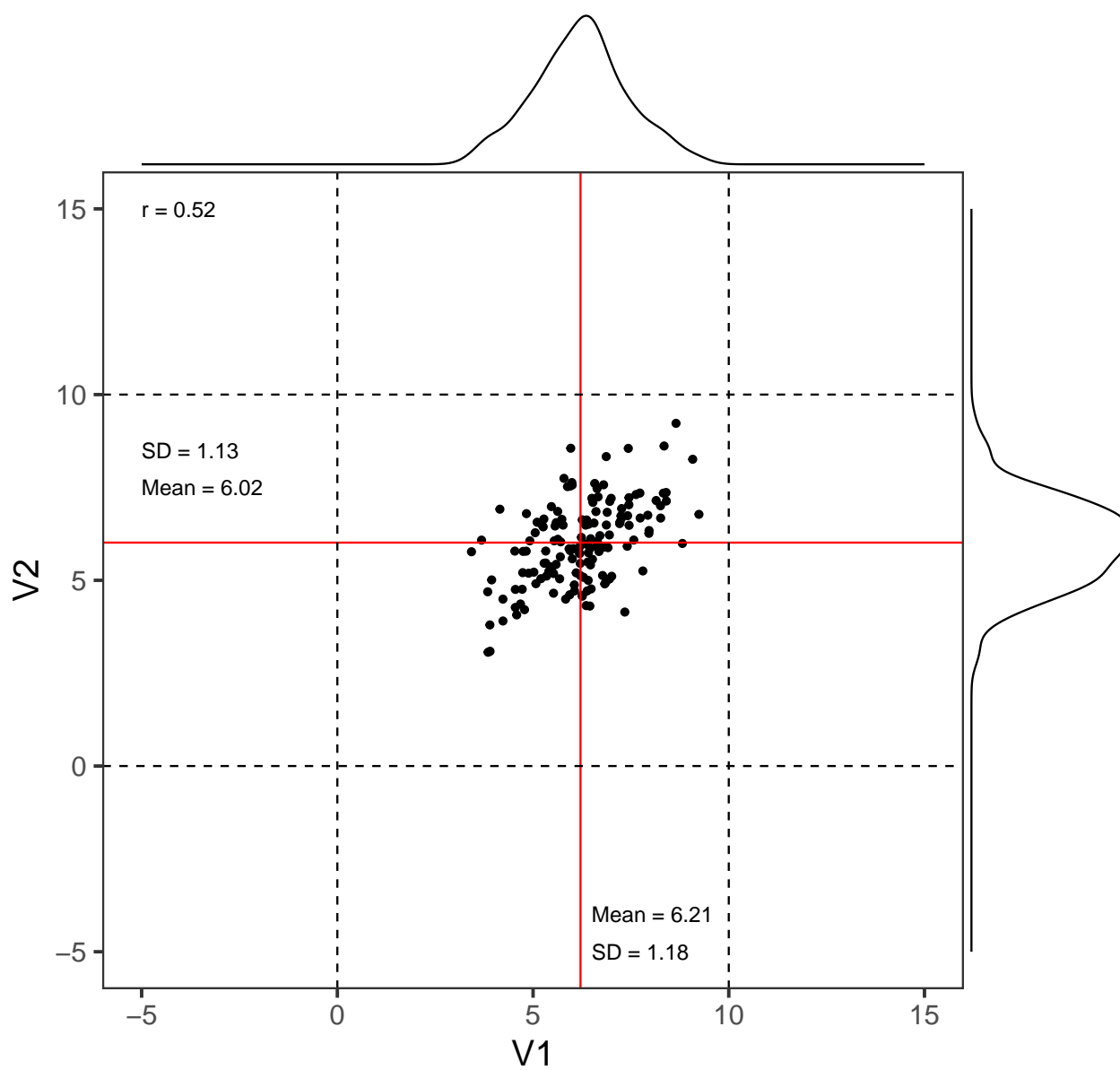
# Plot base data
ggMarginal(ggplot(data = six_1.base) +
            aes(x = V1, y = V2) +
            geom_point() +
            geom_hline(yintercept = mean(six_1.base$V2), colour = 'red') +
            geom_vline(xintercept = mean(six_1.base$V1), colour = 'red') +
            geom_hline(yintercept = 0, linetype = 2) +
            geom_hline(yintercept = 10, linetype = 2) +
            geom_vline(xintercept = 0, linetype = 2) +
```

```

geom_vline(xintercept = 10, linetype = 2) +
annotate(geom = 'text', x = -5, y = 15, hjust = 0,
         label = str_glue("r = {round(cor(six_1.base$V1, six_1.base$V2), 2)}")) +
annotate(geom = 'text', x = -5, y = 7.5, hjust = 0,
         label = str_glue("Mean = {round(mean(six_1.base$V2), 2)}")) +
annotate(geom = 'text', x = -5, y = 8.5, hjust = 0,
         label = str_glue("SD = {round(sd(six_1.base$V2), 2)}")) +
annotate(geom = 'text', x = 6.5, y = -4, hjust = 0,
         label = str_glue("Mean = {round(mean(six_1.base$V1), 2)}")) +
annotate(geom = 'text', x = 6.5, y = -5, hjust = 0,
         label = str_glue("SD = {round(sd(six_1.base$V1), 2)}")) +
labs(subtitle = 'Mean = 6.2, SD = 1.2, cor = 0.5') +
scale_y_continuous(limits = c(-5, 15)) +
scale_x_continuous(limits = c(-5, 15)) +
theme(panel.grid = element_blank(),
      plot.subtitle = element_text(size = 16)))

```

Mean = 6.2, SD = 1.2, cor = 0.5



```
# Linear regression
summary(lm(V2 ~ V1, data = six_1.base))

##
## Call:
## lm(formula = V2 ~ V1, data = six_1.base)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -2.43530 -0.60848 -0.08843  0.63099  2.66504
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  2.91166    0.42123   6.912 1.33e-10 ***
## V1           0.49956    0.06661   7.499 5.47e-12 ***
```

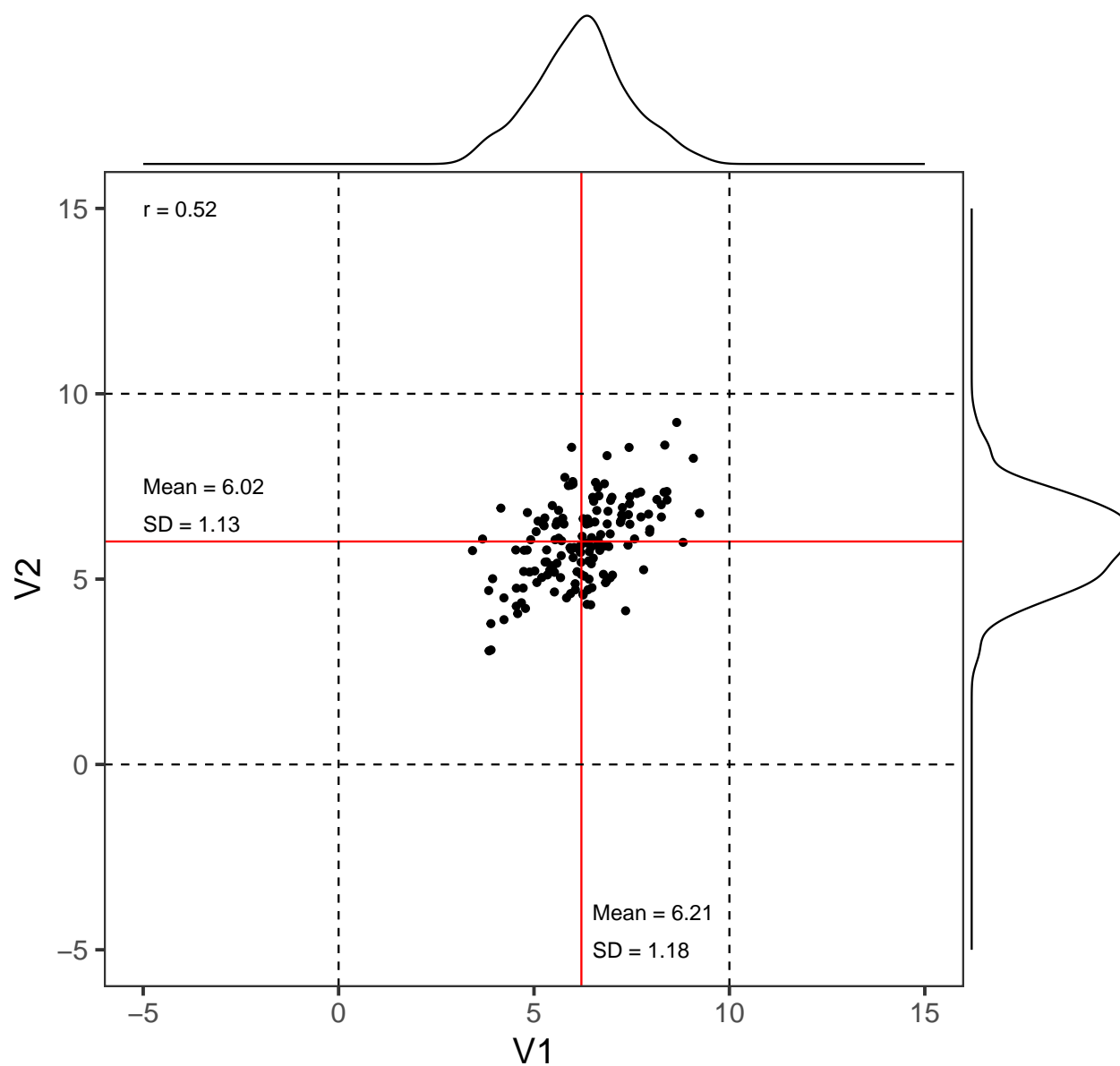
```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.9619 on 148 degrees of freedom
## Multiple R-squared:  0.2754, Adjusted R-squared:  0.2705
## F-statistic: 56.24 on 1 and 148 DF,  p-value: 5.468e-12
```

Constrain values to 0-10 range

```
# Process data
six_1 <- six_1.base %>%
  mutate(V1 = case_when(
    V1 < 1 ~ 1,
    V1 > 10 ~ 10,
    TRUE ~ V1)) %>%
  mutate(V2 = case_when(
    V2 < 0 ~ 0,
    V2 > 10 ~ 10,
    TRUE ~ V2)) %>%
  mutate(group = 'No cut-off')

# Plot processed data
ggMarginal(ggplot(data = six_1) +
  aes(x = V1, y = V2) +
  geom_point() +
  geom_hline(yintercept = mean(six_1$V2), colour = 'red') +
  geom_vline(xintercept = mean(six_1$V1), colour = 'red') +
  geom_hline(yintercept = 0, linetype = 2) +
  geom_hline(yintercept = 10, linetype = 2) +
  geom_vline(xintercept = 0, linetype = 2) +
  geom_vline(xintercept = 10, linetype = 2) +
  annotate(geom = 'text', x = -5, y = 15, hjust = 0,
    label = str_glue("r = {round(cor(six_1$V1, six_1$V2), 2)}")) +
  annotate(geom = 'text', x = -5, y = 7.5, hjust = 0,
    label = str_glue("Mean = {round(mean(six_1$V2), 2)}")) +
  annotate(geom = 'text', x = -5, y = 6.5, hjust = 0,
    label = str_glue("SD = {round(sd(six_1$V2), 2)}")) +
  annotate(geom = 'text', x = 6.5, y = -4, hjust = 0,
    label = str_glue("Mean = {round(mean(six_1$V1), 2)}")) +
  annotate(geom = 'text', x = 6.5, y = -5, hjust = 0,
    label = str_glue("SD = {round(sd(six_1$V1), 2)}")) +
  labs(subtitle = 'Mean = 6.2, SD = 1.2 (constrained), cor = 0.5') +
  scale_y_continuous(limits = c(-5, 15)) +
  scale_x_continuous(limits = c(-5, 15)) +
  theme(panel.grid = element_blank(),
    plot.subtitle = element_text(size = 16)))
```

Mean = 6.2, SD = 1.2 (constrained), cor = 0.5



```
# Linear regression
summary(lm(V2 ~ V1, data = six_1))

##
## Call:
## lm(formula = V2 ~ V1, data = six_1)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -2.43530 -0.60848 -0.08843  0.63099  2.66504
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   2.91166    0.42123   6.912 1.33e-10 ***
## V1            0.49956    0.06661   7.499 5.47e-12 ***
```

```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.9619 on 148 degrees of freedom
## Multiple R-squared:  0.2754, Adjusted R-squared:  0.2705
## F-statistic: 56.24 on 1 and 148 DF,  p-value: 5.468e-12
```

## Model mean of V1 with increasing V1 cut-offs from 0 to 5

```
# Extract visit 1 data
six_1V1 <- six_1$V1

# Generate a vector of cut-off values to iterate over
cutoff <- 0:5

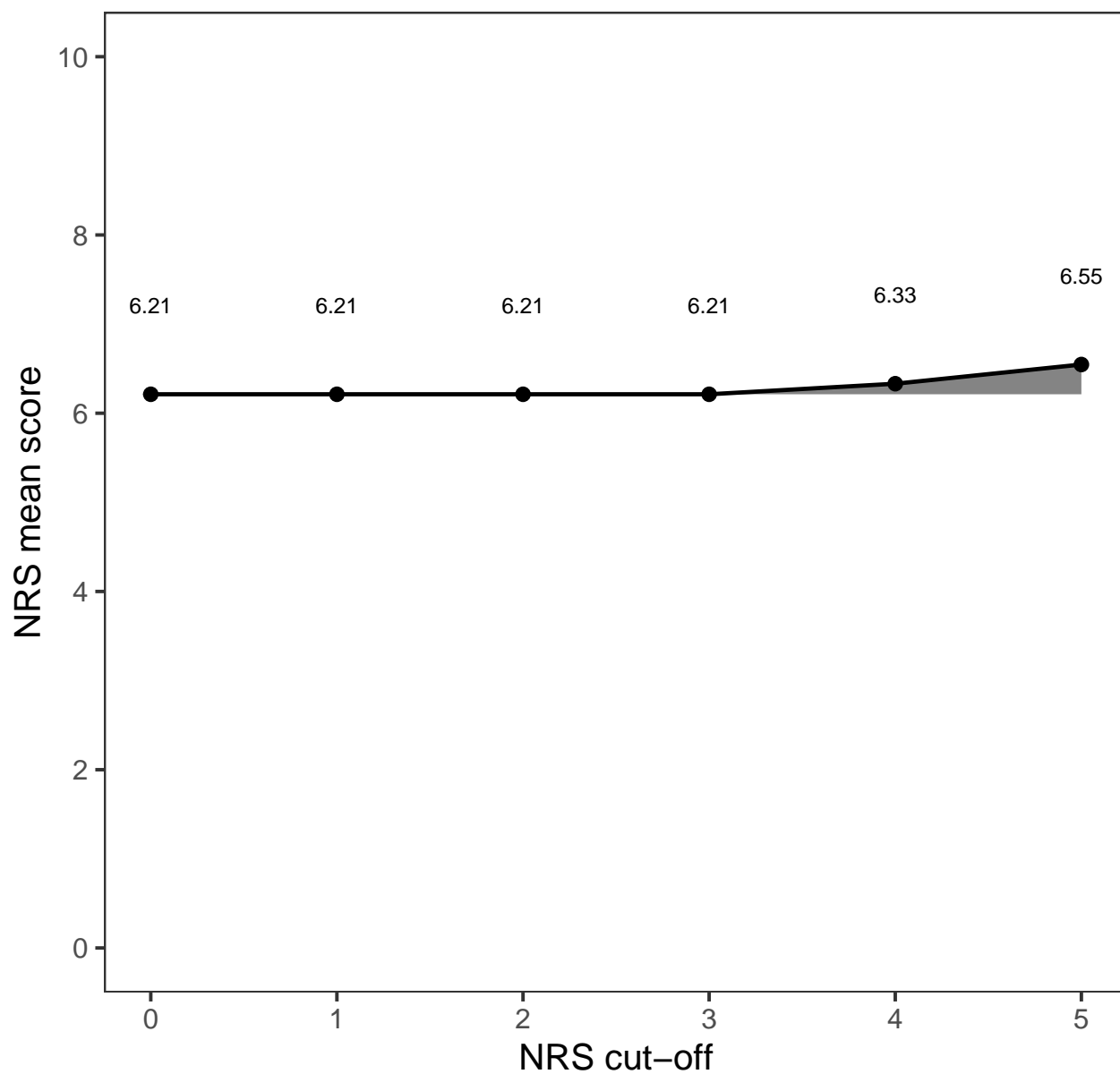
# Generate a vector of V1 means at each V1 cut-off
six_1V1.shift <- sapply(cutoff, function(x){mean(six_1V1[six_1V1 > x])})

# Calculate deviation
(six_1V1.df <- data.frame(cutoff = cutoff,
                          mean = six_1V1.shift) %>%
  mutate(deviation = mean - mean(six_1V1)))

##   cutoff    mean deviation
## 1      0 6.212484 0.0000000
## 2      1 6.212484 0.0000000
## 3      2 6.212484 0.0000000
## 4      3 6.212484 0.0000000
## 5      4 6.330978 0.1184938
## 6      5 6.546297 0.3338133

# Plot data
ggplot(data = six_1V1.df) +
  aes(x = cutoff, y = mean, ymin = mean(six_1V1), ymax = mean) +
  geom_ribbon(alpha = 0.6) +
  geom_point(size = 3) +
  geom_line(size = 1) +
  geom_text(aes(label = round(mean, 2)), nudge_y = 1) +
  scale_y_continuous(limits = c(0, 10),
                     breaks = c(0, 2, 4, 6, 8, 10)) +
  labs(subtitle = 'Shift in V1 mean with increasing V1 NRS cut-off value',
       x = 'NRS cut-off',
       y = 'NRS mean score') +
  theme(panel.grid = element_blank(),
        plot.subtitle = element_text(size = 16))
```

Shift in V1 mean with increasing V1 NRS cut-off value



Model mean of V2 with increasing V1 cut-offs from 0 to 5

```
# Extract visit 2 data
six_1V2 <- six_1$V2

# Generate a vector of cut-off values to iterate over
cutoff <- 0:5

# Generate a vector of V2 means at each V1 cut-off
six_1V2.shift <- map_dbl(.x = cutoff,
  ~ six_1 %>%
    filter(V1 > .x) %>%
    .$V2 %>%
```



```

      mean(.))

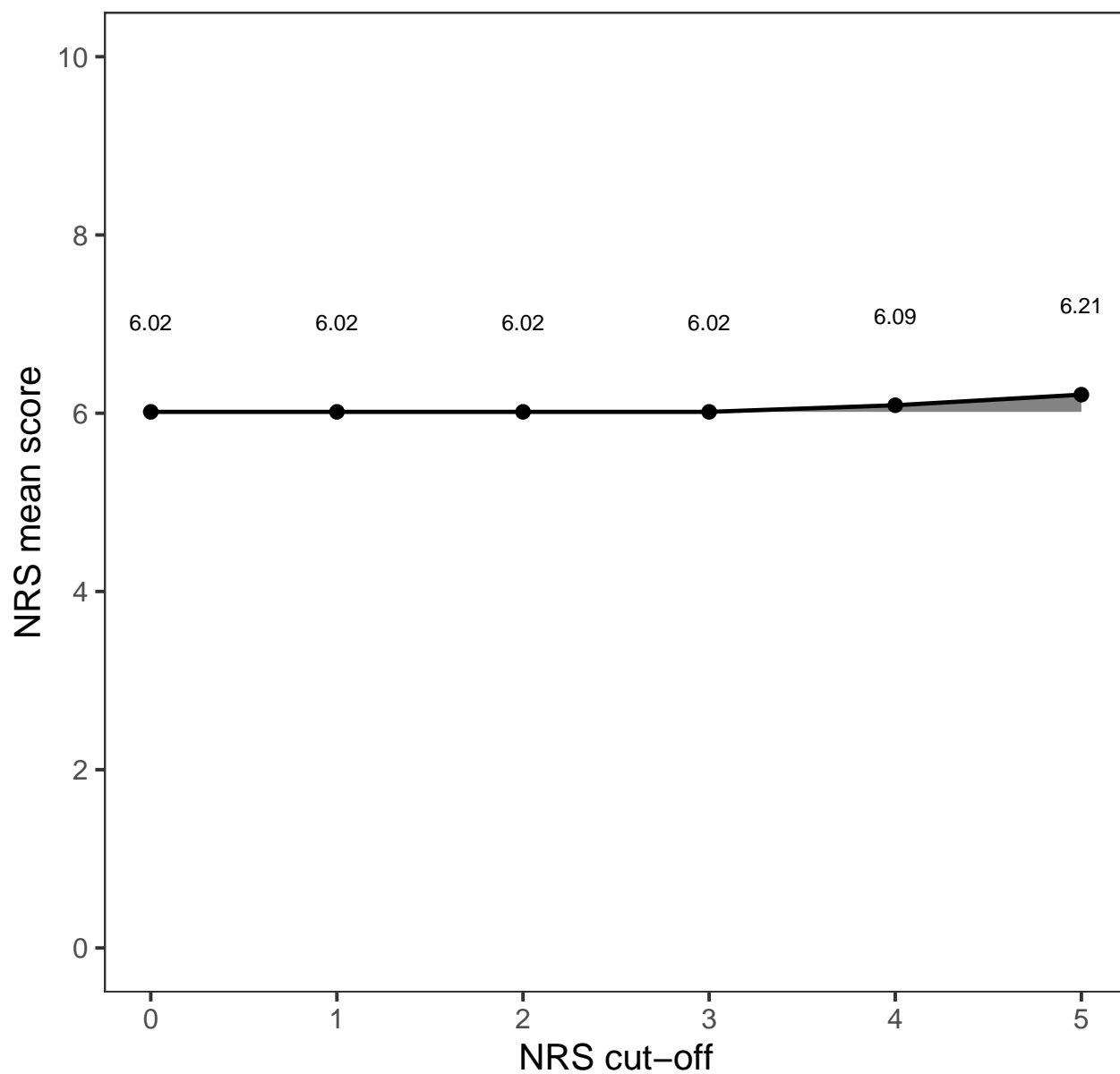
# Calculate deviation
(six_1V2.df <- data.frame(cutoff = cutoff,
                          mean = six_1V2.shift) %>%
  mutate(deviation = mean - mean(six_1V2)))

##   cutoff    mean deviation
## 1      0 6.015193 0.00000000
## 2      1 6.015193 0.00000000
## 3      2 6.015193 0.00000000
## 4      3 6.015193 0.00000000
## 5      4 6.089323 0.07412992
## 6      5 6.208126 0.19293313

# Plot data
ggplot(data = six_1V2.df) +
  aes(x = cutoff, y = mean, ymin = mean(six_1V2), ymax = mean) +
  geom_ribbon(alpha = 0.6) +
  geom_point(size = 3) +
  geom_line(size = 1) +
  geom_text(aes(label = round(mean, 2)), nudge_y = 1) +
  scale_y_continuous(limits = c(0, 10),
                     breaks = c(0, 2, 4, 6, 8, 10)) +
  labs(subtitle = 'Shift in V2 mean with increasing V1 NRS cut-off value',
       x = 'NRS cut-off',
       y = 'NRS mean score') +
  theme(panel.grid = element_blank(),
        plot.subtitle = element_text(size = 16))

```

Shift in V2 mean with increasing V1 NRS cut-off value



## Placebo response

threshold: 0

```
# Process data
placebo_1.0 <- six_1 %>%
  filter(V1 >= 0) %>%
  mutate(difference = V1 - V2) %>%
  mutate(group = 'Cut-off')

# Calculate the mean (95%CI) difference between V1 and V2
groupwiseMean(difference ~ 1,
  data = placebo_1.0,
```

```

R = 5000,
traditional = FALSE,
bca = TRUE) %>%

kable(.)

```

.id	n	Mean	Conf.level	Bca.lower	Bca.upper
NA	150	0.197	0.95	0.00485	0.38

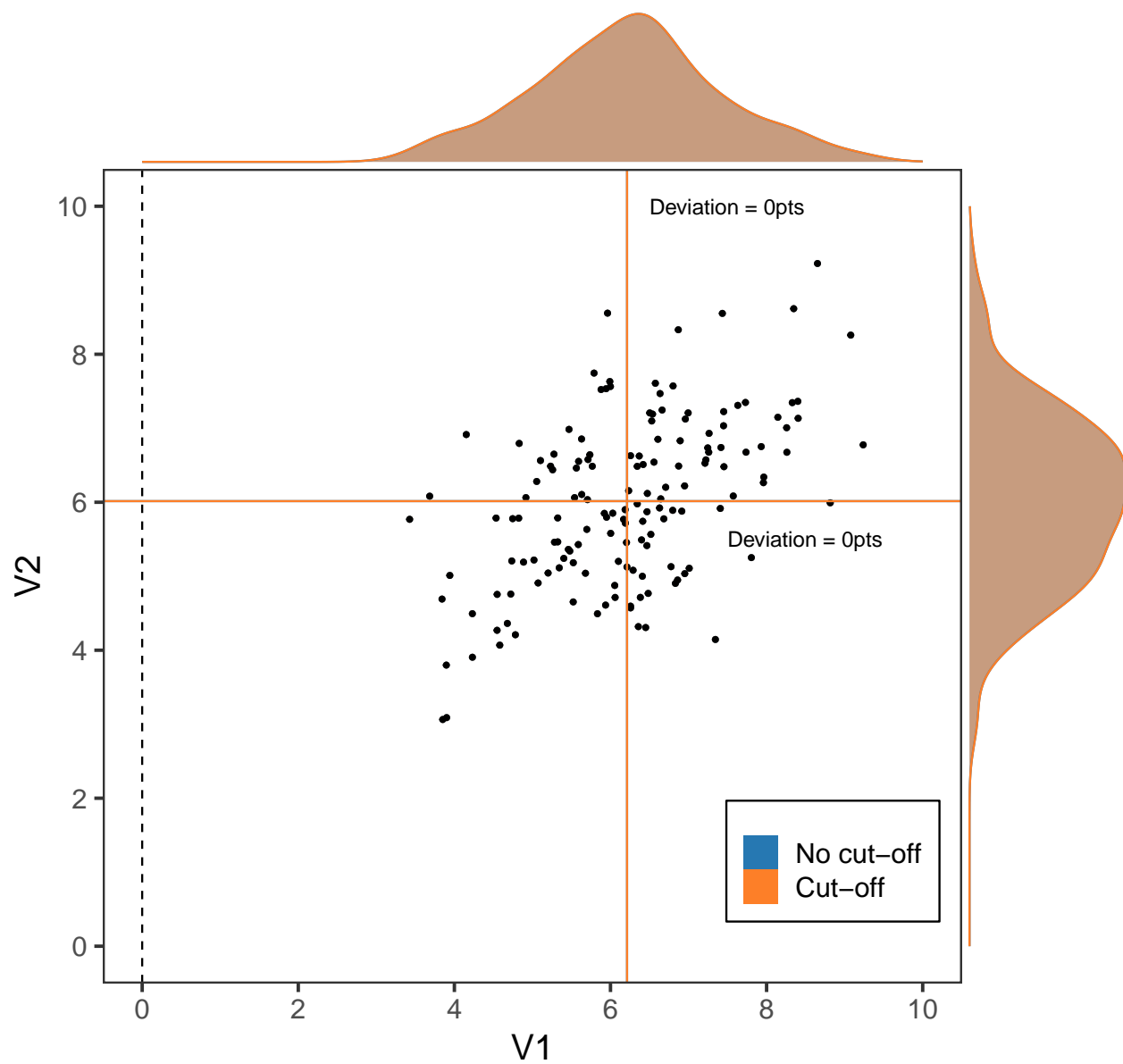
```

# Plot the data
ggMarginal(placebo_1.0[, 1:3] %>%
  bind_rows(six_1) %>%
  mutate(group = factor(group,
                        levels = c('No cut-off', 'Cut-off'),
                        ordered = TRUE)) %>%

  ggplot(data = .) +
  aes(x = V1, y = V2) +
  geom_point(aes(colour = group, fill = group),
    size = 1,
    key_glyph = draw_key_rect) +
  geom_point(data = six_1,
    colour = '#999999',
    size = 1) +
  geom_point(data = placebo_1.0,
    size = 1,
    colour = '#000000') +
  geom_vline(xintercept = mean(six_1$V1),
    colour = pal[1]) +
  geom_vline(xintercept = mean(placebo_1.0$V1),
    colour = pal[2]) +
  geom_vline(xintercept = 0, linetype = 2) +
  geom_hline(yintercept = mean(six_1$V2),
    colour = pal[1]) +
  geom_hline(yintercept = mean(placebo_1.0$V2),
    colour = pal[2]) +
  annotate(geom = 'text', x = 6.5, y = 10, hjust = 0,
    label = str_glue("Deviation = {round(six_1V1.df$deviation[4], 2)}pts")) +
  annotate(geom = 'text', x = 7.5, y = 5.5, hjust = 0,
    label = str_glue("Deviation = {round(six_1V2.df$deviation[4], 2)}pts")) +
  scale_y_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_x_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_fill_manual(values = pal) +
  scale_colour_manual(values = pal) +
  labs(subtitle = 'Base model: Mean = 6.2, SD = 1.2, cor = 0.5; Cut-off: 0') +
  theme(legend.title = element_blank(),
    legend.position = c(0.85, 0.15),
    legend.background = element_rect(colour = '#000000',
      size = 0.5),
    panel.grid = element_blank(),
    plot.subtitle = element_text(size = 16)),
groupColour = TRUE,
groupFill = TRUE)

```

Base model: Mean = 6.2, SD = 1.2, cor = 0.5; Cut-off: 0



threshold: 3

```
# Process data
placebo_1.3 <- six_1 %>%
  filter(V1 >= 3) %>%
  mutate(difference = V1 - V2) %>%
  mutate(group = 'Cut-off')

# Calculate the mean (95%CI) difference between V1 and V2
groupwiseMean(difference ~ 1,
  data = placebo_1.3,
  R = 5000,
  traditional = FALSE,
```

```

kable(.)
bca = TRUE) %>%

```

.id	n	Mean	Conf.level	Bca.lower	Bca.upper
NA	150	0.197	0.95	0.0228	0.373

```

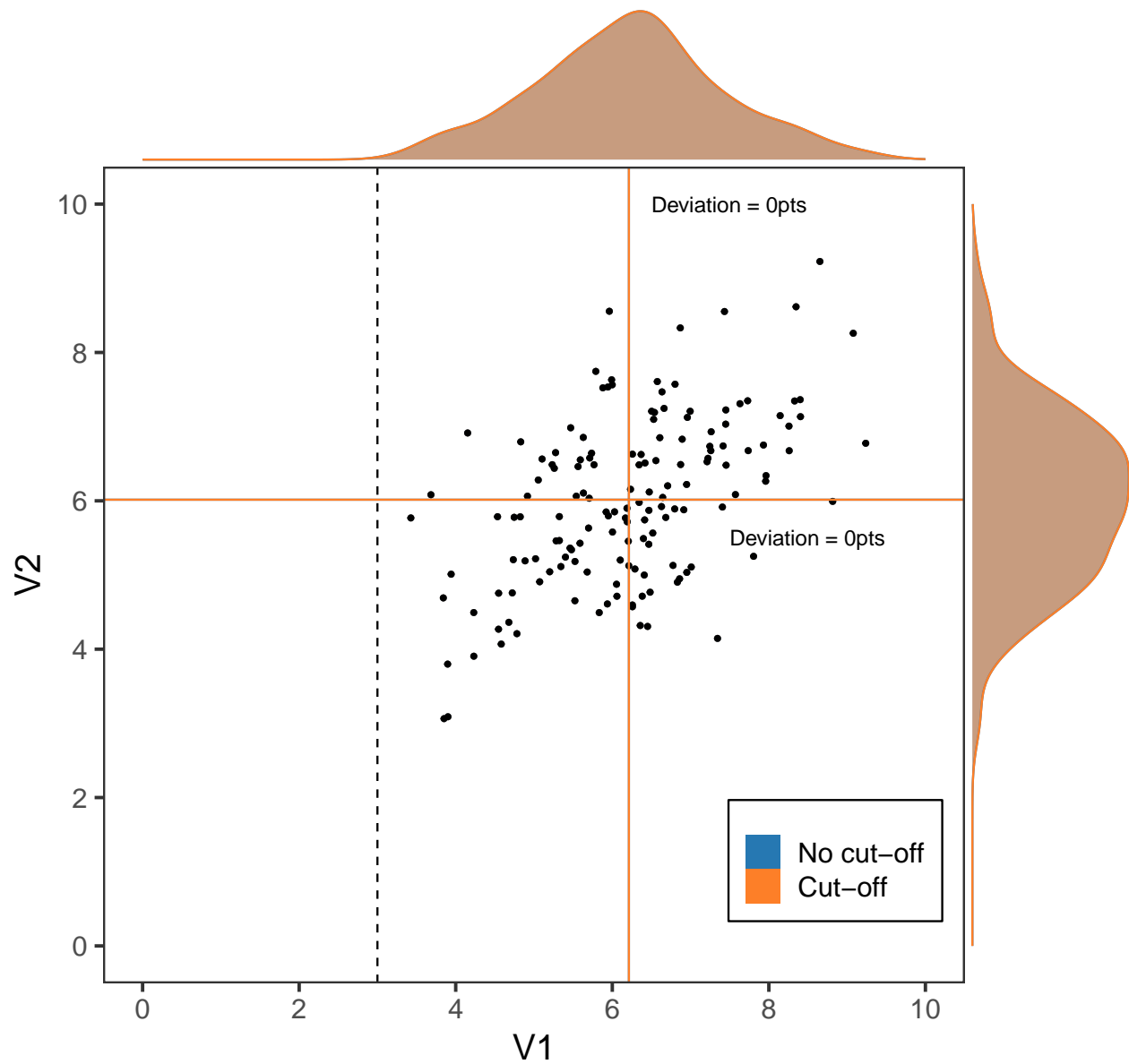
# Plot the data
ggMarginal(placebo_1.3[, 1:3] %>%
  bind_rows(six_1) %>%
  mutate(group = factor(group,
    levels = c('No cut-off', 'Cut-off'),
    ordered = TRUE)) %>%

  ggplot(data = .) +
  aes(x = V1, y = V2) +
  geom_point(aes(colour = group, fill = group),
    size = 1,
    key_glyph = draw_key_rect) +
  geom_point(data = six_1,
    colour = '#999999',
    size = 1) +
  geom_point(data = placebo_1.3,
    size = 1,
    colour = '#000000') +
  geom_vline(xintercept = mean(six_1$V1),
    colour = pal[1]) +
  geom_vline(xintercept = mean(placebo_1.3$V1),
    colour = pal[2]) +
  geom_vline(xintercept = 3, linetype = 2) +
  geom_hline(yintercept = mean(six_1$V2),
    colour = pal[1]) +
  geom_hline(yintercept = mean(placebo_1.3$V2),
    colour = pal[2]) +
  annotate(geom = 'text', x = 6.5, y = 10, hjust = 0,
    label = str_glue("Deviation = {round(six_1V1.df$deviation[4], 2)}pts")) +
  annotate(geom = 'text', x = 7.5, y = 5.5, hjust = 0,
    label = str_glue("Deviation = {round(six_1V2.df$deviation[4], 2)}pts")) +
  scale_y_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_x_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_fill_manual(values = pal) +
  scale_colour_manual(values = pal) +
  labs(subtitle = 'Base model: Mean = 6.2, SD = 1.2, cor = 0.5; Cut-off: 3') +
  theme(legend.title = element_blank(),
    legend.position = c(0.85, 0.15),
    legend.background = element_rect(colour = '#000000',
      size = 0.5),

    panel.grid = element_blank(),
    plot.subtitle = element_text(size = 16)),
groupColour = TRUE,
groupFill = TRUE)

```

Base model: Mean = 6.2, SD = 1.2, cor = 0.5; Cut-off: 3



threshold: 4

```
# Process that data
placebo_1.4 <- six_1 %>%
  filter(V1 >= 4) %>%
  mutate(difference = V1 - V2) %>%
  mutate(group = 'Cut-off')

# Set seed
set.seed(2019)

# Calculate the mean (95%CI) difference between V1 and V2
groupwiseMean(difference ~ 1,
```

```

data = placebo_1.4,
R = 5000,
traditional = FALSE,
bca = TRUE) %>%

kable(.)

```

.id	n	Mean	Conf.level	Bca.lower	Bca.upper
NA	143	0.242	0.95	0.0567	0.418

```

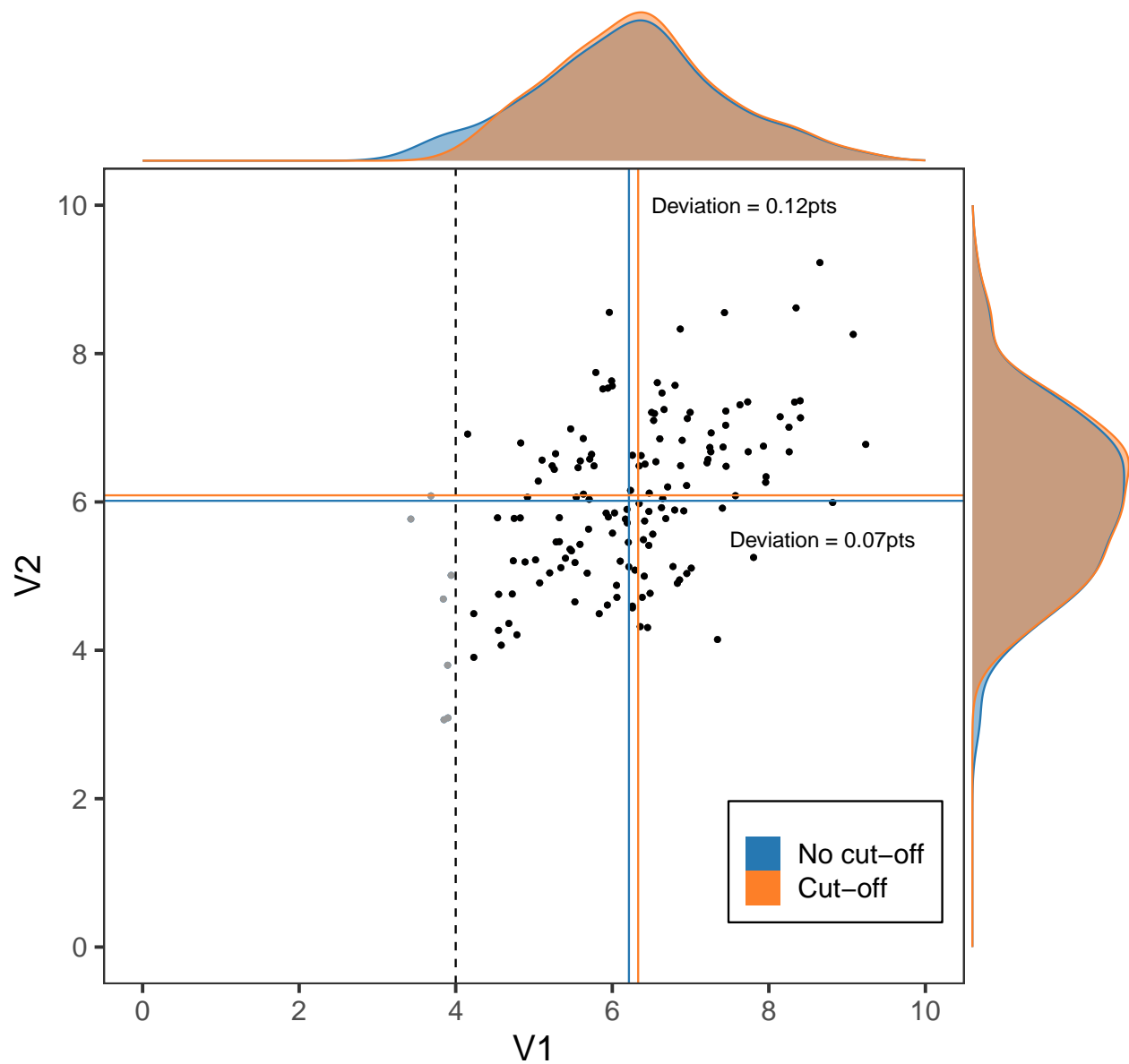
# Plot the data
ggMarginal(placebo_1.4[, 1:3] %>%
  bind_rows(six_1) %>%
  mutate(group = factor(group,
    levels = c('No cut-off', 'Cut-off'),
    ordered = TRUE)) %>%

  ggplot(data = .) +
  aes(x = V1, y = V2) +
  geom_point(aes(colour = group, fill = group),
    size = 1,
    key_glyph = draw_key_rect) +
  geom_point(data = six_1,
    colour = '#999999',
    size = 1) +
  geom_point(data = placebo_1.4,
    size = 1,
    colour = '#000000') +
  geom_vline(xintercept = mean(six_1$V1),
    colour = pal[1]) +
  geom_vline(xintercept = mean(placebo_1.4$V1),
    colour = pal[2]) +
  geom_vline(xintercept = 4, linetype = 2) +
  geom_hline(yintercept = mean(six_1$V2),
    colour = pal[1]) +
  geom_hline(yintercept = mean(placebo_1.4$V2),
    colour = pal[2]) +
  annotate(geom = 'text', x = 6.5, y = 10, hjust = 0,
    label = str_glue("Deviation = {round(six_1V1.df$deviation[5], 2)}pts")) +
  annotate(geom = 'text', x = 7.5, y = 5.5, hjust = 0,
    label = str_glue("Deviation = {round(six_1V2.df$deviation[5], 2)}pts")) +
  scale_y_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_x_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_fill_manual(values = pal) +
  scale_colour_manual(values = pal) +
  labs(subtitle = 'Base model: Mean = 6.2, SD = 1.2, cor = 0.5; Cut-off: 4') +
  theme(legend.title = element_blank(),
    legend.position = c(0.85, 0.15),
    legend.background = element_rect(colour = '#000000',
      size = 0.5),
    panel.grid = element_blank(),
    plot.subtitle = element_text(size = 16)),
groupColour = TRUE,

```

```
groupFill = TRUE)
```

Base model: Mean = 6.2, SD = 1.2, cor = 0.5; Cut-off: 4



threshold: 5

```
# Process that data
placebo_1.5 <- six_1 %>%
  filter(V1 >= 5) %>%
  mutate(difference = V1 - V2) %>%
  mutate(group = 'Cut-off')
```

```
# Set seed
set.seed(2019)
```



```
# Calculate the mean (95%CI) difference between V1 and V2
```

```
groupwiseMean(difference ~ 1,
  data = placebo_1.5,
  R = 5000,
  traditional = FALSE,
  bca = TRUE) %>%
```

```
kable(.)
```

.id	n	Mean	Conf.level	Bca.lower	Bca.upper
NA	127	0.338	0.95	0.146	0.532

```
# Plot the data
```

```
ggMarginal(placebo_1.5[, 1:3] %>%
  bind_rows(six_1) %>%
  mutate(group = factor(group,
    levels = c('No cut-off', 'Cut-off'),
    ordered = TRUE)) %>%

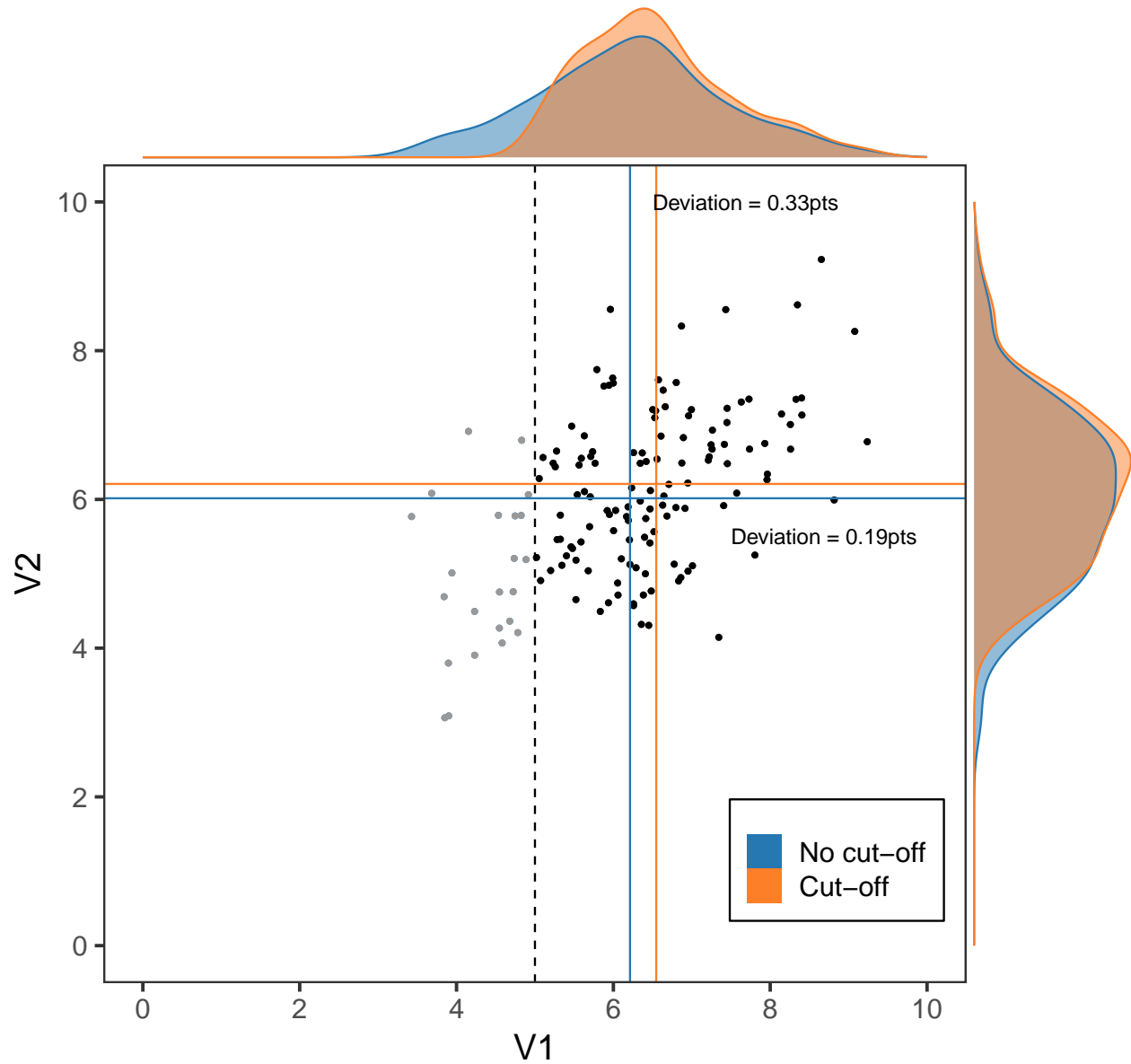
  ggplot(data = .) +
  aes(x = V1, y = V2) +
  geom_point(aes(colour = group, fill = group),
    size = 1,
    key_glyph = draw_key_rect) +
  geom_point(data = six_1,
    colour = '#999999',
    size = 1) +
  geom_point(data = placebo_1.5,
    size = 1,
    colour = '#000000') +
  geom_vline(xintercept = mean(six_1$V1),
    colour = pal[1]) +
  geom_vline(xintercept = mean(placebo_1.5$V1),
    colour = pal[2]) +
  geom_vline(xintercept = 5, linetype = 2) +
  geom_hline(yintercept = mean(six_1$V2),
    colour = pal[1]) +
  geom_hline(yintercept = mean(placebo_1.5$V2),
    colour = pal[2]) +
  annotate(geom = 'text', x = 6.5, y = 10, hjust = 0,
    label = str_glue("Deviation = {round(six_1V1.df$deviation[6], 2)}pts")) +
  annotate(geom = 'text', x = 7.5, y = 5.5, hjust = 0,
    label = str_glue("Deviation = {round(six_1V2.df$deviation[6], 2)}pts")) +
  scale_y_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_x_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_fill_manual(values = pal) +
  scale_colour_manual(values = pal) +
  labs(subtitle = 'Base model: Mean = 6.2, SD = 1.2, cor = 0.5; Cut-off: 5') +
  theme(legend.title = element_blank(),
    legend.position = c(0.85, 0.15),
    legend.background = element_rect(colour = '#000000',
      size = 0.5),
    panel.grid = element_blank(),
```

```

    plot.subtitle = element_text(size = 16)),
    groupColour = TRUE,
    groupFill = TRUE)

```

Base model: Mean = 6.2, SD = 1.2, cor = 0.5; Cut-off: 5



Mean = 6.2, SD = 1.7, cor = 0.5

Generate and summarise data

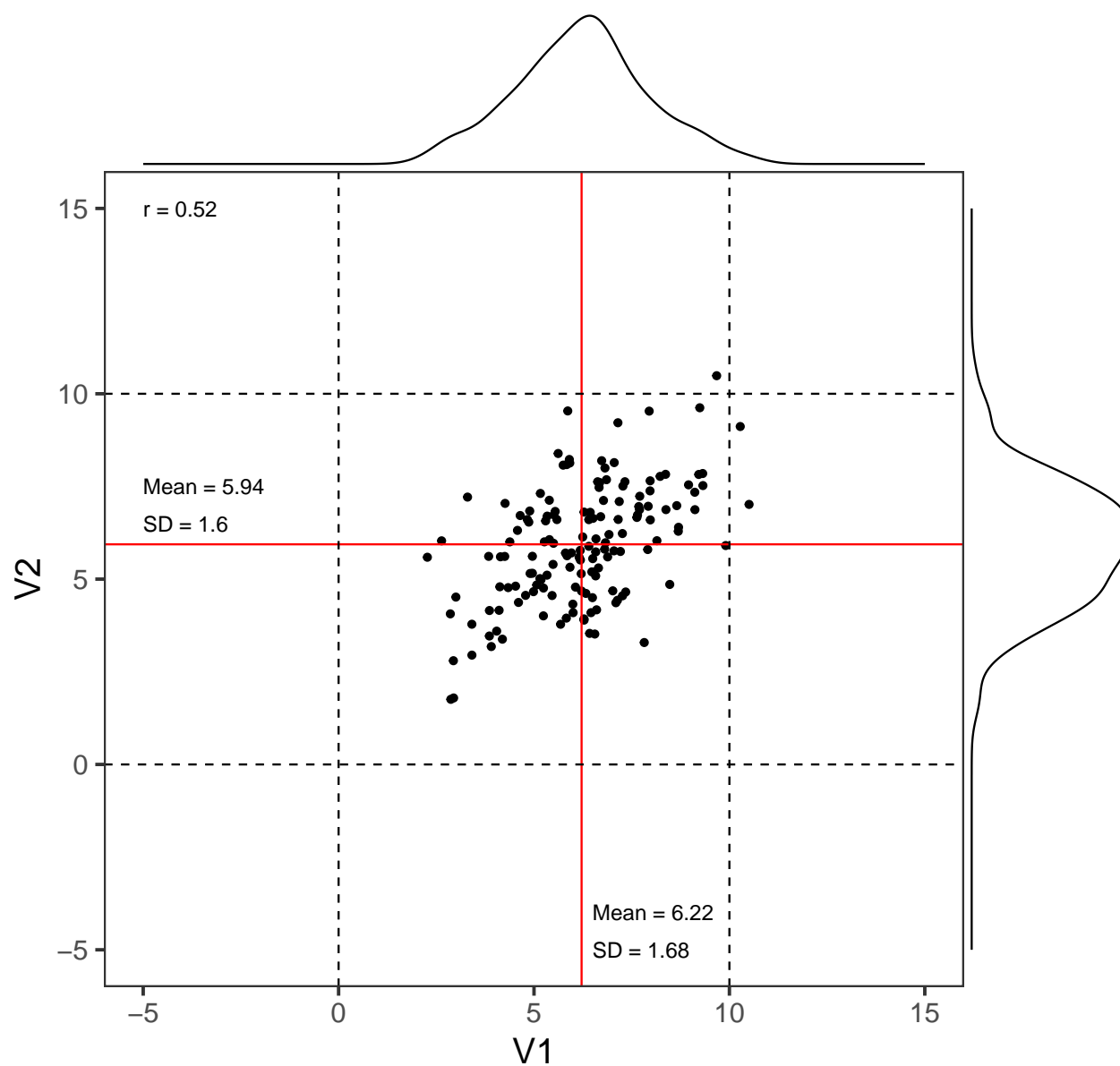
Base data

```
# Set the random seed for reproducibility
set.seed(2019)

# Generate the data
six_2.base <- as.data.frame(mvrnorm(n = 150, mu = c(6.2, 6.2), Sigma = cov_2))

# Plot base data
ggMarginal(ggplot(data = six_2.base) +
  aes(x = V1, y = V2) +
  geom_point() +
  geom_hline(yintercept = mean(six_2.base$V2), colour = 'red') +
  geom_vline(xintercept = mean(six_2.base$V1), colour = 'red') +
  geom_hline(yintercept = 0, linetype = 2) +
  geom_hline(yintercept = 10, linetype = 2) +
  geom_vline(xintercept = 0, linetype = 2) +
  geom_vline(xintercept = 10, linetype = 2) +
  annotate(geom = 'text', x = -5, y = 15, hjust = 0,
    label = str_glue("r = {round(cor(six_2.base$V1, six_2.base$V2), 2)}")) +
  annotate(geom = 'text', x = -5, y = 7.5, hjust = 0,
    label = str_glue("Mean = {round(mean(six_2.base$V2), 2)}")) +
  annotate(geom = 'text', x = -5, y = 6.5, hjust = 0,
    label = str_glue("SD = {round(sd(six_2.base$V2), 2)}")) +
  annotate(geom = 'text', x = 6.5, y = -4, hjust = 0,
    label = str_glue("Mean = {round(mean(six_2.base$V1), 2)}")) +
  annotate(geom = 'text', x = 6.5, y = -5, hjust = 0,
    label = str_glue("SD = {round(sd(six_2.base$V1), 2)}")) +
  labs(subtitle = 'Mean = 6.2, SD = 1.7, cor = 0.5') +
  scale_y_continuous(limits = c(-5, 15)) +
  scale_x_continuous(limits = c(-5, 15)) +
  theme(panel.grid = element_blank(),
    plot.subtitle = element_text(size = 16)))
```

Mean = 6.2, SD = 1.7, cor = 0.5



```
# Linear regression
summary(lm(V2 ~ V1, data = six_2.base))

##
## Call:
## lm(formula = V2 ~ V1, data = six_2.base)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -3.4500 -0.8620 -0.1253  0.8939  3.7755
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  2.83206    0.42887   6.603 6.75e-10 ***
## V1           0.49956    0.06661   7.499 5.47e-12 ***
```

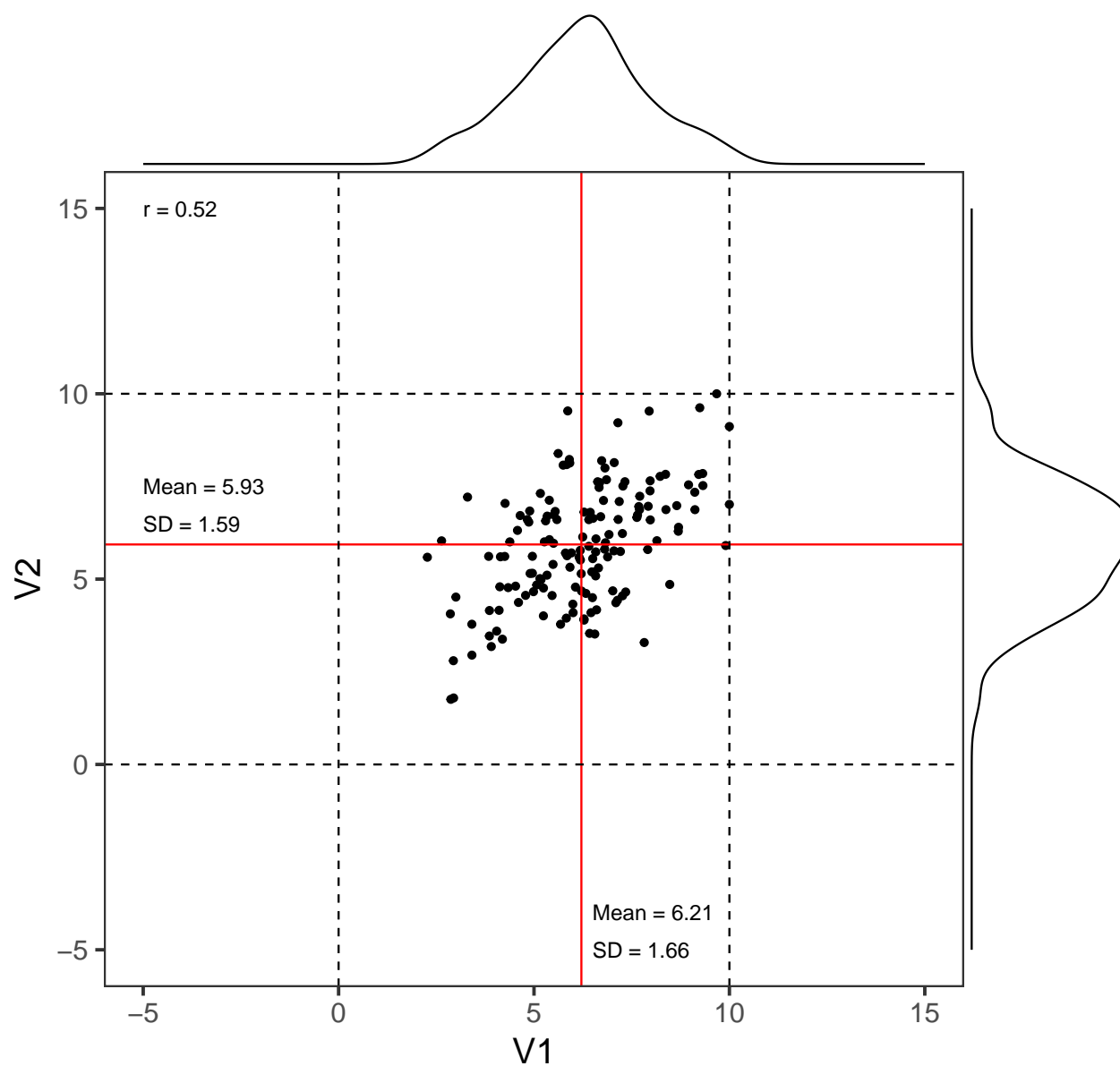
```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.363 on 148 degrees of freedom
## Multiple R-squared:  0.2754, Adjusted R-squared:  0.2705
## F-statistic: 56.24 on 1 and 148 DF,  p-value: 5.468e-12
```

Constrain values to 0-10 range

```
# Process data
six_2 <- six_2.base %>%
  mutate(V1 = case_when(
    V1 < 1 ~ 1,
    V1 > 10 ~ 10,
    TRUE ~ V1)) %>%
  mutate(V2 = case_when(
    V2 < 0 ~ 0,
    V2 > 10 ~ 10,
    TRUE ~ V2)) %>%
  mutate(group = 'No cut-off')

# Plot processed data
ggMarginal(ggplot(data = six_2) +
  aes(x = V1, y = V2) +
  geom_point() +
  geom_hline(yintercept = mean(six_2$V2), colour = 'red') +
  geom_vline(xintercept = mean(six_2$V1), colour = 'red') +
  geom_hline(yintercept = 0, linetype = 2) +
  geom_hline(yintercept = 10, linetype = 2) +
  geom_vline(xintercept = 0, linetype = 2) +
  geom_vline(xintercept = 10, linetype = 2) +
  annotate(geom = 'text', x = -5, y = 15, hjust = 0,
    label = str_glue("r = {round(cor(six_2$V1, six_2$V2), 2)}")) +
  annotate(geom = 'text', x = -5, y = 7.5, hjust = 0,
    label = str_glue("Mean = {round(mean(six_2$V2), 2)}")) +
  annotate(geom = 'text', x = -5, y = 6.5, hjust = 0,
    label = str_glue("SD = {round(sd(six_2$V2), 2)}")) +
  annotate(geom = 'text', x = 6.5, y = -4, hjust = 0,
    label = str_glue("Mean = {round(mean(six_2$V1), 2)}")) +
  annotate(geom = 'text', x = 6.5, y = -5, hjust = 0,
    label = str_glue("SD = {round(sd(six_2$V1), 2)}")) +
  labs(subtitle = 'Mean = 6.2, SD = 1.7 (constrained), cor = 0.5') +
  scale_y_continuous(limits = c(-5, 15)) +
  scale_x_continuous(limits = c(-5, 15)) +
  theme(panel.grid = element_blank(),
    plot.subtitle = element_text(size = 16)))
```

Mean = 6.2, SD = 1.7 (constrained), cor = 0.5



```
# Linear regression
summary(lm(V2 ~ V1, data = six_2))

##
## Call:
## lm(formula = V2 ~ V1, data = six_2)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -3.4494 -0.8419 -0.1246  0.8946  3.7761
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  2.83129    0.42947   6.592 7.15e-10 ***
## V1           0.49958    0.06679   7.480 6.10e-12 ***
```

```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.356 on 148 degrees of freedom
## Multiple R-squared:  0.2743, Adjusted R-squared:  0.2694
## F-statistic: 55.94 on 1 and 148 DF,  p-value: 6.1e-12
```

## Model mean of V1 with increasing V1 cut-offs from 0 to 5

```
# Extract visit 1 data
six_2V1 <- six_2$V1

# Generate a vector of cut-off values to iterate over
cutoff <- 0:5

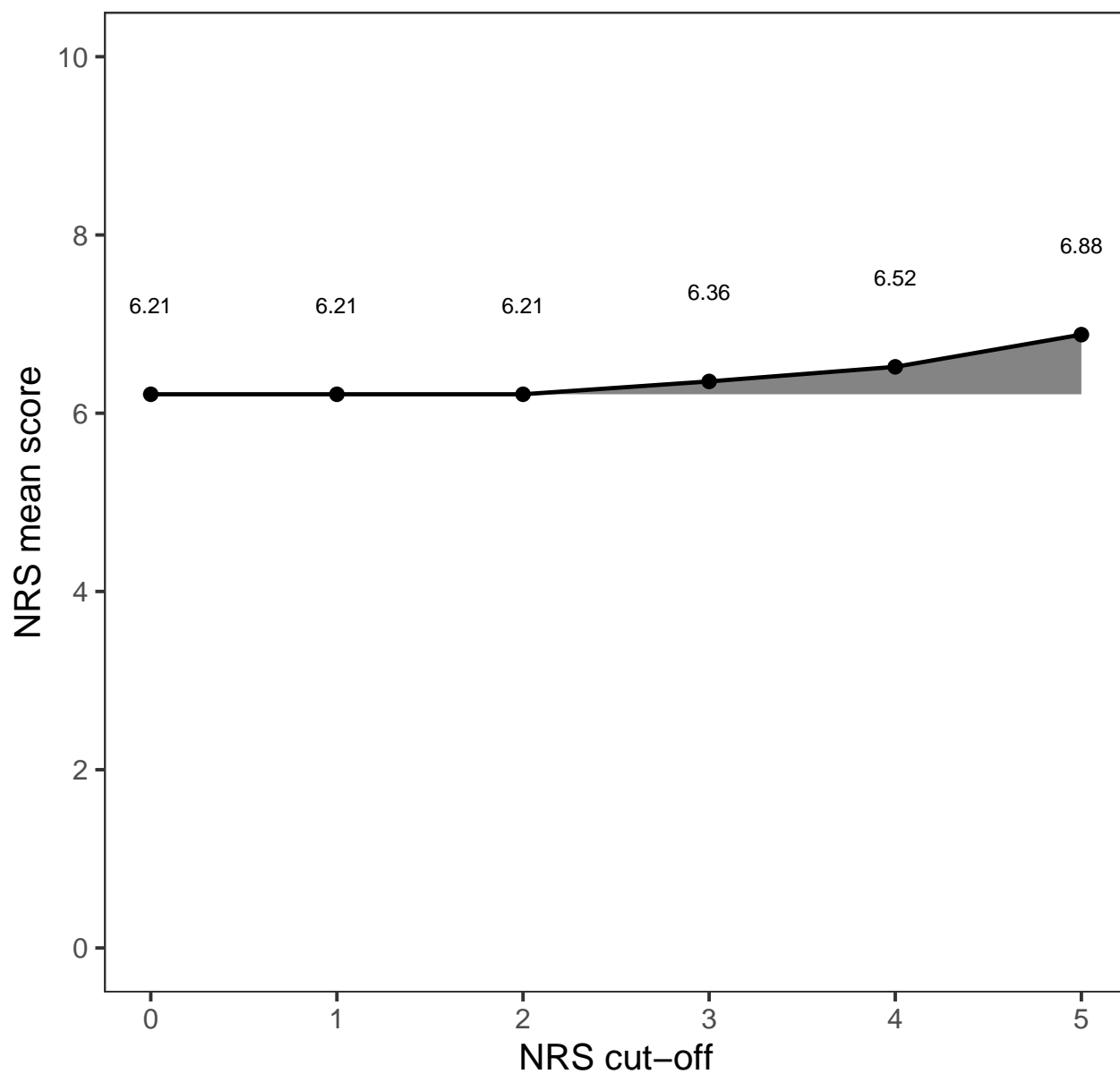
# Generate a vector of V1 means at each V1 cut-off
six_2V1.shift <- sapply(cutoff, function(x){mean(six_2V1[six_2V1 > x])})

# Calculate deviation
(six_2V1.df <- data.frame(cutoff = cutoff,
                          mean = six_2V1.shift) %>%
  mutate(deviation = mean - mean(six_2V1)))

##   cutoff    mean deviation
## 1      0 6.212475 0.0000000
## 2      1 6.212475 0.0000000
## 3      2 6.212475 0.0000000
## 4      3 6.356619 0.1441441
## 5      4 6.520334 0.3078593
## 6      5 6.881636 0.6691615

# Plot data
ggplot(data = six_2V1.df) +
  aes(x = cutoff, y = mean, ymin = mean(six_2V1), ymax = mean) +
  geom_ribbon(alpha = 0.6) +
  geom_point(size = 3) +
  geom_line(size = 1) +
  geom_text(aes(label = round(mean, 2)), nudge_y = 1) +
  scale_y_continuous(limits = c(0, 10),
                     breaks = c(0, 2, 4, 6, 8, 10)) +
  labs(subtitle = 'Shift in V1 mean with increasing V1 NRS cut-off value',
       x = 'NRS cut-off',
       y = 'NRS mean score') +
  theme(panel.grid = element_blank(),
        plot.subtitle = element_text(size = 16))
```

Shift in V1 mean with increasing V1 NRS cut-off value



Model mean of V2 with increasing V1 cut-offs from 0 to 5

```
# Extract visit 2 data
six_2V2 <- six_2$V2

# Generate a vector of cut-off values to iterate over
cutoff <- 0:5

# Generate a vector of V2 means at each V1 cut-off
six_2V2.shift <- map_dbl(.x = cutoff,
  ~ six_2 %>%
    filter(V1 > .x) %>%
    .$V2 %>%
```



```

      mean(.))

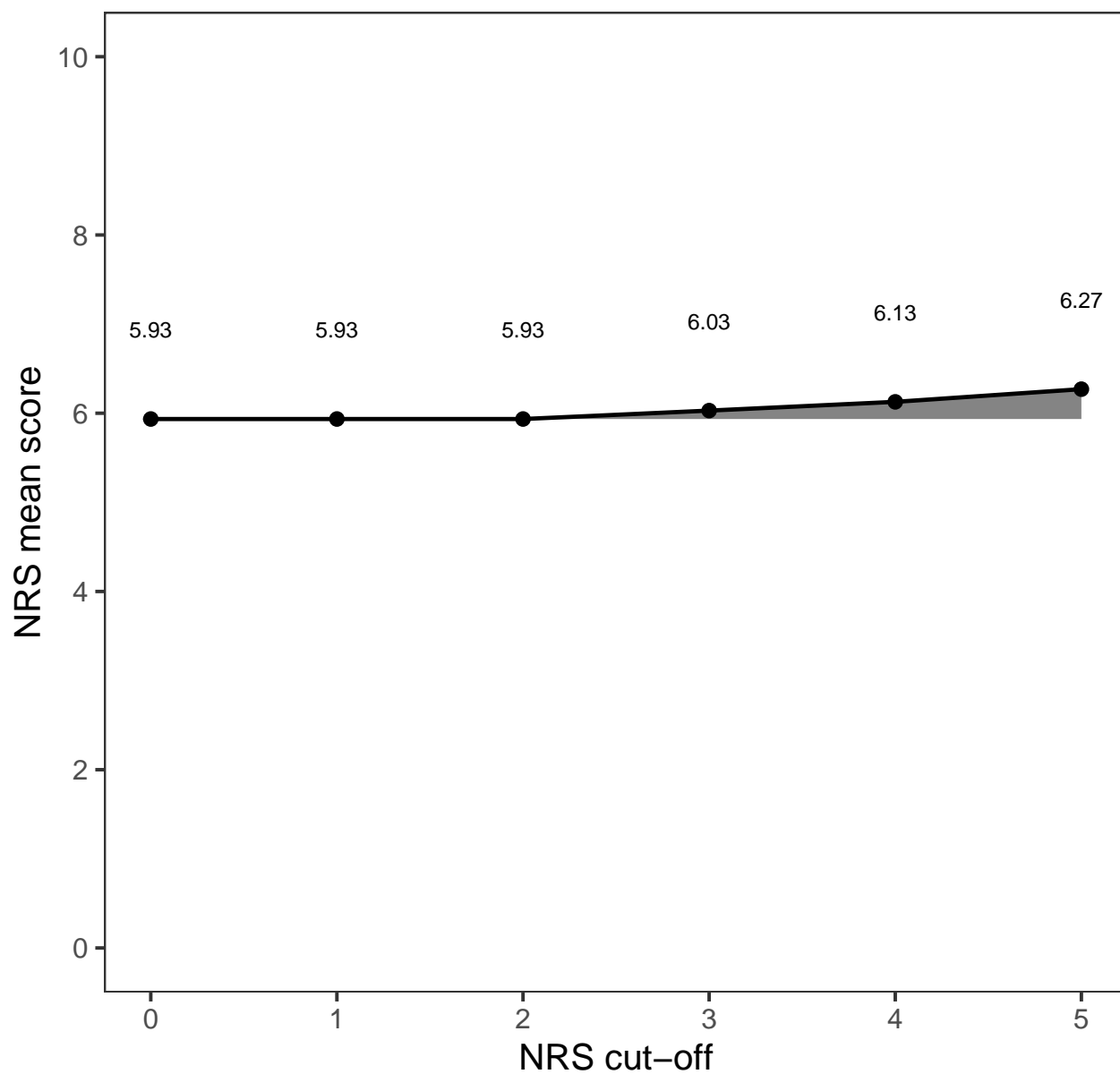
# Calculate deviation
(six_2V2.df <- data.frame(cutoff = cutoff,
                          mean = six_2V2.shift) %>%
  mutate(deviation = mean - mean(six_2V2)))

##   cutoff    mean deviation
## 1      0 5.934941 0.00000000
## 2      1 5.934941 0.00000000
## 3      2 5.934941 0.00000000
## 4      3 6.029214 0.09427288
## 5      4 6.127460 0.19251869
## 6      5 6.269851 0.33490973

# Plot data
ggplot(data = six_2V2.df) +
  aes(x = cutoff, y = mean, ymin = mean(six_2V2), ymax = mean) +
  geom_ribbon(alpha = 0.6) +
  geom_point(size = 3) +
  geom_line(size = 1) +
  geom_text(aes(label = round(mean, 2)), nudge_y = 1) +
  scale_y_continuous(limits = c(0, 10),
                     breaks = c(0, 2, 4, 6, 8, 10)) +
  labs(subtitle = 'Shift in V2 mean with increasing V1 NRS cut-off value',
       x = 'NRS cut-off',
       y = 'NRS mean score') +
  theme(panel.grid = element_blank(),
        plot.subtitle = element_text(size = 16))

```

Shift in V2 mean with increasing V1 NRS cut-off value



## Placebo response

threshold: 0

```
# Process data
placebo_2.0 <- six_2 %>%
  filter(V1 >= 0) %>%
  mutate(difference = V1 - V2) %>%
  mutate(group = 'Cut-off')

# Calculate the mean (95%CI) difference between V1 and V2
groupwiseMean(difference ~ 1,
  data = placebo_2.0,
```

```

R = 5000,
traditional = FALSE,
bca = TRUE) %>%

kable(.)

```

.id	n	Mean	Conf.level	Bca.lower	Bca.upper
NA	150	0.278	0.95	0.0115	0.534

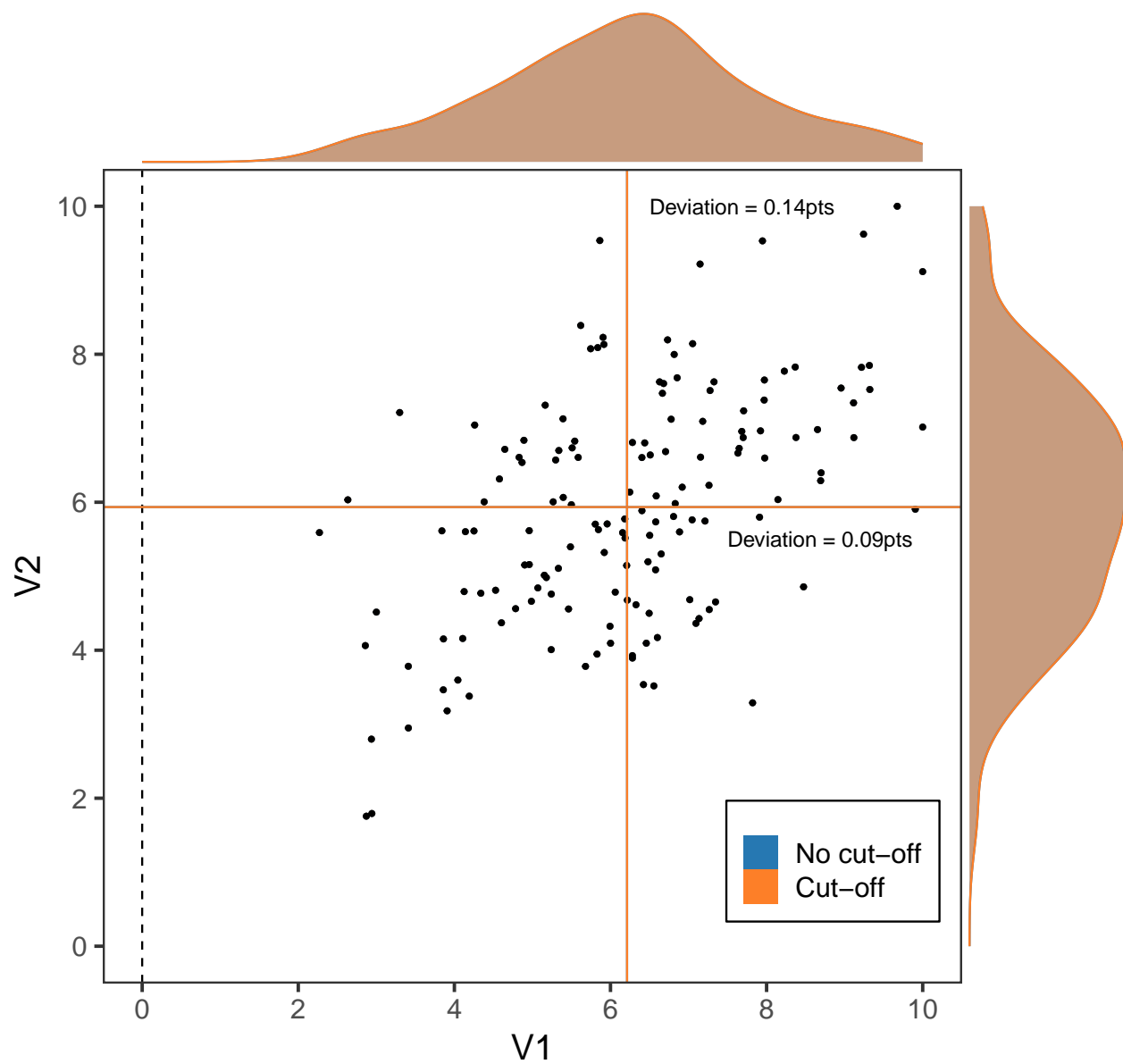
```

# Plot the data
ggMarginal(placebo_2.0[, 1:3] %>%
  bind_rows(six_2) %>%
  mutate(group = factor(group,
                        levels = c('No cut-off', 'Cut-off'),
                        ordered = TRUE)) %>%

  ggplot(data = .) +
  aes(x = V1, y = V2) +
  geom_point(aes(colour = group, fill = group),
    size = 1,
    key_glyph = draw_key_rect) +
  geom_point(data = six_2,
    colour = '#999999',
    size = 1) +
  geom_point(data = placebo_2.0,
    size = 1,
    colour = '#000000') +
  geom_vline(xintercept = mean(six_2$V1),
    colour = pal[1]) +
  geom_vline(xintercept = mean(placebo_2.0$V1),
    colour = pal[2]) +
  geom_vline(xintercept = 0, linetype = 2) +
  geom_hline(yintercept = mean(six_2$V2),
    colour = pal[1]) +
  geom_hline(yintercept = mean(placebo_2.0$V2),
    colour = pal[2]) +
  annotate(geom = 'text', x = 6.5, y = 10, hjust = 0,
    label = str_glue("Deviation = {round(six_2V1.df$deviation[4], 2)}pts")) +
  annotate(geom = 'text', x = 7.5, y = 5.5, hjust = 0,
    label = str_glue("Deviation = {round(six_2V2.df$deviation[4], 2)}pts")) +
  scale_y_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_x_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_fill_manual(values = pal) +
  scale_colour_manual(values = pal) +
  labs(subtitle = 'Base model: Mean = 6.2, SD = 1.2, cor = 0.5; Cut-off: 0') +
  theme(legend.title = element_blank(),
    legend.position = c(0.85, 0.15),
    legend.background = element_rect(colour = '#000000',
      size = 0.5),
    panel.grid = element_blank(),
    plot.subtitle = element_text(size = 16)),
groupColour = TRUE,
groupFill = TRUE)

```

Base model: Mean = 6.2, SD = 1.2, cor = 0.5; Cut-off: 0



threshold: 3

```
# Process data
placebo_2.3 <- six_2 %>%
  filter(V1 >= 3) %>%
  mutate(difference = V1 - V2) %>%
  mutate(group = 'Cut-off')

# Set seed
set.seed(2019)

# Calculate the mean (95%CI) difference between V1 and V2
groupwiseMean(difference ~ 1,
```

```

data = placebo_2.3,
R = 5000,
traditional = FALSE,
bca = TRUE) %>%

kable(.)

```

.id	n	Mean	Conf.level	Bca.lower	Bca.upper
NA	144	0.327	0.95	0.0797	0.571

```

# Plot the data
ggMarginal(placebo_2.3[, 1:3] %>%
  bind_rows(six_2) %>%
  mutate(group = factor(group,
                        levels = c('No cut-off', 'Cut-off'),
                        ordered = TRUE)) %>%

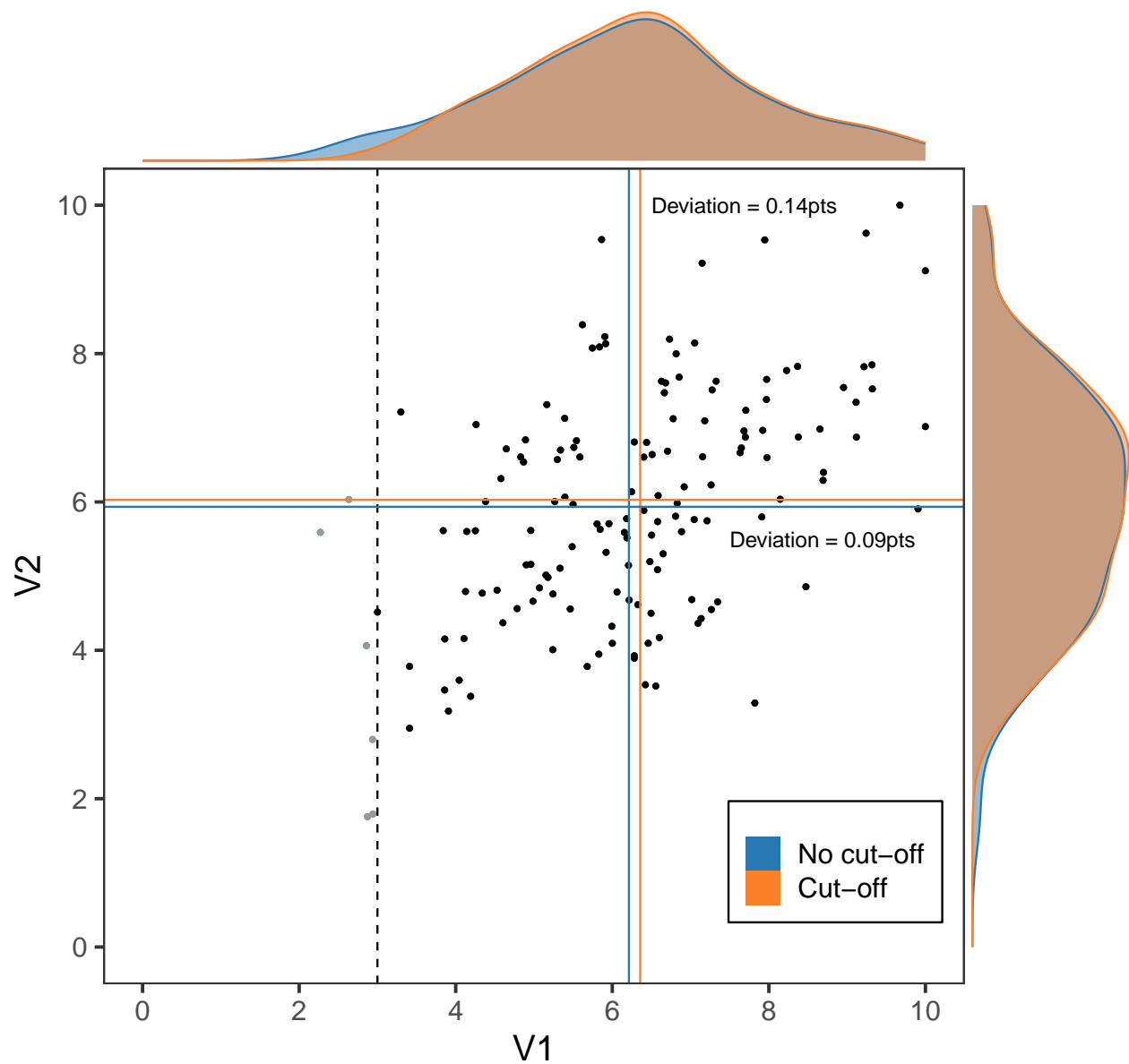
  ggplot(data = .) +
  aes(x = V1, y = V2) +
  geom_point(aes(colour = group, fill = group),
            size = 1,
            key_glyph = draw_key_rect) +
  geom_point(data = six_2,
            colour = '#999999',
            size = 1) +
  geom_point(data = placebo_2.3,
            size = 1,
            colour = '#000000') +
  geom_vline(xintercept = mean(six_2$V1),
            colour = pal[1]) +
  geom_vline(xintercept = mean(placebo_2.3$V1),
            colour = pal[2]) +
  geom_vline(xintercept = 3, linetype = 2) +
  geom_hline(yintercept = mean(six_2$V2),
            colour = pal[1]) +
  geom_hline(yintercept = mean(placebo_2.3$V2),
            colour = pal[2]) +
  annotate(geom = 'text', x = 6.5, y = 10, hjust = 0,
            label = str_glue("Deviation = {round(six_2V1.df$deviation[4], 2)}pts")) +
  annotate(geom = 'text', x = 7.5, y = 5.5, hjust = 0,
            label = str_glue("Deviation = {round(six_2V2.df$deviation[4], 2)}pts")) +
  scale_y_continuous(limits = c(0, 10),
                    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_x_continuous(limits = c(0, 10),
                    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_fill_manual(values = pal) +
  scale_colour_manual(values = pal) +
  labs(subtitle = 'Base model: Mean = 6.2, SD = 1.7, cor = 0.5; Cut-off: 3') +
  theme(legend.title = element_blank(),
        legend.position = c(0.85, 0.15),
        legend.background = element_rect(colour = '#000000',
                                          size = 0.5),

        panel.grid = element_blank(),
        plot.subtitle = element_text(size = 16)),
groupColour = TRUE,

```

```
groupFill = TRUE)
```

Base model: Mean = 6.2, SD = 1.7, cor = 0.5; Cut-off: 3



threshold: 4

```
# Process that data
placebo_2.4 <- six_2 %>%
  filter(V1 >= 4) %>%
  mutate(difference = V1 - V2) %>%
  mutate(group = 'Cut-off')
```

```
# Set seed
set.seed(2019)
```

```
# Calculate the mean (95%CI) difference between V1 and V2
```

```
groupwiseMean(difference ~ 1,
  data = placebo_2.4,
  R = 5000,
  traditional = FALSE,
  bca = TRUE) %>%
```

```
kable(.)
```

.id	n	Mean	Conf.level	Bca.lower	Bca.upper
NA	136	0.393	0.95	0.13	0.654

```
# Plot the data
```

```
ggMarginal(placebo_2.4[, 1:3] %>%
  bind_rows(six_2) %>%
  mutate(group = factor(group,
    levels = c('No cut-off', 'Cut-off'),
    ordered = TRUE)) %>%

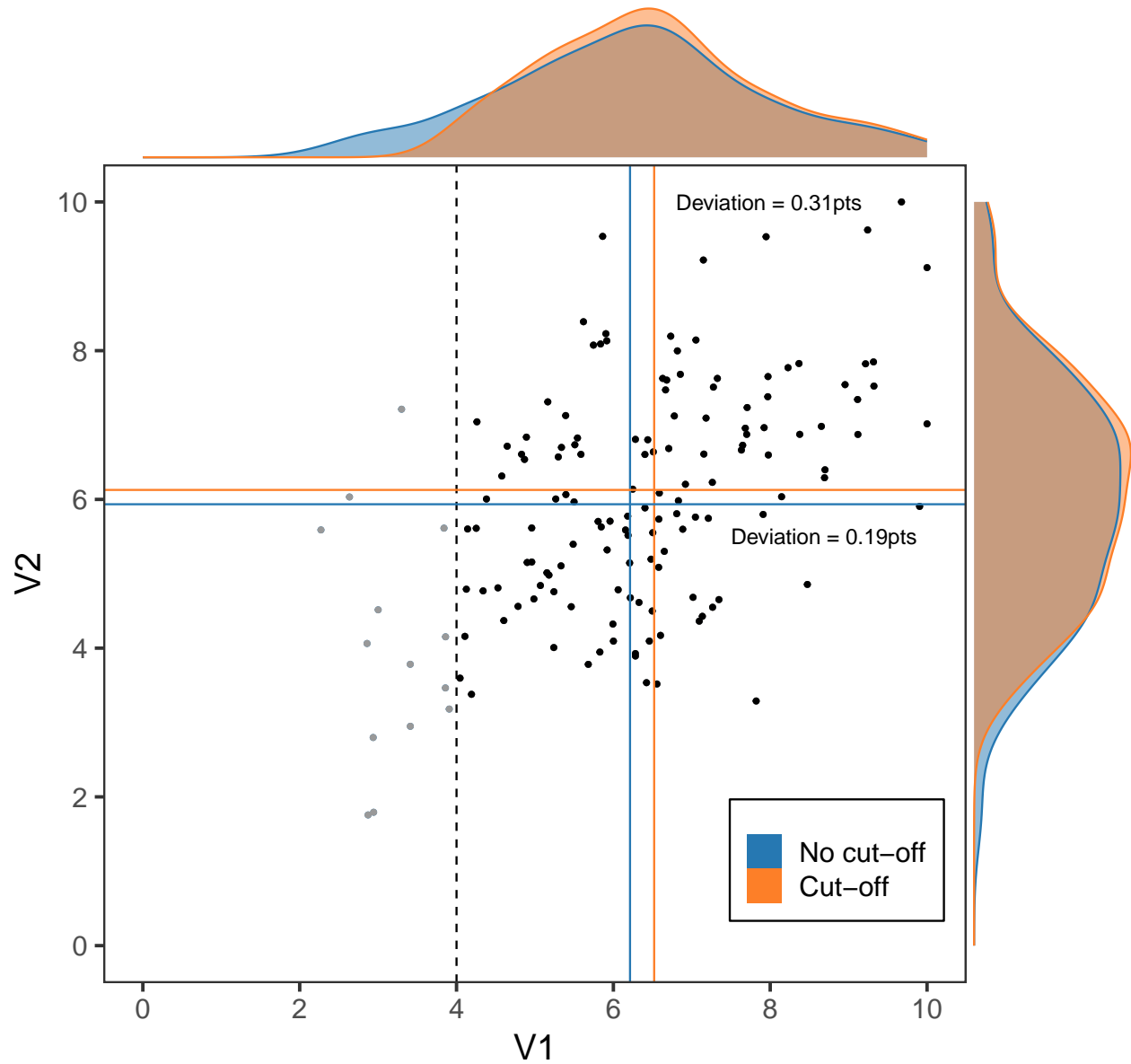
  ggplot(data = .) +
  aes(x = V1, y = V2) +
  geom_point(aes(colour = group, fill = group),
    size = 1,
    key_glyph = draw_key_rect) +
  geom_point(data = six_2,
    colour = '#999999',
    size = 1) +
  geom_point(data = placebo_2.4,
    size = 1,
    colour = '#000000') +
  geom_vline(xintercept = mean(six_2$V1),
    colour = pal[1]) +
  geom_vline(xintercept = mean(placebo_2.4$V1),
    colour = pal[2]) +
  geom_vline(xintercept = 4, linetype = 2) +
  geom_hline(yintercept = mean(six_2$V2),
    colour = pal[1]) +
  geom_hline(yintercept = mean(placebo_2.4$V2),
    colour = pal[2]) +
  annotate(geom = 'text', x = 6.8, y = 10, hjust = 0,
    label = str_glue("Deviation = {round(six_2V1.df$deviation[5], 2)}pts")) +
  annotate(geom = 'text', x = 7.5, y = 5.5, hjust = 0,
    label = str_glue("Deviation = {round(six_2V2.df$deviation[5], 2)}pts")) +
  scale_y_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_x_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_fill_manual(values = pal) +
  scale_colour_manual(values = pal) +
  labs(subtitle = 'Base model: Mean = 6.2, SD = 1.7, cor = 0.5; Cut-off: 4') +
  theme(legend.title = element_blank(),
    legend.position = c(0.85, 0.15),
    legend.background = element_rect(colour = '#000000',
      size = 0.5),
    panel.grid = element_blank(),
```

```

    plot.subtitle = element_text(size = 16)),
    groupColour = TRUE,
    groupFill = TRUE)

```

Base model: Mean = 6.2, SD = 1.7, cor = 0.5; Cut-off: 4



threshold: 5

```

# Process that data
placebo_2.5 <- six_2 %>%
  filter(V1 >= 5) %>%
  mutate(difference = V1 - V2) %>%
  mutate(group = 'Cut-off')

```

```

# Set seed

```



```
set.seed(2019)
```

```
# Calculate the mean (95%CI) difference between V1 and V2
```

```
groupwiseMean(difference ~ 1,
  data = placebo_2.5,
  R = 5000,
  traditional = FALSE,
  bca = TRUE) %>%
```

```
kable(.)
```

.id	n	Mean	Conf.level	Bca.lower	Bca.upper
NA	115	0.612	0.95	0.33	0.885

```
# Plot the data
```

```
ggMarginal(placebo_2.5[, 1:3] %>%
  bind_rows(six_2) %>%
  mutate(group = factor(group,
    levels = c('No cut-off', 'Cut-off'),
    ordered = TRUE)) %>%

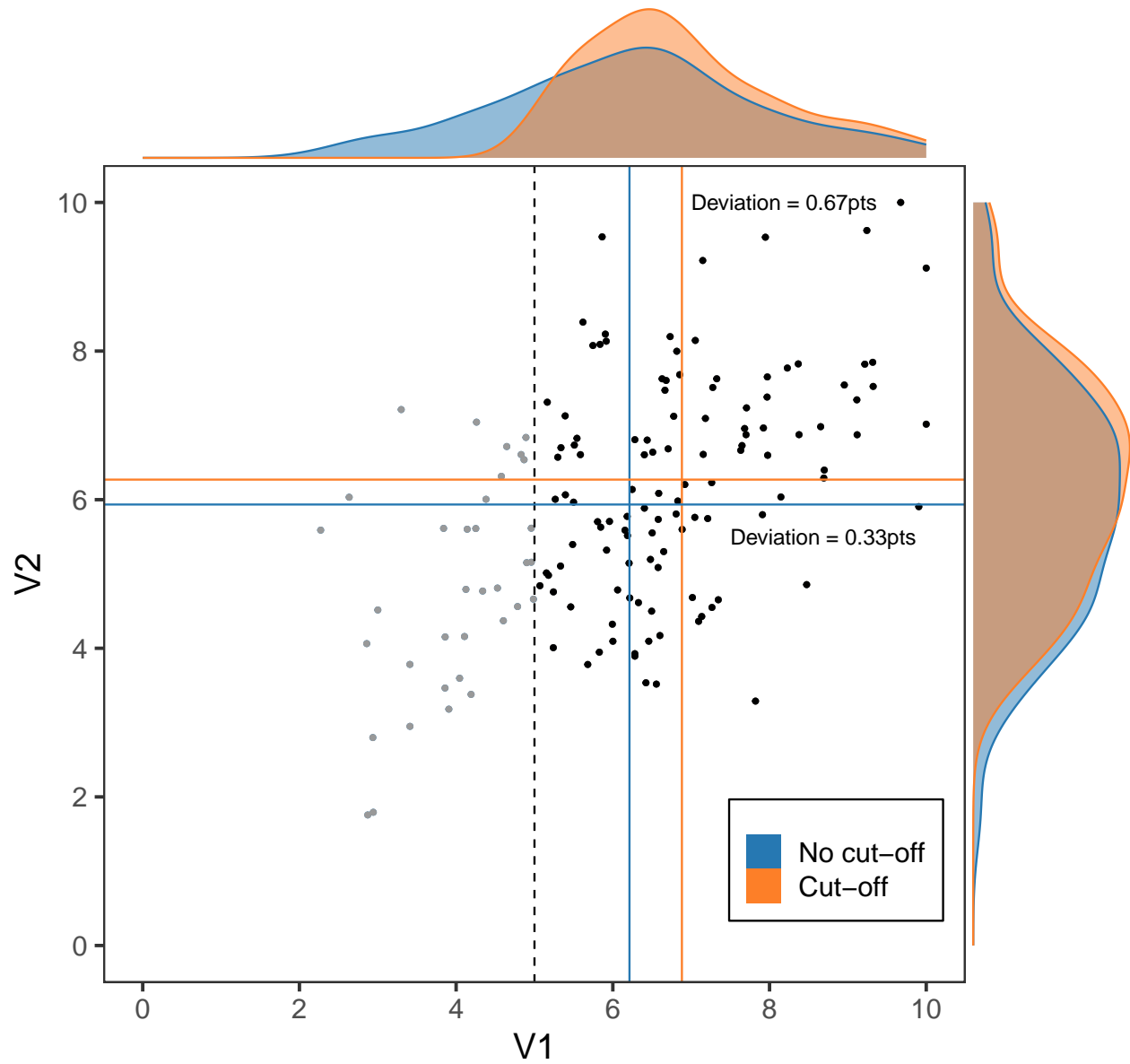
  ggplot(data = .) +
  aes(x = V1, y = V2) +
  geom_point(aes(colour = group, fill = group),
    size = 1,
    key_glyph = draw_key_rect) +
  geom_point(data = six_2,
    colour = '#999999',
    size = 1) +
  geom_point(data = placebo_2.5,
    size = 1,
    colour = '#000000') +
  geom_vline(xintercept = mean(six_2$V1),
    colour = pal[1]) +
  geom_vline(xintercept = mean(placebo_2.5$V1),
    colour = pal[2]) +
  geom_vline(xintercept = 5, linetype = 2) +
  geom_hline(yintercept = mean(six_2$V2),
    colour = pal[1]) +
  geom_hline(yintercept = mean(placebo_2.5$V2),
    colour = pal[2]) +
  annotate(geom = 'text', x = 7, y = 10, hjust = 0,
    label = str_glue("Deviation = {round(six_2V1.df$deviation[6], 2)}pts")) +
  annotate(geom = 'text', x = 7.5, y = 5.5, hjust = 0,
    label = str_glue("Deviation = {round(six_2V2.df$deviation[6], 2)}pts")) +
  scale_y_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_x_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_fill_manual(values = pal) +
  scale_colour_manual(values = pal) +
  labs(subtitle = 'Base model: Mean = 6.2, SD = 1.7, cor = 0.5; Cut-off: 5') +
  theme(legend.title = element_blank(),
    legend.position = c(0.85, 0.15),
    legend.background = element_rect(colour = '#000000',
```

```

size = 0.5),
panel.grid = element_blank(),
plot.subtitle = element_text(size = 16)),
groupColour = TRUE,
groupFill = TRUE)

```

Base model: Mean = 6.2, SD = 1.7, cor = 0.5; Cut-off: 5



Mean = 6.2, SD = 2.2, cor = 0.5

Generate and summarise data

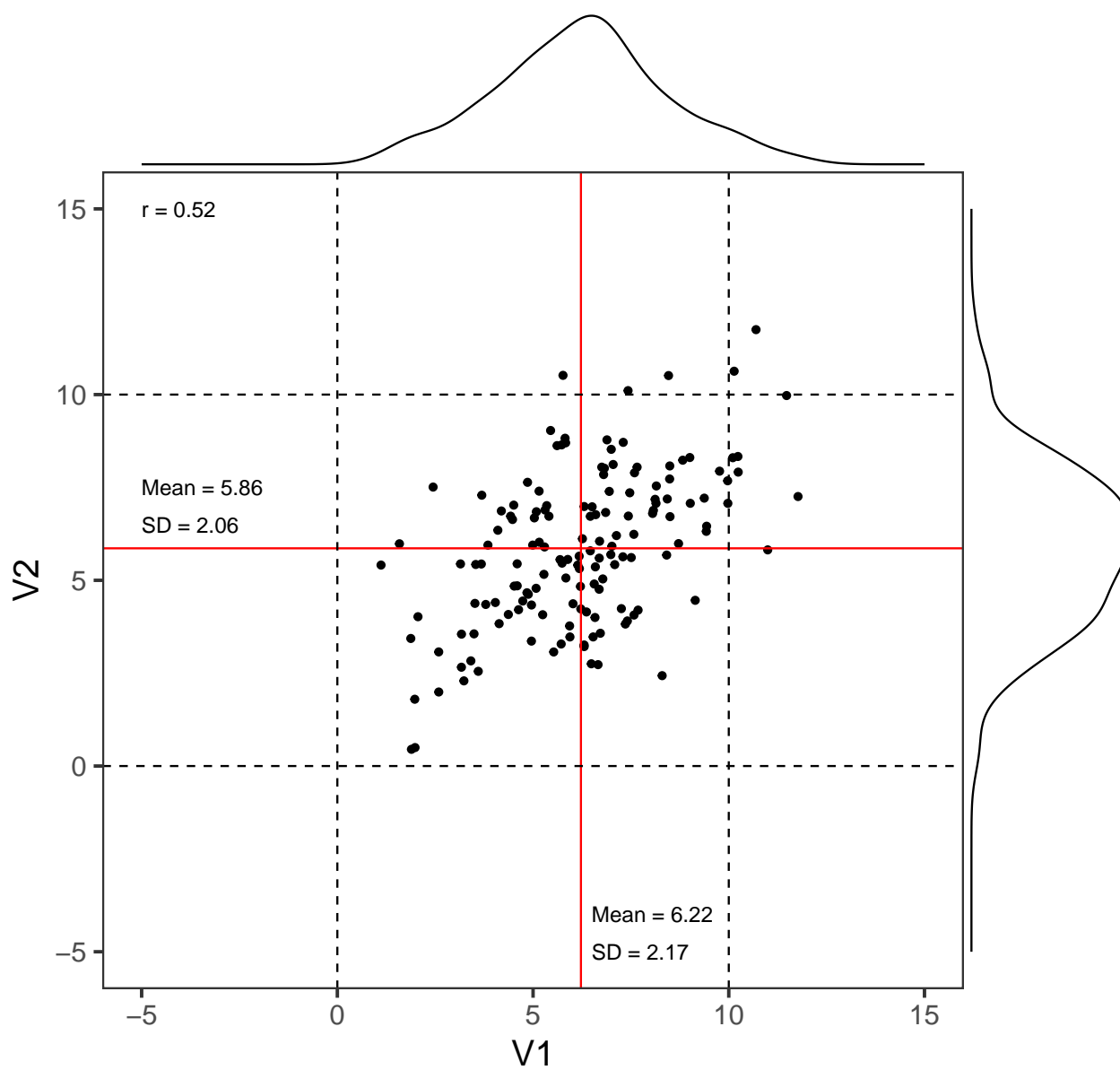
Base data

```
# Set the random seed for reproducibility
set.seed(2019)

# Generate the data
six_3.base <- as.data.frame(mvrnorm(n = 150, mu = c(6.2, 6.2), Sigma = cov_3))

# Plot base data
ggMarginal(ggplot(data = six_3.base) +
  aes(x = V1, y = V2) +
  geom_point() +
  geom_hline(yintercept = mean(six_3.base$V2), colour = 'red') +
  geom_vline(xintercept = mean(six_3.base$V1), colour = 'red') +
  geom_hline(yintercept = 0, linetype = 2) +
  geom_hline(yintercept = 10, linetype = 2) +
  geom_vline(xintercept = 0, linetype = 2) +
  geom_vline(xintercept = 10, linetype = 2) +
  annotate(geom = 'text', x = -5, y = 15, hjust = 0,
    label = str_glue("r = {round(cor(six_3.base$V1, six_3.base$V2), 2)}")) +
  annotate(geom = 'text', x = -5, y = 7.5, hjust = 0,
    label = str_glue("Mean = {round(mean(six_3.base$V2), 2)}")) +
  annotate(geom = 'text', x = -5, y = 6.5, hjust = 0,
    label = str_glue("SD = {round(sd(six_3.base$V2), 2)}")) +
  annotate(geom = 'text', x = 6.5, y = -4, hjust = 0,
    label = str_glue("Mean = {round(mean(six_3.base$V1), 2)}")) +
  annotate(geom = 'text', x = 6.5, y = -5, hjust = 0,
    label = str_glue("SD = {round(sd(six_3.base$V1), 2)}")) +
  labs(subtitle = 'Mean = 6.2, SD = 2.2, cor = 0.5') +
  scale_y_continuous(limits = c(-5, 15)) +
  scale_x_continuous(limits = c(-5, 15)) +
  theme(panel.grid = element_blank(),
    plot.subtitle = element_text(size = 16)))
```

Mean = 6.2, SD = 2.2, cor = 0.5



```
# Linear regression
summary(lm(V2 ~ V1, data = six_3.base))

##
## Call:
## lm(formula = V2 ~ V1, data = six_3.base)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -4.4647 -1.1155 -0.1621  1.1568  4.8859
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  2.75246    0.43883   6.272 3.71e-09 ***
## V1           0.49956    0.06661   7.499 5.47e-12 ***
```

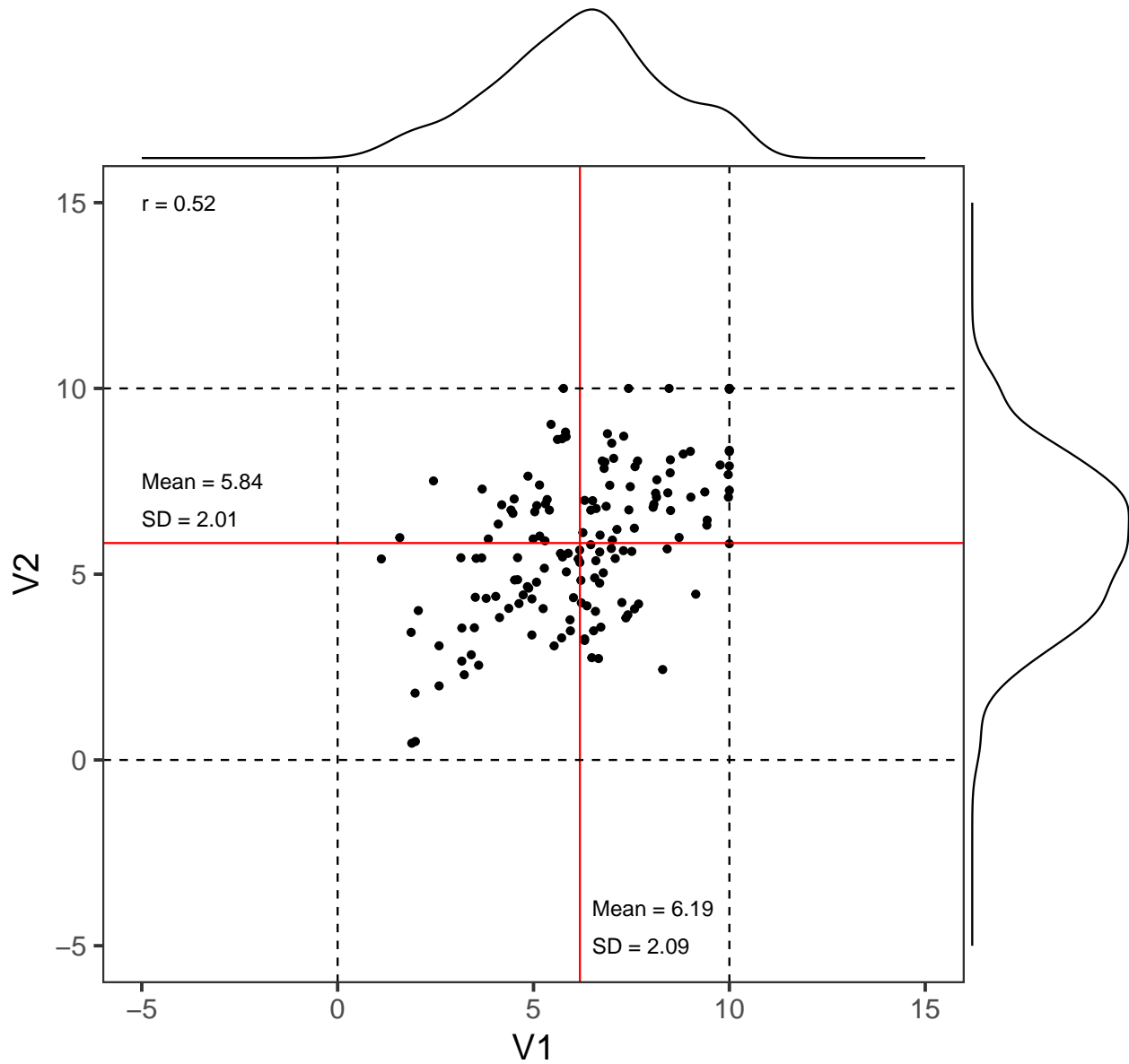
```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.764 on 148 degrees of freedom
## Multiple R-squared:  0.2754, Adjusted R-squared:  0.2705
## F-statistic: 56.24 on 1 and 148 DF,  p-value: 5.468e-12
```

Constrain values to 0-10 range

```
# Process data
six_3 <- six_3.base %>%
  mutate(V1 = case_when(
    V1 < 1 ~ 1,
    V1 > 10 ~ 10,
    TRUE ~ V1)) %>%
  mutate(V2 = case_when(
    V2 < 0 ~ 0,
    V2 > 10 ~ 10,
    TRUE ~ V2)) %>%
  mutate(group = 'No cut-off')

# Plot processed data
ggMarginal(ggplot(data = six_3) +
  aes(x = V1, y = V2) +
  geom_point() +
  geom_hline(yintercept = mean(six_3$V2), colour = 'red') +
  geom_vline(xintercept = mean(six_3$V1), colour = 'red') +
  geom_hline(yintercept = 0, linetype = 2) +
  geom_hline(yintercept = 10, linetype = 2) +
  geom_vline(xintercept = 0, linetype = 2) +
  geom_vline(xintercept = 10, linetype = 2) +
  annotate(geom = 'text', x = -5, y = 15, hjust = 0,
    label = str_glue("r = {round(cor(six_3$V1, six_3$V2), 2)}")) +
  annotate(geom = 'text', x = -5, y = 7.5, hjust = 0,
    label = str_glue("Mean = {round(mean(six_3$V2), 2)}")) +
  annotate(geom = 'text', x = -5, y = 6.5, hjust = 0,
    label = str_glue("SD = {round(sd(six_3$V2), 2)}")) +
  annotate(geom = 'text', x = 6.5, y = -4, hjust = 0,
    label = str_glue("Mean = {round(mean(six_3$V1), 2)}")) +
  annotate(geom = 'text', x = 6.5, y = -5, hjust = 0,
    label = str_glue("SD = {round(sd(six_3$V1), 2)}")) +
  labs(subtitle = 'Mean = 6.2, SD = 2.2 (constrained), cor = 0.5') +
  scale_y_continuous(limits = c(-5, 15)) +
  scale_x_continuous(limits = c(-5, 15)) +
  theme(panel.grid = element_blank(),
    plot.subtitle = element_text(size = 16)))
```

Mean = 6.2, SD = 2.2 (constrained), cor = 0.5



```
# Linear regression
summary(lm(V2 ~ V1, data = six_3))

##
## Call:
## lm(formula = V2 ~ V1, data = six_3)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -4.4607 -1.0781 -0.1572  1.1615  4.3723
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  2.74618    0.44075   6.231 4.58e-09 ***
## V1           0.49984    0.06753   7.402 9.37e-12 ***
```

```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.723 on 148 degrees of freedom
## Multiple R-squared:  0.2702, Adjusted R-squared:  0.2652
## F-statistic: 54.78 on 1 and 148 DF,  p-value: 9.372e-12
```

## Model mean of V1 with increasing V1 cut-offs from 0 to 5

```
# Extract visit 1 data
six_3V1 <- six_3$V1

# Generate a vector of cut-off values to iterate over
cutoff <- 0:5

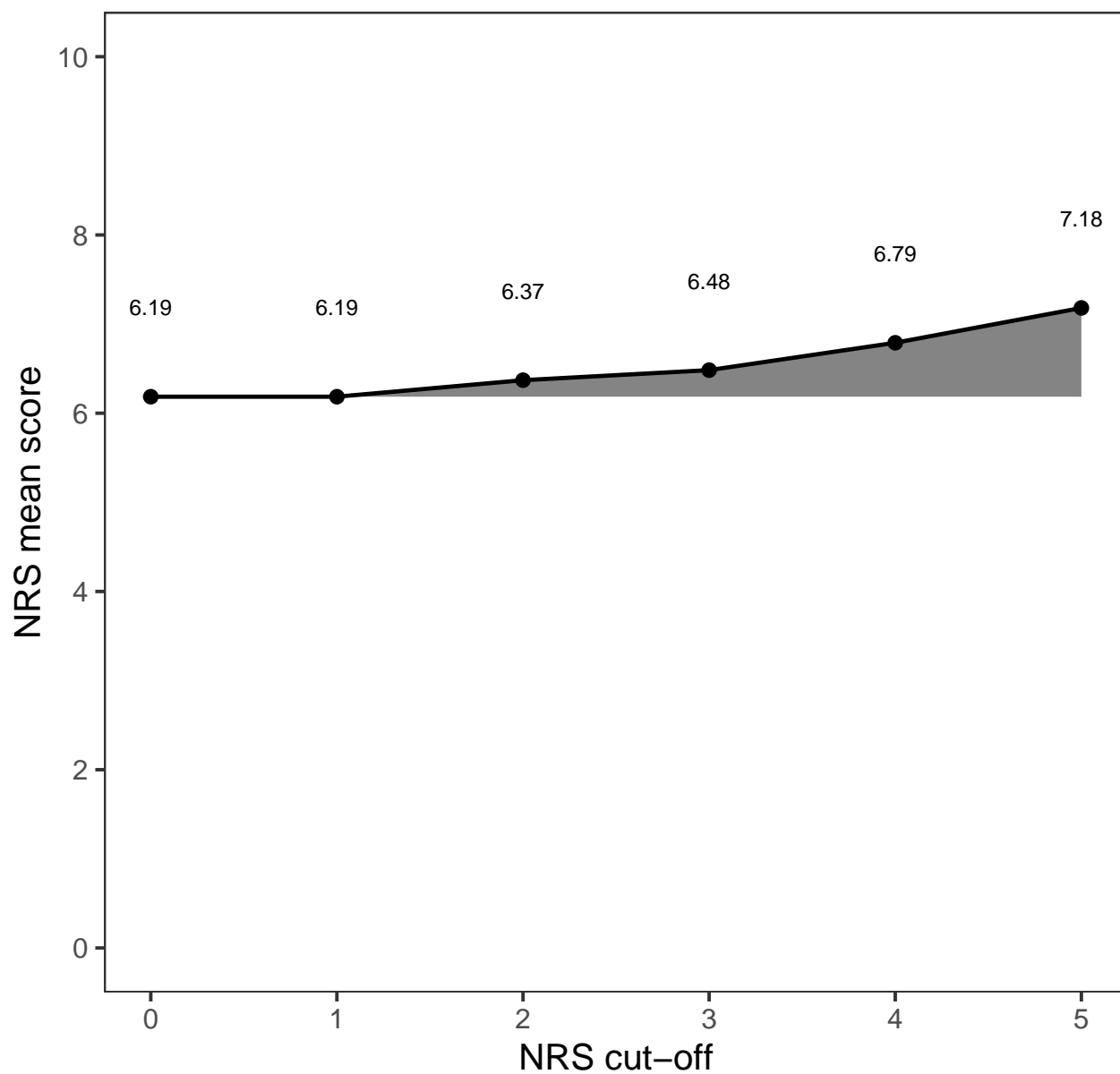
# Generate a vector of V1 means at each V1 cut-off
six_3V1.shift <- sapply(cutoff, function(x){mean(six_3V1[six_3V1 > x])})

# Calculate deviation
(six_3V1.df <- data.frame(cutoff = cutoff,
                          mean = six_3V1.shift) %>%
  mutate(deviation = mean - mean(six_3V1)))

##   cutoff    mean deviation
## 1      0 6.185189 0.0000000
## 2      1 6.185189 0.0000000
## 3      2 6.370438 0.1852496
## 4      3 6.483263 0.2980742
## 5      4 6.790352 0.6051637
## 6      5 7.181210 0.9960212

# Plot data
ggplot(data = six_3V1.df) +
  aes(x = cutoff, y = mean, ymin = mean(six_3V1), ymax = mean) +
  geom_ribbon(alpha = 0.6) +
  geom_point(size = 3) +
  geom_line(size = 1) +
  geom_text(aes(label = round(mean, 2)), nudge_y = 1) +
  scale_y_continuous(limits = c(0, 10),
                     breaks = c(0, 2, 4, 6, 8, 10)) +
  labs(subtitle = 'Shift in V1 mean with increasing V1 NRS cut-off value',
       x = 'NRS cut-off',
       y = 'NRS mean score') +
  theme(panel.grid = element_blank(),
        plot.subtitle = element_text(size = 16))
```

Shift in V1 mean with increasing V1 NRS cut-off value



Model mean of V2 with increasing V1 cut-offs from 0 to 5

```
# Extract visit 2 data
six_3V2 <- six_3$V2

# Generate a vector of cut-off values to iterate over
cutoff <- 0:5

# Generate a vector of V2 means at each V1 cut-off
six_3V2.shift <- map_dbl(.x = cutoff,
  ~ six_3 %>%
    filter(V1 > .x) %>%
    .$V2 %>%
```



```

mean(.))

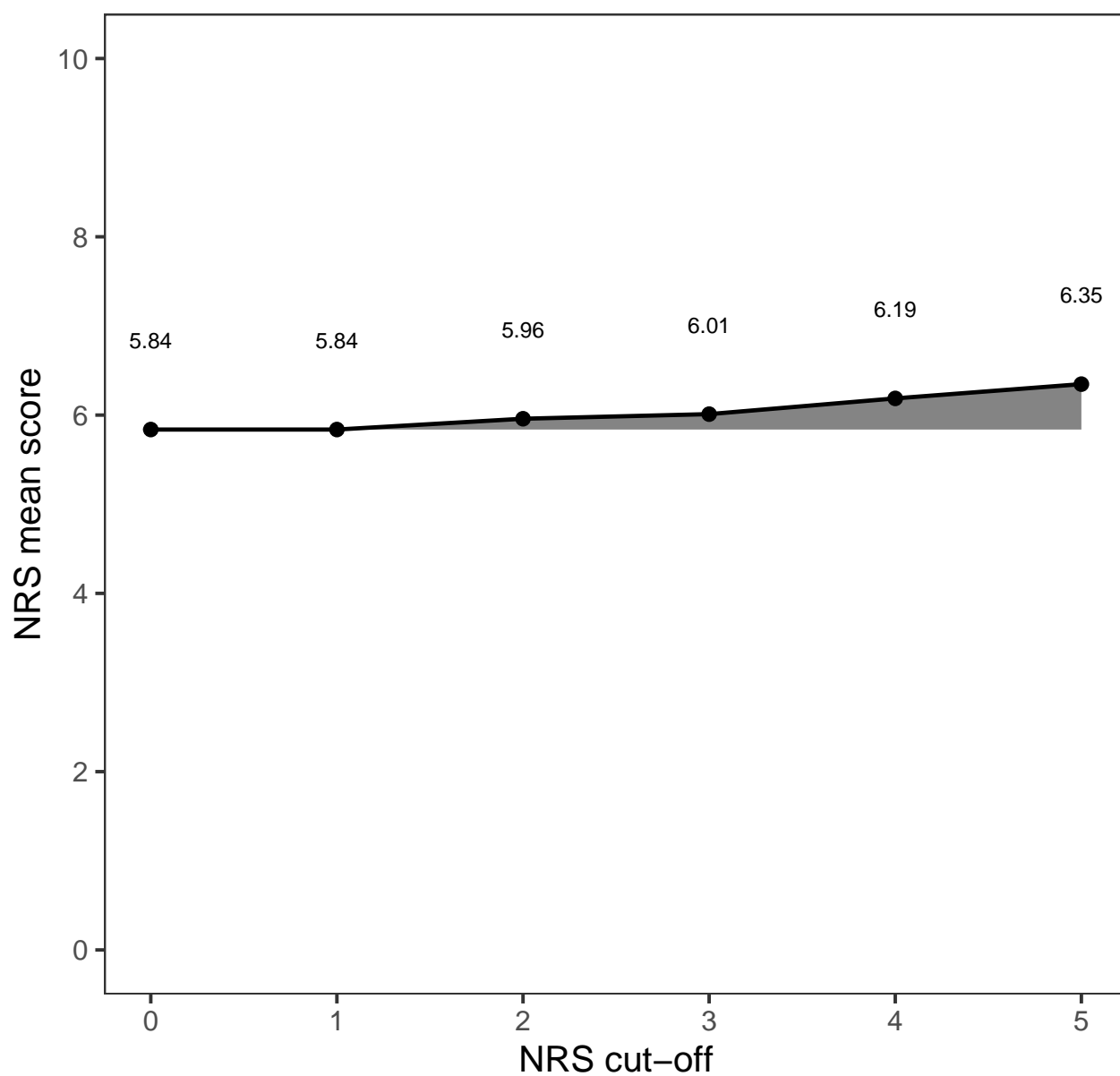
# Calculate deviation
(six_3V2.df <- data.frame(cutoff = cutoff,
                          mean = six_3V2.shift) %>%
  mutate(deviation = mean - mean(six_3V2)))

##   cutoff    mean deviation
## 1      0 5.837755 0.0000000
## 2      1 5.837755 0.0000000
## 3      2 5.958954 0.1211991
## 4      3 6.010669 0.1729136
## 5      4 6.187222 0.3494671
## 6      5 6.347169 0.5094139

# Plot data
ggplot(data = six_3V2.df) +
  aes(x = cutoff, y = mean, ymin = mean(six_3V2), ymax = mean) +
  geom_ribbon(alpha = 0.6) +
  geom_point(size = 3) +
  geom_line(size = 1) +
  geom_text(aes(label = round(mean, 2)), nudge_y = 1) +
  scale_y_continuous(limits = c(0, 10),
                    breaks = c(0, 2, 4, 6, 8, 10)) +
  labs(subtitle = 'Shift in V2 mean with increasing V1 NRS cut-off value',
       x = 'NRS cut-off',
       y = 'NRS mean score') +
  theme(panel.grid = element_blank(),
        plot.subtitle = element_text(size = 16))

```

Shift in V2 mean with increasing V1 NRS cut-off value



## Placebo response

threshold: 0

```
# Process data
placebo_3.0 <- six_3 %>%
  filter(V1 >= 0) %>%
  mutate(difference = V1 - V2) %>%
  mutate(group = 'Cut-off')

# Calculate the mean (95%CI) difference between V1 and V2
groupwiseMean(difference ~ 1,
  data = placebo_3.0,
```

```

R = 5000,
traditional = FALSE,
bca = TRUE) %>%

kable(.)

```

.id	n	Mean	Conf.level	Bca.lower	Bca.upper
NA	150	0.347	0.95	0.0097	0.677

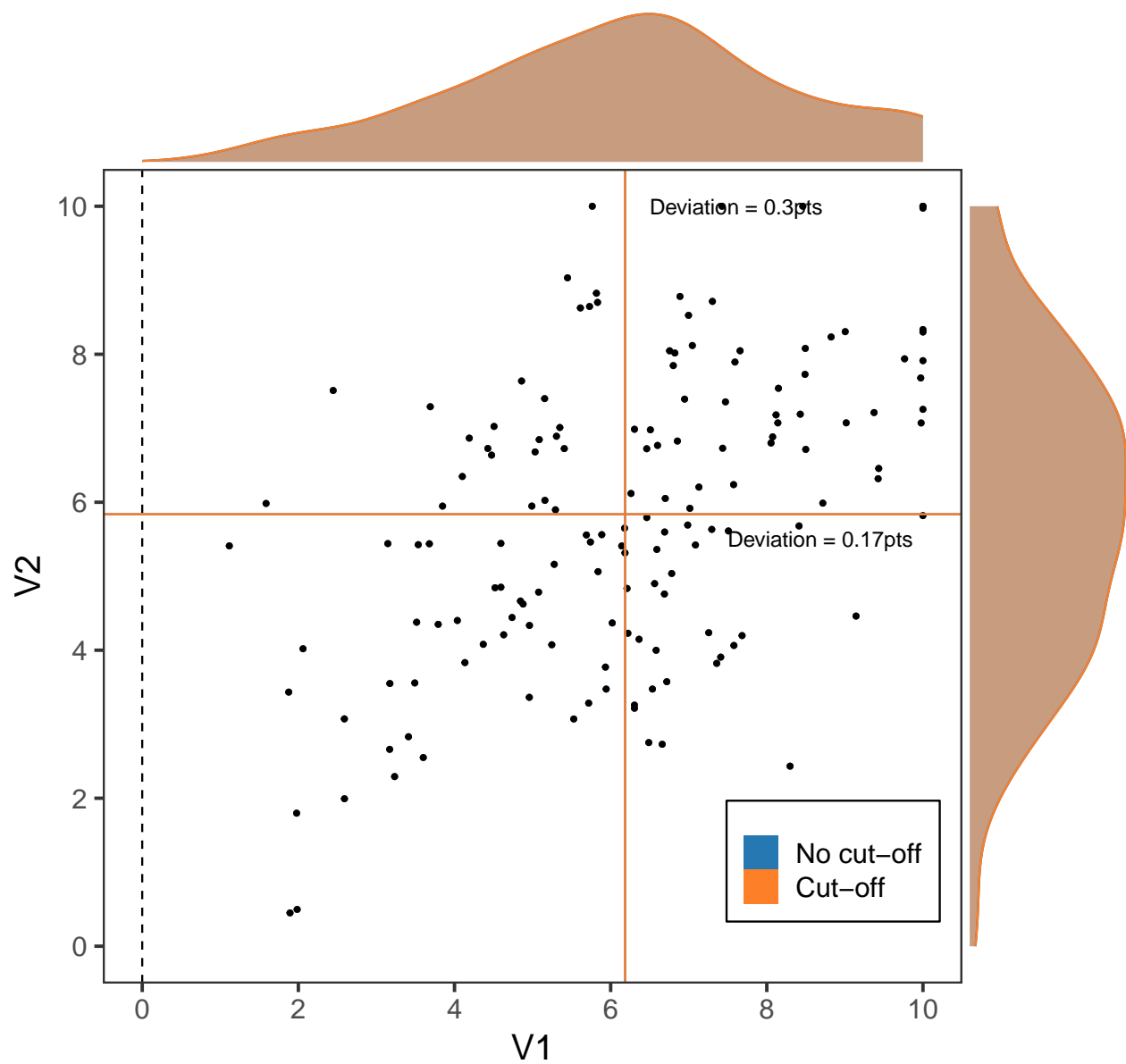
```

# Plot the data
ggMarginal(placebo_3.0[, 1:3] %>%
  bind_rows(six_3) %>%
  mutate(group = factor(group,
                        levels = c('No cut-off', 'Cut-off'),
                        ordered = TRUE)) %>%

  ggplot(data = .) +
  aes(x = V1, y = V2) +
  geom_point(aes(colour = group, fill = group),
            size = 1,
            key_glyph = draw_key_rect) +
  geom_point(data = six_3,
            colour = '#999999',
            size = 1) +
  geom_point(data = placebo_3.0,
            size = 1,
            colour = '#000000') +
  geom_vline(xintercept = mean(six_3$V1),
            colour = pal[1]) +
  geom_vline(xintercept = mean(placebo_3.0$V1),
            colour = pal[2]) +
  geom_vline(xintercept = 0, linetype = 2) +
  geom_hline(yintercept = mean(six_3$V2),
            colour = pal[1]) +
  geom_hline(yintercept = mean(placebo_3.0$V2),
            colour = pal[2]) +
  annotate(geom = 'text', x = 6.5, y = 10, hjust = 0,
            label = str_glue("Deviation = {round(six_3V1.df$deviation[4], 2)}pts")) +
  annotate(geom = 'text', x = 7.5, y = 5.5, hjust = 0,
            label = str_glue("Deviation = {round(six_3V2.df$deviation[4], 2)}pts")) +
  scale_y_continuous(limits = c(0, 10),
                    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_x_continuous(limits = c(0, 10),
                    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_fill_manual(values = pal) +
  scale_colour_manual(values = pal) +
  labs(subtitle = 'Base model: Mean = 6.2, SD = 1.2, cor = 0.5; Cut-off: 0') +
  theme(legend.title = element_blank(),
        legend.position = c(0.85, 0.15),
        legend.background = element_rect(colour = '#000000',
                                          size = 0.5),
        panel.grid = element_blank(),
        plot.subtitle = element_text(size = 16)),
groupColour = TRUE,
groupFill = TRUE)

```

Base model: Mean = 6.2, SD = 1.2, cor = 0.5; Cut-off: 0



threshold: 3

```
# Process data
placebo_3.3 <- six_3 %>%
  filter(V1 >= 3) %>%
  mutate(difference = V1 - V2) %>%
  mutate(group = 'Cut-off')

# Set seed
set.seed(2019)

# Calculate the mean (95%CI) difference between V1 and V2
groupwiseMean(difference ~ 1,
```

```

data = placebo_3.3,
R = 5000,
traditional = FALSE,
bca = TRUE) %>%

kable(.)

```

.id	n	Mean	Conf.level	Bca.lower	Bca.upper
NA	140	0.473	0.95	0.158	0.799

```

# Plot the data
ggMarginal(placebo_3.3[, 1:3] %>%
  bind_rows(six_3) %>%
  mutate(group = factor(group,
                        levels = c('No cut-off', 'Cut-off'),
                        ordered = TRUE)) %>%

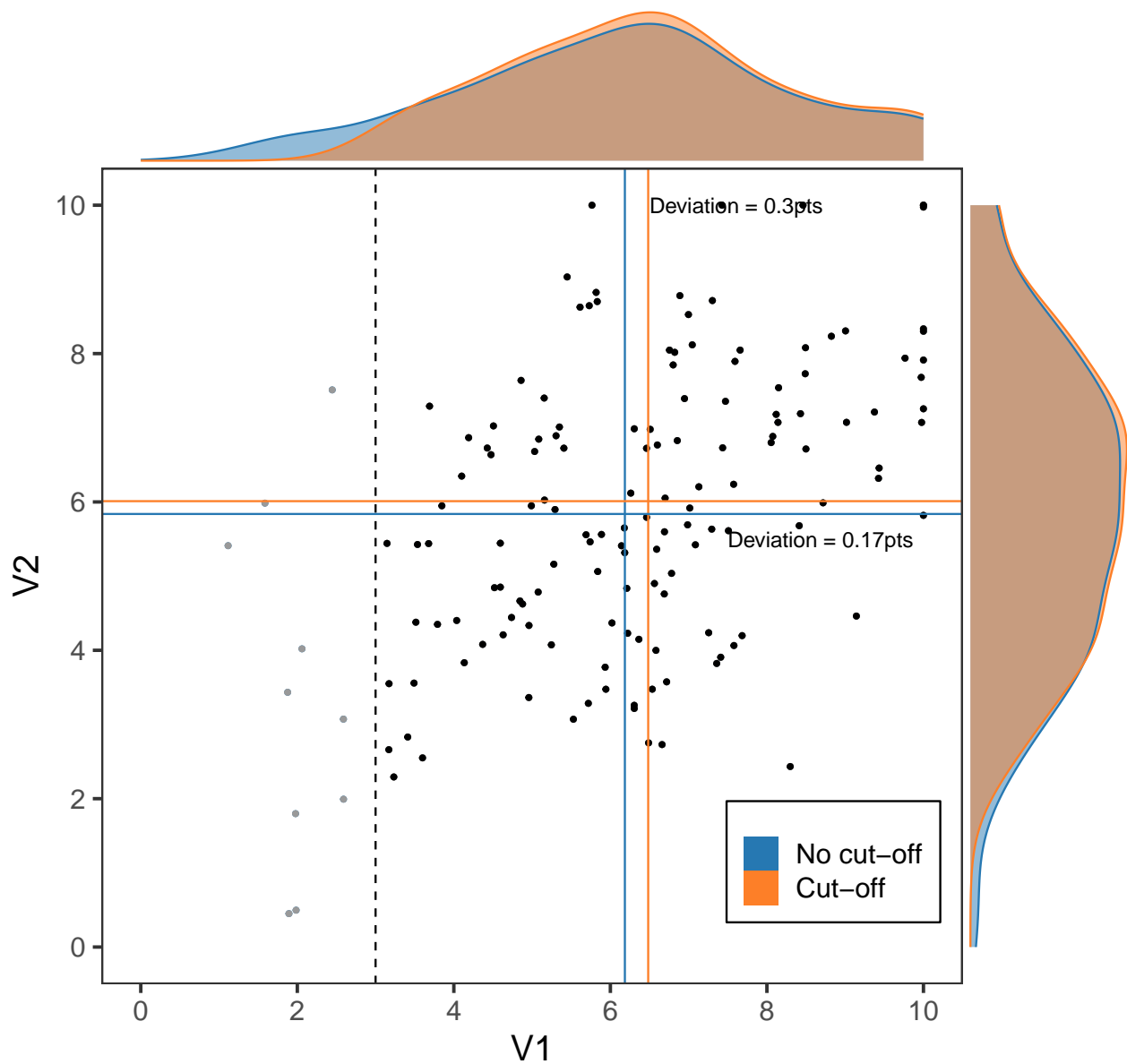
  ggplot(data = .) +
  aes(x = V1, y = V2) +
  geom_point(aes(colour = group, fill = group),
            size = 1,
            key_glyph = draw_key_rect) +
  geom_point(data = six_3,
            colour = '#999999',
            size = 1) +
  geom_point(data = placebo_3.3,
            size = 1,
            colour = '#000000') +
  geom_vline(xintercept = mean(six_3$V1),
            colour = pal[1]) +
  geom_vline(xintercept = mean(placebo_3.3$V1),
            colour = pal[2]) +
  geom_vline(xintercept = 3, linetype = 2) +
  geom_hline(yintercept = mean(six_3$V2),
            colour = pal[1]) +
  geom_hline(yintercept = mean(placebo_3.3$V2),
            colour = pal[2]) +
  annotate(geom = 'text', x = 6.5, y = 10, hjust = 0,
            label = str_glue("Deviation = {round(six_3V1.df$deviation[4], 2)}pts")) +
  annotate(geom = 'text', x = 7.5, y = 5.5, hjust = 0,
            label = str_glue("Deviation = {round(six_3V2.df$deviation[4], 2)}pts")) +
  scale_y_continuous(limits = c(0, 10),
                    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_x_continuous(limits = c(0, 10),
                    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_fill_manual(values = pal) +
  scale_colour_manual(values = pal) +
  labs(subtitle = 'Base model: Mean = 6.2, SD = 2.2, cor = 0.5; Cut-off: 3') +
  theme(legend.title = element_blank(),
        legend.position = c(0.85, 0.15),
        legend.background = element_rect(colour = '#000000',
                                          size = 0.5),

        panel.grid = element_blank(),
        plot.subtitle = element_text(size = 16)),
groupColour = TRUE,

```

```
groupFill = TRUE)
```

Base model: Mean = 6.2, SD = 2.2, cor = 0.5; Cut-off: 3



threshold: 4

```
# Process that data
placebo_3.4 <- six_3 %>%
  filter(V1 >= 4) %>%
  mutate(difference = V1 - V2) %>%
  mutate(group = 'Cut-off')
```

```
# Set seed
set.seed(2019)
```

```
# Calculate the mean (95%CI) difference between V1 and V2
```

```
groupwiseMean(difference ~ 1,
  data = placebo_3.4,
  R = 5000,
  traditional = FALSE,
  bca = TRUE) %>%
```

```
kable(.)
```

.id	n	Mean	Conf.level	Bca.lower	Bca.upper
NA	127	0.603	0.95	0.258	0.945

```
# Plot the data
```

```
ggMarginal(placebo_3.4[, 1:3] %>%
  bind_rows(six_3) %>%
  mutate(group = factor(group,
    levels = c('No cut-off', 'Cut-off'),
    ordered = TRUE)) %>%

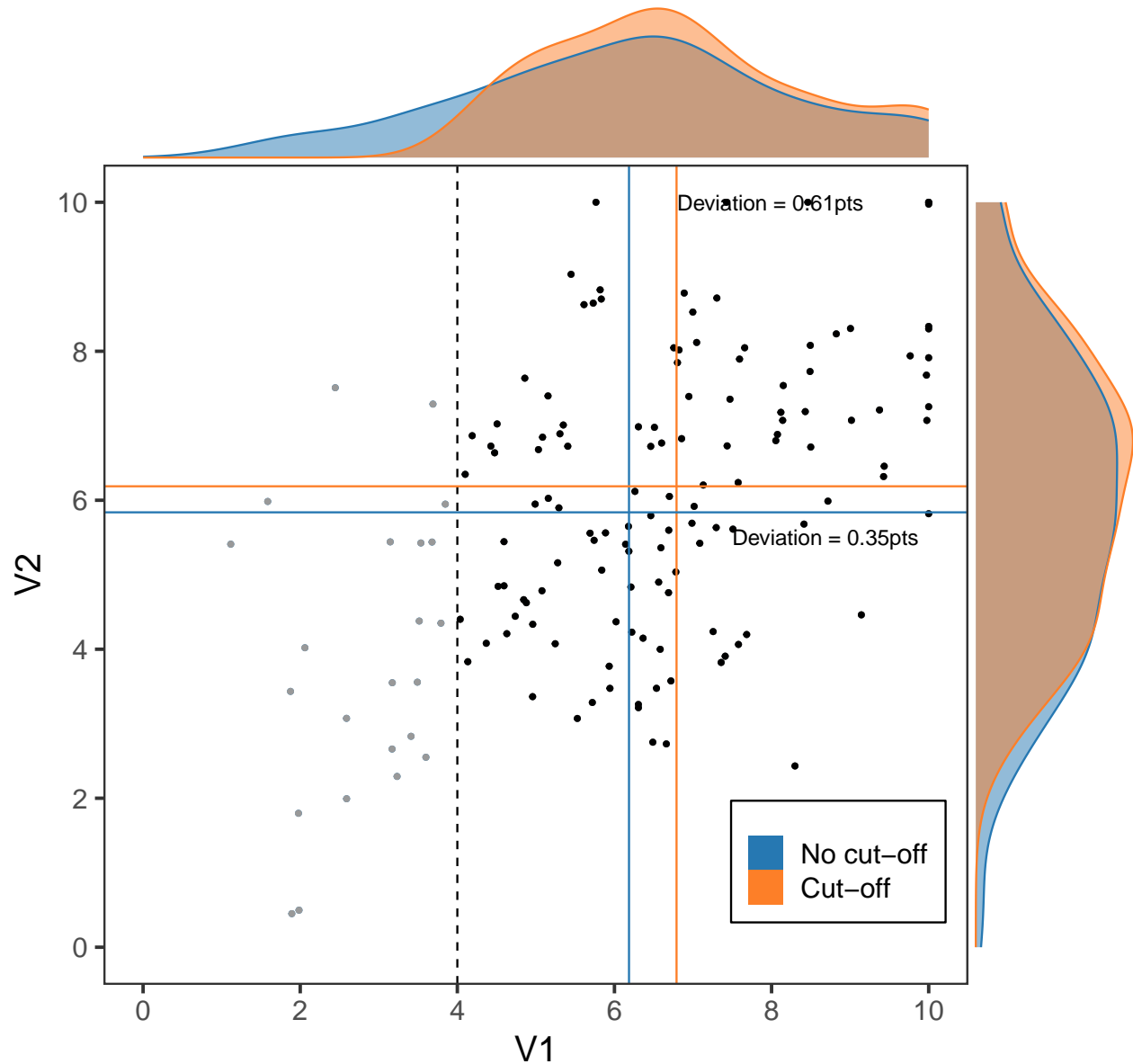
  ggplot(data = .) +
  aes(x = V1, y = V2) +
  geom_point(aes(colour = group, fill = group),
    size = 1,
    key_glyph = draw_key_rect) +
  geom_point(data = six_3,
    colour = '#999999',
    size = 1) +
  geom_point(data = placebo_3.4,
    size = 1,
    colour = '#000000') +
  geom_vline(xintercept = mean(six_3$V1),
    colour = pal[1]) +
  geom_vline(xintercept = mean(placebo_3.4$V1),
    colour = pal[2]) +
  geom_vline(xintercept = 4, linetype = 2) +
  geom_hline(yintercept = mean(six_3$V2),
    colour = pal[1]) +
  geom_hline(yintercept = mean(placebo_3.4$V2),
    colour = pal[2]) +
  annotate(geom = 'text', x = 6.8, y = 10, hjust = 0,
    label = str_glue("Deviation = {round(six_3V1.df$deviation[5], 2)}pts")) +
  annotate(geom = 'text', x = 7.5, y = 5.5, hjust = 0,
    label = str_glue("Deviation = {round(six_3V2.df$deviation[5], 2)}pts")) +
  scale_y_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_x_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_fill_manual(values = pal) +
  scale_colour_manual(values = pal) +
  labs(subtitle = 'Base model: Mean = 6.2, SD = 2.2, cor = 0.5; Cut-off: 4') +
  theme(legend.title = element_blank(),
    legend.position = c(0.85, 0.15),
    legend.background = element_rect(colour = '#000000',
      size = 0.5),
    panel.grid = element_blank(),
```

```

    plot.subtitle = element_text(size = 16)),
    groupColour = TRUE,
    groupFill = TRUE)

```

Base model: Mean = 6.2, SD = 2.2, cor = 0.5; Cut-off: 4



threshold: 5

```

# Process that data
placebo_3.5 <- six_3 %>%
  filter(V1 >= 5) %>%
  mutate(difference = V1 - V2) %>%
  mutate(group = 'Cut-off')

```

```

# Set seed

```



```
set.seed(2019)
```

```
# Calculate the mean (95%CI) difference between V1 and V2
```

```
groupwiseMean(difference ~ 1,
  data = placebo_3.5,
  R = 5000,
  traditional = FALSE,
  bca = TRUE) %>%
```

```
kable(.)
```

.id	n	Mean	Conf.level	Bca.lower	Bca.upper
NA	108	0.834	0.95	0.458	1.18

```
# Plot the data
```

```
ggMarginal(placebo_3.5[, 1:3] %>%
  bind_rows(six_3) %>%
  mutate(group = factor(group,
    levels = c('No cut-off', 'Cut-off'),
    ordered = TRUE)) %>%

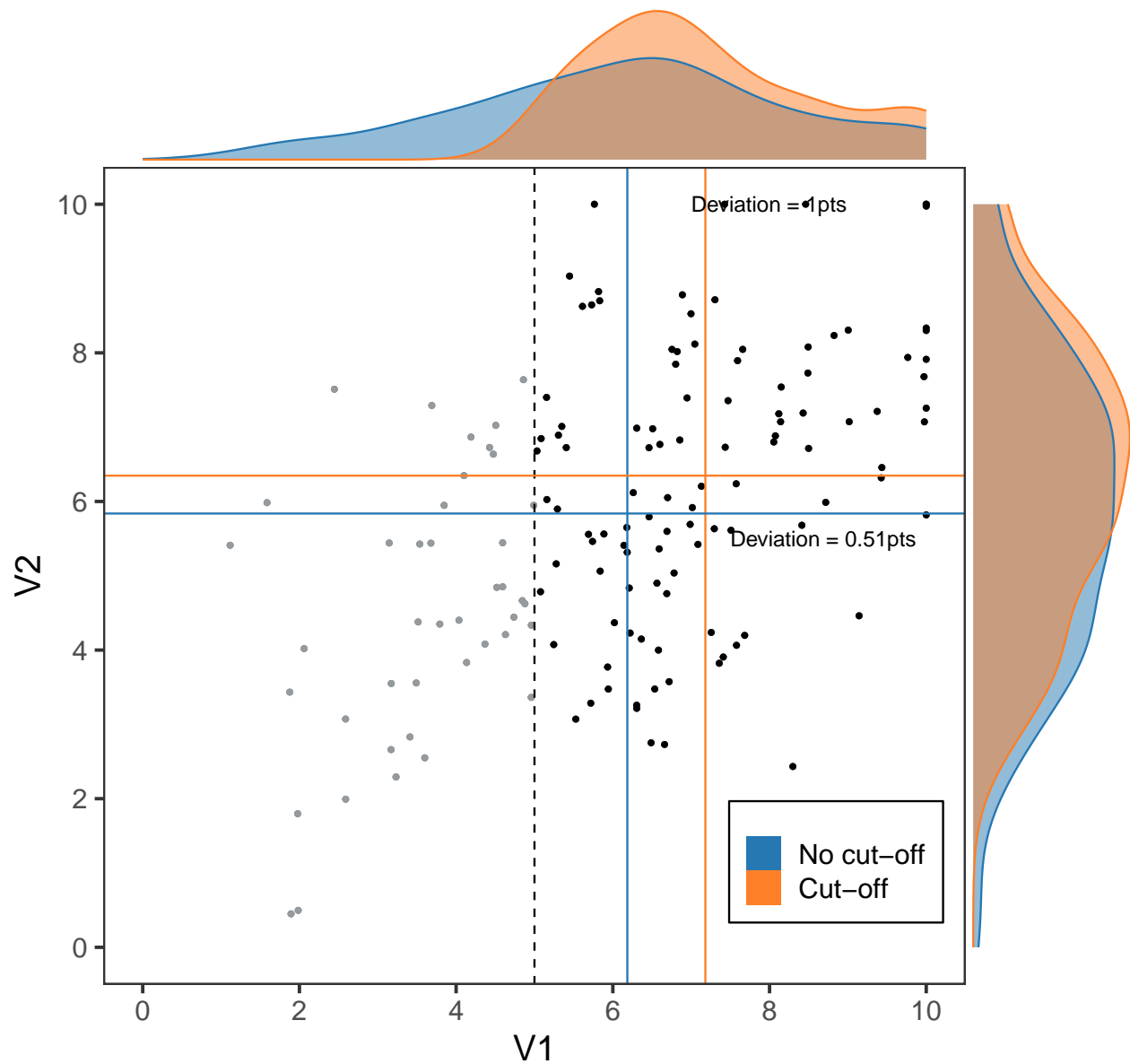
  ggplot(data = .) +
  aes(x = V1, y = V2) +
  geom_point(aes(colour = group, fill = group),
    size = 1,
    key_glyph = draw_key_rect) +
  geom_point(data = six_3,
    colour = '#999999',
    size = 1) +
  geom_point(data = placebo_3.5,
    size = 1,
    colour = '#000000') +
  geom_vline(xintercept = mean(six_3$V1),
    colour = pal[1]) +
  geom_vline(xintercept = mean(placebo_3.5$V1),
    colour = pal[2]) +
  geom_vline(xintercept = 5, linetype = 2) +
  geom_hline(yintercept = mean(six_3$V2),
    colour = pal[1]) +
  geom_hline(yintercept = mean(placebo_3.5$V2),
    colour = pal[2]) +
  annotate(geom = 'text', x = 7, y = 10, hjust = 0,
    label = str_glue("Deviation = {round(six_3V1.df$deviation[6], 2)}pts")) +
  annotate(geom = 'text', x = 7.5, y = 5.5, hjust = 0,
    label = str_glue("Deviation = {round(six_3V2.df$deviation[6], 2)}pts")) +
  scale_y_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_x_continuous(limits = c(0, 10),
    breaks = c(0, 2, 4, 6, 8, 10)) +
  scale_fill_manual(values = pal) +
  scale_colour_manual(values = pal) +
  labs(subtitle = 'Base model: Mean = 6.2, SD = 2.2, cor = 0.5; Cut-off: 5') +
  theme(legend.title = element_blank(),
    legend.position = c(0.85, 0.15),
    legend.background = element_rect(colour = '#000000',
```

```

size = 0.5),
panel.grid = element_blank(),
plot.subtitle = element_text(size = 16)),
groupColour = TRUE,
groupFill = TRUE)

```

Base model: Mean = 6.2, SD = 2.2, cor = 0.5; Cut-off: 5



## Session information

```

sessionInfo()
## R version 3.6.0 (2019-04-26)
## Platform: x86_64-apple-darwin15.6.0 (64-bit)

```

```

## Running under: macOS Mojave 10.14.5
##
## Matrix products: default
## BLAS:   /Library/Frameworks/R.framework/Versions/3.6/Resources/lib/libRblas.0.dylib
## LAPACK: /Library/Frameworks/R.framework/Versions/3.6/Resources/lib/libRlapack.dylib
##
## locale:
## [1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8
##
## attached base packages:
## [1] stats      graphics  grDevices  utils      datasets  methods   base
##
## other attached packages:
## [1] knitr_1.23      MBESS_4.6.0      ggExtra_0.8      rcompanion_2.2.2
## [5] MASS_7.3-51.4   magrittr_1.5     forcats_0.4.0    stringr_1.4.0
## [9] dplyr_0.8.3     purrr_0.3.2     readr_1.3.1      tidyr_0.8.3.9000
## [13] tibble_2.1.3    ggplot2_3.2.0    tidyverse_1.2.1
##
## loaded via a namespace (and not attached):
## [1] nlme_3.1-140      matrixStats_0.54.0 lubridate_1.7.4
## [4] httr_1.4.0        tools_3.6.0        backports_1.1.4
## [7] R6_2.4.0          nortest_1.0-4      lazyeval_0.2.2
## [10] colorspace_1.4-1 withr_2.1.2.9000    tidyselect_0.2.5
## [13] compiler_3.6.0    cli_1.1.0          rvest_0.3.4
## [16] expm_0.999-4      xml2_1.2.1         sandwich_2.5-1
## [19] labeling_0.3      scales_1.0.0       lmtest_0.9-37
## [22] mvtnorm_1.0-11    multcompView_0.1-7 digest_0.6.20
## [25] foreign_0.8-71    rmarkdown_1.14     pkgconfig_2.0.2
## [28] htmltools_0.3.6  manipulate_1.0.1   highr_0.8
## [31] rlang_0.4.0       readxl_1.3.1       rstudioapi_0.10
## [34] shiny_1.3.2       generics_0.0.2     zoo_1.8-6
## [37] jsonlite_1.6      modeltools_0.2-22  Matrix_1.2-17
## [40] Rcpp_1.0.2        DescTools_0.99.28  munsell_0.5.0
## [43] stringi_1.4.3     multcomp_1.4-10    yaml_2.2.0
## [46] plyr_1.8.4        grid_3.6.0         parallel_3.6.0
## [49] promises_1.0.1    crayon_1.3.4       miniUI_0.1.1.1
## [52] lattice_0.20-38   haven_2.1.1        splines_3.6.0
## [55] hms_0.5.0         zeallot_0.1.0      pillar_1.4.2
## [58] EMT_1.1           boot_1.3-23        codetools_0.2-16
## [61] stats4_3.6.0      glue_1.3.1         evaluate_0.14
## [64] modelr_0.1.4      vctrs_0.2.0        httpuv_1.5.1
## [67] cellranger_1.1.0  gtable_0.3.0       assertthat_0.2.1
## [70] xfun_0.8          mime_0.7            coin_1.3-0
## [73] libcoin_1.0-4     xtable_1.8-4       broom_0.5.2
## [76] later_0.8.0       survival_2.44-1.1  TH.data_1.0-10

```