

Challenges

Overview

Initial analysis of survey Q9: “How frequently have you encountered the following challenges while working on open-source projects?”

Import packages and utilities

```
project_root <- here::here() # requires that you be somewhere in the
# project directory (not above it)
# packages
suppressMessages(source(file.path(project_root, "scripts/packages.R")))
# functions and objects used across scripts
suppressMessages(source(file.path(project_root, "scripts/utils.R")))
```

Set seed

```
set.seed(42)
```

Define functions

multiple_plots

- Arguments:
 - `df`: In this script, this will always be the `to_plot` data frame. Must contain (at least) three columns: `challenge`, `challenge_level` (a character column), and `total`.

- `title_codes`: In this script, this will always be the `titles` list. Keys are shorthand codes for each challenge, and values are the full challenge from the survey.
- `challenges_of_interest`: A character vector of the challenges you want to plot.
- Details:
 - A simple function to call my `basic_bar_chart` function (from `scripts/utls.R`) on multiple challenges, producing multiple plots.
- Outputs:
 - Prints `n` plots, where `n` is the number of challenges of interest.

```
multiple_plots <- function(df, title_codes, challenges_of_interest) {
  for (ch in challenges_of_interest) {
    df_ch <- filter(df, challenge == ch)
    plot_title <- title_codes[[ch]]
    p <- basic_bar_chart(
      df_ch,
      x_var = "challenge_level",
      y_var = "total",
      title = plot_title,
      show_grid = TRUE
    )
    print(p)
  }
}
```

Load data

```
challenges <- load_qualtrics_data("clean_data/challenges_Q9.tsv")
```

Wrangle data

Remove empty rows, i.e. rows from respondents who didn't receive this question. As with many questions in this survey, we can cut some corners in the code because the question was mandatory. For example, no need to worry about incomplete answers.

```
nrow(challenges)
```

```
[1] 332
```

```
challenges <- exclude_empty_rows(challenges) # from scripts/utils.R
nrow(challenges)
```

```
[1] 233
```

Let's reshape the data from wide to long format for easier plotting later. We'll also recode the Likert values to integers, so we can get descriptive statistics of the responses. ("Never" = 0, "Non-applicable" = 0, "Rarely" = 1, "Occasionally" = 2, "Frequently" = 3, "Always" = 4)

```
long_data <- challenges %>%
  pivot_longer(
    cols = everything(),
    names_to = "challenge",
    values_to = "challenge_level"
  )

long_data <- long_data %>%
  mutate(
    challenge_score = recode(
      challenge_level,
      "Never" = 0L,
      "Non-applicable" = 0L,
      "Rarely" = 1L,
      "Occasionally" = 2L,
      "Frequently" = 3L,
      "Always" = 4L
    )
  )
# Using interger literals 0L, 1L, etc., ensures that
# the new column will be integers, not doubles.

long_data
```

```
# A tibble: 3,262 x 3
  challenge      challenge_level challenge_score
  <chr>          <chr>             <int>
1 Coding time    Always                4
2 Documentation time Always                4
3 Managing issues Always                4
4 Attracting users Always                4
5 Recognition    Always                4
```

```

6 Hiring                Always                4
7 Security               Always                4
8 Finding peers          Always                4
9 Finding mentors        Always                4
10 Education time        Always                4
# i 3,252 more rows

```

Next, let's calculate some simple descriptive statistics. I will choose:

* The total "score", that is, the total number of "points" a challenge received * The mean (which might be misleading if 0s drag it down, and also, who's to say what a 2.5 really means? Are the distances between the Likert points equal? We assume so, but this is hand-wavy.) * The median * The mode * The standard deviation

```

# Helper to compute the (numeric) mode
get_mode <- function(x) {
  ux <- unique(x)
  ux[which.max(tabulate(match(x, ux)))]
}

summary_df <- long_data %>%
  group_by(challenge) %>%
  summarise(
    total  = sum(challenge_score),
    mean   = mean(challenge_score, na.rm = TRUE),
    median = median(challenge_score),
    mode   = get_mode(challenge_score),
    st_dev = sd(challenge_score, na.rm = TRUE)
  ) %>%
  ungroup()

# Order by highest total "score"
summary_df <- summary_df %>%
  arrange(desc(total))

summary_df

```

```

# A tibble: 14 x 6
  challenge          total  mean median  mode st_dev
  <chr>          <int> <dbl>   <int> <int> <dbl>
1 Documentation time    686  2.94     3     3  1.08
2 Coding time           606  2.60     3     3  1.24
3 Education time        539  2.31     2     3  1.26

```

4	Managing issues	451	1.94	2	2	1.29
5	Attracting users	442	1.90	2	0	1.45
6	Securing funding	438	1.88	2	0	1.74
7	Finding funding	432	1.85	2	0	1.68
8	Educational resources	369	1.58	2	1	1.19
9	Recognition	334	1.43	1	0	1.35
10	Legal	333	1.43	1	0	1.24
11	Finding mentors	323	1.39	1	0	1.31
12	Security	307	1.32	1	0	1.31
13	Hiring	291	1.25	0	0	1.53
14	Finding peers	267	1.15	1	0	1.13

Cool! It looks like finding the time for documentation, coding, and self-education are the challenges encountered most frequently. These are the only responses that had a mode of 3 (“Frequently”) and a mean of **greater** than 2 (“Occasionally”).

```
# Prettify
summary_df_to_print <- summary_df %>%
  rename(
    `Challenge` = `challenge`,
    `Total` = `total`,
    `Mean` = `mean`,
    `Median` = `median`,
    `Mode` = `mode`,
    `Standard Deviation` = `st_dev`
  )

write_df_to_file(summary_df_to_print, "supplementary_tables/challenges.tsv")
```

Out of curiosity, how does it look when we order by variability?

```
sd_df <- summary_df %>%
  arrange(desc(st_dev))

sd_df
```

```
# A tibble: 14 x 6
  challenge      total  mean median  mode st_dev
  <chr>      <int> <dbl>  <int> <int> <dbl>
1 Securing funding    438  1.88     2     0  1.74
2 Finding funding    432  1.85     2     0  1.68
```

3	Hiring	291	1.25	0	0	1.53
4	Attracting users	442	1.90	2	0	1.45
5	Recognition	334	1.43	1	0	1.35
6	Security	307	1.32	1	0	1.31
7	Finding mentors	323	1.39	1	0	1.31
8	Managing issues	451	1.94	2	2	1.29
9	Education time	539	2.31	2	3	1.26
10	Legal	333	1.43	1	0	1.24
11	Coding time	606	2.60	3	3	1.24
12	Educational resources	369	1.58	2	1	1.19
13	Finding peers	267	1.15	1	0	1.13
14	Documentation time	686	2.94	3	3	1.08

Fascinating! The greatest standard deviations are from securing funding, finding funding, and hiring. This makes sense, as these are, at least in my perception, “manager tasks”—tasks that only some people face, but they’re likely to be a big challenge for those who face them. I would guess that these might show a bimodal distribution. Let’s plot them and find out!

Plot the distributions

Prepare data for plotting

```
ordered_levels <- c(
  "Non-applicable",
  "Never",
  "Rarely",
  "Occasionally",
  "Frequently",
  "Always"
)

to_plot <- long_data %>%
  mutate(
    challenge_level = factor(
      challenge_level,
      levels = ordered_levels
    )
  ) %>%
  count( # Count number of occurrences,
    #i.e. number of people who selected that challenge level
    challenge,
```

```

    challenge_level,
    name = "total"
  ) %>%
  ungroup()

```

```
to_plot
```

```

# A tibble: 84 x 3
  challenge challenge_level total
  <chr>      <fct>      <int>
1 Attracting users Non-applicable 50
2 Attracting users Never 15
3 Attracting users Rarely 24
4 Attracting users Occasionally 53
5 Attracting users Frequently 52
6 Attracting users Always 39
7 Coding time Non-applicable 21
8 Coding time Never 4
9 Coding time Rarely 13
10 Coding time Occasionally 54
# i 74 more rows

```

Create a plot for each “challenge”. After inspecting the plots, I attempted to order them into groups based on the shape of their distribution. These are the shapes I observed (this is extremely subjective):

- * Right-skewed: Documentation time, coding time, education time * Interpretation: Common tasks that are frequently challenging
- * Highly bimodal: Securing funding, identifying funding, hiring * Interpretation: Tasks that are not as common, but they are frequently challenging for the people tasked with them.
- * Normal: Educational resources, Legal * Interpretation: Moderately common tasks that are challenging with moderate frequency.
- * NA-skewed but otherwise normal: Attracting users, Receiving recognition, finding mentors, managing security risks, managing issues * Interpretation: Less-common tasks that are challenging with moderate frequency.
- * Left-skewed: Finding peers * Interpretation: Moderately common tasks that are infrequently challenging.

```

titles <- list(
  "Coding time" = "Limited time for writing new code",
  "Documentation time" = "Limited time for writing documentation",
  "Managing issues" = "Managing issues and pull requests",
  "Attracting users" = "Attracting users and/or contributors",
  "Recognition" = "Receiving recognition for my contributions",

```

```

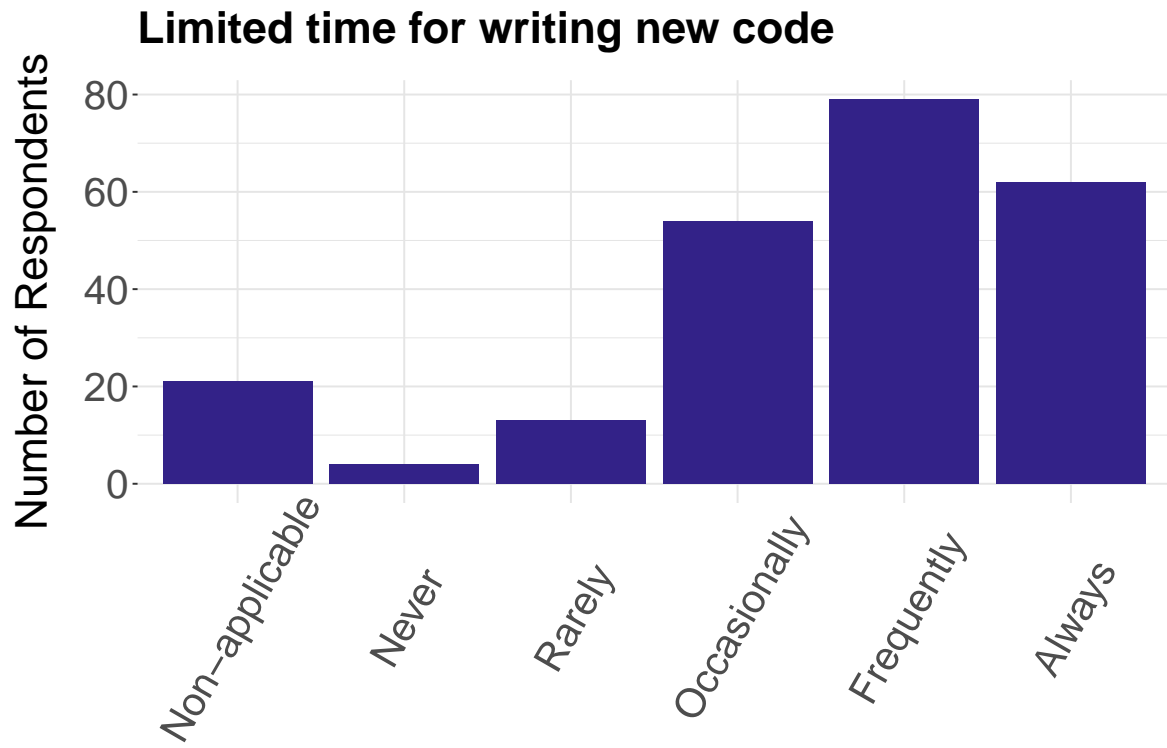
"Hiring" = "Finding and hiring qualified personnel",
"Security" = "Managing security risks",
"Finding peers" = "Finding a community of peers who share my interests",
"Finding mentors" = "Finding mentors",
"Education time" = "Finding time to educate myself",
"Educational resources" = "Identifying helpful educational resources",
"Legal" = "Navigating licensing and other legal issues",
"Finding funding" = "Identifying potential funding sources\nfor my open source projects"
"Securing funding" = "Securing funding for my open source projects"
)

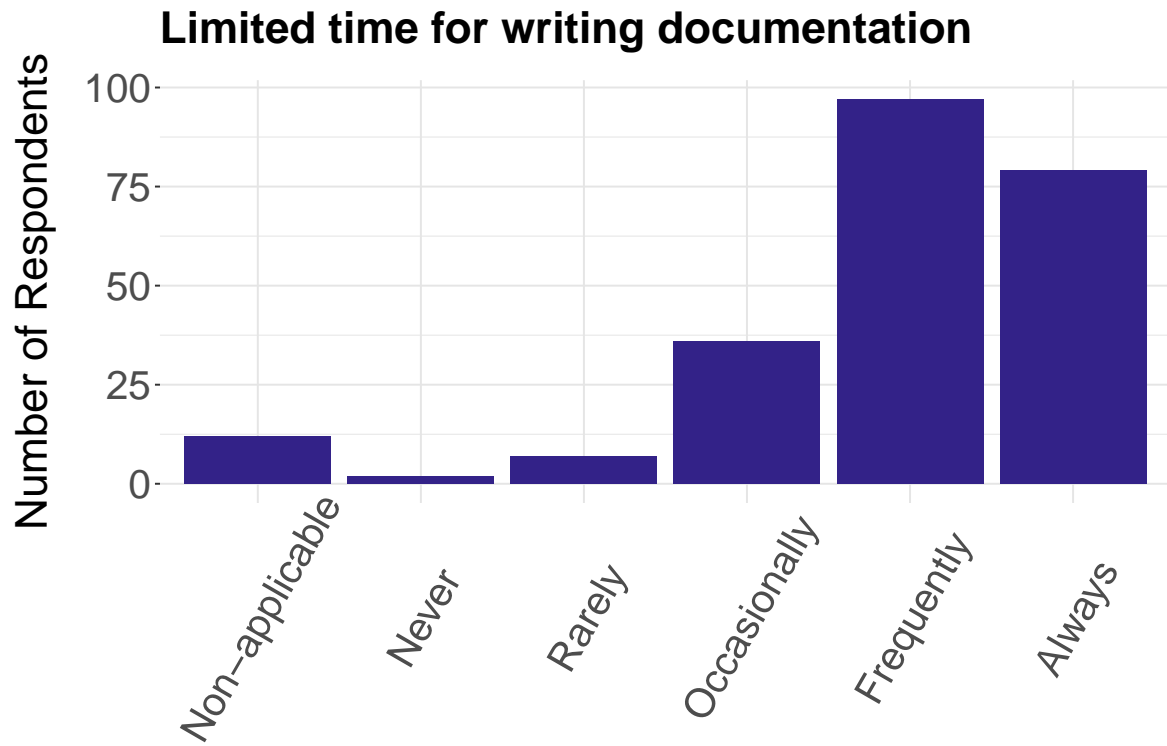
right_skewed <- c(
  "Coding time",
  "Documentation time",
  "Education time"
)
bimodal <- c(
  "Finding funding",
  "Securing funding",
  "Hiring"
)
normal <- c(
  "Educational resources",
  "Legal"
)
na_skewed <- c(
  "Managing issues",
  "Attracting users",
  "Recognition",
  "Security",
  "Finding mentors"
)
left_skewed <- c(
  "Finding peers"
)

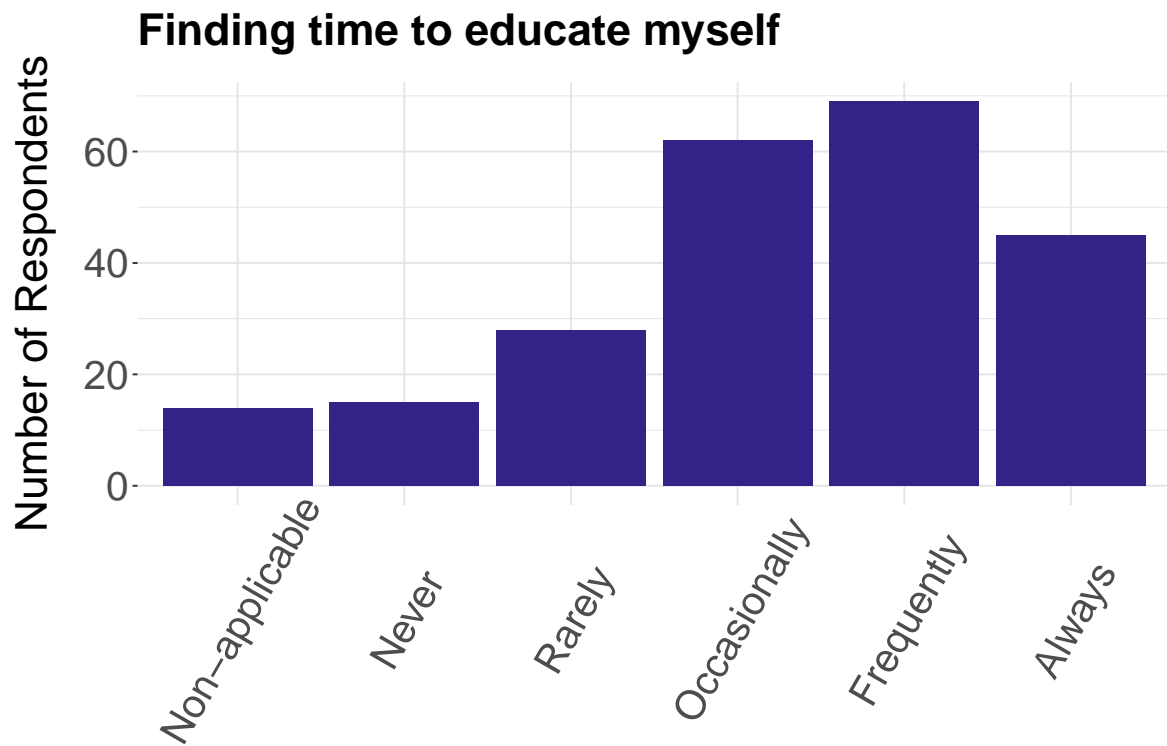
```

“right-skewed”

```
multiple_plots(to_plot, titles, right_skewed)
```

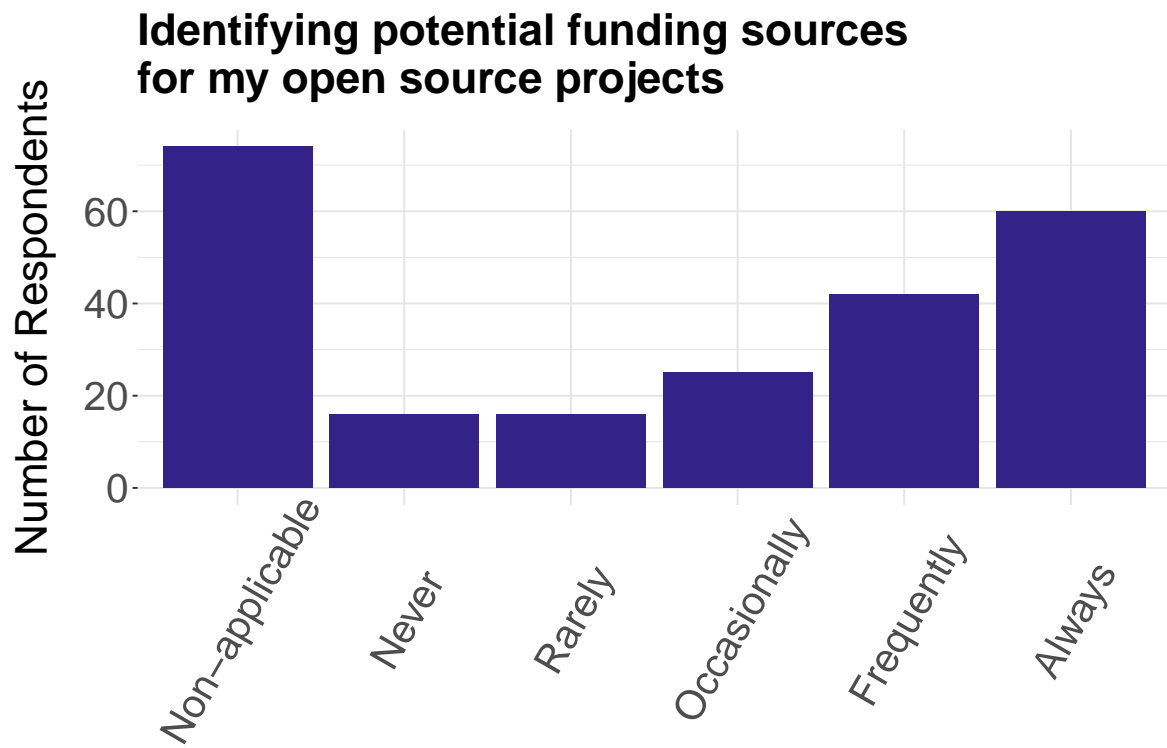



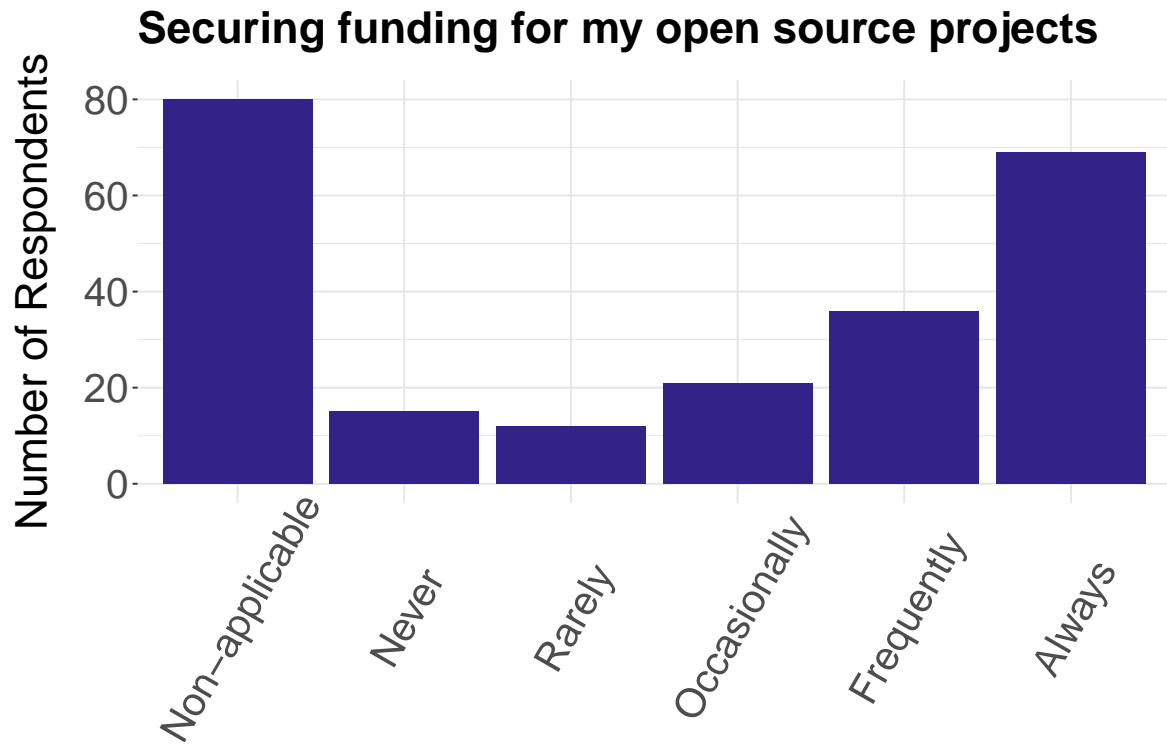


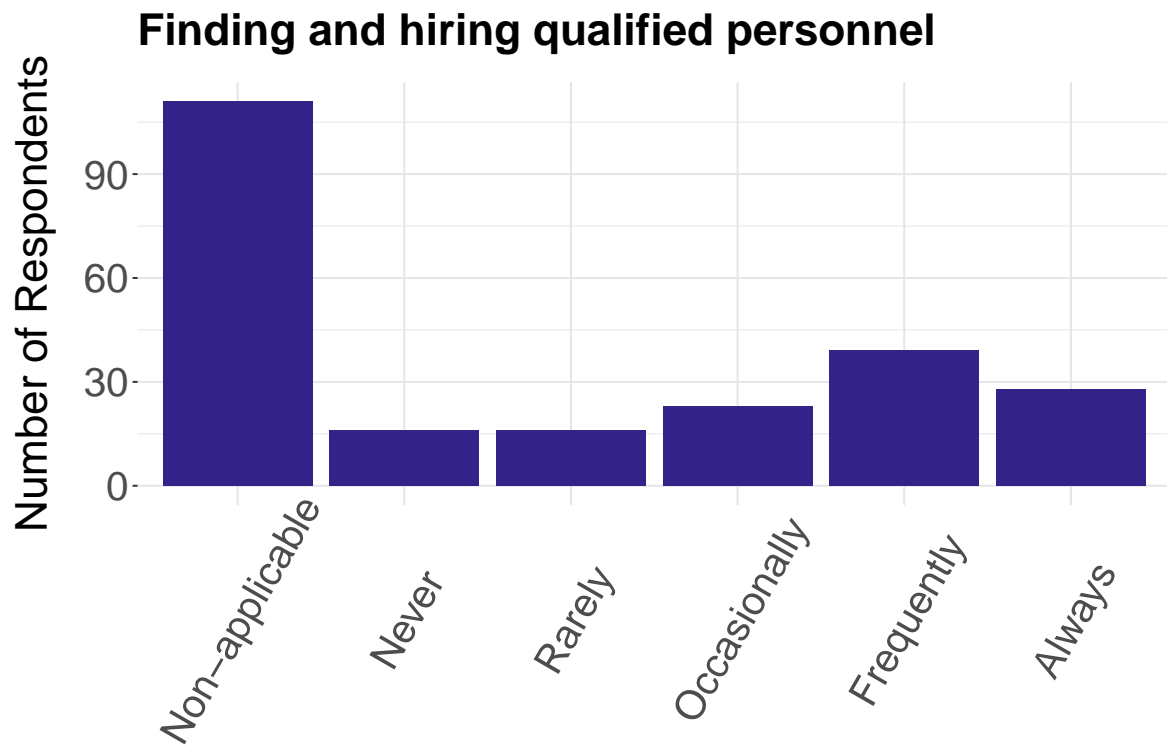


“highly bimodal”

```
multiple_plots(to_plot, titles, bimodal)
```

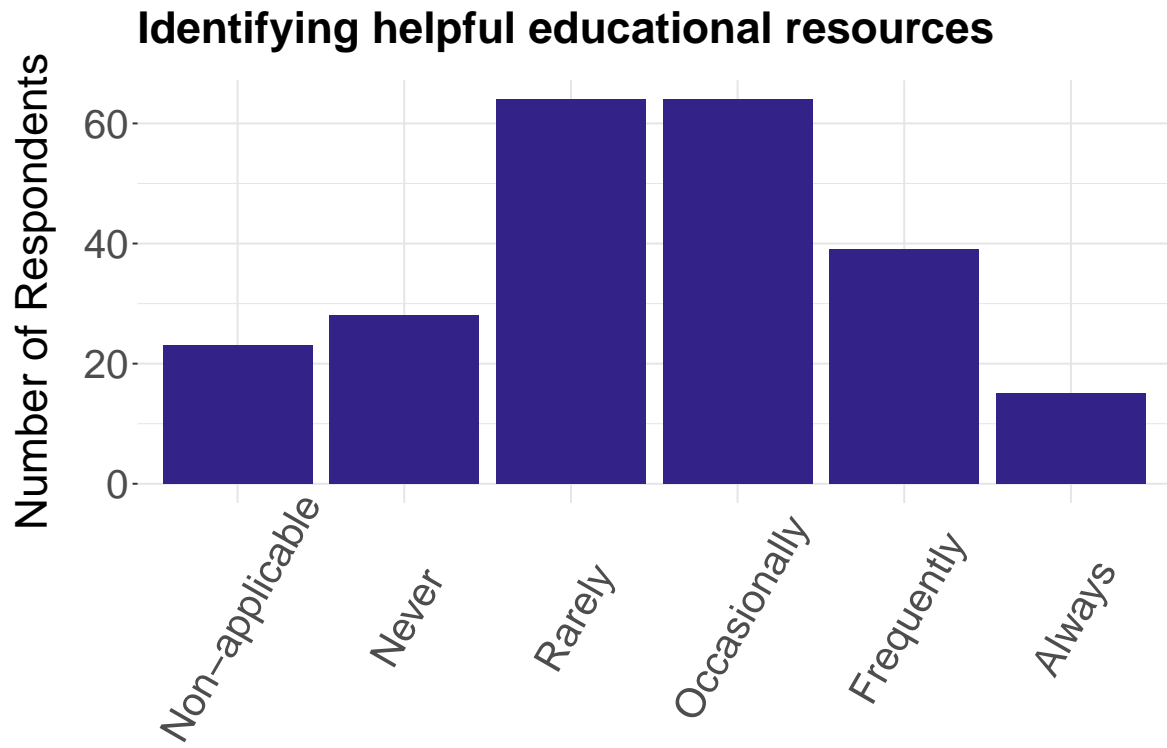


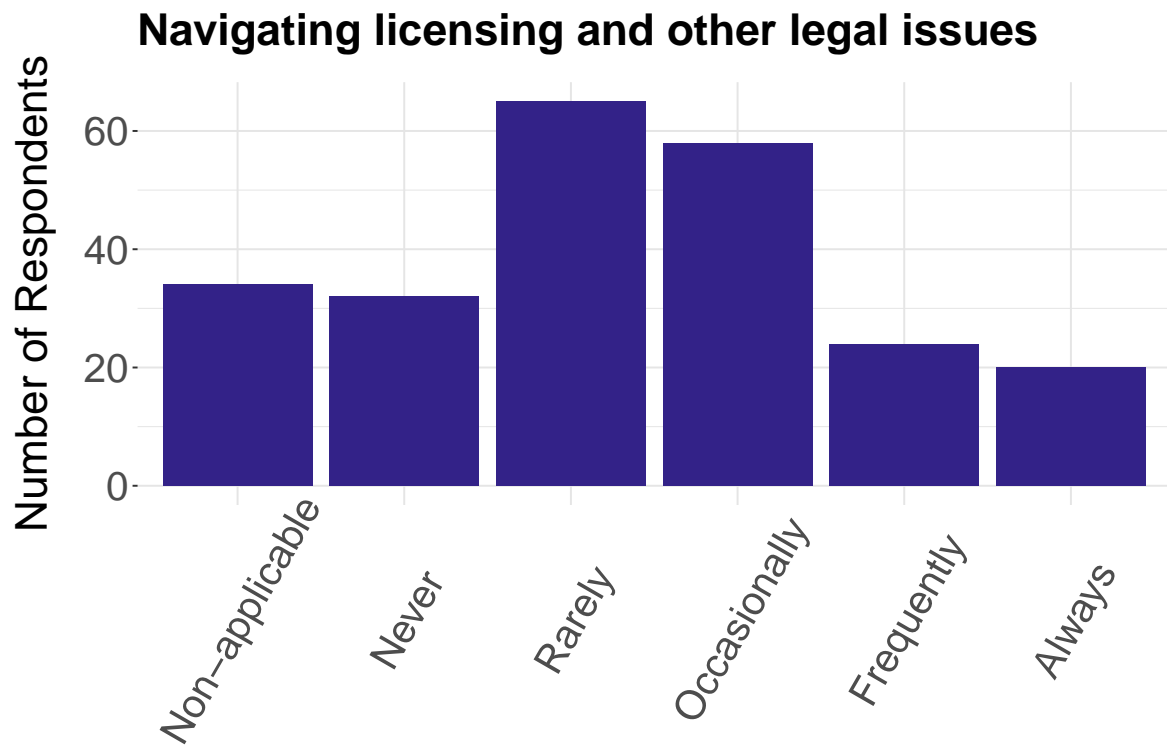




“normal”

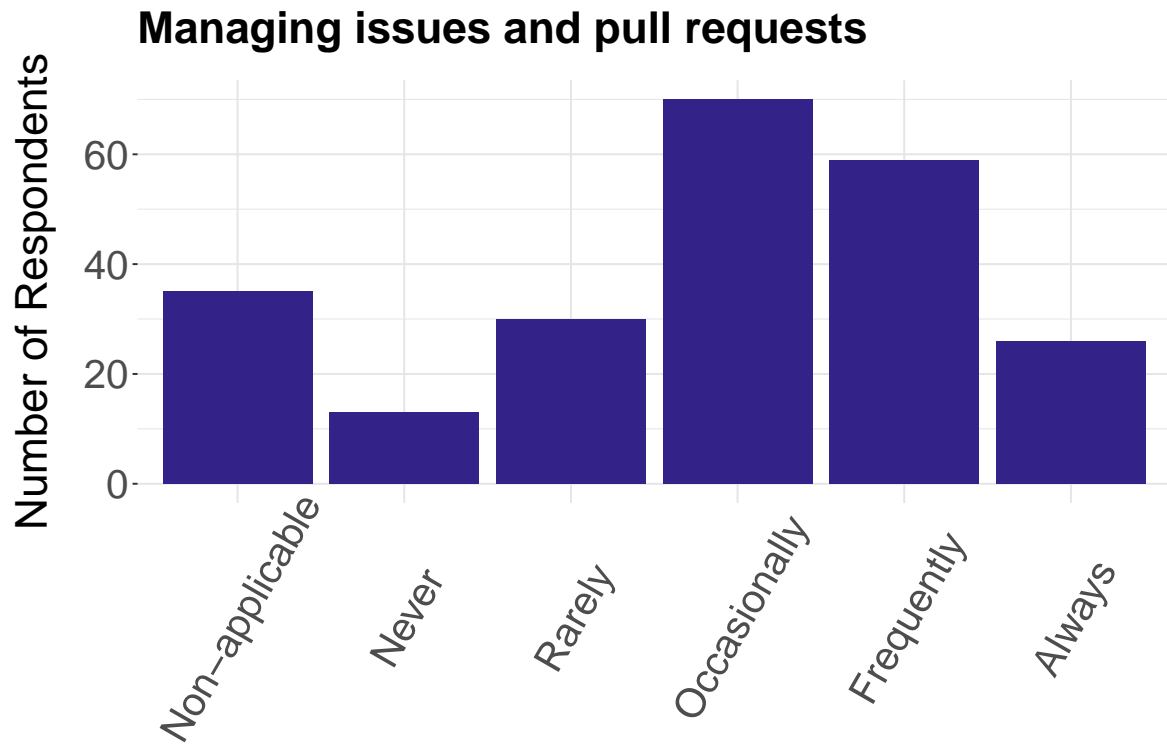
```
multiple_plots(to_plot, titles, normal)
```

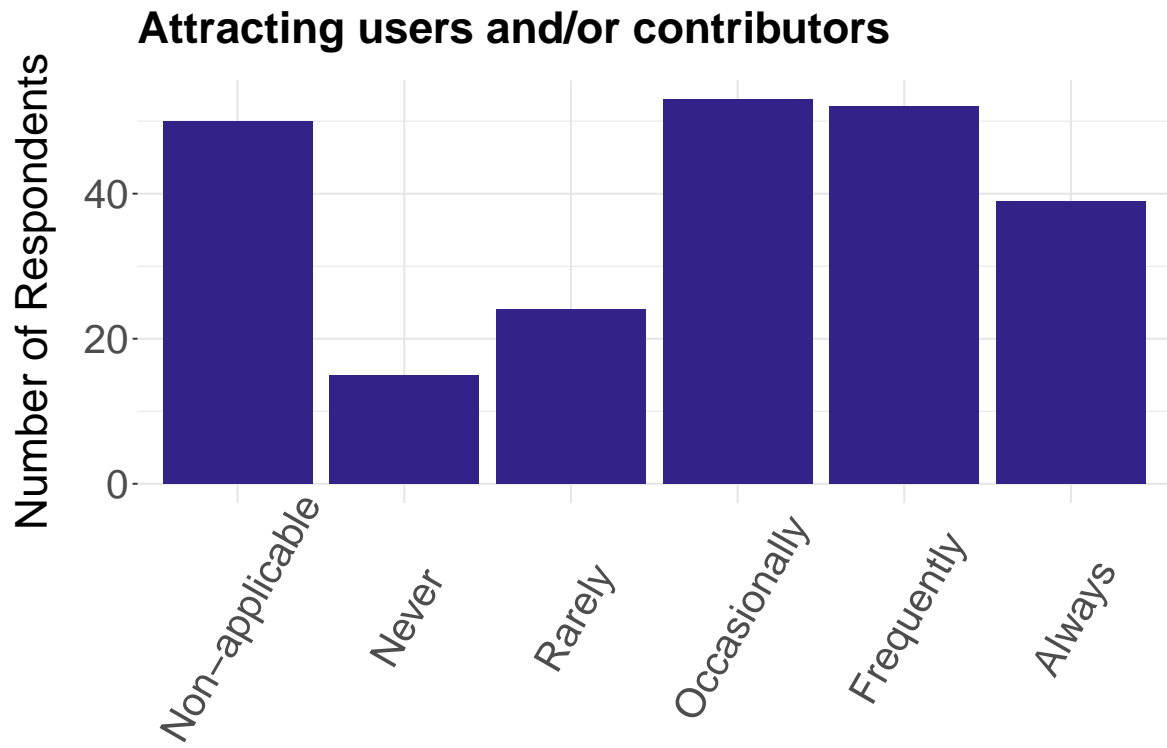


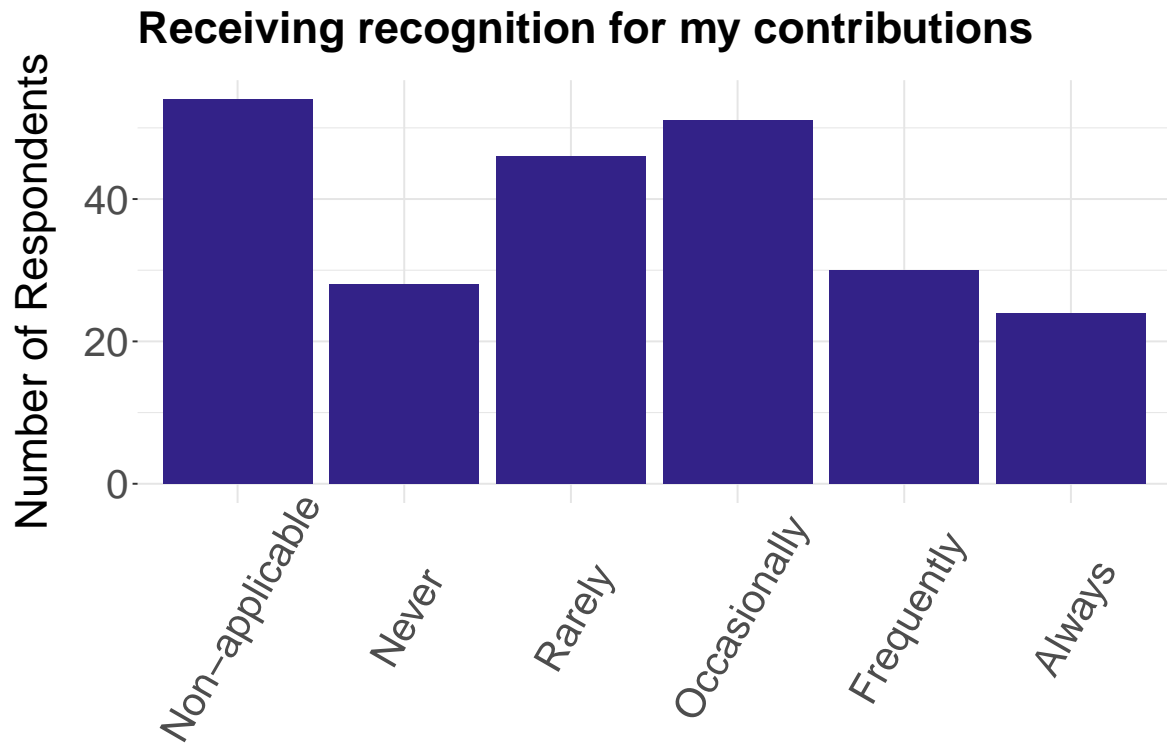


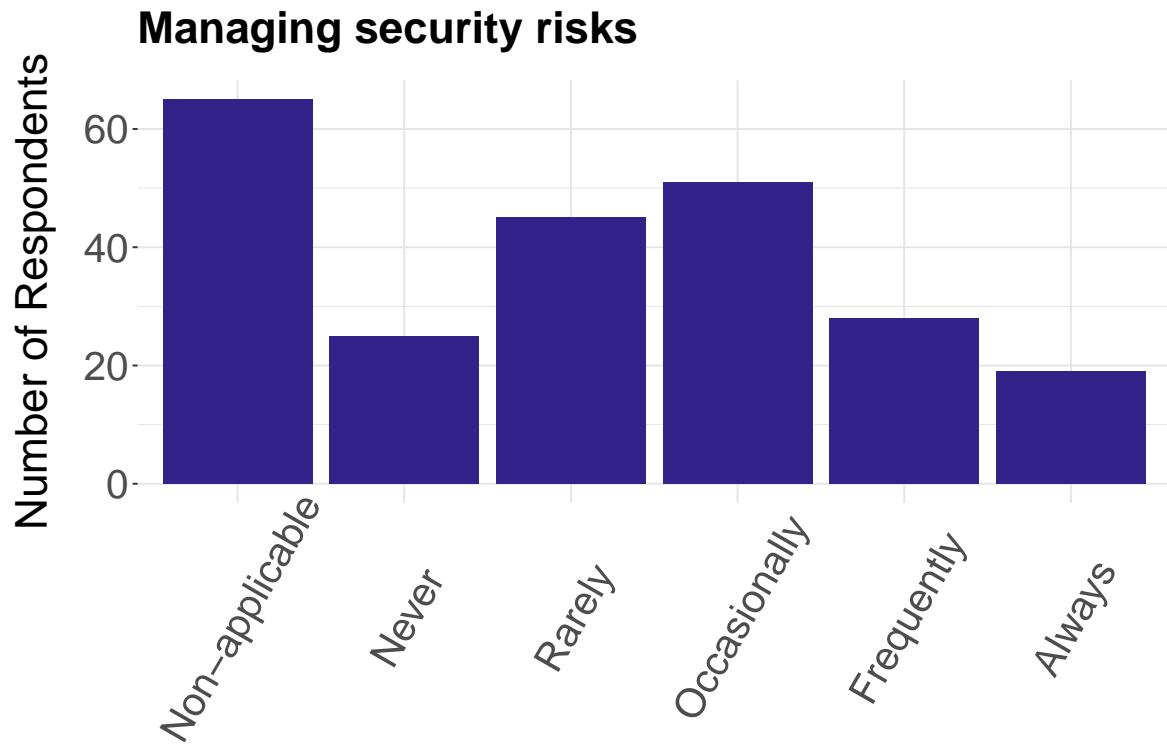
“na-skewed”

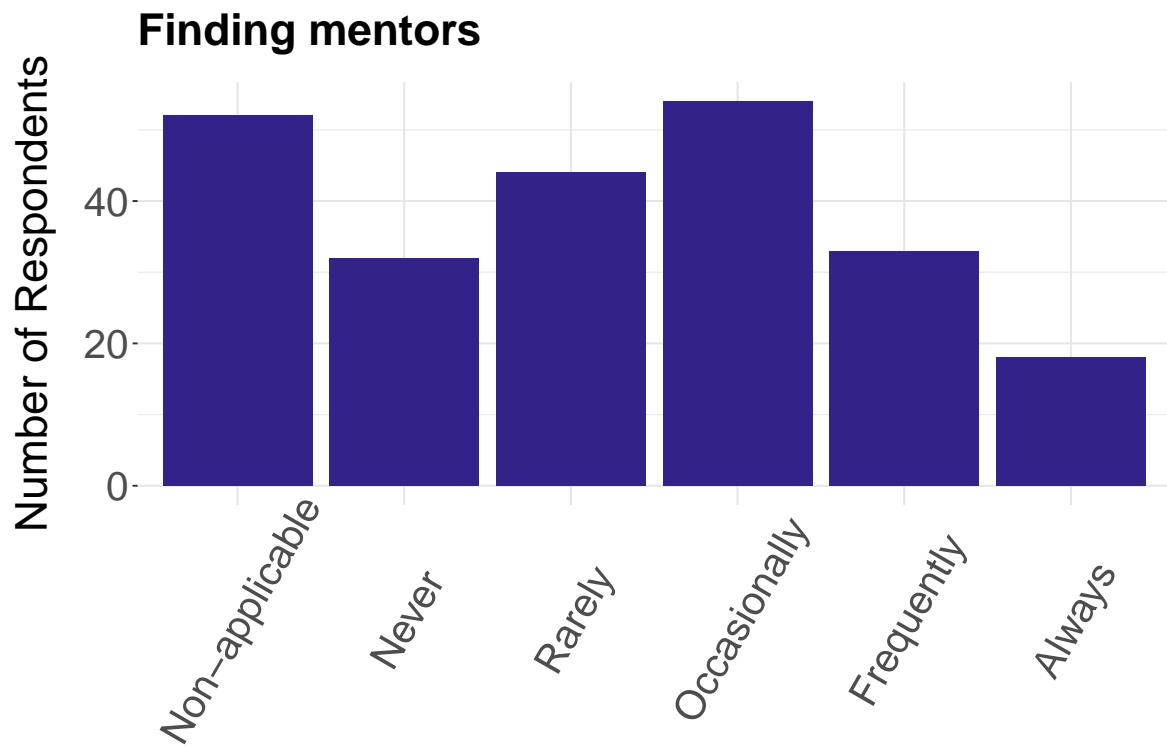
```
multiple_plots(to_plot, titles, na_skewed)
```



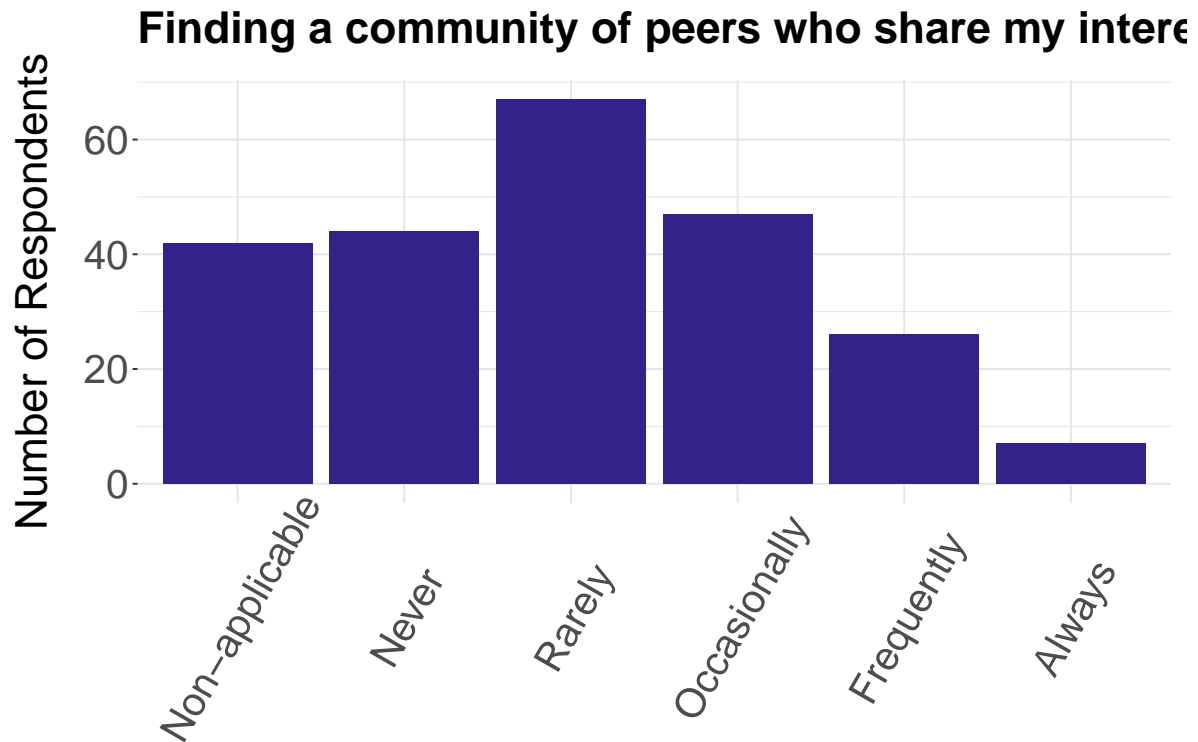






“left-skewed”

```
multiple_plots(to_plot, titles, left_skewed)
```



K-means clustering of distributions

This seems like an interesting line of inquiry. Let's make it a little more rigorous by clustering the challenges based on the response rates (actually, the absolute response numbers).

Wrangle data

```
wide_counts <- to_plot %>%  
  pivot_wider(  
    names_from = challenge_level,  
    values_from = total,  
    values_fill = 0  
  )  
  
wide_counts <- data.frame(wide_counts)  
# Turn this categorical column into row names
```

```
rownames(wide_counts) <- wide_counts$challenge
wide_counts <- wide_counts[,2:(ncol(wide_counts))]
head(wide_counts)
```

	Non.applicable	Never	Rarely	Occasionally	Frequently
Attracting users	50	15	24	53	52
Coding time	21	4	13	54	79
Documentation time	12	2	7	36	97
Education time	14	15	28	62	69
Educational resources	23	28	64	64	39
Finding funding	74	16	16	25	42

	Always
Attracting users	39
Coding time	62
Documentation time	79
Education time	45
Educational resources	15
Finding funding	60

```
# Scaling probably isn't necessary?
# We have the same number of responses throughout,
# so the units for each challenge are the same
# (number of responses).
scaled <- scale(wide_counts)
scaled
```

	Non.applicable	Never	Rarely	Occasionally
Attracting users	0.08407151	-0.4615675	-0.4780961	0.33347644
Coding time	-0.95026278	-1.4093196	-1.0002719	0.39743082
Documentation time	-1.27126308	-1.5816381	-1.2850951	-0.75374811
Education time	-1.19992968	-0.4615675	-0.2882139	0.90906590
Educational resources	-0.87892938	0.6585030	1.4207252	1.03697467
Finding funding	0.94007229	-0.3754083	-0.8578603	-1.45724635
Finding mentors	0.15540491	1.0031401	0.4713146	0.39743082
Finding peers	-0.20126209	2.0370514	1.5631368	-0.05024987
Hiring	2.25974018	-0.3754083	-0.8578603	-1.58515512
Legal	-0.48659569	1.0031401	1.4681957	0.65324836
Managing issues	-0.45092899	-0.6338861	-0.1932729	1.42070098
Recognition	0.22673830	0.6585030	0.5662556	0.20556767
Securing funding	1.15407249	-0.4615675	-1.0477424	-1.71306389
Security	0.61907200	0.4000252	0.5187851	0.20556767

	Frequently	Always
Attracting users	0.2440265	0.1098315
Coding time	1.4739199	1.1202809
Documentation time	2.2938488	1.8671349
Education time	1.0184038	0.3734270
Educational resources	-0.3481444	-0.9445506
Finding funding	-0.2114896	1.0324157
Finding mentors	-0.6214541	-0.8127528
Finding peers	-0.9403153	-1.2960113
Hiring	-0.3481444	-0.3734270
Legal	-1.0314186	-0.7248876
Managing issues	0.5628877	-0.4612921
Recognition	-0.7581089	-0.5491573
Securing funding	-0.4847993	1.4278090
Security	-0.8492121	-0.7688202

attr("scaled:center")

Non.applicable	Never	Rarely	Occasionally	Frequently
47.64286	20.35714	34.07143	47.78571	46.64286

Always
36.50000

attr("scaled:scale")

Non.applicable	Never	Rarely	Occasionally	Frequently
28.03736	11.60641	21.06570	15.63614	21.95312

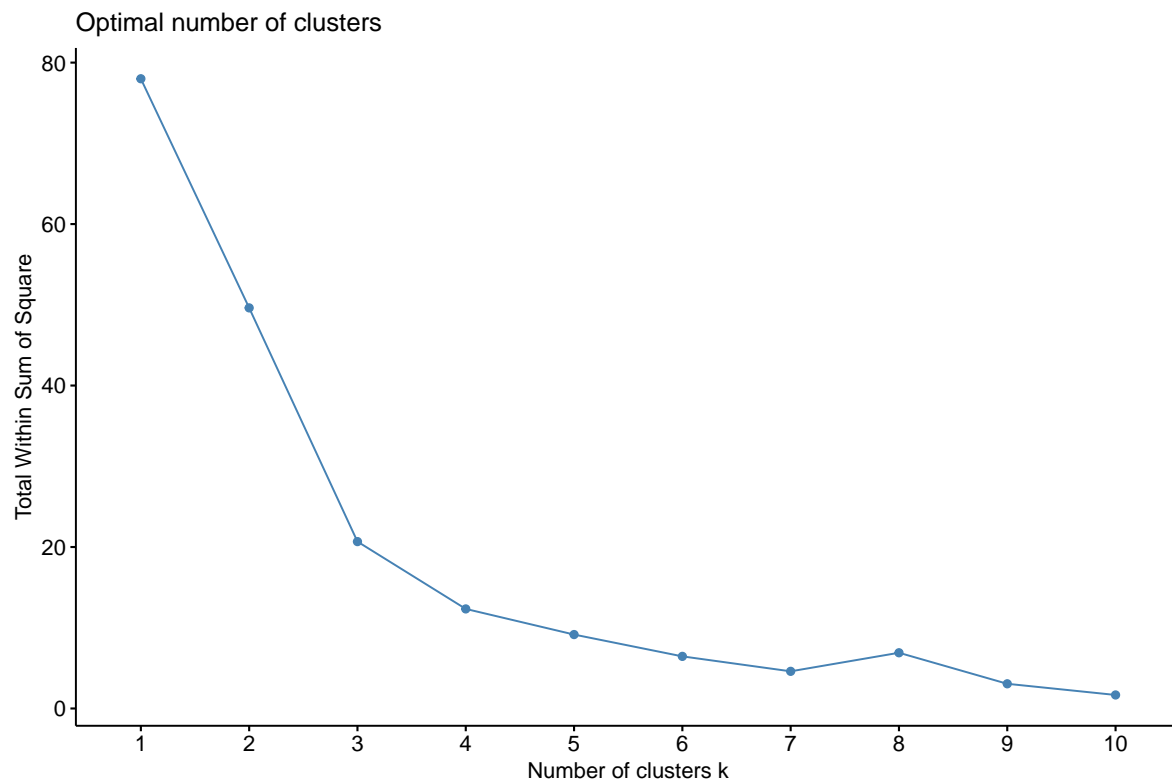
Always
22.76215

Plot an elbow plot to find the point of diminishing returns.

```
factoextra::fviz_nbclust(scaled, kmeans, method = "wss")
```

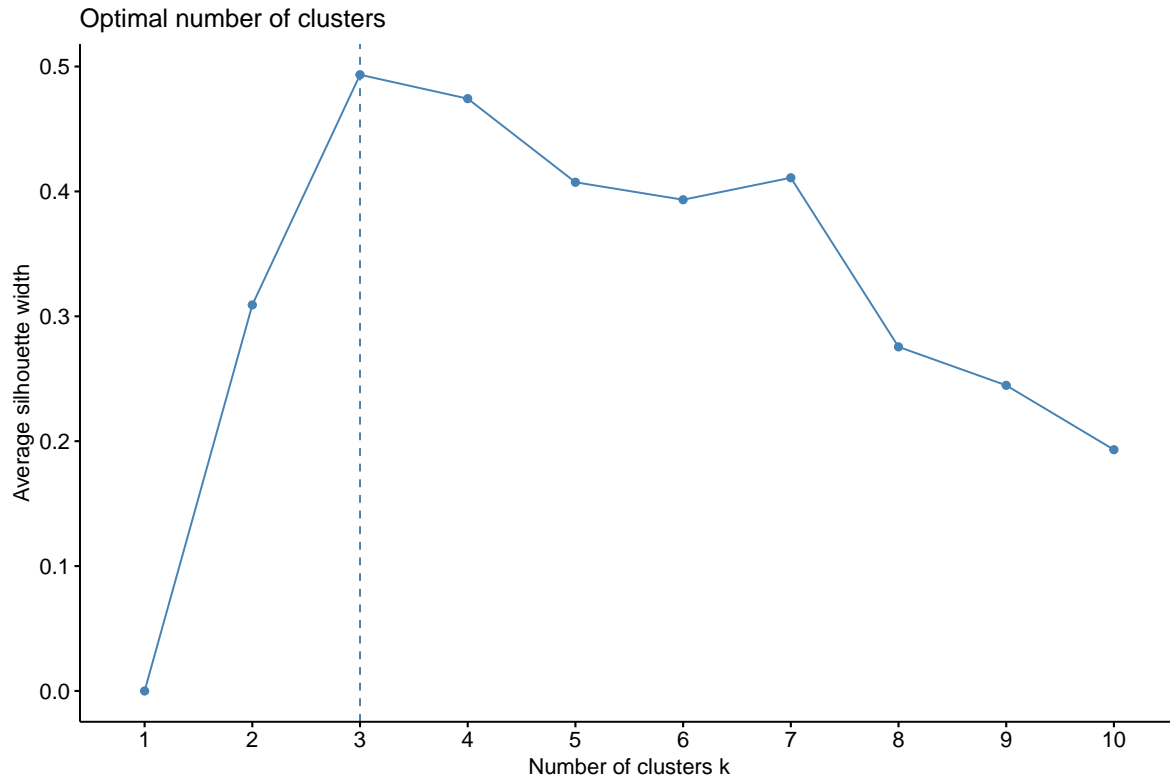
Registered S3 methods overwritten by 'car':

```
method      from
hist.boot   FSA
confint.boot FSA
```

I seem to get diminishing returns around $k=4$.

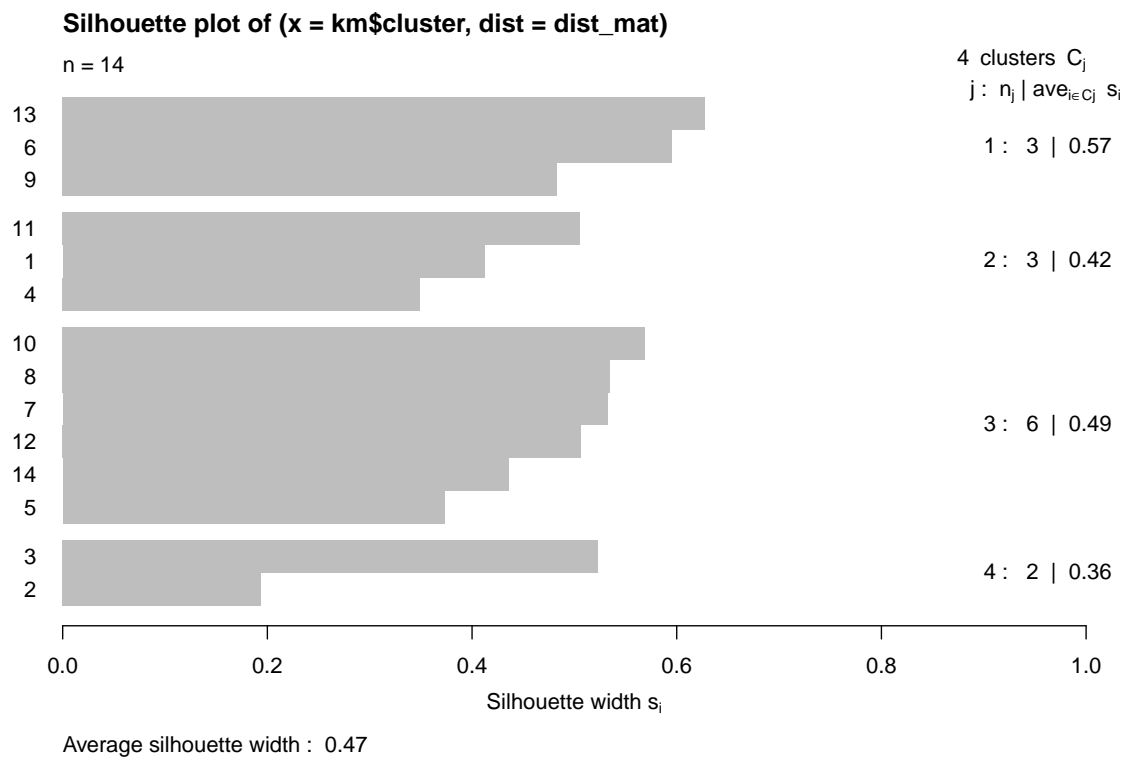
```
factoextra::fviz_nbclust(scaled, kmeans, method = "silhouette")
```



Hm. The silhouette plot indicates I should use $k=3$.

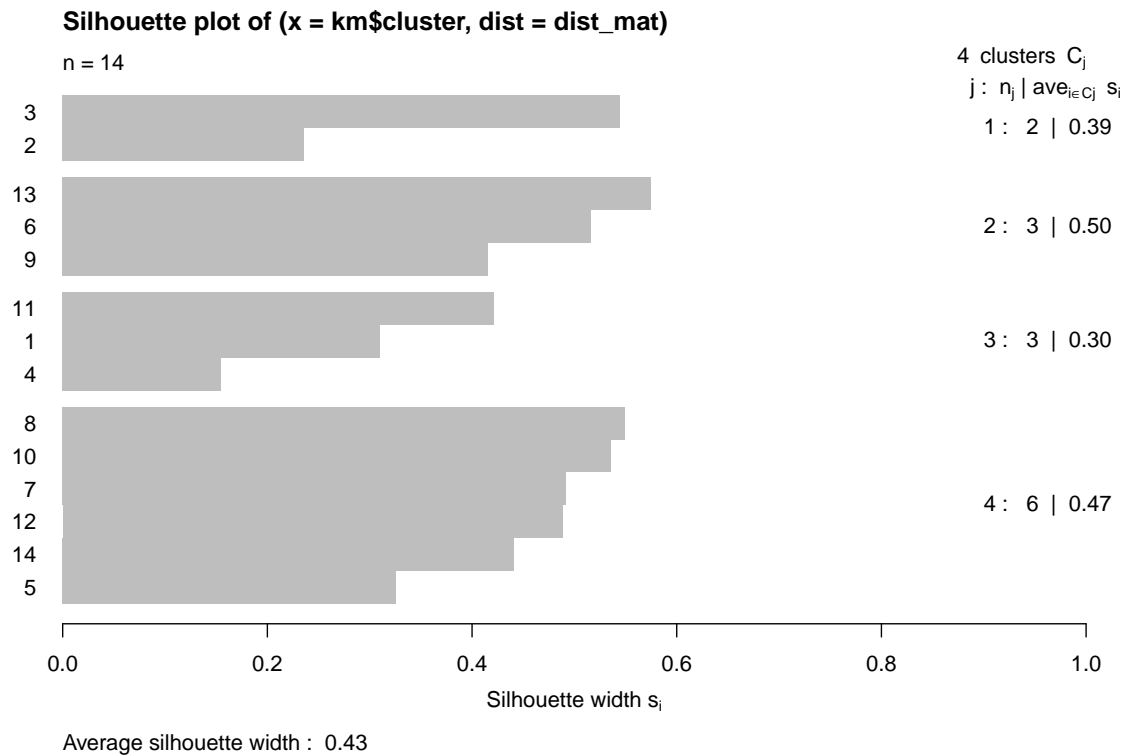
I think I'll try $k=4$ first, since it's closer to the number I got from eyeballing. Let's look at a different type of silhouette plot, which shows us the silhouette width of each cluster and on average across the clusters.

```
km <- stats::kmeans(scaled, centers = 4, nstart = 25)
dist_mat <- dist(scaled)
sil <- cluster::silhouette(km$cluster, dist_mat)
plot(sil)
```



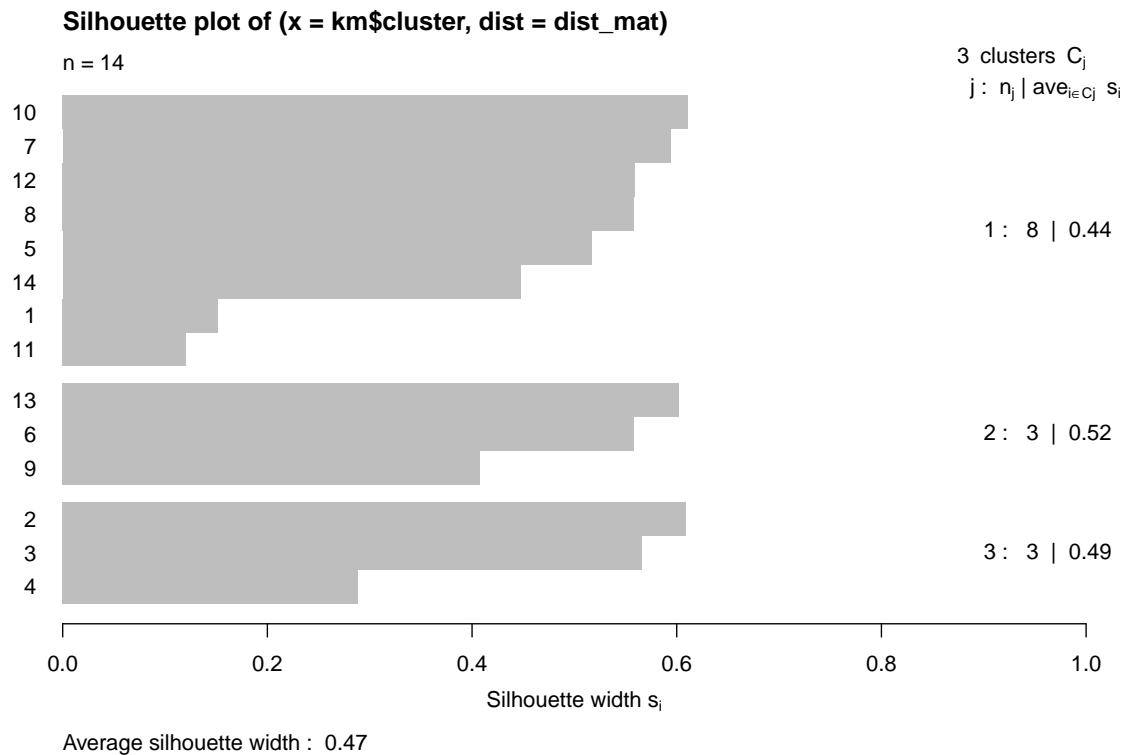
Hm. Looks... acceptable. Average silhouette width of 0.47. From Wikipedia: “A clustering with an average silhouette width of over 0.7 is considered to be “strong”, a value over 0.5 “reasonable” and over 0.25 “weak.” Let’s try unscaled data.

```
km <- stats::kmeans(wide_counts, centers = 4, nstart = 25)
dist_mat <- dist(wide_counts)
sil <- cluster::silhouette(km$cluster, dist_mat)
plot(sil)
```



Looks slightly worse: average silhouette width = 0.43. Still, I think we should probably stick with unscaled data because it's simpler, and I don't think we should add extra unnecessary procedures. What if we try 3 clusters?

```
km <- stats::kmeans(wide_counts, centers = 3, nstart = 25)
dist_mat <- dist(wide_counts)
sil <- cluster::silhouette(km$cluster, dist_mat)
plot(sil)
```



With an average silhouette width of 0.44-0.52, our clusters aren't looking amazing (unscaled data, 3 clusters). But they're not terrible, either. I prefer to use unscaled data with $k=3$, which results in an average silhouette width of 0.47. I think these results are consistent with my hunch that the data for the challenges are not all drawn from the same distribution. These are the cluster assignments:

```
# A little extra code to achieve prettier printing
cluster_df <- data.frame(sort(km$cluster))
cluster_df$challenge <- rownames(cluster_df)
clusters <- unique(cluster_df[,1])
for (cl in clusters) {
  print(cluster_df[cluster_df[,1] == cl,], row.names = FALSE)
  cat("\n")
}
```

```
sort.km.cluster.      challenge
1      Attracting users
1 Educational resources
1      Finding mentors
1      Finding peers
```

```

1          Legal
1    Managing issues
1      Recognition
1      Security

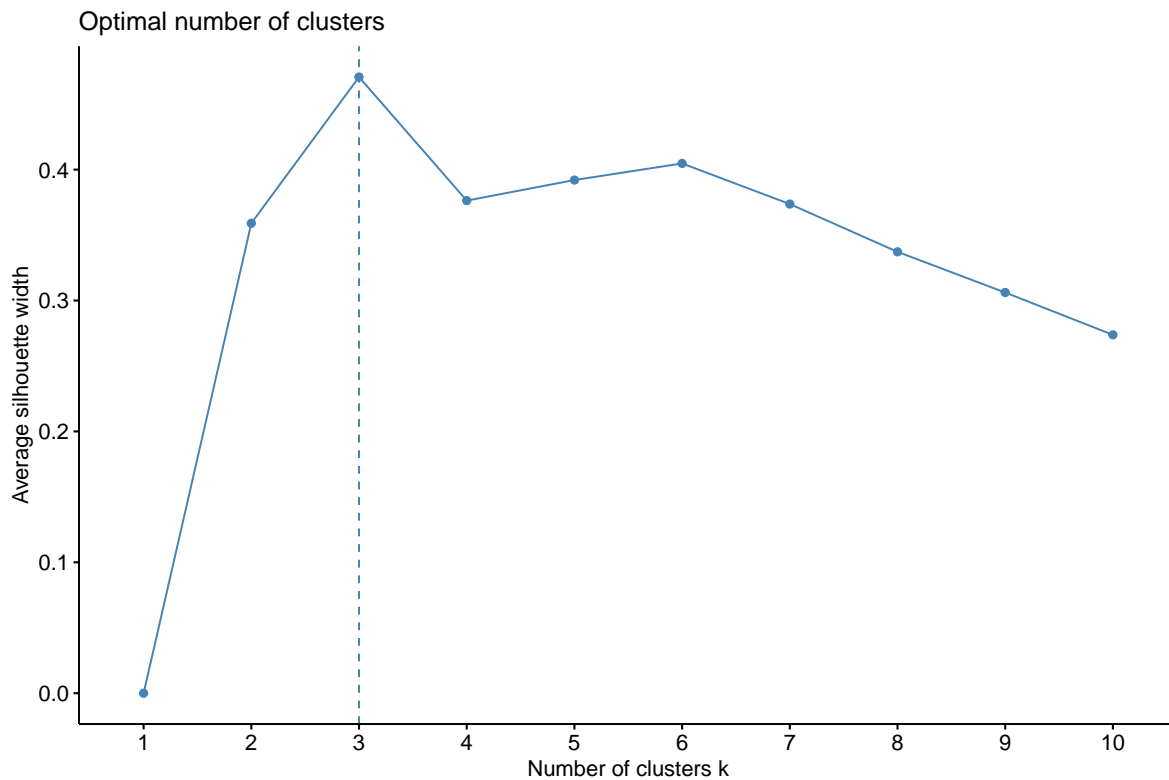
sort.km.cluster.    challenge
2  Finding funding
2      Hiring
2  Securing funding

sort.km.cluster.    challenge
3      Coding time
3  Documentation time
3      Education time

```

Let's look at a silhouette plot for the PAM method, too.

```
factoextra::fviz_nbclust(wide_counts, FUNcluster = pam, method = "silhouette")
```



This also says that 3 clusters is ideal.

Let's try PAM clustering on the unscaled data with k=3.

```
pm <- cluster::pam(wide_counts, k=3)
```

Print the clusters in a more readable format.

```
cluster_df <- data.frame(sort(pm$cluster))
cluster_df$challenge <- rownames(cluster_df)
clusters <- unique(cluster_df[,1])
for (cl in clusters) {
  print(cluster_df[cluster_df[,1] == cl,], row.names = FALSE)
  cat("\n")
}
```

```
sort.pm.cluster.      challenge
      1      Attracting users
      1 Educational resources
      1      Finding mentors
      1      Finding peers
      1           Legal
      1      Managing issues
      1      Recognition
      1      Security
```

```
sort.pm.cluster.      challenge
      2      Coding time
      2 Documentation time
      2      Education time
```

```
sort.pm.cluster.      challenge
      3 Finding funding
      3      Hiring
      3 Securing funding
```

We see the same groups we saw with k-means clustering. Good!

One last check: what about a stability assessment by bootstrap resampling?

```
# Note I'm hiding the printed status update from each iteration
boot_res <- fpc::clusterboot(
  wide_counts,
  clustermethod = fpc::kmeansCBI,
```

```

    krange = 3
)
# Annoyingly, the documentation doesn't explain 'krange',
# but I'm pretty sure that this argument lets you specify
# a desired k or range of k values (e.g. 5:7)

```

```
boot_res
```

```

* Cluster stability assessment *
Cluster method:  kmeans
Full clustering results are given as parameter result
of the clusterboot object, which also provides further statistics
of the resampling results.
Number of resampling runs:  100

Number of clusters found in data:  3

Clusterwise Jaccard bootstrap (omitting multiple points) mean:
[1] 0.7761190 0.8803333 0.8630714
dissolved:
[1] 30 16  1
recovered:
[1] 61 80 71

```

```
mean(boot_res$bootmean)
```

```
[1] 0.8398413
```

The clusterwise Jaccard bootstrap means are around 0.8-0.9, which is pretty respectable. Although this analysis was brief, I think we can conclude that these three clusters are reasonably stable and meaningful.

Stacked bar plots

A request from Amber: how about color-coded stacked bar plots, instead of the bar charts I made above?


```

# Rename this column, poorly named by the cluster package
# From `sort.km.cluster.` to sort_km_cluster
names(cluster_df)[1] <- "sort_km_cluster"

temp <- to_plot
temp$challenge_level <- factor(temp$challenge_level, levels = rev(ordered_levels))

data_cluster1 <- temp %>%
  filter(
    challenge %in% rownames(subset(cluster_df, sort_km_cluster == 1))
  )

data_cluster2 <- temp %>%
  filter(
    challenge %in% rownames(subset(cluster_df, sort_km_cluster == 2))
  )

data_cluster3 <- temp %>%
  filter(
    challenge %in% rownames(subset(cluster_df, sort_km_cluster == 3))
  )

```

WARNING! I am changing the cluster names ONLY on the plot, not in the rest of the code. The plot looks better when the two small clusters—clusters 1 and 3—are next to each other (we want to visually compare them, and it's easier to do so when they're adjacent). However, the plot looks dumb when I have them in order 1, 3, 2. It looks like I made a mistake. So I am renaming the plot panel titles ONLY.

```

# Paul Tol's sunset theme
#https://sronpersonalpages.nl/~pault/
sunset <- c(
  "#A50026",
  "#F67E4B",
  "#FEDA8B",
  "#C2E4EF",
  "#6EA6CD",
  "#364B9A"
)

```

```

p1 <- stacked_bar_chart(
  df = data_cluster2,
  x_var = "challenge",
  y_var = "total",
  fill = "challenge_level",
  title = "Cluster 1", # NAME CHANGE, SEE TEXT ABOVE
  ylabel = NULL,
  show_axis_title_y = FALSE,
  show_x_axis_text = FALSE,
  show_grid = TRUE,
  show_legend = FALSE, # don't show legend
  horizontal = TRUE,
  proportional = TRUE,
  cpalette = sunset
)

p2 <- stacked_bar_chart(
  df = data_cluster3,
  x_var = "challenge",
  y_var = "total",
  fill = "challenge_level",
  title = "Cluster 2", # NAME CHANGE, SEE TEXT ABOVE
  ylabel = NULL,
  legend_left_margin = 45, # show legend, with a wide margin
  show_axis_title_y = FALSE,
  show_x_axis_text = FALSE,
  show_grid = TRUE,
  horizontal = TRUE,
  proportional = TRUE,
  cpalette = sunset
)

p3 <- stacked_bar_chart(
  df = data_cluster1,
  x_var = "challenge",
  y_var = "total",
  fill = "challenge_level",
  title = "Cluster 3", # NAME CHANGE, SEE TEXT ABOVE
  ylabel = NULL,
  show_axis_title_y = FALSE,
  show_x_axis_text = FALSE,
  show_grid = TRUE,

```

```

    show_legend = FALSE, # don't show legend
    horizontal = TRUE,
    proportional = TRUE,
    cpalette = sunset
)

```

Save the data for tweaking the final figure. NOTE CLUSTER NAME CHANGE.

```

write_df_to_file(data_cluster1, "data_for_plots/chall_cluster3.tsv")
write_df_to_file(data_cluster3, "data_for_plots/chall_cluster2.tsv")
write_df_to_file(data_cluster2, "data_for_plots/chall_cluster1.tsv")

```

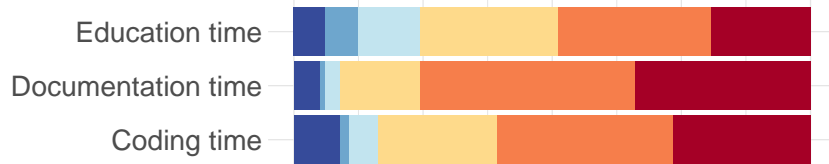
```

combined <- patchwork::wrap_plots(p1, p2, p3, ncol = 1) +
  patchwork::plot_layout(heights = c(1, 1, 2)) +
  patchwork::plot_annotation(
    title = "Frequency of Open Source Challenges",
    theme = theme(plot.title = element_text(
      size = 24,
      margin = margin(t = 15),
      hjust = 0.5)
    )
  )
combined

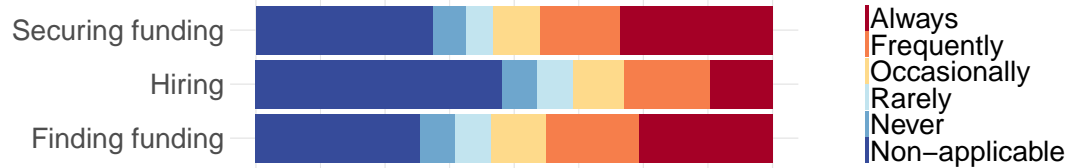
```

Frequency of Open Source Challenges

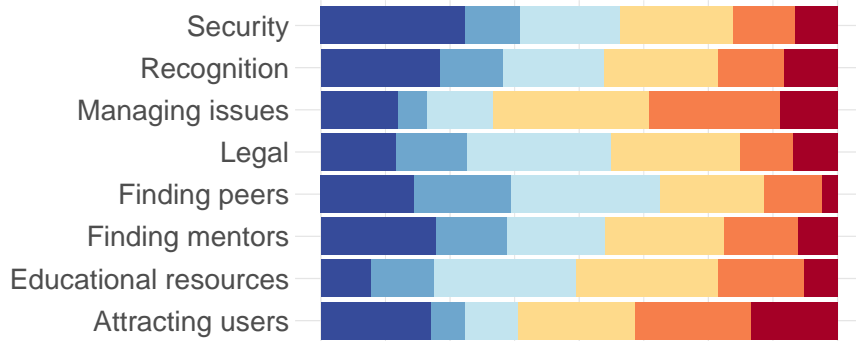
Cluster 1



Cluster 2



Cluster 3



```
save_plot("challenge_stacks.tiff", 14, 12, p=combined)
```

How does it look if we arrange these plots horizontally? We'll need to put the legend on the right-most plot in this case.

```
p2_hor <- stacked_bar_chart(
  df = data_cluster3,
  x_var = "challenge",
  y_var = "total",
  fill = "challenge_level",
  title = "Cluster 2", # NAME CHANGE, SEE TEXT ABOVE
  ylabel = NULL,
  show_legend = FALSE, # don't show legend
```

```

    show_axis_title_y = FALSE,
    show_x_axis_text = FALSE,
    show_grid = TRUE,
    horizontal = TRUE,
    proportional = TRUE,
    cpalette = sunset
  )

p3_hor <- stacked_bar_chart(
  df = data_cluster1,
  x_var = "challenge",
  y_var = "total",
  fill = "challenge_level",
  title = "Cluster 3", # NAME CHANGE, SEE TEXT ABOVE
  ylabel = NULL,
  show_axis_title_y = FALSE,
  show_x_axis_text = FALSE,
  show_grid = TRUE,
  legend_left_margin = 45, # show legend, with a wide margin
  horizontal = TRUE,
  proportional = TRUE,
  cpalette = sunset
)

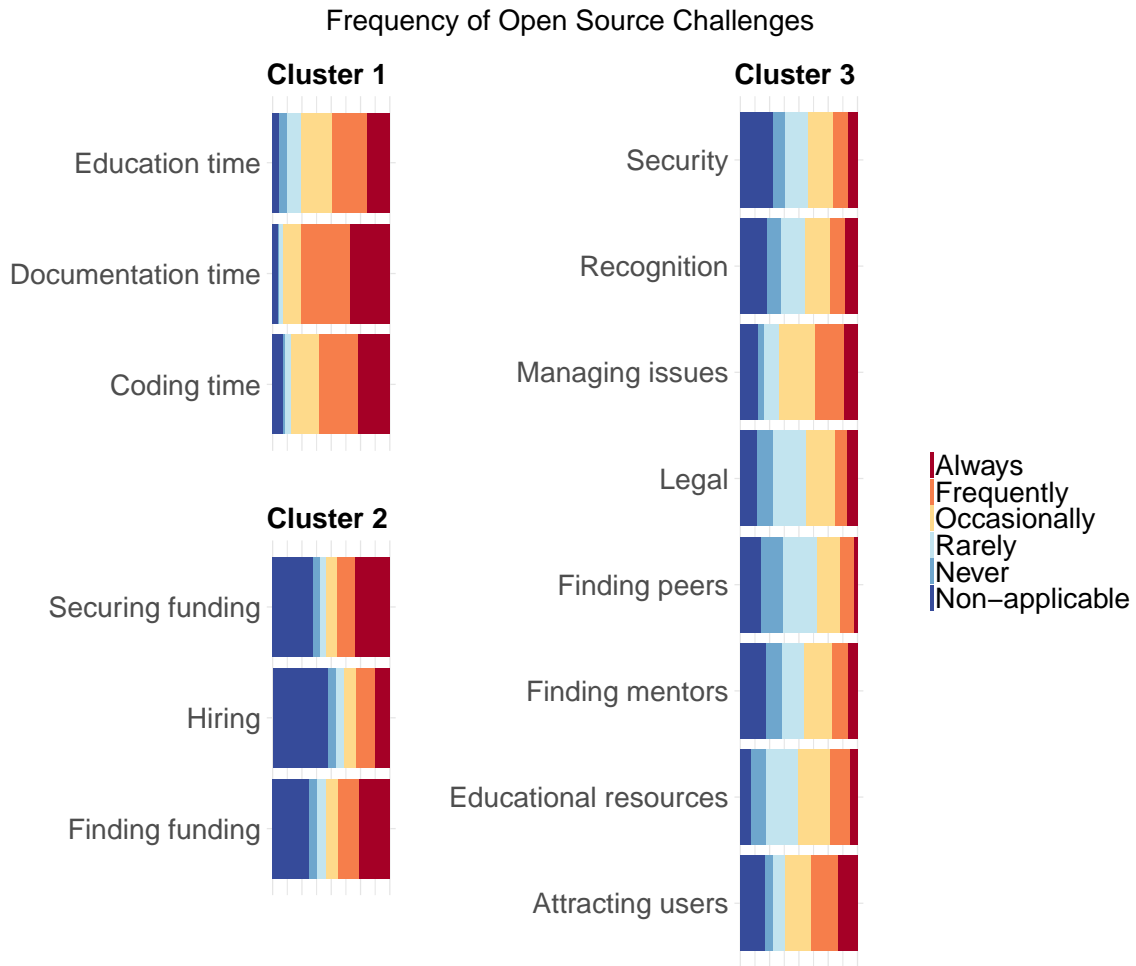
```

Tried plotting horizontally a couple different ways. Sticking with the vertical version.

```

# left column: p1 over p2; right column: p3 spanning full height
combined_hor <- patchwork::wrap_plots(
  (p1 / p2_hor / patchwork::plot_spacer()) + plot_layout(heights = c(1,1,0.1)) | p3_hor
) +
  #patchwork::plot_layout(heights = c(1, 1, 1)) +
  patchwork::plot_annotation(
    title = "Frequency of Open Source Challenges",
    theme = theme(
      plot.title = element_text(
        size = 24,
        margin = margin(t = 15),
        hjust = 0.5
      )
    )
  )
combined_hor

```

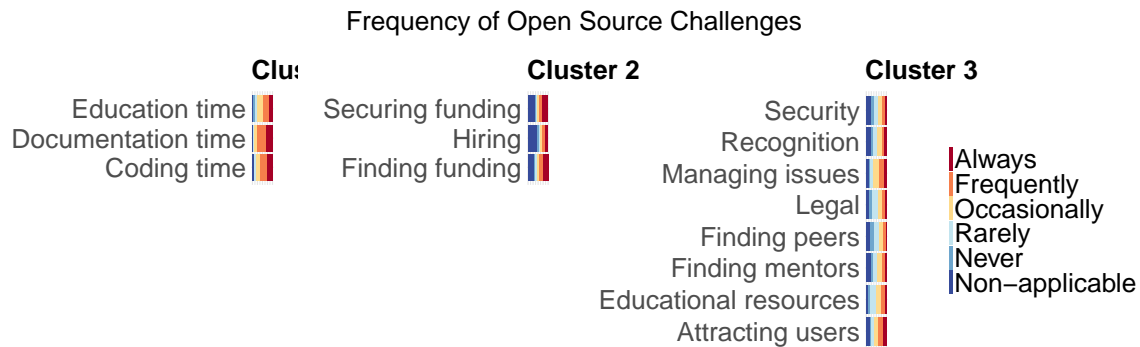


```
# three plots in a row
combined_hor <- patchwork::wrap_plots(
  (p1 / patchwork::plot_spacer() ) + plot_layout(heights = c(1,1.4)) |
  (p2_hor / patchwork::plot_spacer() ) + plot_layout(heights = c(1,1.4)) |
  p3_hor
) +
#patchwork::plot_layout(heights = c(1, 1, 1)) +
patchwork::plot_annotation(
  title = "Frequency of Open Source Challenges",
  theme = theme(
    plot.title = element_text(
      size = 24,
      margin = margin(t = 15),
```

```

    hjust = 0.5
  )
)
combined_hor

```



```

save_plot("challenge_stacks_horizontal.tiff", 36, 8, p=combined_hor)

```

```

sessionInfo()

```

```

R version 4.4.2 (2024-10-31)
Platform: aarch64-apple-darwin20
Running under: macOS 26.1

```

```

Matrix products: default

```

```

BLAS:   /Library/Frameworks/R.framework/Versions/4.4-arm64/Resources/lib/libRblas.0.dylib
LAPACK: /Library/Frameworks/R.framework/Versions/4.4-arm64/Resources/lib/libRlapack.dylib;

```

```

locale:

```

```

[1] C.UTF-8/C.UTF-8/C.UTF-8/C/C.UTF-8/C.UTF-8

```

```

time zone: America/Los_Angeles

```

```

tzcode source: internal

```

```

attached base packages:

```

```

[1] tools      grid        stats      graphics  grDevices  datasets  utils
[8] methods    base

```

```

other attached packages:

```

[1] treemapify_2.5.6	tidyr_1.3.1	svglite_2.2.1
[4] stringr_1.5.1	scales_1.4.0	readr_2.1.5
[7] pwr_1.3-0	patchwork_1.3.2	ordinal_2023.12-4.1
[10] lme4_1.1-37	Matrix_1.7-1	languageserver_0.3.16
[13] here_1.0.1	gtools_3.9.5	ggforce_0.5.0
[16] FSA_0.10.0	fpc_2.2-13	forcats_1.0.0
[19] factoextra_1.0.7	ggplot2_3.5.2	emmeans_1.11.2
[22] dplyr_1.1.4	corrplot_0.95	ComplexHeatmap_2.22.0
[25] cluster_2.1.8.1	BiocManager_1.30.26	

loaded via a namespace (and not attached):

[1] Rdpack_2.6.4	rlang_1.1.6	magrittr_2.0.3
[4] clue_0.3-66	GetoptLong_1.0.5	matrixStats_1.5.0
[7] compiler_4.4.2	flexmix_2.3-20	systemfonts_1.2.3
[10] png_0.1-8	callr_3.7.6	vctrs_0.6.5
[13] pkgconfig_2.0.3	shape_1.4.6.1	crayon_1.5.3
[16] fastmap_1.2.0	backports_1.5.0	labeling_0.4.3
[19] utf8_1.2.6	rmarkdown_2.29	ggfitttext_0.10.2
[22] tzdb_0.5.0	ps_1.9.1	nloptr_2.2.1
[25] purrr_1.1.0	xfun_0.53	modeltools_0.2-24
[28] jsonlite_2.0.0	tweenr_2.0.3	broom_1.0.9
[31] parallel_4.4.2	prabclus_2.3-4	R6_2.6.1
[34] stringi_1.8.7	RColorBrewer_1.1-3	car_3.1-3
[37] boot_1.3-31	diptest_0.77-2	numDeriv_2016.8-1.1
[40] estimability_1.5.1	Rcpp_1.1.0	iterators_1.0.14
[43] knitr_1.50	IRanges_2.40.1	splines_4.4.2
[46] nnet_7.3-19	tidyselect_1.2.1	abind_1.4-8
[49] yaml_2.3.10	doParallel_1.0.17	codetools_0.2-20
[52] processx_3.8.6	lattice_0.22-6	tibble_3.3.0
[55] withr_3.0.2	evaluate_1.0.4	polyclip_1.10-7
[58] xml2_1.4.0	circlize_0.4.16	mclust_6.1.1
[61] kernlab_0.9-33	ggpubr_0.6.1	pillar_1.11.0
[64] carData_3.0-5	renv_1.1.5	foreach_1.5.2
[67] stats4_4.4.2	reformulas_0.4.1	generics_0.1.4
[70] rprojroot_2.1.1	S4Vectors_0.44.0	hms_1.1.3
[73] minqa_1.2.8	xtable_1.8-4	class_7.3-22
[76] glue_1.8.0	robustbase_0.99-4-1	ggsignif_0.6.4
[79] mvtnorm_1.3-3	rbibutils_2.3	colorspace_2.1-1
[82] nlme_3.1-166	Formula_1.2-5	cli_3.6.5
[85] textshaping_1.0.1	gtable_0.3.6	DEoptimR_1.1-4
[88] rstatix_0.7.2	digest_0.6.37	BiocGenerics_0.52.0
[91] ucminf_1.2.2	ggrepel_0.9.6	rjson_0.2.23
[94] farver_2.1.2	htmltools_0.5.8.1	lifecycle_1.0.4

[97] GlobalOptions_0.1.2 MASS_7.3-61