

# 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 correspond to 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

```
data <- load_qualtrics_data("deidentified_no_qual.tsv")
```

## Wrangle data

```
challenges <- data %>%
  select(
    starts_with("challenges")
  )

head(challenges)
```

	challenges_1	challenges_2	challenges_3	challenges_4	challenges_5	challenges_6
1	Always	Always	Always	Always	Always	Always
2	Frequently	Occasionally	Occasionally	Occasionally	Occasionally	Rarely
3	Frequently	Always	Occasionally	Always	Occasionally	Frequently
4	Always	Always	Frequently	Occasionally	Frequently	Always
5	Always	Always	Rarely	Occasionally	Frequently	Never
6						

	challenges_7	challenges_8	challenges_9	challenges_10	challenges_11
1	Always	Always	Always	Always	Always
2	Frequently	Occasionally	Frequently	Frequently	Frequently
3	Frequently	Occasionally	Occasionally	Rarely	Rarely
4	Occasionally	Rarely	Rarely	Frequently	Rarely
5	Never	Never	Never	Always	Occasionally
6					

	challenges_12	challenges_13	challenges_14
1	Always	Always	Always
2	Frequently	Frequently	Occasionally
3	Always	Always	Always
4	Occasionally	Frequently	Frequently
5	Occasionally	Rarely	Always
6			

**STOP!!** Presumably, “challenges\_1” corresponds to the first option, “challenges\_2” corresponds to the second option, etc., but we still need to check. I am manually comparing the answers in this data frame to those in the Qualtrics interface, which shows the whole response, i.e. “Limited time for writing new code”, not just “challenges\_1”. To be extra confident that I am comparing the same rows between the two tables, I am looking at responses associated with a particular email. After this code chunk, I go back to using the data frame that doesn’t contain the emails.

Since this code only needed to be run once, I’ve commented it out.

```
# pii <- load_qualtrics_data("pii.tsv")
# emails <- pii %>%
#   select(starts_with("stay_in_touch_email"))

# t <- cbind(emails, challenges)

# # Next, I run this line repeatedly with different emails,
# # to make sure that this person's response to "challenges_1"
# # matches their response to "Limited time for writing new code", etc.
# subset(t, startsWith(stay_in_touch_email, "PERSON_EMAIL_HERE"))
```

My assumption above was correct; the options are ordered as expected. Let's rename the columns accordingly.

```
challenge_codes <- c(
  "Coding time" = "challenges_1",
  "Documentation time" = "challenges_2",
  "Managing issues" = "challenges_3",
  "Attracting users" = "challenges_4",
  "Recognition" = "challenges_5",
  "Hiring" = "challenges_6",
  "Security" = "challenges_7",
  "Finding peers" = "challenges_8",
  "Finding mentors" = "challenges_9",
  "Education time" = "challenges_10",
  "Educational resources" = "challenges_11",
  "Legal" = "challenges_12",
  "Finding funding" = "challenges_13",
  "Securing funding" = "challenges_14"
)
challenges <- rename(challenges, any_of(challenge_codes))
```

Next, 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.

```
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 ("Never" = 0, "Non-applicable" = 0, "Rarely" = 1, "Occasionally" = 2, "Frequently" = 3, "Always" = 4) \* 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 don't know.) \* 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),
    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 5
  challenge          total mean  mode st_dev
  <chr>          <int> <dbl> <int> <dbl>
1 Documentation time     686  2.94     3  1.08
2 Coding time           606  2.60     3  1.24
3 Education time         539  2.31     3  1.26
4 Managing issues        451  1.94     2  1.29
5 Attracting users       442  1.90     0  1.45
6 Securing funding       438  1.88     0  1.74
7 Finding funding       432  1.85     0  1.68
8 Educational resources  369  1.58     1  1.19
9 Recognition           334  1.43     0  1.35
10 Legal                333  1.43     0  1.24
11 Finding mentors       323  1.39     0  1.31
12 Security             307  1.32     0  1.31
13 Hiring               291  1.25     0  1.53
14 Finding peers        267  1.15     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”).

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 5  
  challenge      total  mean  mode st_dev  
  <chr>      <int> <dbl> <int> <dbl>  
1 Securing funding    438  1.88     0  1.74  
2 Finding funding    432  1.85     0  1.68  
3 Hiring              291  1.25     0  1.53  
4 Attracting users   442  1.90     0  1.45  
5 Recognition        334  1.43     0  1.35  
6 Security           307  1.32     0  1.31  
7 Finding mentors    323  1.39     0  1.31  
8 Managing issues    451  1.94     2  1.29  
9 Education time     539  2.31     3  1.26  
10 Legal             333  1.43     0  1.24  
11 Coding time       606  2.60     3  1.24  
12 Educational resources 369  1.58     1  1.19  
13 Finding peers     267  1.15     0  1.13  
14 Documentation time 686  2.94     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(
    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"
)

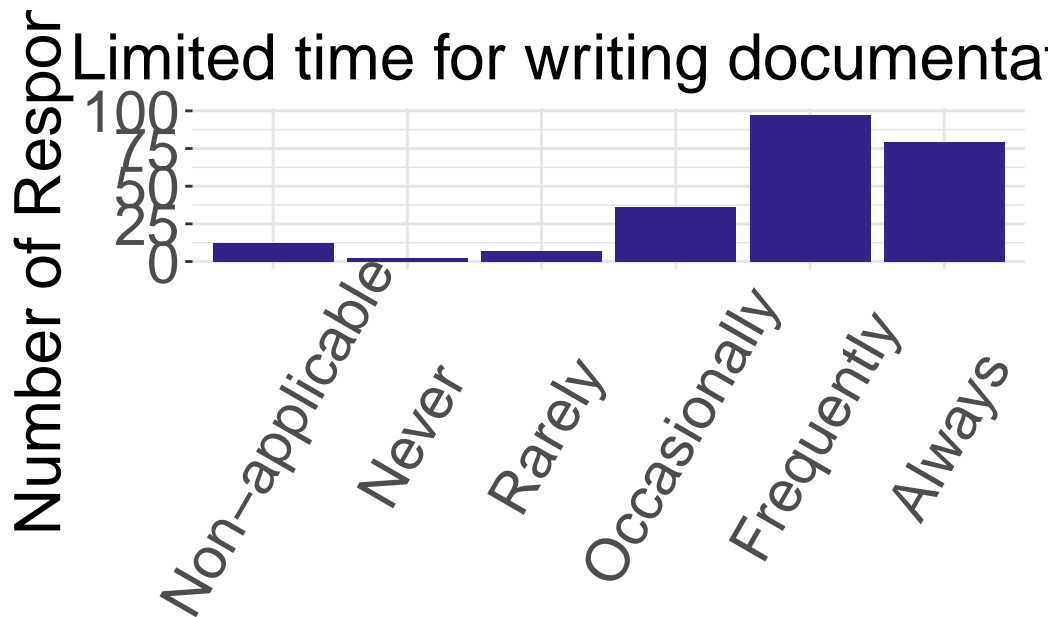
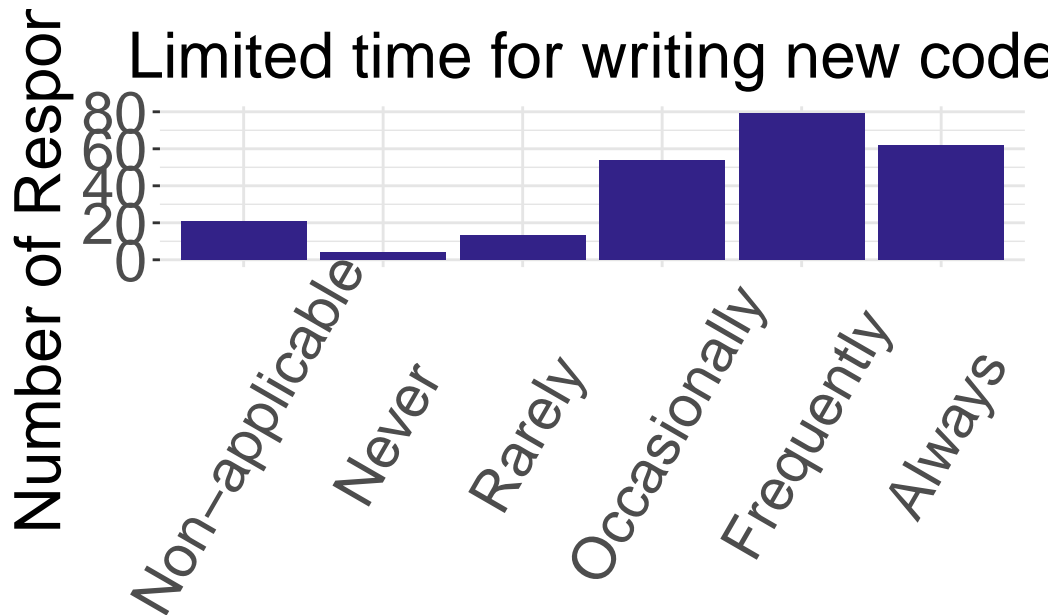
na_skewed <- c(
  "Managing issues",
  "Attracting users",
  "Recognition",
  "Security",
  "Finding mentors"
)

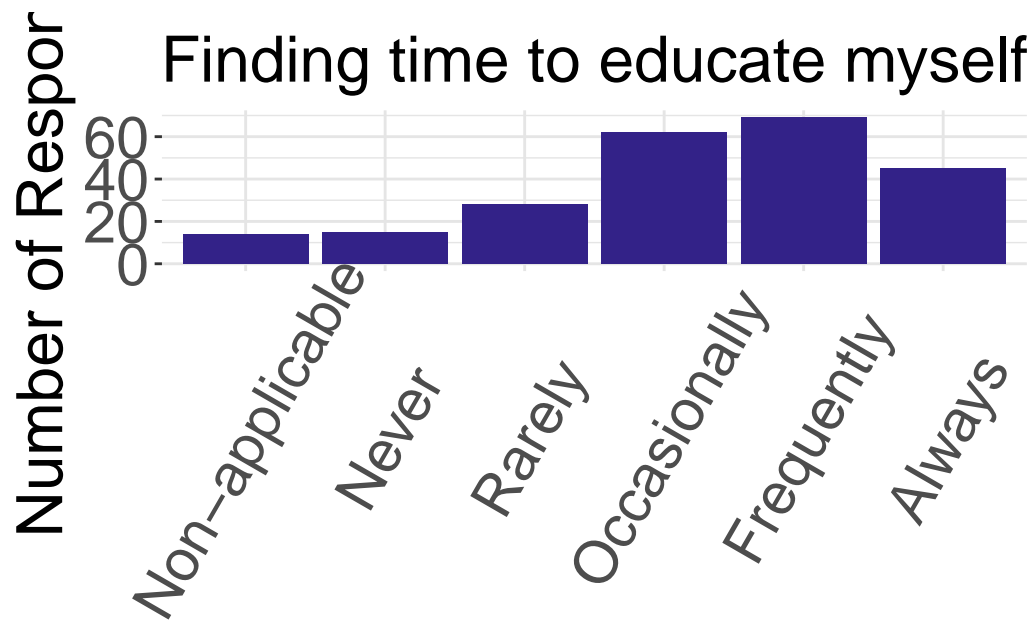
left_skewed <- c(
  "Finding peers"
)

```

“right-skewed”

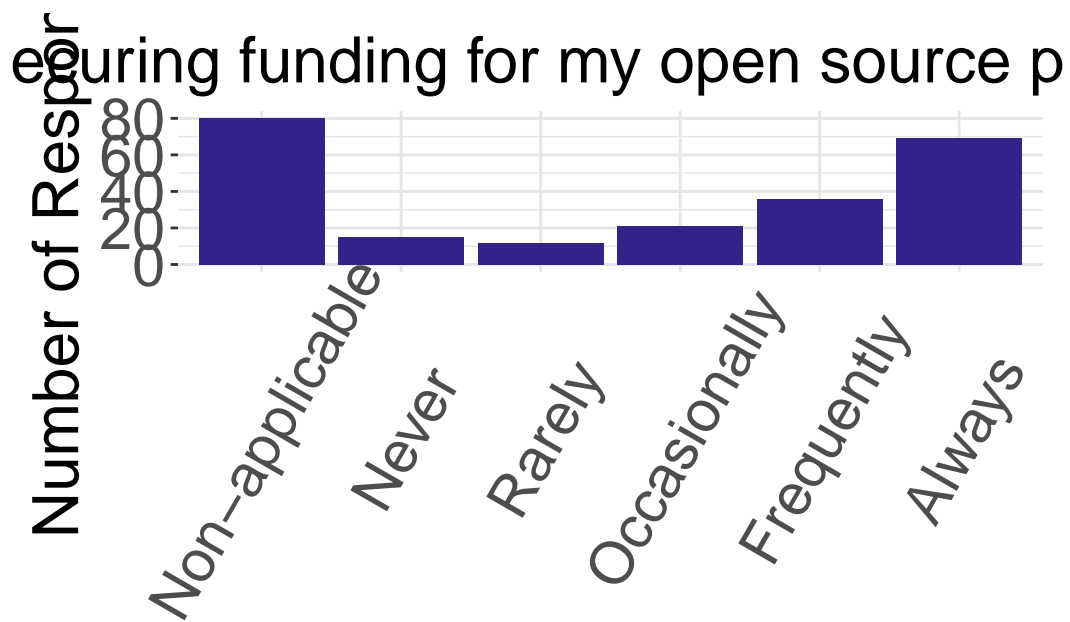
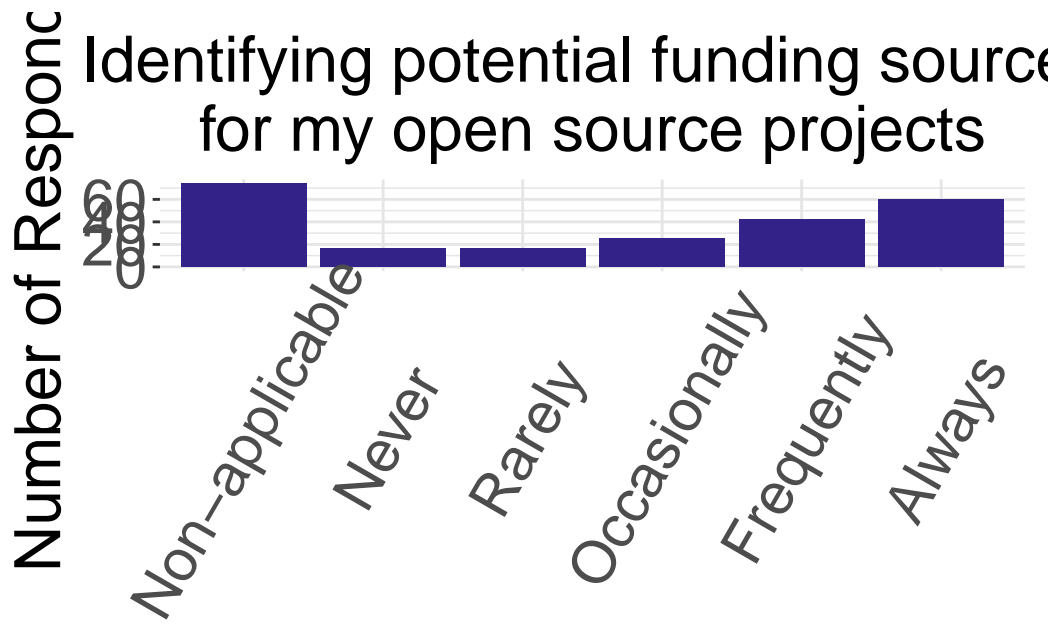
```
multiple_plots(to_plot, titles, right_skewed)
```

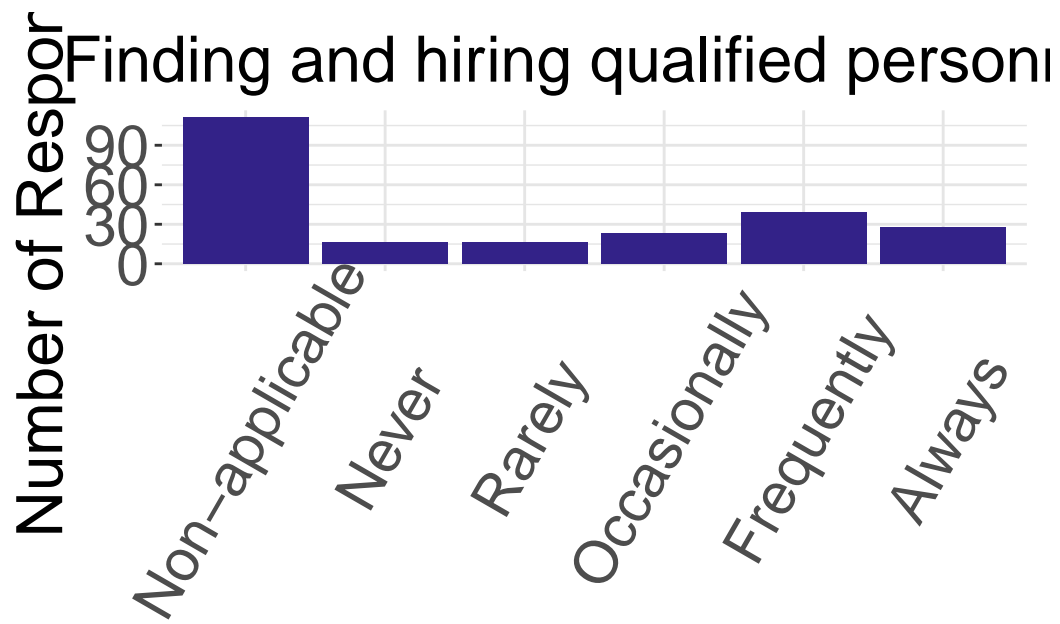




“highly bimodal”

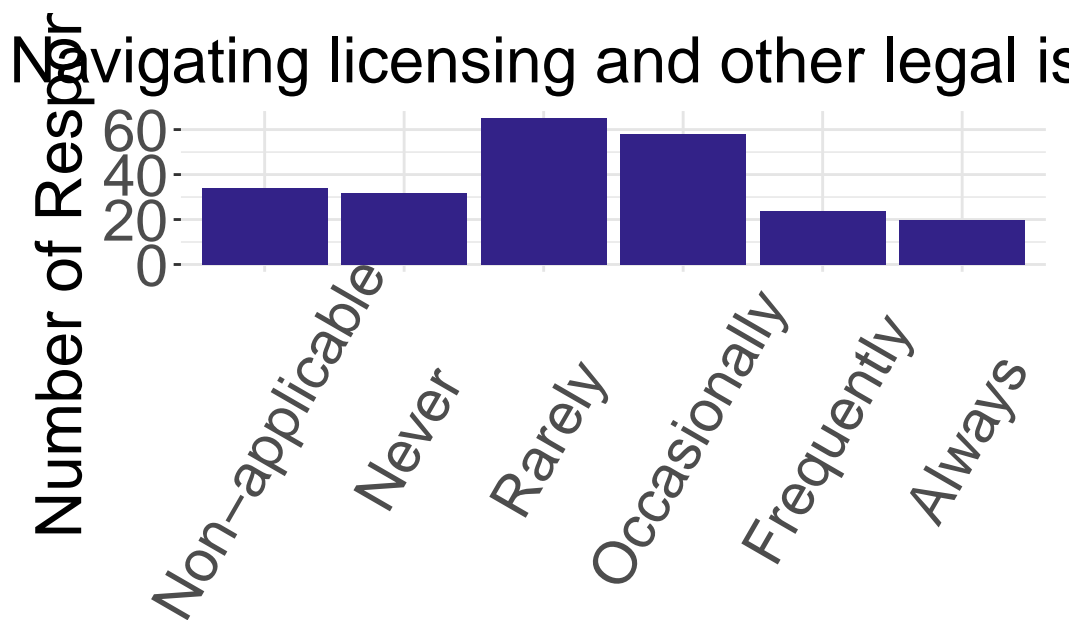
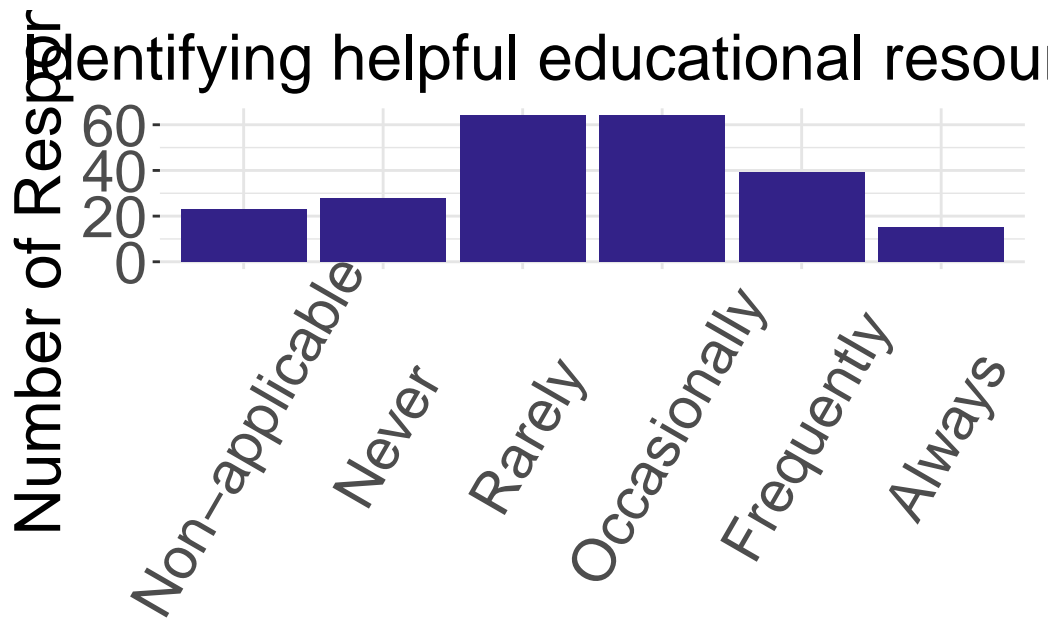
```
multiple_plots(to_plot, titles, bimodal)
```





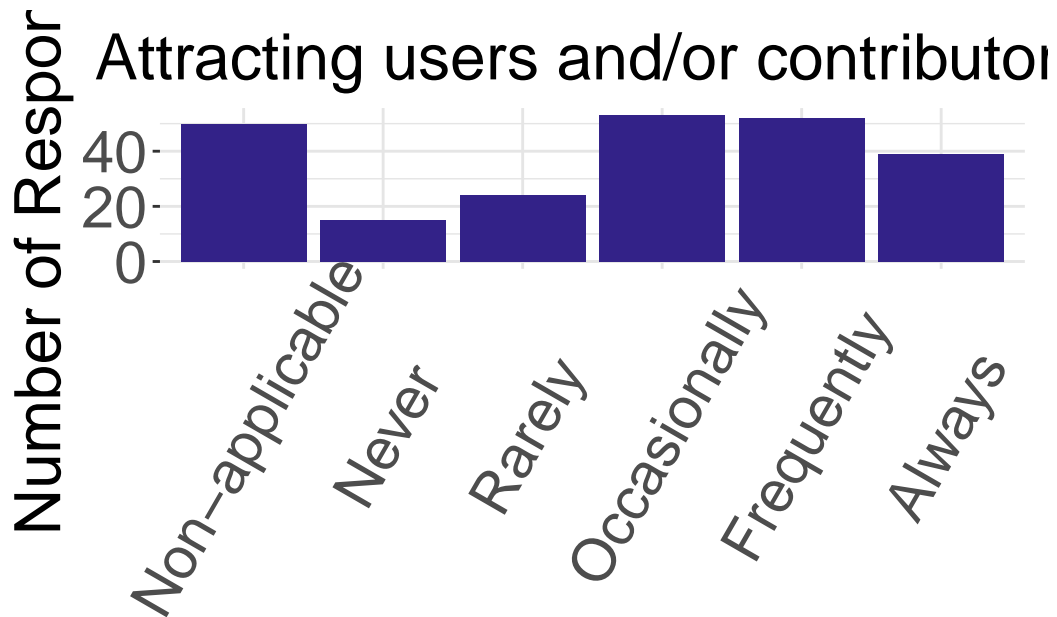
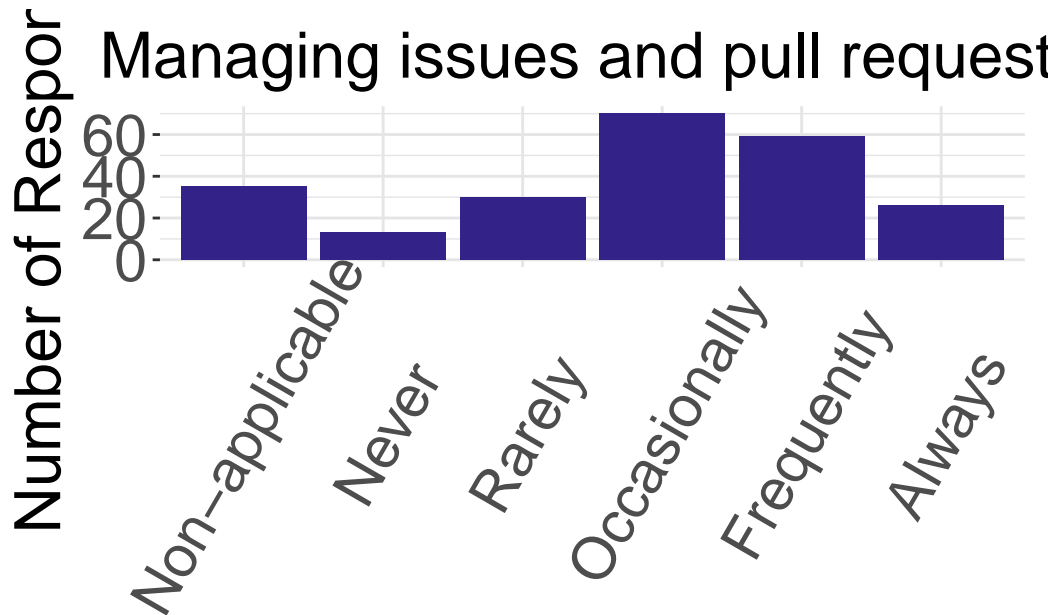
“normal”

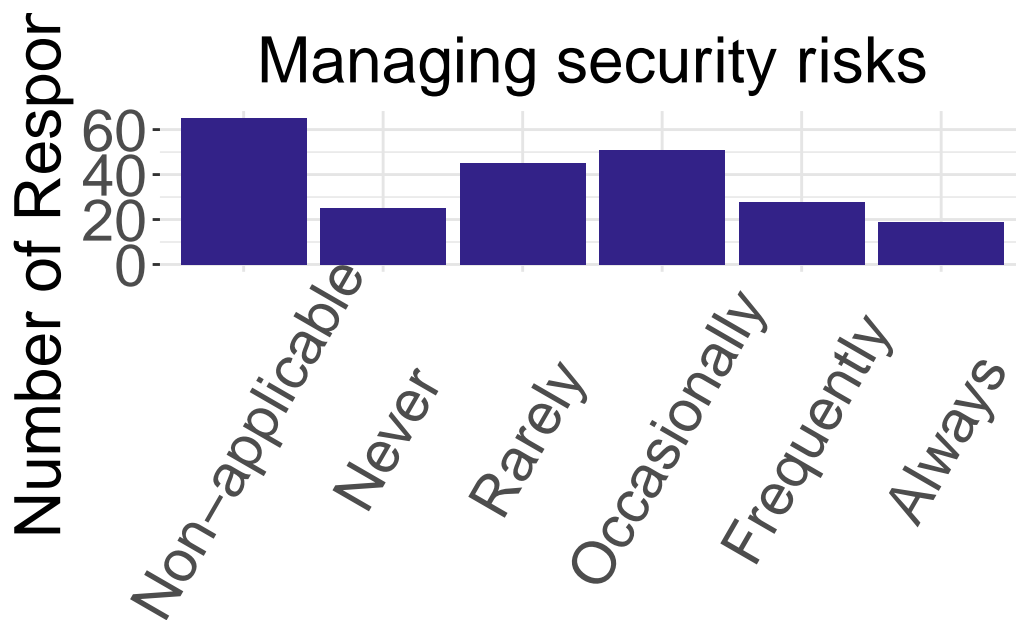
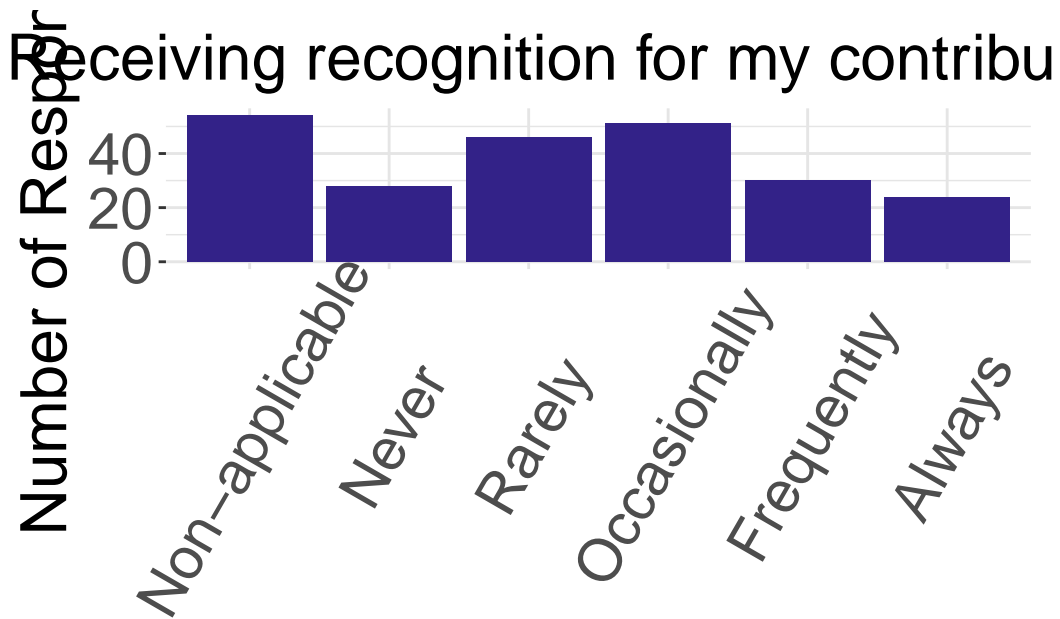
```
multiple_plots(to_plot, titles, normal)
```



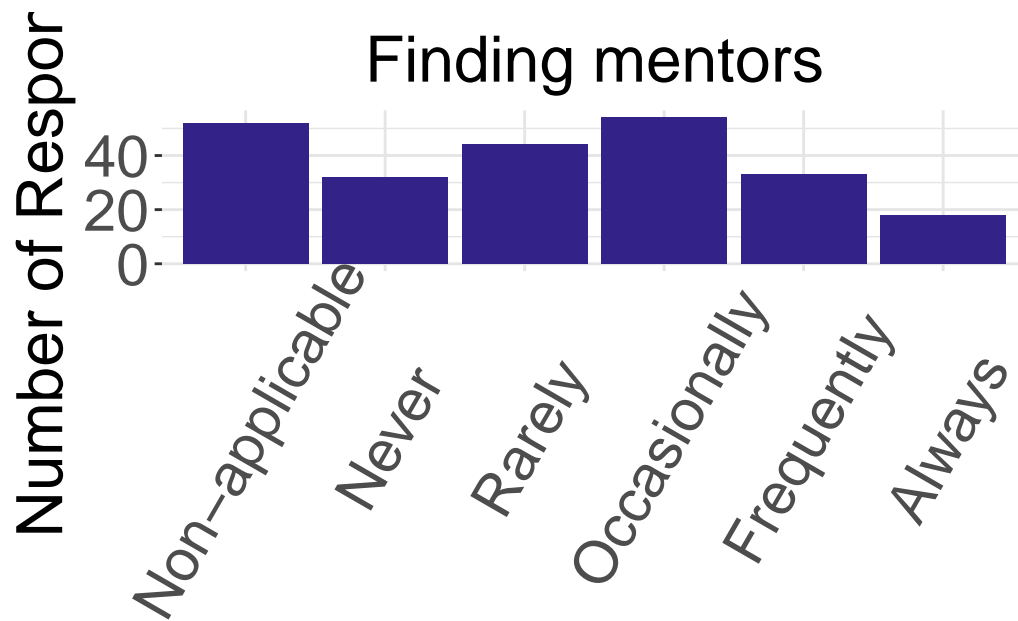
“na-skewed”

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



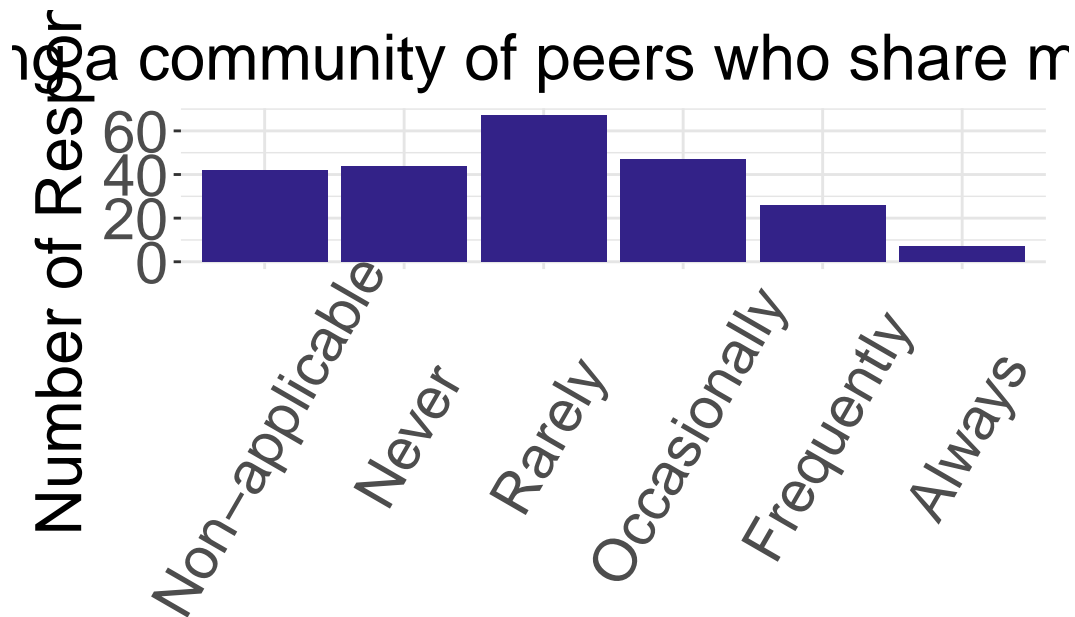






“left-skewed”

```
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))]

# 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
```

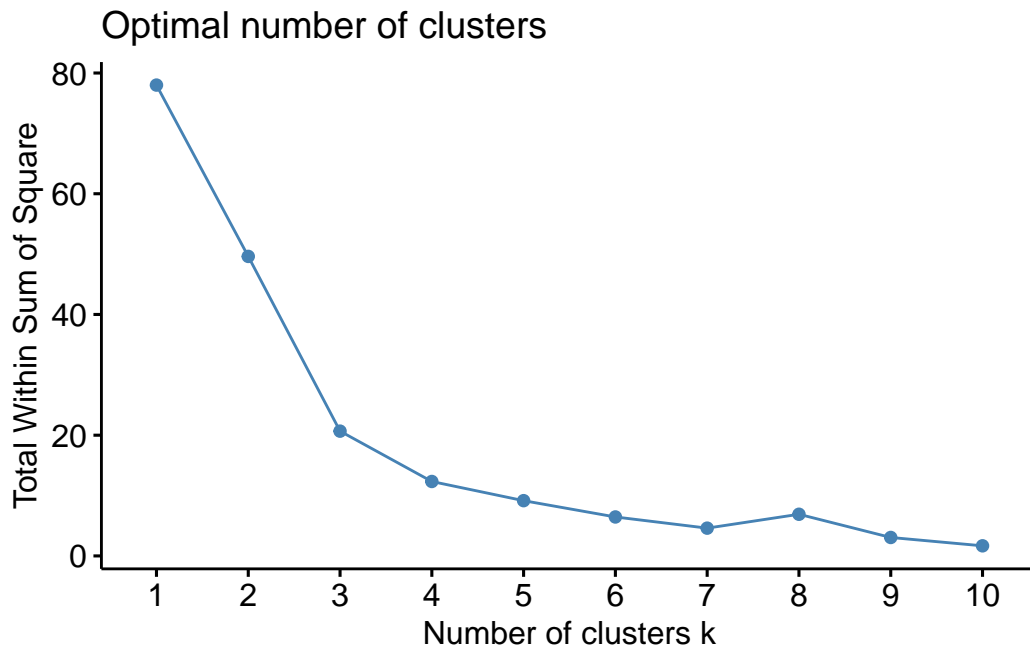
```
attr(,"scaled:scale")
```

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

Always  
22.76215

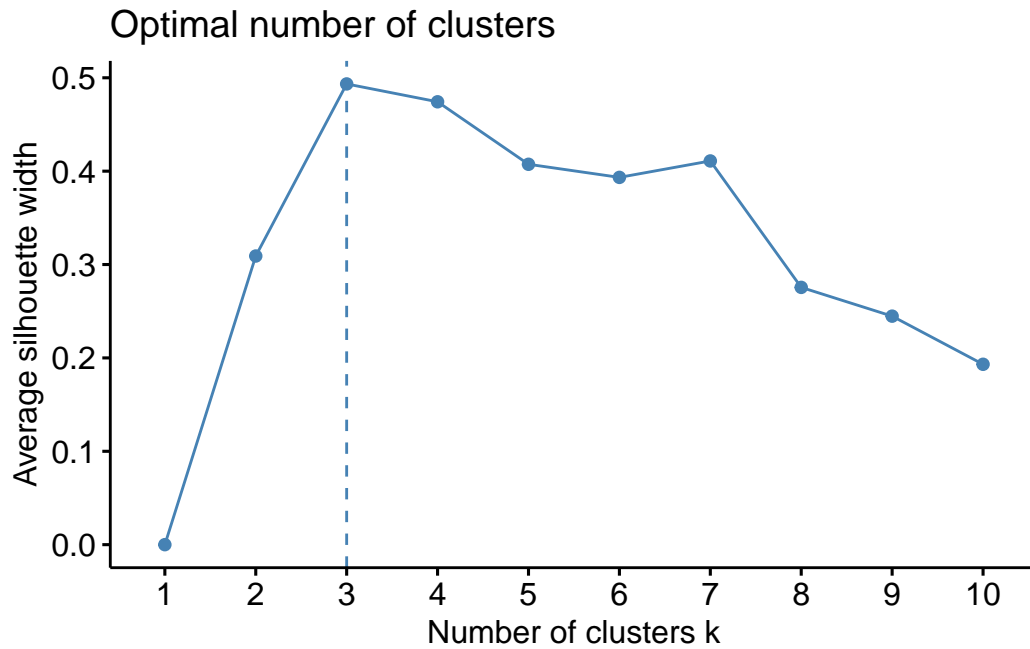
Plot an elbow plot to find the point of diminishing returns.

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



I seem to get diminishing returns around k=4.

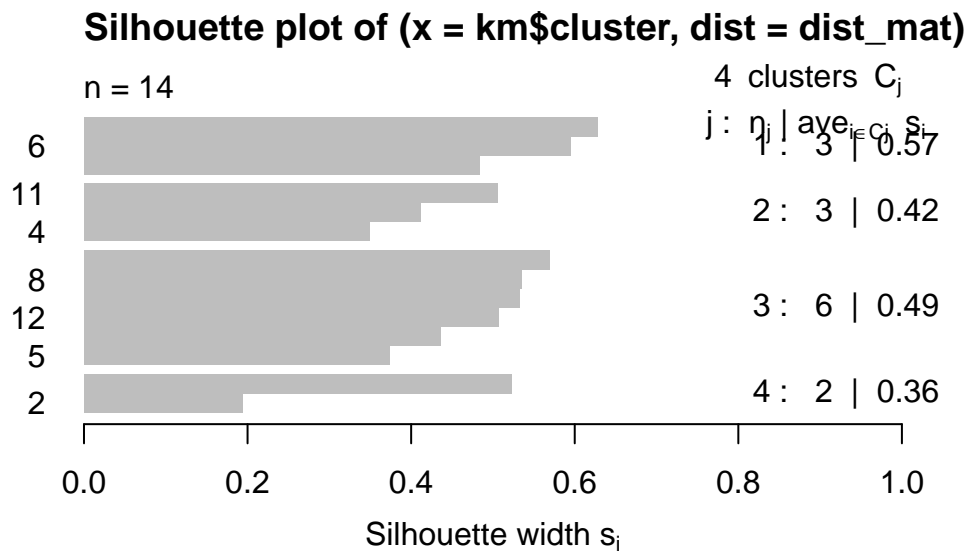
```
factoextra::fviz_nbclust(scaled, kmeans, method = "silhouette")
```



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.

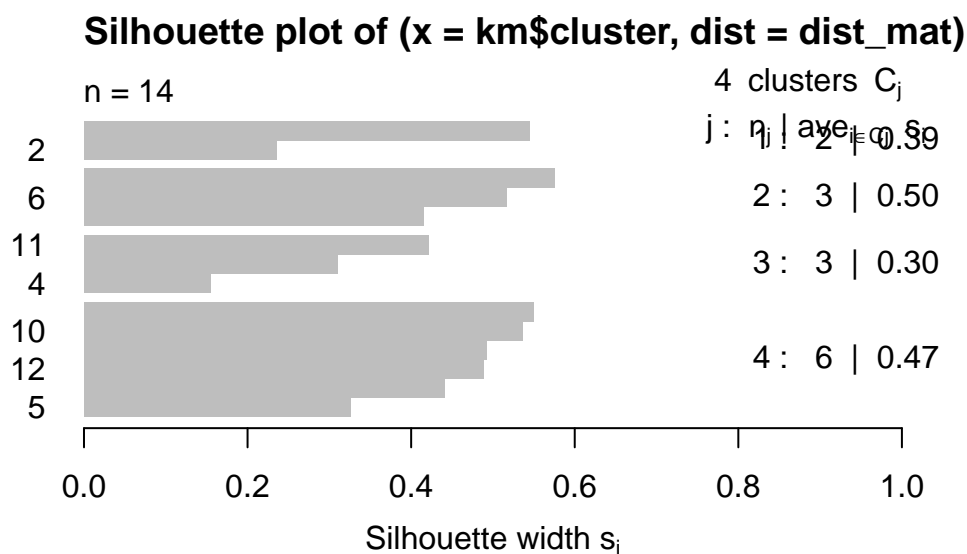
```
km <- kmeans(scaled, centers = 4, nstart = 25)
dist_mat <- dist(scaled)
sil <- cluster::silhouette(km$cluster, dist_mat)
plot(sil)
```



Average silhouette width : 0.47

Hm. Looks... acceptable. 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.

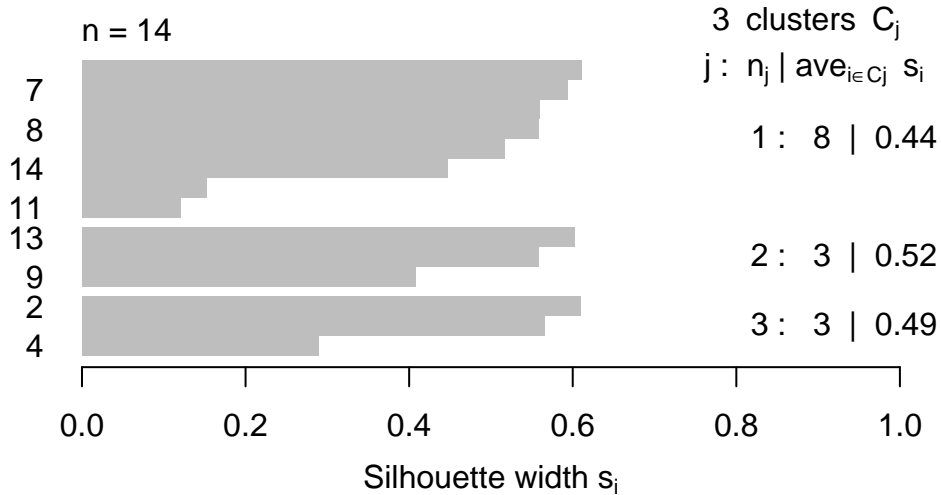
```
km <- kmeans(wide_counts, centers = 4, nstart = 25)
dist_mat <- dist(wide_counts)
sil <- cluster::silhouette(km$cluster, dist_mat)
plot(sil)
```



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

### Silhouette plot of (x = km\$cluster, dist = dist\_mat)



Average silhouette width : 0.47

With an average silhouette width of 0.43-0.47, our clusters aren't looking amazing. Still, the average silhouette score is 0.47 with unscaled data and  $k=3$ . This is not terrible. I think it's 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
```

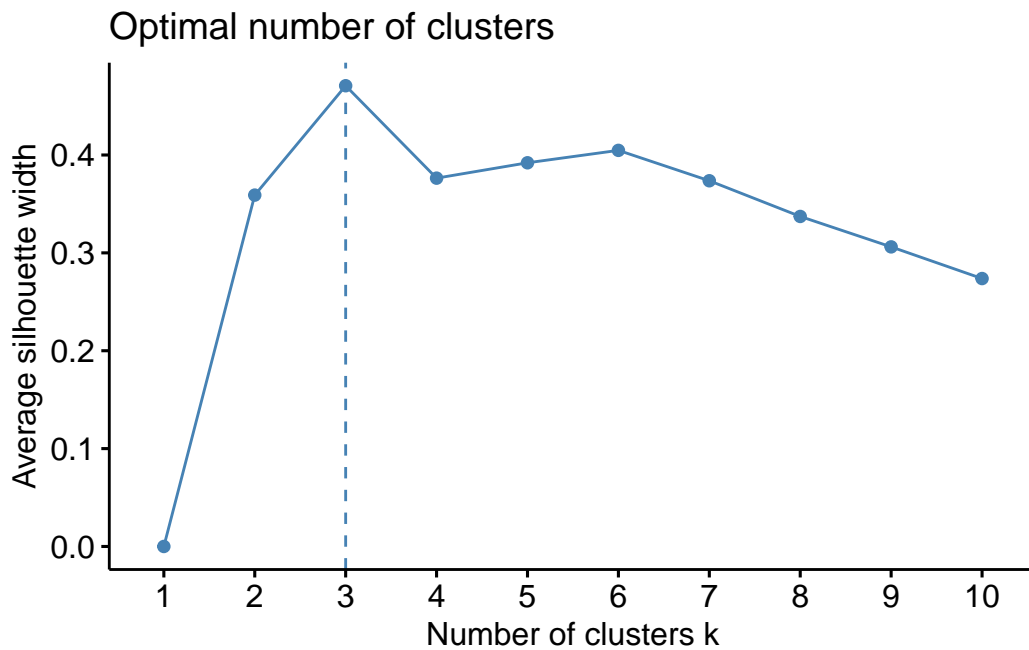


```
sort.km.cluster.      challenge
      2 Finding funding
      2 Hiring
      2 Securing funding

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.

## PCA (Abandoned)

In the above sections, I was clustering challenges into groups. Here, I started clustering people into groups. However, this didn’t seem too promising, and I don’t know if I care enough to pursue it. NOTE THE DIFFERENT CODING SCHEME

```
challnumeric <- challenges %>%
  mutate(
    across(
      everything(),
      ~ recode(
        .x,
        "Never" = 0L,
        "Non-applicable" = -1L, # THIS IS DIFFERENT (-1, not 0)
        "Rarely" = 1L,
        "Occasionally" = 2L,
        "Frequently" = 3L,
```

```

    "Always" = 4L
  )
)
)

pca <- prcomp(challnumeric, scale = TRUE)
summary(pca)

```

Importance of components:

	PC1	PC2	PC3	PC4	PC5	PC6	PC7
Standard deviation	2.1561	1.3366	1.1930	1.06788	0.9642	0.8606	0.79525
Proportion of Variance	0.3321	0.1276	0.1017	0.08146	0.0664	0.0529	0.04517
Cumulative Proportion	0.3321	0.4597	0.5613	0.64279	0.7092	0.7621	0.80726
	PC8	PC9	PC10	PC11	PC12	PC13	PC14
Standard deviation	0.74086	0.70184	0.67486	0.64866	0.62517	0.56317	0.26958
Proportion of Variance	0.03921	0.03518	0.03253	0.03005	0.02792	0.02265	0.00519
Cumulative Proportion	0.84647	0.88165	0.91418	0.94424	0.97215	0.99481	1.00000

```
biplot(pca)
```

