Script 3a

Pain progression: group data

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Analysis notes

Definitions of missingness

Data were regarded as **missing** when *pain in the last week* data were not present for one or more of weeks 0, 12, 24, 36, 48. Data also were classified as **missing** when there were inconsistencies in the data across the variables collected within a week.

Definition of data inconsistencies

Pain was defined as pain in the last week being 'Yes', and pain at its worst being > 0. These two measurements were then the "gatekeeper" measurements, such that the two measurements both had to be positive ('Yes' and '> 0', respectively) in order for there to be any entries for site of pain and site of worst pain. Were the data were inconsistent (e.g., when

there was no pain in the last week and pain at its worst = 0, but there were entries for site of pain and site of worst pain), then the site of pain and site of worst pain entries were marked as **inconsistent**.

Data also were considered **inconsistent** when pain in the last week = 'Yes', but site of worst pain = 'None'.

Lastly, data were considered **inconsistent** when *site of worst pain* was not listed as one of the pain locations for a given measurement week.

For analysis purposes, missing data in the *site of pain* columns were changed to 'No' (pain not present in the site). This approach was conservative, but we believed that the approach would have the least effect on the outcome, while still retaining as many participants as possible.

Import data

Quick look

```
head(df)
## # A tibble: 6 x 4
##
     ranid
             interval_name pain_in_the_last_week any_missing
##
     <chr>>
             <ord>
                            <chr>>
                                                   <chr>>
## 1 01-0001 0 weeks
                            No
                                                   No
## 2 01-0001 12 weeks
                            No
                                                   No
## 3 01-0001 24 weeks
                            No
                                                   No
## 4 01-0001 36 weeks
                            No
                                                   No
## 5 01-0001 48 weeks
                            No
                                                   No
## 6 01-0002 0 weeks
                            No
                                                   No
glimpse(df)
## Observations: 5,265
## Variables: 4
## $ ranid
                            <chr> "01-0001", "01-0001", "01-0001", "01-0001", "...
## $ interval_name
                            <ord> 0 weeks, 12 weeks, 24 weeks, 36 weeks, 48 wee...
## $ pain_in_the_last_week <chr> "No", "No", "No", "No", "No", "No", "No", "Yes", "Y...
                            <chr> "No", "No", "No", "No", "No", "No", "No", "No...
## $ any_missing
```

Basic clean

Quick tabulation

Analysis data set for the period 0 to 48 weeks

```
# Tabulate data
xtabs(~interval_name, data = df)

## interval_name
## 0 weeks 12 weeks 24 weeks 36 weeks 48 weeks
## 787 787 787 787 787
```

Analysis

Prepare data

Add a dummy frequency column for sankey diagram.

```
# Add dummy freq
df_sankey <- df %>%
  mutate(freq = 1)
```

Numeric summary: All

```
# Tabulate data
tab_prop1 <- as.data.frame(xtabs(~time_weeks + pain_in_the_last_week, data = df)) %>%
    pivot_wider(names_from = pain_in_the_last_week,
                values_from = Freq,
                names_prefix = 'pain_') %>%
    mutate(total_cases = rowSums(.[2:3])) %>%
    mutate(point_estimate = pain_Yes / total_cases) %>%
    select(time_weeks, total_cases, everything()) %>%
    mutate(time_weeks = as.numeric(as.character(time_weeks)))
# Calculate 95%CI
# Boot function
booted <- function(d, i, prop_factor = 'Yes'){</pre>
    data <- d[i, ]
    data2 <- data$pain_in_the_last_week</pre>
    prop <- mean(data2 == prop_factor)</pre>
    prop
}
# Set the seed
```

```
set.seed(2019)
# Bootstrap values
tab_prop2 <- df %>%
   group_by(time_weeks) %>%
   nest() %>%
   mutate(data_boot = map(.x = data,
                           ~ boot(data = .x,
                                  R = 5000,
                                  statistic = booted,
                                  stype = 'i'))) %>%
   mutate(data_ci = map(.x = data_boot,
                         ~ boot.ci(.x,
                                   type = 'basic'))) %>%
   mutate(data_point = map(.x = data_boot,
                            ~ .x$t0),
           data_ci.lower = map(.x = data_ci,
                               ~ .x$basic[[4]]),
           data_ci.upper = map(.x = data_ci,
                               ~ .x$basic[[5]]))
# Extract bootstrapped data
tab_prop2 %<>%
   select(time_weeks, data_point, data_ci.lower, data_ci.upper) %>%
   unnest(cols = c('data_point', 'data_ci.lower', 'data_ci.upper')) %>%
   rename(point_estimate = data_point,
           lower_CI = data_ci.lower,
           upper_CI = data_ci.upper)
# Join tab_prop1 and tab_prop2
tab_prop3 <- tab_prop1 %>%
   left_join(tab_prop2) %>%
   mutate(time_weeks = factor(time_weeks))
# Tabulate
kable(tab_prop3,
      caption = 'Estimate and 95% CI of the proportion with pain by week')
```

Table 1: Estimate and 95% CI of the proportion with pain by week

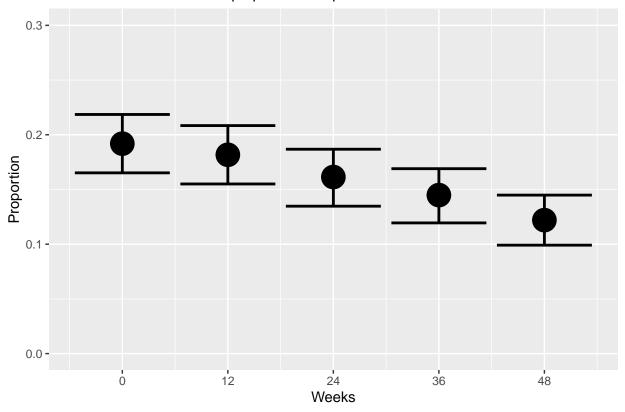
time_weeks	total_cases	pain_No	pain_Yes	point_estimate	lower_CI	upper_CI
0	787	636	151	0.1918679	0.1651842	0.2185515
12	787	644	143	0.1817027	0.1550191	0.2083863
24	787	660	127	0.1613723	0.1347206	0.1867853
36	787	673	114	0.1448539	0.1194409	0.1689962
48	787	691	96	0.1219822	0.0991105	0.1448539

Point plot with 95% CI: All

```
# Plot bootstrapped data
ggplot(data = tab_prop3) +
   aes(x = as.numeric(as.character(time_weeks)),
        y = point_estimate,
        ymin = lower_CI,
        ymax = upper_CI) +
        geom_errorbar(size = 1) +
```

```
geom_point(size = 8) +
scale_y_continuous(limits = c(0, 0.3)) +
scale_x_continuous(breaks = c(0, 12, 24, 36, 48)) +
labs(subtitle = 'Estimate and 95% CI of the proportion with pain',
        x = 'Weeks',
        y = 'Proportion')
```

Estimate and 95% CI of the proportion with pain

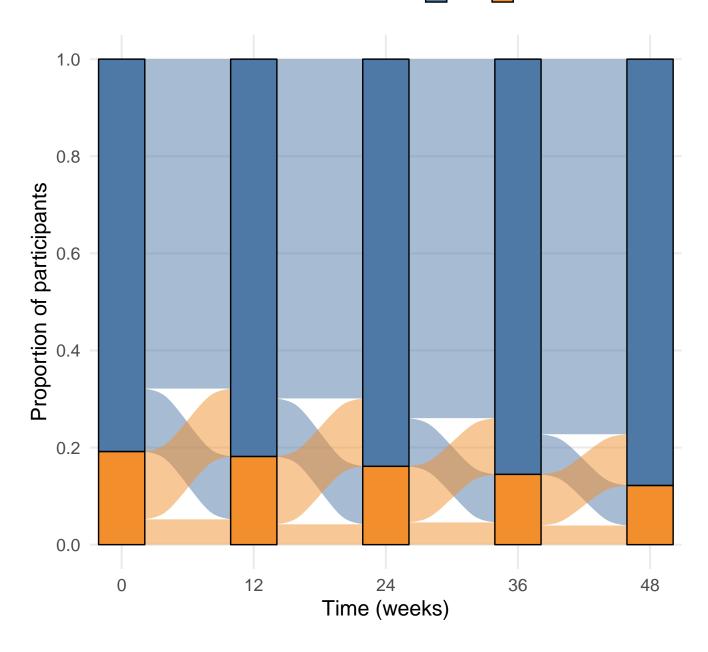


Sankey plot 1: All

```
# Breaks
n <- length(unique(df$ranid))</pre>
# Generate plot for publication
sankey_plot1 <- ggplot(data = df_sankey) +</pre>
    aes(x = factor(time_weeks),
        stratum = pain_in_the_last_week,
        alluvium = ranid,
        y = freq,
        fill = pain_in_the_last_week,
        label = pain_in_the_last_week) +
    geom_flow() +
    geom_stratum(width = 0.35) +
    scale_fill_tableau(name = 'Pain in the last week') +
    scale_y_continuous(breaks = c(n * 0, n * 0.2, n * 0.4, n * 0.6, n * 0.8, n),
                       labels = c('0.0', '0.2', '0.4', '0.6', '0.8', '1.0')) +
    scale_x_discrete(expand = c(0.06, 0)) +
    labs(x = 'Time (weeks)',
         y = 'Proportion of participants') +
    theme_minimal(base_size = 20) +
    theme(legend.position = 'top',
```

```
panel.grid.minor = element_blank())
# Save plot
ggsave(filename = 'figures/figure-2.png', sankey_plot1,
       width = 10.2, height = 7.72)
# Generate plot for knitr output
ggplot(data = df_sankey) +
   aes(x = factor(time_weeks),
        stratum = pain_in_the_last_week,
        alluvium = ranid,
       y = freq,
       fill = pain_in_the_last_week,
       label = pain_in_the_last_week) +
   geom_flow() +
   geom_stratum(width = 0.35) +
   scale_fill_tableau(name = 'Pain in the last week') +
   scale_y = continuous(breaks = c(n * 0, n * 0.2, n * 0.4, n * 0.6, n * 0.8, n),
                       labels = c('0.0', '0.2', '0.4', '0.6', '0.8', '1.0')) +
   scale_x_discrete(expand = c(0.06, 0)) +
   labs(x = 'Time (weeks)',
        y = 'Proportion of participants') +
   theme_minimal(base_size = 16) +
   theme(legend.position = 'top',
         panel.grid.minor = element_blank())
```



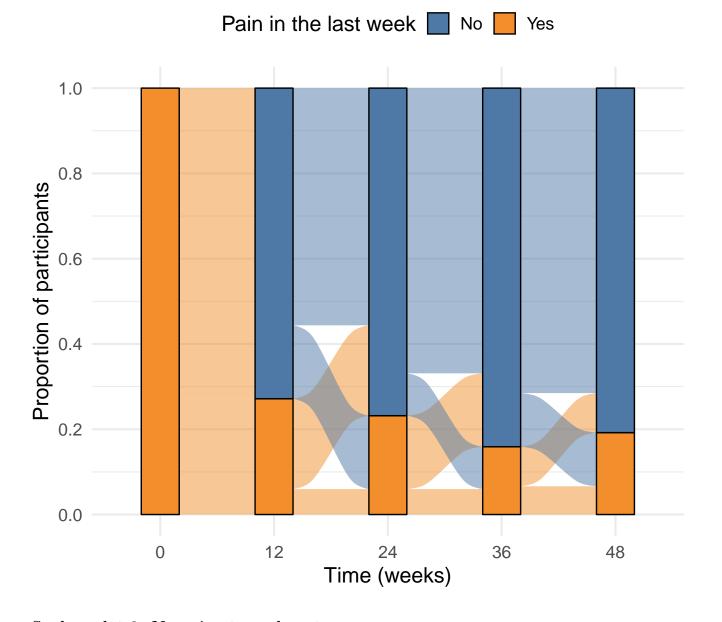


Sankey plot 2: Pain at enrolment

```
# Breaks
n2 <- length(unique(df$ranid[df$pain_at_enrolment == 'Yes']))

# Generate plot
sankey_plot2 <- df_sankey %>%
    select(-interval_name) %>%
    filter(pain_at_enrolment == 'Yes') %>%
    ggplot(data = .) +
    aes(x = factor(time_weeks),
        stratum = pain_in_the_last_week,
        alluvium = ranid,
        y = freq,
        fill = pain_in_the_last_week,
        label = pain_in_the_last_week) +
```

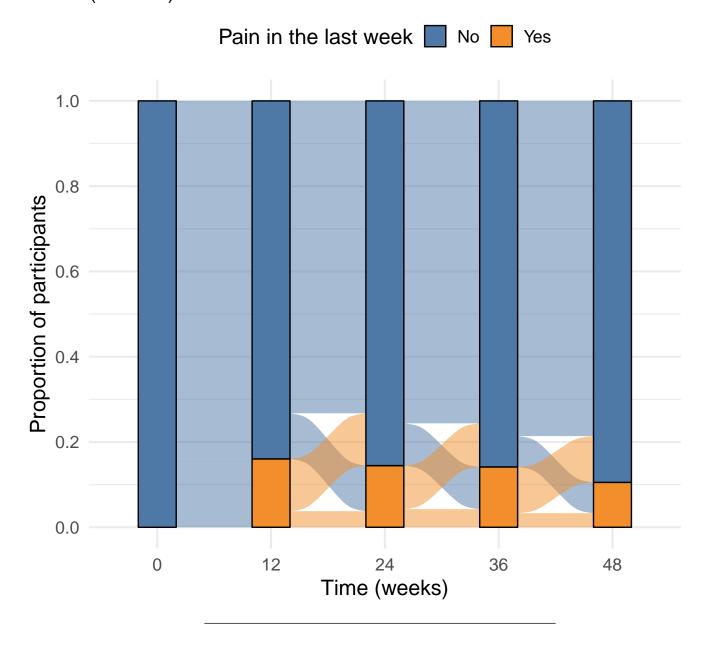
Participants with pain at enrolment (n = 151)



Sankey plot 3: No pain at enrolment

```
# Breaks
n3 <- length(unique(df$ranid[df$pain_at_enrolment == 'No']))
# Generate plot
sankey_plot3 <- df_sankey %>%
    select(-interval_name) %>%
    filter(pain_at_enrolment == 'No') %>%
    ggplot(data = .) +
    aes(x = factor(time_weeks),
        stratum = pain_in_the_last_week,
        alluvium = ranid,
        y = freq,
        fill = pain_in_the_last_week,
        label = pain_in_the_last_week) +
    geom_flow() +
    geom_stratum() +
    scale_fill_tableau(name = 'Pain in the last week') +
    scale_y = continuous(breaks = c(n3 * 0, n3 * 0.2, n3 * 0.4, n3 * 0.6, n3 * 0.8, n3),
                       labels = c('0.0', '0.2', '0.4', '0.6', '0.8', '1.0')) +
    labs(title = 'Participants with no pain at enrolment',
         subtitle = str_glue("(n = {length(unique(df$ranid[df$pain_at_enrolment == 'No']))})"),
         x = 'Time (weeks)',
         y = 'Proportion of participants') +
    theme_minimal(base_size = 16) +
    theme(legend.position = 'top'); sankey_plot3
```

Participants with no pain at enrolment (n = 636)



Number of people transitioning between pain states at each time period Separate time points

```
# Week 0 to 12
t0.12 <- df %>%
    filter(time_weeks %in% c(0, 12)) %>%
    select(-interval_name, -pain_at_enrolment)

# Week 12 to 24
t12.24 <- df %>%
    filter(time_weeks %in% c(12, 24)) %>%
    select(-interval_name, -pain_at_enrolment)
```

```
# Week 24 to 36
t24.36 <- df %>%
    filter(time_weeks %in% c(24, 36)) %>%
    select(-interval_name, -pain_at_enrolment)

# Week 36 to 48
t36.48 <- df %>%
    filter(time_weeks %in% c(36, 48)) %>%
    select(-interval_name, -pain_at_enrolment)
```

Calculate percent transitioning pain state

Table 2: Pain transitions: week 0 to 12

pain_transition	count	prop	percent
No pain to pain	102	0.1296061	13
Pain to no pain	110	0.1397713	14

Table 3: Pain transitions: week 12 to 24

pain_transition	count	prop	percent
No pain to pain	94	0.1194409	11.9
Pain to no pain	110	0.1397713	14.0

Table 4: Pain transitions: week 24 to 36

pain_transition	count	prop	percent
No pain to pain	78	0.0991105	9.9
Pain to no pain	91	0.1156290	11.6

Table 5: Pain transitions: week 36 to 38

pain_transition	count	prop	percent
No pain to pain	65	0.0825921	8.3
Pain to no pain	83	0.1054638	10.5

Session information

```
## R version 3.6.1 (2019-07-05)
## Platform: x86_64-apple-darwin15.6.0 (64-bit)
## Running under: macOS Mojave 10.14.6
##
## Matrix products: default
```

```
/Library/Frameworks/R.framework/Versions/3.6/Resources/lib/libRblas.0.dylib
## BLAS:
## 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:
                          patchwork_1.0.0
##
   [1] knitr_1.27
                                            boot_1.3-24
                                                              ggthemes_4.2.0
##
   [5] ggalluvial_0.11.1 magrittr_1.5
                                            forcats_0.4.0
                                                               stringr_1.4.0
   [9] dplyr_0.8.3
                          purrr_0.3.3
                                            readr_1.3.1
                                                              tidyr_1.0.0
## [13] tibble_2.1.3
                          ggplot2_3.2.1
                                            tidyverse_1.3.0
##
## loaded via a namespace (and not attached):
   [1] tidyselect_0.2.5 xfun_0.12
                                          haven_2.2.0
                                                            lattice_0.20-38
   [5] colorspace_1.4-1 vctrs_0.2.1
##
                                          generics_0.0.2
                                                           htmltools_0.4.0
## [9] yaml_2.2.0
                         utf8_1.1.4
                                          rlang_0.4.2
                                                           pillar_1.4.3
## [13] withr 2.1.2
                         glue 1.3.1
                                          DBI 1.1.0
                                                            dbplyr 1.4.2
## [17] modelr_0.1.5
                         readxl_1.3.1
                                          plyr_1.8.5
                                                            lifecycle_0.1.0
## [21] munsell_0.5.0
                         gtable_0.3.0
                                          cellranger_1.1.0 rvest_0.3.5
## [25] evaluate_0.14
                         labeling_0.3
                                          fansi_0.4.1
                                                           highr_0.8
## [29] broom_0.5.3
                         Rcpp_1.0.3
                                          scales_1.1.0
                                                            backports_1.1.5
## [33] jsonlite_1.6
                         farver_2.0.3
                                          fs_1.3.1
                                                           hms_0.5.3
## [37] digest_0.6.23
                         stringi_1.4.5
                                          grid_3.6.1
                                                            cli_2.0.1
## [41] tools_3.6.1
                         lazyeval_0.2.2
                                          crayon_1.3.4
                                                            pkgconfig_2.0.3
## [45] zeallot_0.1.0
                         xm12_1.2.2
                                          reprex_0.3.0
                                                           lubridate_1.7.4
## [49] assertthat_0.2.1 rmarkdown_2.1
                                          httr_1.4.1
                                                            rstudioapi_0.10
## [53] R6_2.4.1
                         nlme_3.1-143
                                          compiler_3.6.1
```