Supplement 1

Demographics

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1 Import and check data

```
# Import
demo <- read_rds('data-cleaned/data-demographics.rds')%>%
    # Fix site names:
   ## DD = Site 1, NM = Site 2, RESI = Site 3, RP = Site 4, STIG = Site 5
   mutate(Site = case_when(
       Site == 'DD' ~ 'Site 1',
       Site == 'NM' ~ 'Site 2',
       Site == 'RESI' ~ 'Site 3',
       Site == 'RP' ~ 'Site 4',
       Site == 'STIG' ~ 'Site 5'
   ))
# Check
## Demographics
dim(demo)
## [1] 596
names (demo)
## [1] "ID"
                                                "Sex"
## [4] "Age"
                           "Employment_status" "CD4_recent"
## [7] "ART_currently"
                           "Education"
```

glimpse(demo)

```
## Rows: 596
## Columns: 8
## $ ID
                      <chr> "RPB73", "RPB74", "RPB75", "RPB76", "RPB77", "RPB...
                      <chr> "Site 4", "Site 4", "Site 4", "Site 4", "Site 4", ...
## $ Site
                      <chr> "Female", "Female", "Female", "Female", "Female", ...
## $ Sex
                      <dbl> 36, 27, 39, 36, 31, 32, 28, 37, 31, 25, 31, 24, 3...
## $ Age
## $ Employment_status <chr> "Other", "Unemployed", "Other", "Unemployed", "Un...
                      <dbl> 391, 571, 591, 207, 126, 225, 543, 410, 74, 212, ...
## $ CD4 recent
                      <chr> "Yes", "Yes", "Yes", "Yes", "Yes", "Yes", "Yes", ...
## $ ART_currently
## $ Education
                      <chr> "Tertiary", "Secondary", "Secondary", "Primary", ...
```

2 Basic descriptive statistics

2.1 Full cohort

2.1.1 Descriptives

```
demo %>%
    # Remove ID column
select(-ID, -Site) %>%
    # Convert character to factor
mutate_if(is.character, factor) %>%
my_skim()
```

Table 1: Data summary

| Name Number of rows | Piped data 596 |
|------------------------|-------------------|
| Number of columns | 6 |
| Column type frequency: | |
| factor | 4 |
| numeric | 2 |
| Group variables | None |

Variable type: factor

| skim_variable | n_missing | complete_rate | n_unique | top_counts |
|-------------------|-----------|---------------|----------|--------------------------------------|
| Sex | 0 | 1.00 | 2 | Fem: 481, Mal: 115 |
| Employment_status | 49 | 0.92 | 4 | Une: 330, Ful: 131, Par: 52, Oth: 34 |
| ART_currently | 5 | 0.99 | 2 | Yes: 460, No: 131 |
| Education | 37 | 0.94 | 3 | Sec: 395, Pri: 99, Ter: 65 |

Variable type: numeric

| skim_variable | n_missing | complete_rate | mean | sd | p0 | p25 | p50 | p75 | p100 |
|---------------|-----------|---------------|--------|---------------------|----|-----|-----|-----|------|
| Age | 8 | 0.99 | 37.28 | 9.06 | 19 | 31 | 36 | 42 | 76 |
| CD4_recent | 99 | 0.83 | 320.71 | 238.92 | 1 | 155 | 261 | 432 | 1232 |

2.2 By study site

2.2.1 Sample size

```
demo %>%
    # Select ID
    select(Site) %>%
    # Group by study site
    group_by(Site) %>%
    summarise(Count = n()) %>%
    kable(caption = 'Sample size by study site')
```

Table 4: Sample size by study site

| Site | Count |
|--------|-------|
| Site 1 | 60 |
| Site 2 | 239 |
| Site 3 | 99 |
| Site 4 | 148 |
| Site 5 | 50 |
| | |

2.2.2 Descriptives

```
demo %>%
    # Convert character to factor
    mutate_if(is.character, factor) %>%
    # Remove ID
    select(-ID) %>%
    # Group by site
    group_by(Site) %>%
    my_skim()
```

Table 5: Data summary

| Name | Piped data |
|------------------------|------------|
| Number of rows | 596 |
| Number of columns | 7 |
| Column type frequency: | |
| factor | 4 |
| numeric | 2 |
| Group variables | Site |

Variable type: factor

| skim_variable | Site | n_missing | $complete_rate$ | n_unique | top_counts |
|----------------------|--------|-----------|------------------|----------|-----------------------------------|
| Sex | Site 1 | 0 | 1.00 | 2 | Fem: 39, Mal: 21 |
| Sex | Site 2 | 0 | 1.00 | 2 | Fem: 184, Mal: 55 |
| Sex | Site 3 | 0 | 1.00 | 2 | Fem: 66, Mal: 33 |
| Sex | Site 4 | 0 | 1.00 | 1 | Fem: 148, Mal: 0 |
| Sex | Site 5 | 0 | 1.00 | 2 | Fem: 44, Mal: 6 |
| $Employment_status$ | Site 1 | 0 | 1.00 | 4 | Une: 36, Ful: 17, Par: 6, Oth: 1 |
| $Employment_status$ | Site 2 | 49 | 0.79 | 3 | Une: 126, Ful: 55, Par: 9, Oth: 0 |
| $Employment_status$ | Site 3 | 0 | 1.00 | 4 | Une: 47, Ful: 35, Par: 16, Oth: 1 |
| $Employment_status$ | Site 4 | 0 | 1.00 | 4 | Une: 99, Oth: 31, Par: 14, Ful: 4 |
| $Employment_status$ | Site 5 | 0 | 1.00 | 4 | Une: 22, Ful: 20, Par: 7, Oth: 1 |
| ART_currently | Site 1 | 0 | 1.00 | 2 | Yes: 58, No: 2 |
| ART_currently | Site 2 | 3 | 0.99 | 2 | Yes: 130, No: 106 |
| ART_currently | Site 3 | 1 | 0.99 | 2 | Yes: 95, No: 3 |
| ART_currently | Site 4 | 0 | 1.00 | 2 | Yes: 128, No: 20 |
| ART_currently | Site 5 | 1 | 0.98 | 1 | Yes: 49, No: 0 |
| Education | Site 1 | 2 | 0.97 | 3 | Sec: 40, Pri: 9, Ter: 9 |
| Education | Site 2 | 27 | 0.89 | 3 | Sec: 160, Pri: 44, Ter: 8 |
| Education | Site 3 | 4 | 0.96 | 3 | Sec: 61, Pri: 19, Ter: 15 |
| Education | Site 4 | 2 | 0.99 | 3 | Sec: 108, Ter: 20, Pri: 18 |
| Education | Site 5 | 2 | 0.96 | 3 | Sec: 26, Ter: 13, Pri: 9 |

Variable type: numeric

| skim_variable | Site | n_missing | $complete_rate$ | mean | sd | p0 | p25 | p50 | p75 | p100 |
|------------------------|--------|-----------|------------------|--------|---------------------|----|--------|-------|--------|------|
| Age | Site 1 | 0 | 1.00 | 38.63 | 5.87 | 27 | 34.00 | 38.5 | 42.00 | 58 |
| Age | Site 2 | 6 | 0.97 | 36.73 | 8.54 | 20 | 31.00 | 36.0 | 41.00 | 76 |
| Age | Site 3 | 0 | 1.00 | 43.40 | 9.32 | 19 | 35.00 | 42.0 | 49.00 | 66 |
| Age | Site 4 | 2 | 0.99 | 30.82 | 4.72 | 20 | 27.00 | 31.0 | 34.00 | 40 |
| Age | Site 5 | 0 | 1.00 | 44.90 | 9.67 | 25 | 37.25 | 44.5 | 52.50 | 65 |
| $CD4_recent$ | Site 1 | 18 | 0.70 | 428.60 | 255.93 | 16 | 256.00 | 374.5 | 577.25 | 1095 |
| $CD4_recent$ | Site 2 | 37 | 0.85 | 222.10 | 180.15 | 1 | 103.75 | 185.5 | 308.75 | 963 |
| $CD4$ _recent | Site 3 | 16 | 0.84 | 452.76 | 278.07 | 11 | 256.50 | 405.0 | 651.00 | 1232 |
| $CD4$ _recent | Site 4 | 4 | 0.97 | 322.19 | 205.43 | 26 | 174.00 | 270.0 | 415.00 | 1040 |
| $\mathrm{CD4}$ _recent | Site 5 | 24 | 0.52 | 482.88 | 292.90 | 50 | 291.00 | 434.5 | 582.00 | 1138 |

3 Session information

sessionInfo()

```
## R version 3.6.3 (2020-02-29)
## Platform: x86_64-pc-linux-gnu (64-bit)
## Running under: Debian GNU/Linux 10 (buster)
##
## Matrix products: default
## BLAS/LAPACK: /usr/lib/x86_64-linux-gnu/libopenblasp-r0.3.5.so
##
## locale:
   [1] LC CTYPE=en US.UTF-8
                                   LC NUMERIC=C
##
                                   LC_COLLATE=en_US.UTF-8
  [3] LC TIME=en US.UTF-8
   [5] LC MONETARY=en US.UTF-8
                                   LC MESSAGES=C
##
   [7] LC_PAPER=en_US.UTF-8
                                   LC_NAME=C
##
  [9] LC_ADDRESS=C
                                   LC_TELEPHONE=C
## [11] LC MEASUREMENT=en US.UTF-8 LC IDENTIFICATION=C
## attached base packages:
## [1] stats
                 graphics grDevices utils
                                               datasets methods
                                                                    base
##
## other attached packages:
   [1] knitr_1.28
                        skimr_2.1.1
                                        forcats_0.5.0
                                                         stringr_1.4.0
##
   [5] dplyr_0.8.5
                        purrr_0.3.4
                                        readr_1.3.1
                                                         tidyr_1.0.2
   [9] tibble_3.0.1
                        ggplot2_3.3.0
                                        tidyverse_1.3.0
##
## loaded via a namespace (and not attached):
  [1] tidyselect_1.0.0 xfun_0.13
                                          repr_1.1.0
                                                           haven_2.2.0
  [5] lattice 0.20-38 colorspace 1.4-1 vctrs 0.2.4
                                                           generics 0.0.2
## [9] htmltools_0.4.0 base64enc_0.1-3 yaml_2.2.1
                                                           utf8_1.1.4
## [13] rlang_0.4.5
                         pillar_1.4.3
                                          glue_1.4.0
                                                           withr 2.2.0
## [17] DBI_1.1.0
                         dbplyr_1.4.3
                                          modelr_0.1.6
                                                           readxl_1.3.1
## [21] lifecycle_0.2.0 munsell_0.5.0
                                          gtable_0.3.0
                                                           cellranger_1.1.0
                                          fansi 0.4.1
## [25] rvest 0.3.5
                         evaluate 0.14
                                                           highr 0.8
                                                           backports_1.1.6
## [29] broom_0.5.6
                         Rcpp_1.0.4.6
                                          scales_1.1.0
## [33] jsonlite_1.6.1
                         fs_1.4.1
                                          hms_0.5.3
                                                           digest_0.6.25
## [37] stringi_1.4.6
                         grid_3.6.3
                                          cli_2.0.2
                                                           tools_3.6.3
## [41] magrittr_1.5
                         crayon_1.3.4
                                          pkgconfig_2.0.3
                                                           ellipsis_0.3.0
## [45] xml2_1.3.2
                         reprex_0.3.0
                                          lubridate_1.7.8
                                                           assertthat_0.2.1
## [49] rmarkdown_2.1
                         httr_1.4.1
                                          rstudioapi_0.11
                                                           R6_2.4.1
## [53] nlme_3.1-144
                         compiler_3.6.3
```