# Geriatric Syndrome in Older Adults with AF

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#### **SETUP**

- load packages
- · set figure style

```
knitr::opts_chunk$set(echo = TRUE)

## Load the libraries we will be using
library(tidyverse)
```

```
## -- Attaching packages ------ tidyverse 1.3.0 --
```

```
## v ggplot2 3.2.1 v purrr 0.3.3

## v tibble 2.1.3 v dplyr 0.8.3

## v tidyr 1.0.2 v stringr 1.4.0

## v readr 1.3.1 v forcats 0.4.0
```

```
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag() masks stats::lag()
```

```
library(ggthemes)
library(haven)
      Color Palette by Paletton.com 30degrees
##### Palette URL: http://paletton.com/#uid=7070u0ksyu7hHCkmJvyuMo9zZj4
##### Color Palette by Paletton.com 150 degrees
##### Palette URL: http://paletton.com/#uid=7072m0ksyu7hHCkmJvyuMo9zZj4
## FIGURE SETTINGS
sachin_theme <- theme_bw()+</pre>
        theme(
          panel.background = element_blank(),
          axis.line = element_line(colour = "grey75"),
          panel.border = element_blank(),
          panel.grid.major.x = element blank(),
          panel.grid.minor.x = element_blank(),
          panel.grid.major.y = element line(color = "grey", linetype = "dotted" ),
          panel.grid.minor.y = element blank(),
          legend.position = "none",
          axis.title=element_text(size=10)
          )
sachin_theme_leg <- theme_bw()+</pre>
        theme(
          panel.background = element_blank(),
          axis.line = element line(colour = "grey75"),
          panel.border = element blank(),
          panel.grid.major.x = element blank(),
          panel.grid.minor.x = element_blank(),
          panel.grid.major.y = element_line(color = "grey", linetype = "dotted" ),
          panel.grid.minor.y = element_blank(),
          legend.position = "top",
          legend.title = element blank(),
          axis.title=element_text(size=10)
          )
```

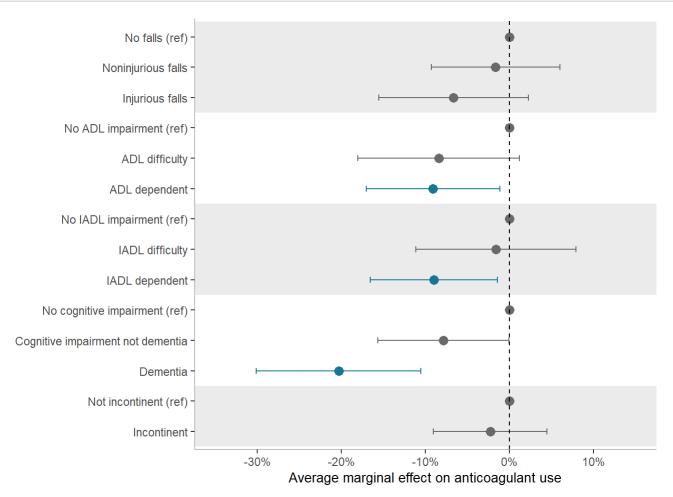
## Figure: Relationship between AC use and geriatric syndrome count

- · analysis done in SAS and saved in dataset thinner ame.sas7dbat
- imported to R to create figure

```
### read in data
library(haven)
ame <- read sas("C:/Users/sachi/Box Sync/AF frailty/tables and figures/thinner ame.sas7bdat",
    NULL)
### Add in reference rows
library(tibble)
toadd_names <- c("Falls", "ADL", "IADL", "Cognitive function", "Incontinence")</pre>
toadd_levels <- c("No falls (ref)", "No ADL impairment (ref)", "No IADL impairment (ref)", "No c
ognitive impairment (ref)", "Not incontinent (ref)")
ame <- add row(ame, syndrome = toadd names, Diff = 0, LL = 0, UL = 0, level = toadd levels)
### create factors for syndrome and lables so that we get the right order
ame$syndrome <- as.factor(ame$syndrome)</pre>
ame$syndrome <- factor(ame$syndrome, levels = c("Falls", "ADL", "IADL", "Cognitive function", "I</pre>
ncontinence"))
color level5 <- c("#FB6549", "#35B862", "#32829D", "#D53E80", "#FBA449")
color sig <- c("#696969", "#157696")
ame$level <- as.factor(ame$level)</pre>
ame$level <- factor(ame$level, levels = c("No falls (ref)", "Noninjurious falls", "Injurious fall</pre>
s", "No ADL impairment (ref)", "ADL difficulty", "ADL dependent", "No IADL impairment (ref)", "I
ADL difficulty", "IADL dependent", "No cognitive impairment (ref)", "Cognitive impairment not de
mentia", "Dementia", "Not incontinent (ref)", "Incontinent"))
### Add columns to aid in the shading of the figure
ame <- ame %>% mutate(Xn=as.numeric(fct rev(level)))
shade_cat <- c("Falls", "IADL", "Incontinence")</pre>
ame <- mutate(ame, shade = ifelse(syndrome %in% shade_cat, "gray", "white"))</pre>
ame <- mutate(ame, significant = ifelse(round(UL,2) < 0, "sig", "notsig"))</pre>
### Create figure
library(stringr)
fig_ac_x_gs_ame <- ggplot(data=ame, aes(y=Diff, x=fct_rev(level), group = syndrome)) +</pre>
        geom_rect(aes(xmin=Xn-0.5, xmax = Xn+0.5, ymin = -Inf, ymax= Inf, fill = shade),
                  alpha = 0.3, stat="identity", show.legend = FALSE) +
                   scale_fill_manual(values = alpha(c("gray", "white"), 0.3)) +
        geom point(aes(color = significant), size=3)+
        geom_errorbar(aes(ymin=LL, ymax=UL, width = 0.2, color=significant)) +
        scale x discrete(name="") +
        scale_y_continuous(name = "Average marginal effect on anticoagulant use",
                            limits = c(-0.35, 0.15),
                           breaks = c(seq(from = -0.3, to = 0.1, by = 0.1)),
                            labels = scales::percent format(accuracy = 1)) +
        scale_color_manual(values = color_sig)+
        theme(
              panel.background = element_blank(),
```

```
axis.line = element_line(colour = "grey75"),
    panel.border = element_blank(),
    panel.grid.major.y = element_blank(),
    panel.grid.minor.x = element_blank(),
    panel.grid.major.x = element_blank(),
    panel.grid.minor.y = element_blank(),
    legend.position = "none",
    legend.title = element_blank(),
    axis.title=element_text(size=10),
    plot.caption = element_text(hjust = +0, face = "italic"),
    plot.title = element_text(hjust = +1)
    ) +

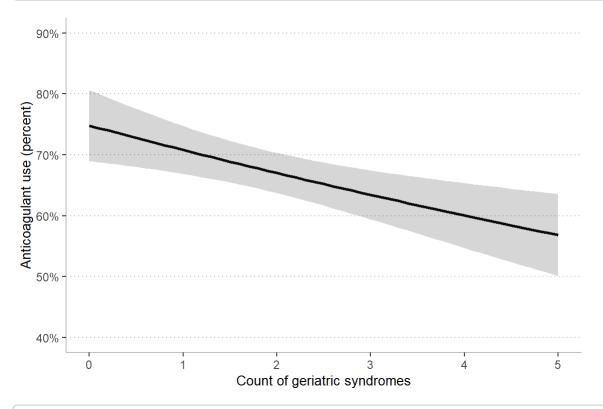
    geom_hline(yintercept = 0, linetype = "dashed") +
    coord_flip()
```



ggsave("C:/Users/sachi/Box Sync/AF frailty/tables and figures/fig gs x ac ame.png", width = 7, h eight = 4)

### Figure AC use by geriatric count

- analysis done in sas and saved to gs\_count\_x\_ac.sas7dbat
- sas output saved to dataset and imported to R



ggsave("C:/Users/sachi/Box Sync/AF frailty/tables and figures/fig\_ac\_x\_gscount.png", width = 6,
height = 4)

### Cohort flow diagram

```
library(DiagrammeR)
# Define some sample data
data \leftarrow list(a=10364, b=9249, c=4882, d=779, e = 776)
flow <- DiagrammeR::grViz("</pre>
digraph graph2 {
graph [layout = dot, fontsize = 11]
# node definitions with substituted label text
node [shape = rectangle, width = 1, fontname = Arial]
a [label = '@@1']
b [label = '@@2']
c [label = '@@3']
d [label = '@@4']
1 [label = '@@5']
2 [label = '@@6']
3 [label = '@@7']
4 [label = '@@8']
5 [label = '@@9']
a -> b -> c -> d
d \rightarrow \{1,2,3,4,5\}
}
[1]: paste0('65+ at 2014 HRS interview\\n', '(n = ', data$a, ')')
[2]: paste0('Agreed to Medicare claims linkage\\n', '(n = ', data$b, ')')
[3]: paste0('Enrolled in Medicare Part A and B for 24\\n', 'consecutive months before 2014 inter
view\\n', '(n = ', data$c, ')')
[4]: paste0('Met claims definition for atrial fibrillation\\n', '(n = ', data$d, ')')
[5]: paste0('Falls prevalence\\n', 'measure excludes\\n', '3 missing falls data\\n', '(n=776)
\\n', '\\n', 'Falls and AC\\n', 'analysis excludes\\n', 'an additional\\n', '3 missing
AC data\\n','(n=773)')
[6]: paste0('ADL prevalence\\n', 'measure excludes\\n', '3 missing ALD data\\n', '(n=776)\\n', '
\n', ' \n', 'ADL and AC\n', 'analysis excludes\n', 'an additional\n', '3 missing AC data
\\n','(n=773)')
[7]: paste0('IADL prevalence\n', 'measure\n', '(n=779)\n', ' \n', ' \n', ' \n', ' IADL and
AC\\n', 'analysis excludes\\n', '3 missing AC data\\n', '(n=776)\\n', '\\n')
[8]: paste0('Cognition prevalence\n', 'measure\n', '(n=779)\n', ' \n', ' \n', '\n', 'Cogni
tion and AC\\n', 'anlaysis excludes\\n', '3 missing AC data\\n', '(n=776)\\n', '\\n')
[9]: paste0('Incontinence prevalence\\n', 'measure excludes\\n', '3 missing\\n', 'incontinence d
ata\n', '(n=776)\n', ' \\n', 'Incontinence and AC\n', 'analysis excludes\n', 'an additional
\\n', '2 missing AC data\\n', '(n=774)')
")
flow
```

