$all_sites_sex_scatters$

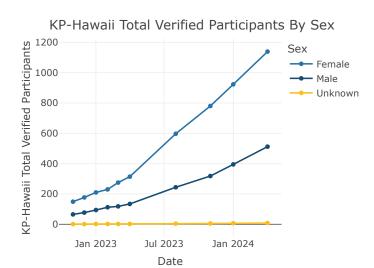
2024-10-03

Refactor Site Names

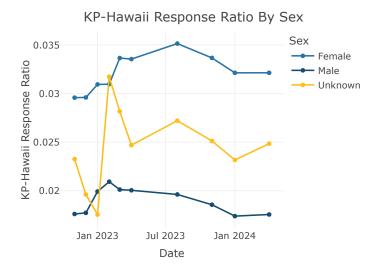
```
data$site <- gsub("^Marshfield.*", "Marshfield", data$site)</pre>
unique(data$site)
                                                         "KP-Northwest"
## [1] "KP-Hawaii"
                                "KP-Colorado"
## [4] "KP-Georgia"
                                "Marshfield"
                                                         "Sanford Health"
## [7] "Henry Ford"
                                "University of Chicago"
```

Write list of sites to be used

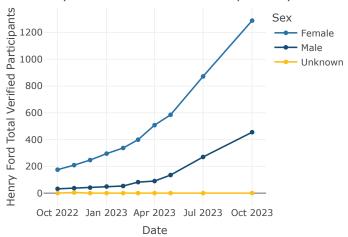
```
items_list = c("KP-Hawaii", "KP-Colorado", "KP-Northwest", "KP-Georgia", "Marshfield", "Sanford Health"
```

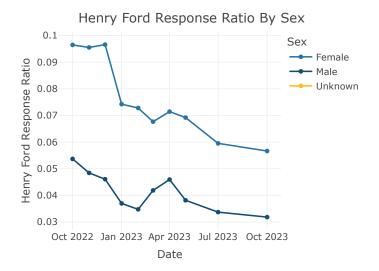


Call by $_sex_scatter$ function for all sites

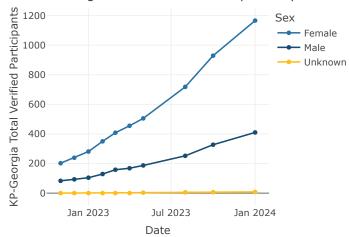


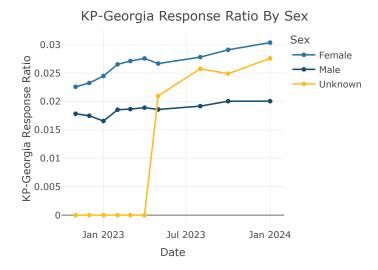
Henry Ford Total Verified Participants By Sex

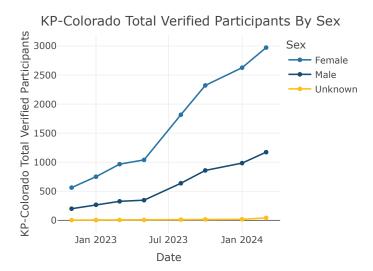


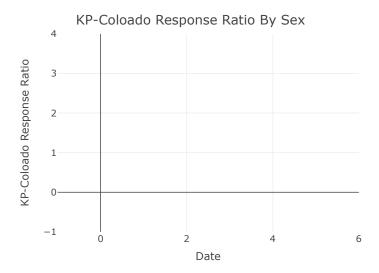


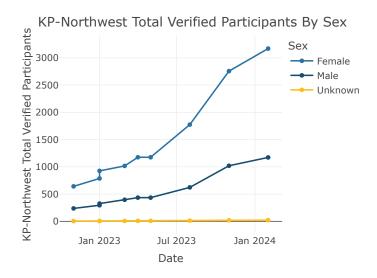
KP-Georgia Total Verified Participants By Sex

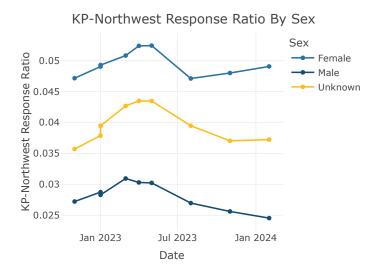


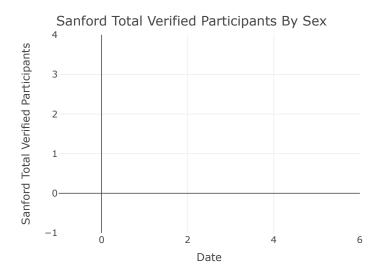


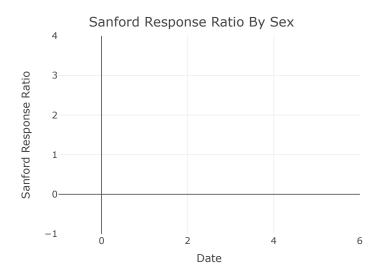


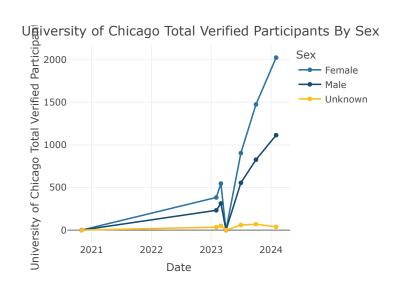




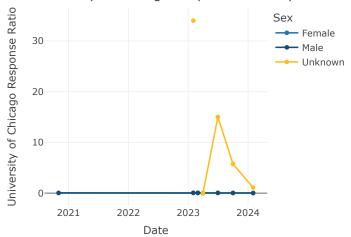




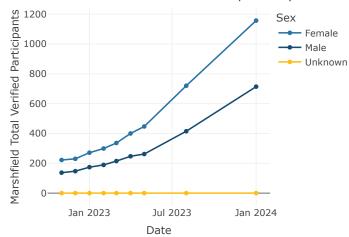


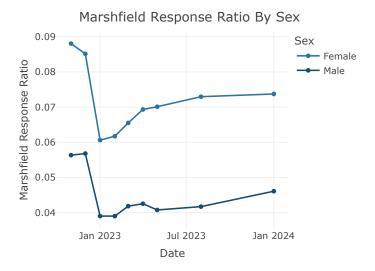


University of Chicago Response Ratio By Sex



Marshfield Total Verified Participants By Sex





race_plots(data, items_list) ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race ethnicity'. You can override using ## the '.groups' argument. ## 'summarise()' has grouped output by 'race_ethnicity'. You can override using ## the '.groups' argument.

'summarise()' has grouped output by 'race_ethnicity'. You can override using

'summarise()' has grouped output by 'race ethnicity'. You can override using

```
ins_plots(data, items_list)
```

```
ruca_plots(data, items_list)
```

the '.groups' argument.

the '.groups' argument.

```
ses_plots(data, items_list)
```