

all_sites_sex_scatters

2024-10-03

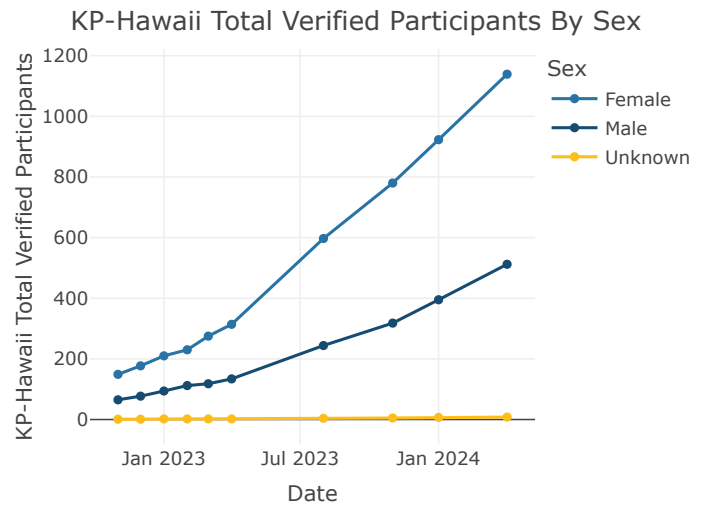
Refactor Site Names

```
data$site <- gsub("^Marshfield.*", "Marshfield", data$site)
unique(data$site)
```

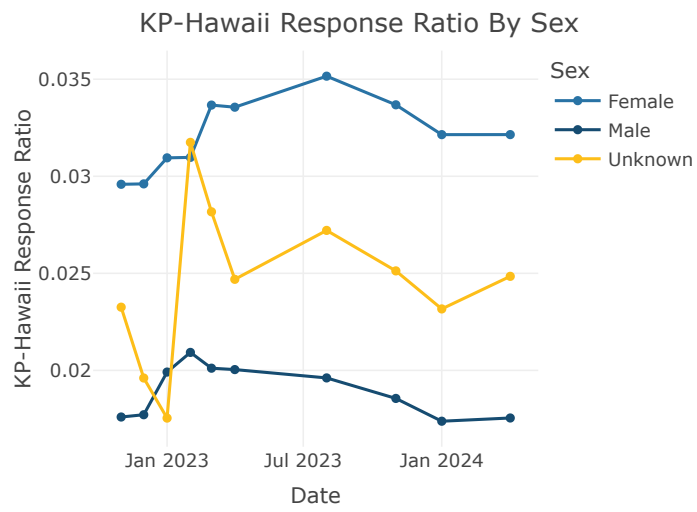
```
## [1] "KP-Hawaii"           "KP-Colorado"         "KP-Northwest"
## [4] "KP-Georgia"          "Marshfield"          "Sanford Health"
## [7] "Henry Ford"          "University of Chicago"
```

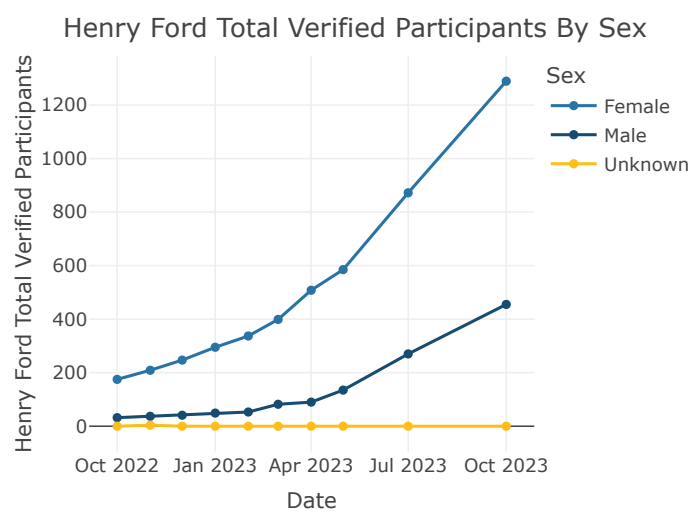
Write list of sites to be used

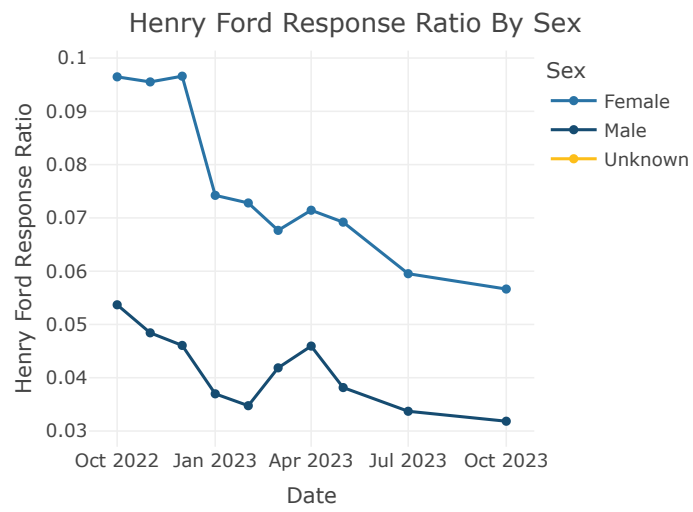
```
items_list = c("KP-Hawaii", "KP-Colorado", "KP-Northwest", "KP-Georgia", "Marshfield", "Sanford Health")
```

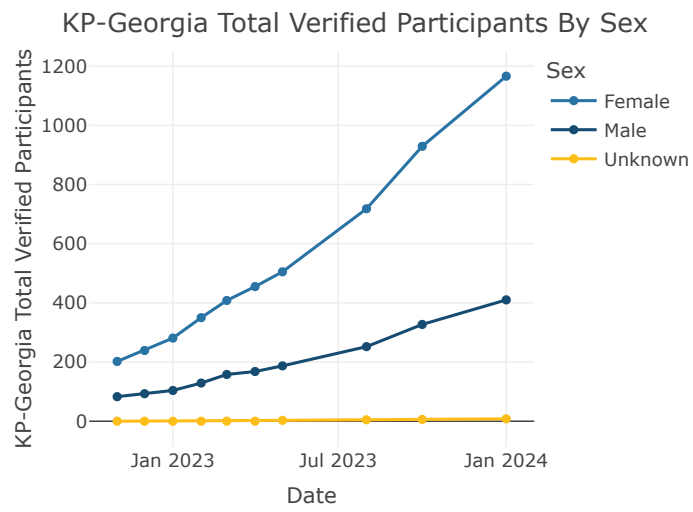


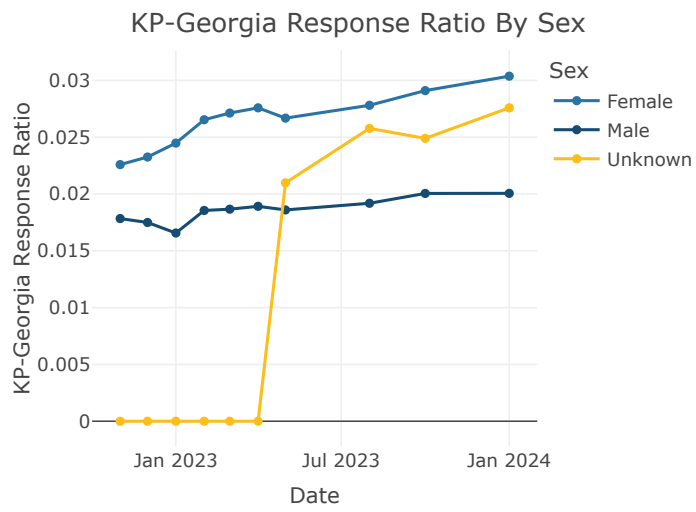
Call by `_sex_scatter` function for all sites

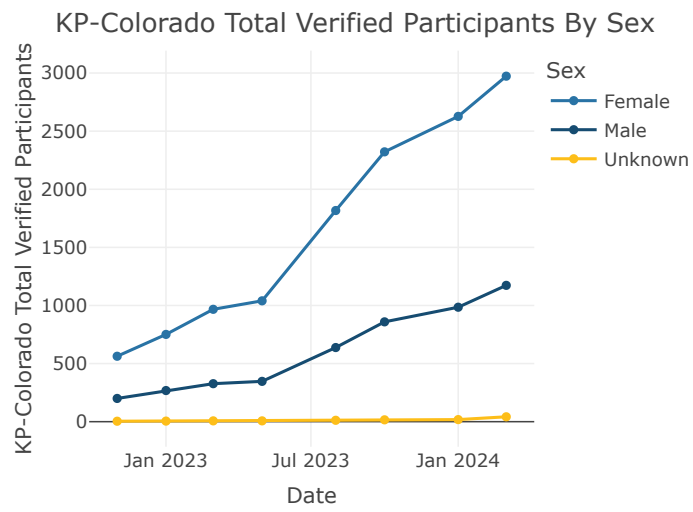


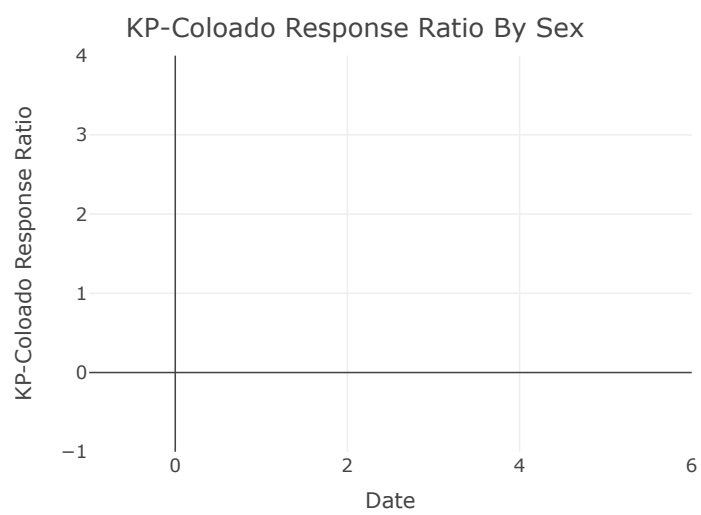


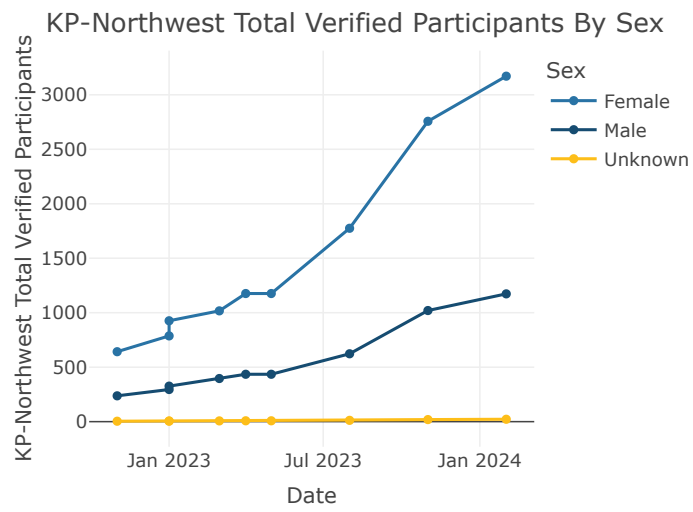


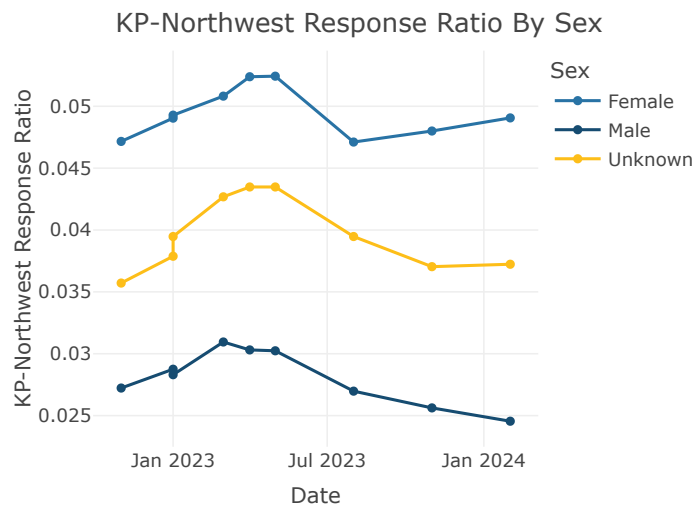


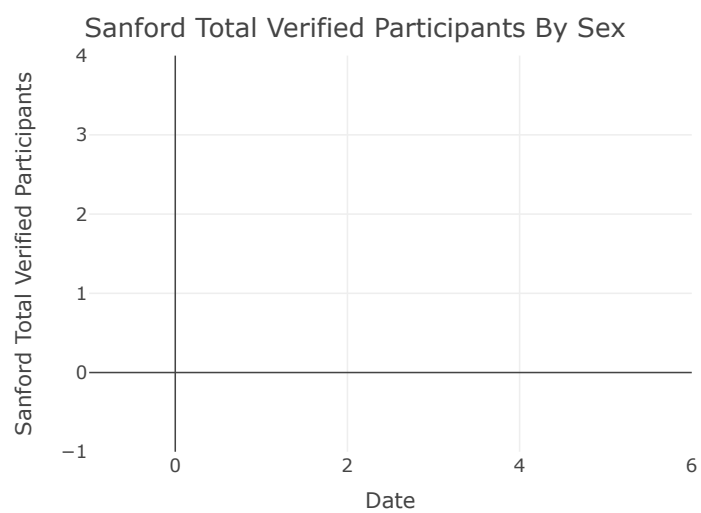


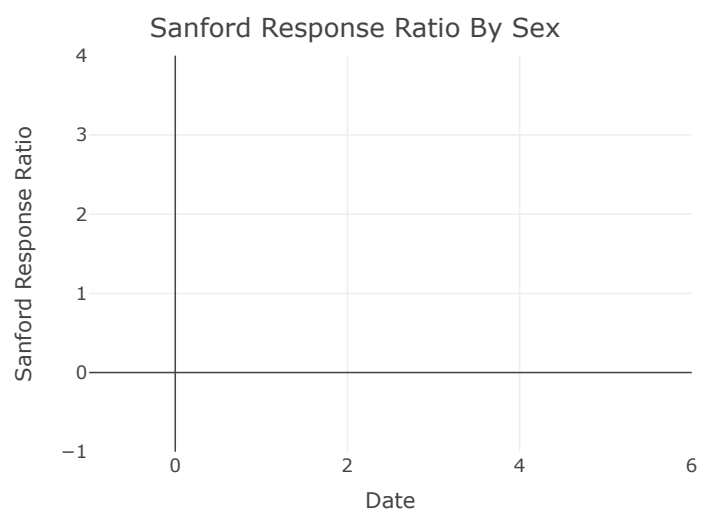












University of Chicago Total Verified Participants By Sex

