

# poa\_health\_8-13

Arielle Herman

3/29/2022

## Contents

<b>Demographics</b>	<b>2</b>
<b>4.1 - 4.2</b>	<b>2</b>
4.1)Households with a case of COVID-19 [36] . . . . .	2
4.2)Individuals with a case of case of COVID-19[36] . . . . .	3
<b>4.3)People who have received a booster vaccine dose [38]</b>	<b>10</b>
<b>4.4)People who returned to work in person are more likely to be boosted against COVID-19</b>	<b>15</b>
<b>4.5, 4.14</b>	<b>16</b>
4.5)People who ranked the government response as average or above average were more likely to be boosted or planning to get boosted against COVID-19 . . . . .	16
4.14) People who rated government services (response of government) above average were more likely to be boosted . . . . .	17
<b>4.6)People who have been discriminated against or are worried about discrimination due     to COVID-19 [35]]</b>	<b>18</b>
<b>4.7)People who have experienced abuse or violence due to COVID-19 [34]</b>	<b>23</b>
<b>4.8)People who score more than average on the quantified mental health quantified ques-     tions [38]</b>	<b>28</b>
<b>4.9 - 4.12</b>	<b>34</b>
4.9) People who rated government response poorly were more likely to score higher than average on the mental health score. . . . .	34
4.10)People who are currently unemployed are more likely to score higher than average on the mental health score . . . . .	35
4.11)People who have experienced abuse or violence are more likely to score higher than average on the mental health score . . . . .	35
4.12)People who have been discriminated against are more likely to score higher than average on the mental health score . . . . .	35

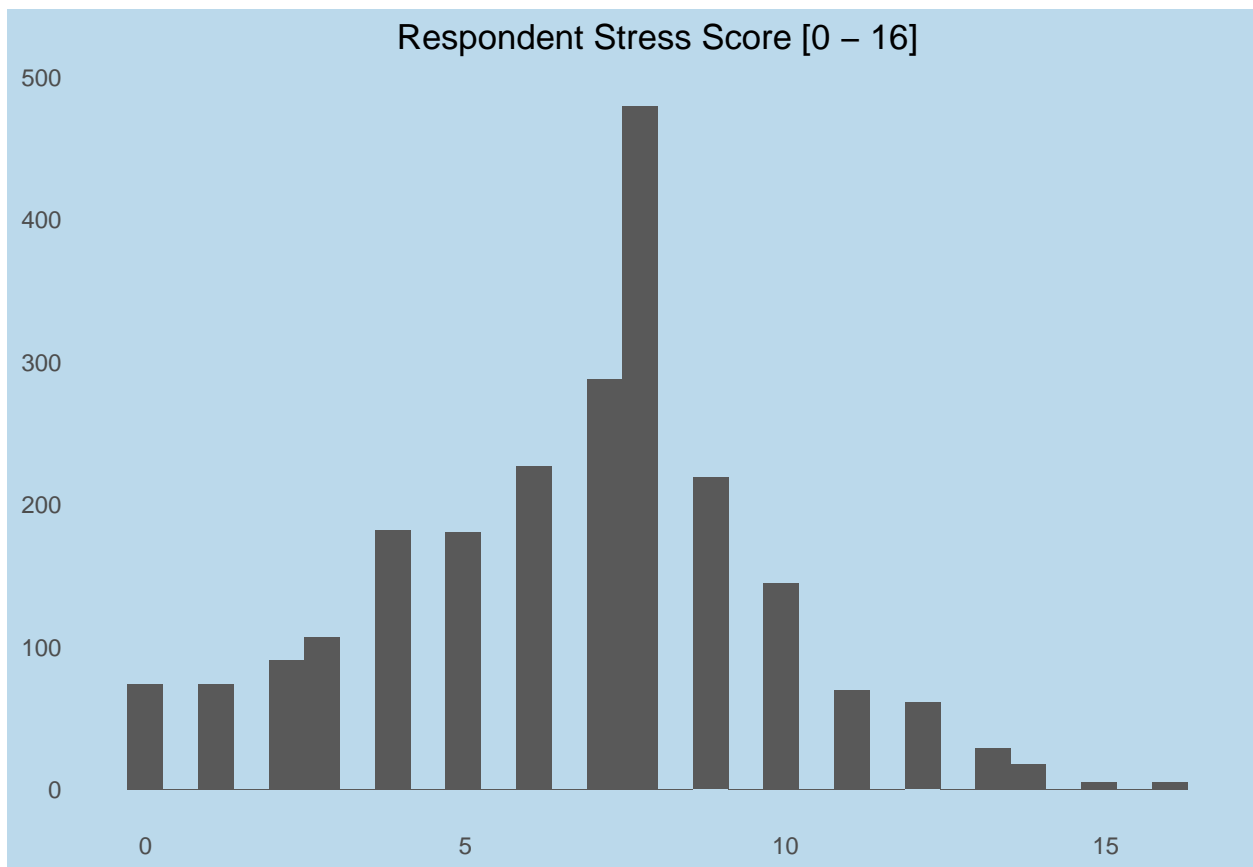
4.13) Run distribution over each activity which has been avoided because of Covid-19	41
4.15) Distribution over frequency of testing	62
4.16) Distribution over each difficulty faced while testing	74

## Demographics

```
wrangled %>% ggplot(aes(x = stress_score)) + geom_histogram() +
  ggtitle("Respondent Stress Score [0 - 16]") +
  xlab(NULL) + ylab(NULL)
```

```
## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.
```

```
## Warning: Removed 59 rows containing non-finite values (stat_bin).
```



## 4.1 - 4.2

### 4.1) Households with a case of COVID-19 [36]

1. Run distribution over population

2. Run distribution by sub-demographics
  - a. Compare and find gaps (test unequal proportions)

## 4.2) Individuals with a case of case of COVID-19[36]

1. Run distribution over population
2. Run distribution by sub-demographics
  - a. Compare and find gaps (test unequal proportions)

```
mean(wrangled$posi_all, na.rm = TRUE)
```

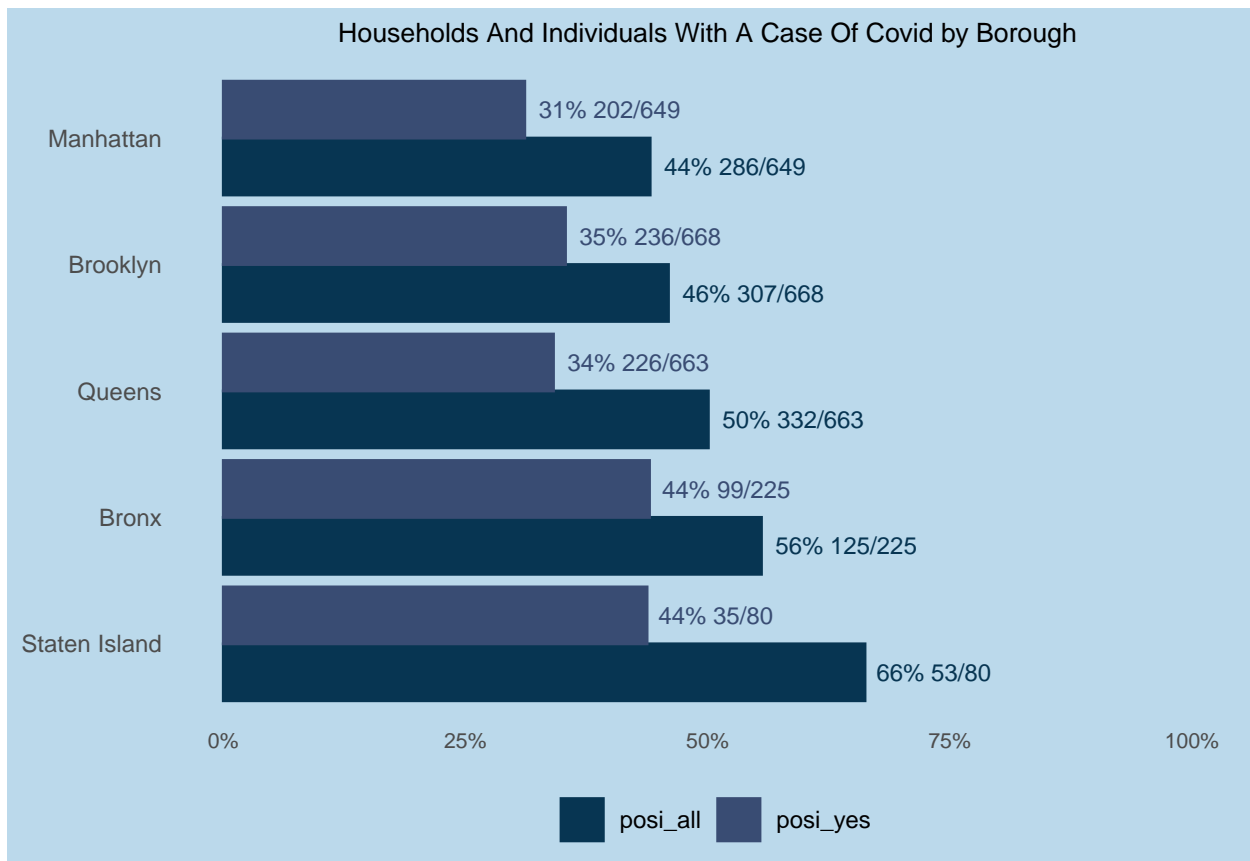
```
## [1] 0.4827133
```

```
mean(wrangled$posi_yes, na.rm = TRUE)
```

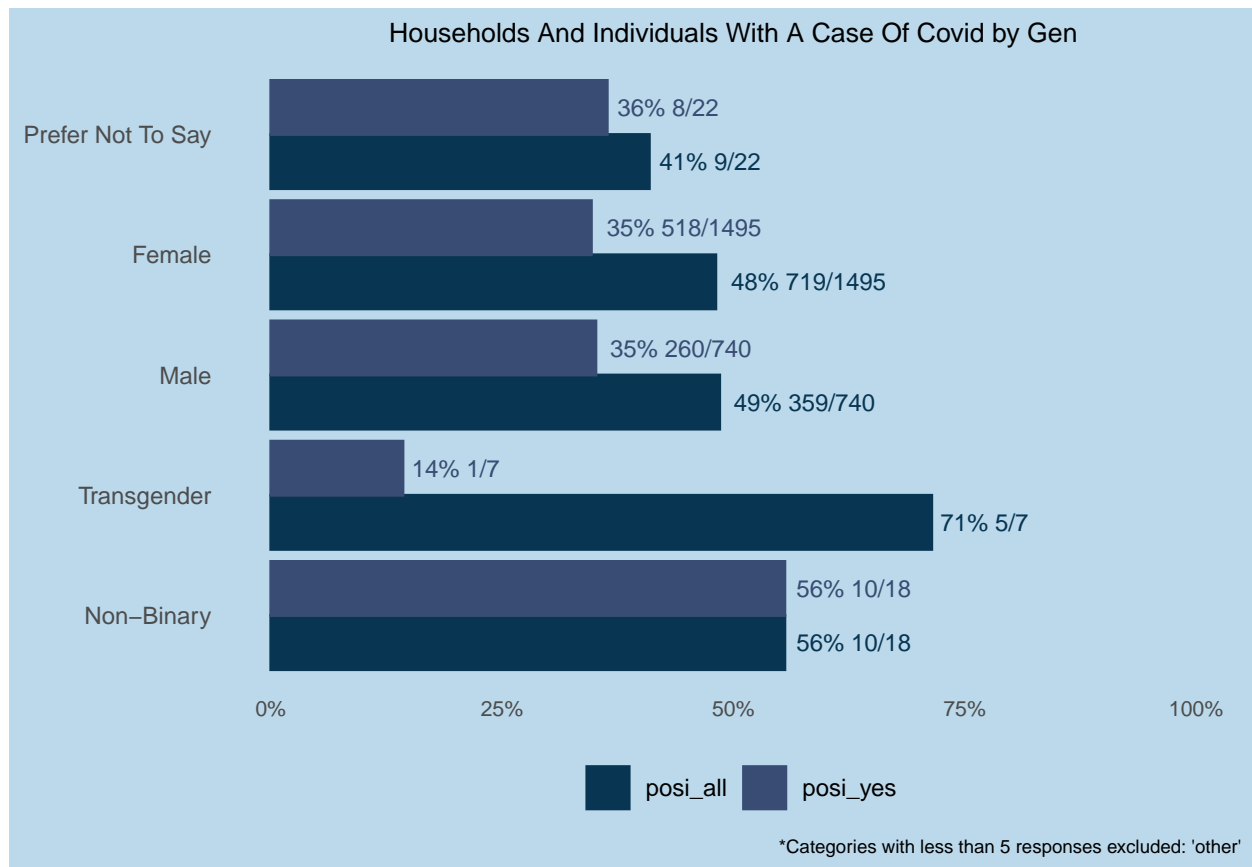
```
## [1] 0.3492341
```

```
make_plots(wrangled, by_vars = demographics,
            hyp_var = c("posi_all", "posi_yes"), title = "Households and Individuals with a case of COVID"
```

```
## $borough
## $borough$plot
```

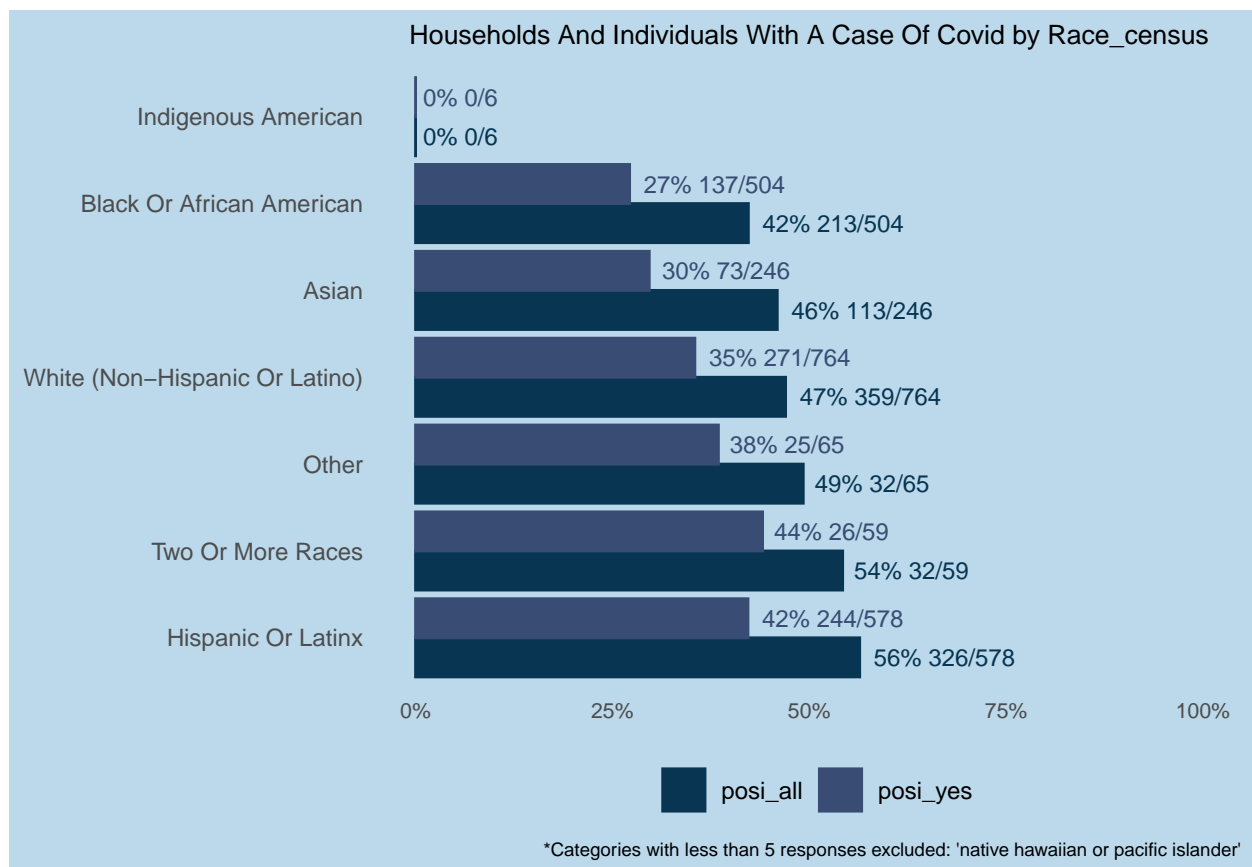


```
##
## $borough$p.values
## $borough$p.values$posi_all
##           manhattan brooklyn queens  bronx staten island
## manhattan           NA         NA    NA 0.0038      0.00028
## brooklyn            NA         NA    NA    NA      0.00092
## queens              NA         NA    NA    NA      0.00890
## bronx               0.00380      NA    NA    NA         NA
## staten island       0.00028 0.00092 0.0089    NA         NA
##
## $borough$p.values$posi_yes
##           manhattan queens brooklyn staten island  bronx
## manhattan           NA     NA      NA           NA 0.00062
## queens              NA     NA      NA           NA 0.00970
## brooklyn            NA     NA      NA           NA    NA
## staten island       NA     NA      NA           NA    NA
## bronx               0.00062 0.0097      NA           NA    NA
##
##
##
## $gen
## $gen$plot
```



```
##
## $gen$p.values
```

```
## $gen$p.values$posi_all
##           prefer not to say female male non-binary transgender
## prefer not to say           NA      NA      NA           NA      NA
## female                     NA      NA      NA           NA      NA
## male                       NA      NA      NA           NA      NA
## non-binary                 NA      NA      NA           NA      NA
## transgender                 NA      NA      NA           NA      NA
##
## $gen$p.values$posi_yes
##           transgender female male prefer not to say non-binary
## transgender           NA      NA      NA           NA      NA
## female                 NA      NA      NA           NA      NA
## male                   NA      NA      NA           NA      NA
## prefer not to say      NA      NA      NA           NA      NA
## non-binary              NA      NA      NA           NA      NA
##
##
##
## $race_census
## $race_census$plot
```

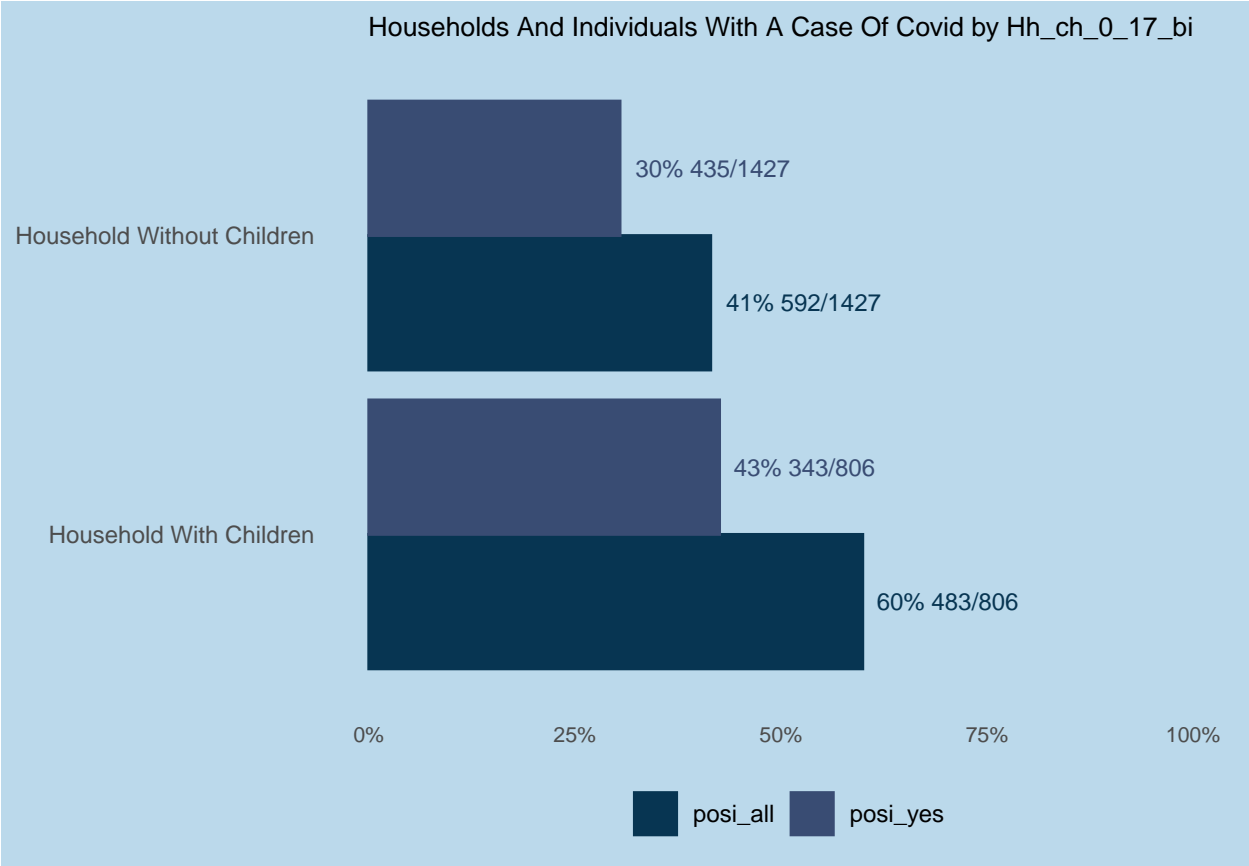


```
##
## $race_census$p.values
## $race_census$p.values$posi_all
##           Indigenous American black or african american
```

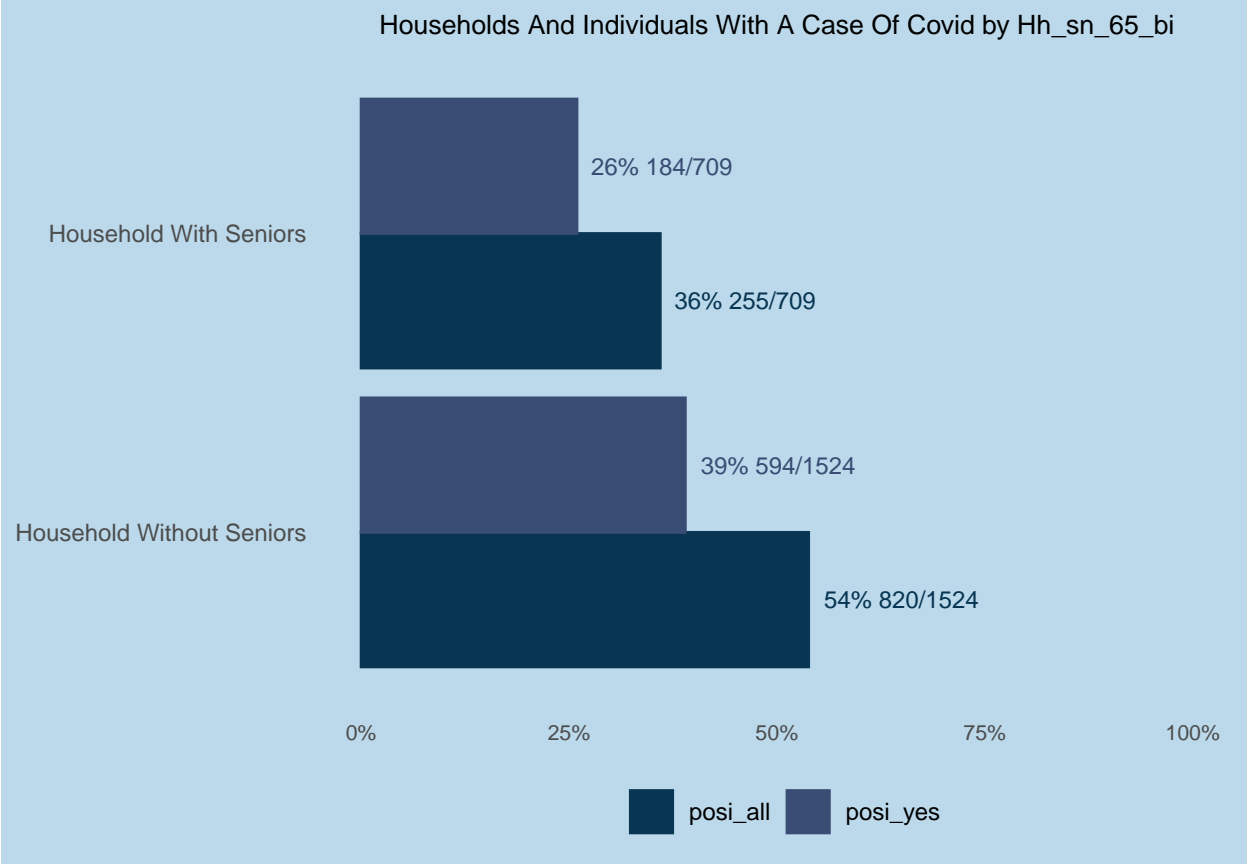
```

## Indigenous American          NA          NA
## black or african american    NA          NA
## asian                        NA          NA
## white (non-hispanic or latino) NA        NA
## other                        NA          NA
## two or more races            NA          NA
## hispanic or latinx           NA        4.7e-06
##
##               asian white (non-hispanic or latino) other
## Indigenous American          NA          NA
## black or african american    NA          NA
## asian                        NA          NA
## white (non-hispanic or latino) NA        NA
## other                        NA          NA
## two or more races            NA          NA
## hispanic or latinx           0.0074      0.00078    NA
##
##               two or more races hispanic or latinx
## Indigenous American          NA          NA
## black or african american    NA        4.7e-06
## asian                        NA        7.4e-03
## white (non-hispanic or latino) NA        7.8e-04
## other                        NA          NA
## two or more races            NA          NA
## hispanic or latinx           NA          NA
##
## $race_census$p.values$posi_yes
##
##               Indigenous American black or african american
## Indigenous American          NA          NA
## black or african american    NA          NA
## asian                        NA          NA
## white (non-hispanic or latino) NA        2.4e-03
## other                        NA          NA
## hispanic or latinx           NA        3.4e-07
## two or more races            NA          NA
##
##               asian white (non-hispanic or latino) other
## Indigenous American          NA          NA
## black or african american    NA        0.0024    NA
## asian                        NA          NA
## white (non-hispanic or latino) NA        NA
## other                        NA          NA
## hispanic or latinx           0.00094      NA        NA
## two or more races            NA          NA
##
##               hispanic or latinx two or more races
## Indigenous American          NA          NA
## black or african american    3.4e-07      NA
## asian                        9.4e-04      NA
## white (non-hispanic or latino) NA        NA
## other                        NA          NA
## hispanic or latinx           NA          NA
## two or more races            NA          NA
##
##
##
## $hh_ch_0_17_bi
## $hh_ch_0_17_bi$plot

```

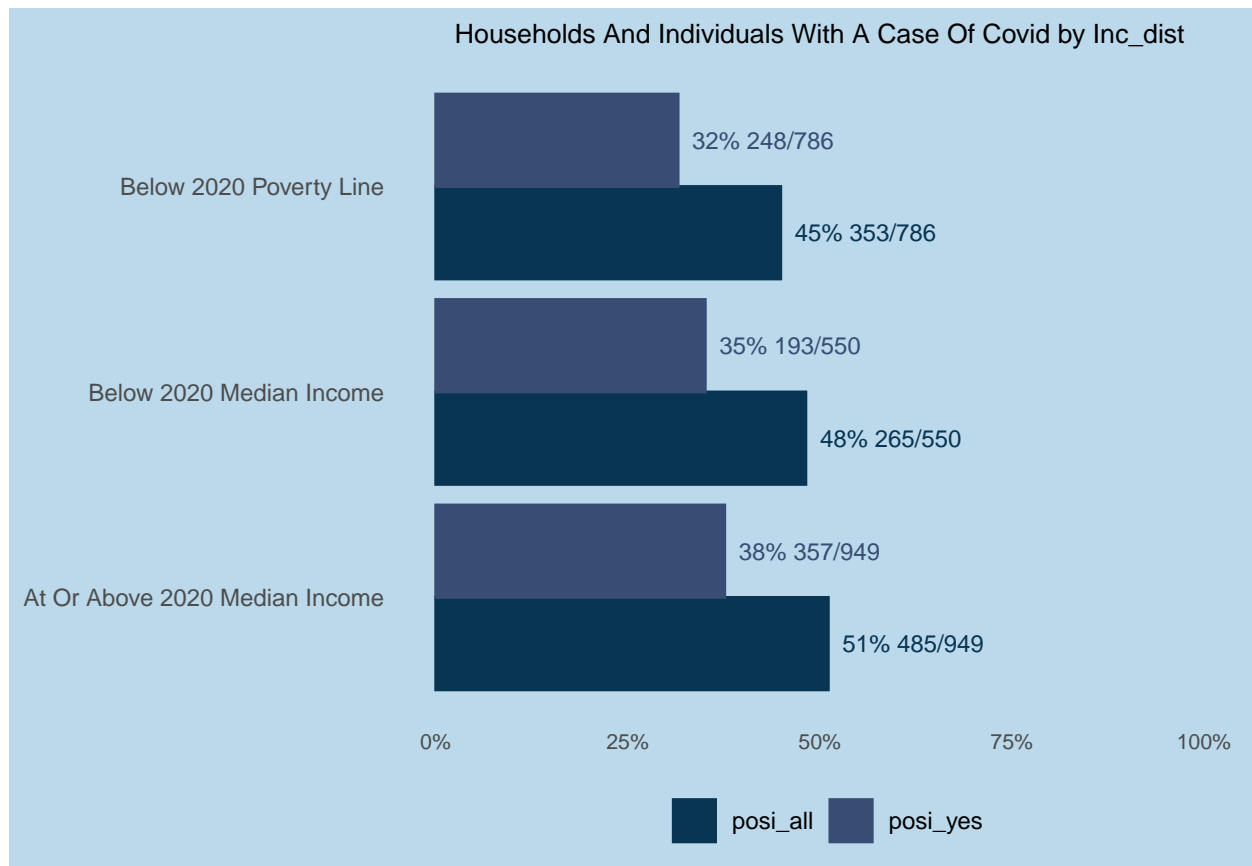


```
##
## $hh_ch_0_17_bi$p.values
## $hh_ch_0_17_bi$p.values$posi_all
##           household without children household with children
## household without children           NA           8e-17
## household with children           8e-17           NA
##
## $hh_ch_0_17_bi$p.values$posi_yes
##           household without children household with children
## household without children           NA           1.2e-08
## household with children           1.2e-08           NA
##
##
##
## $hh_sn_65_bi
## $hh_sn_65_bi$plot
```



```
##
## $hh_sn_65_bi$p.values
## $hh_sn_65_bi$p.values$posi_all
##           household with seniors household without seniors
## household with seniors                NA                5.8e-15
## household without seniors            5.8e-15                NA
##
## $hh_sn_65_bi$p.values$posi_yes
##           household with seniors household without seniors
## household with seniors                NA                2.4e-09
## household without seniors            2.4e-09                NA
##
##
##
## $inc_dist
## $inc_dist$plot
```





```
##
## $inc_dist$p.values
## $inc_dist$p.values$posi_all
##
## below 2020 poverty line below 2020 median income
## below 2020 poverty line NA NA
## below 2020 median income NA NA
## at or above 2020 median income NA NA
##
## at or above 2020 median income
## below 2020 poverty line NA
## below 2020 median income NA
## at or above 2020 median income NA
##
## $inc_dist$p.values$posi_yes
##
## below 2020 poverty line below 2020 median income
## below 2020 poverty line NA NA
## below 2020 median income NA NA
## at or above 2020 median income 0.0096 NA
##
## at or above 2020 median income
## below 2020 poverty line 0.0096
## below 2020 median income NA
## at or above 2020 median income NA

cat("Plots for by gen and inc_dist are not showing at least one statistically significant result")

## Plots for by gen and inc_dist are not showing at least one statistically significant result
```

## 4.3) People who have received a booster vaccine dose [38]

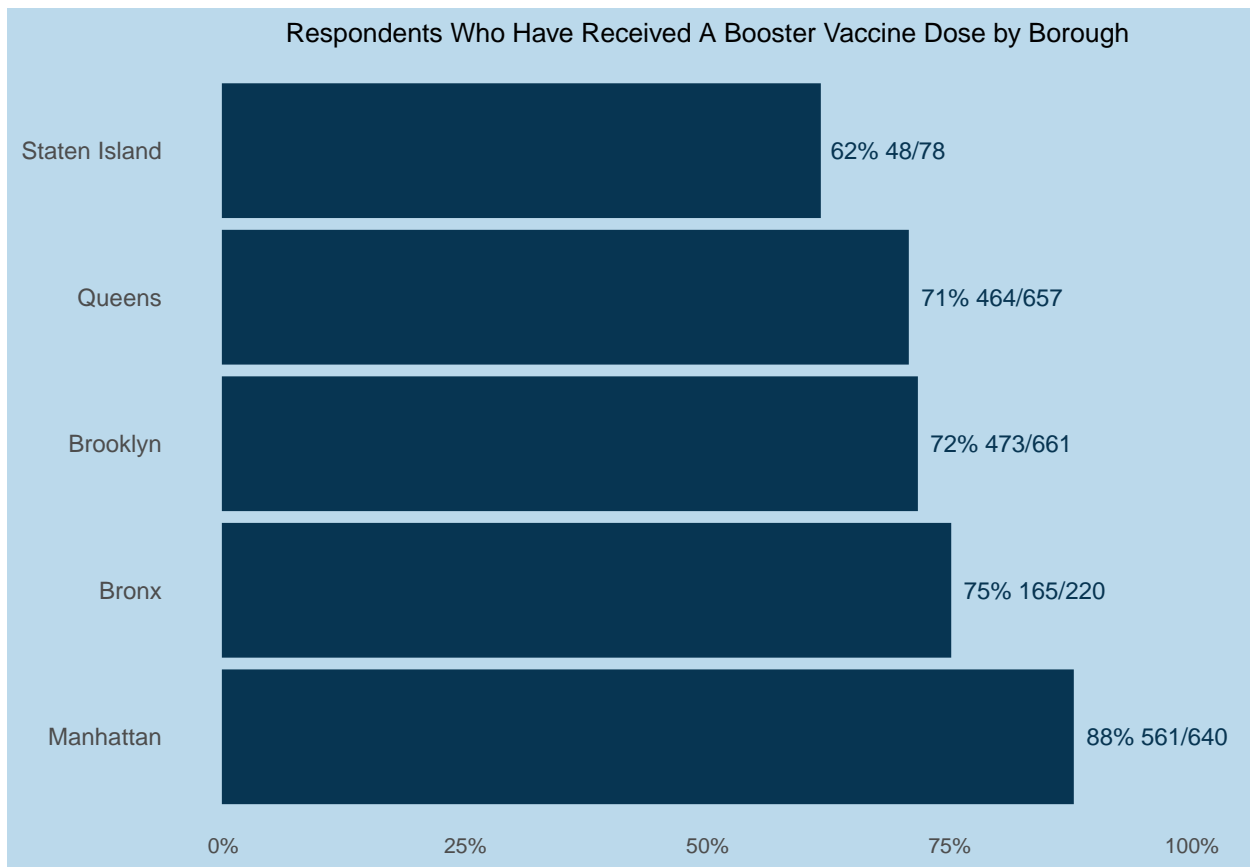
1. Run distribution over population
2. Run distribution by sub-demographics (a-k)
  - a. Compare and find gaps (test unequal proportions)

```
mean(wrangled$boost_bi, na.rm = TRUE)
```

```
## [1] 0.758422
```

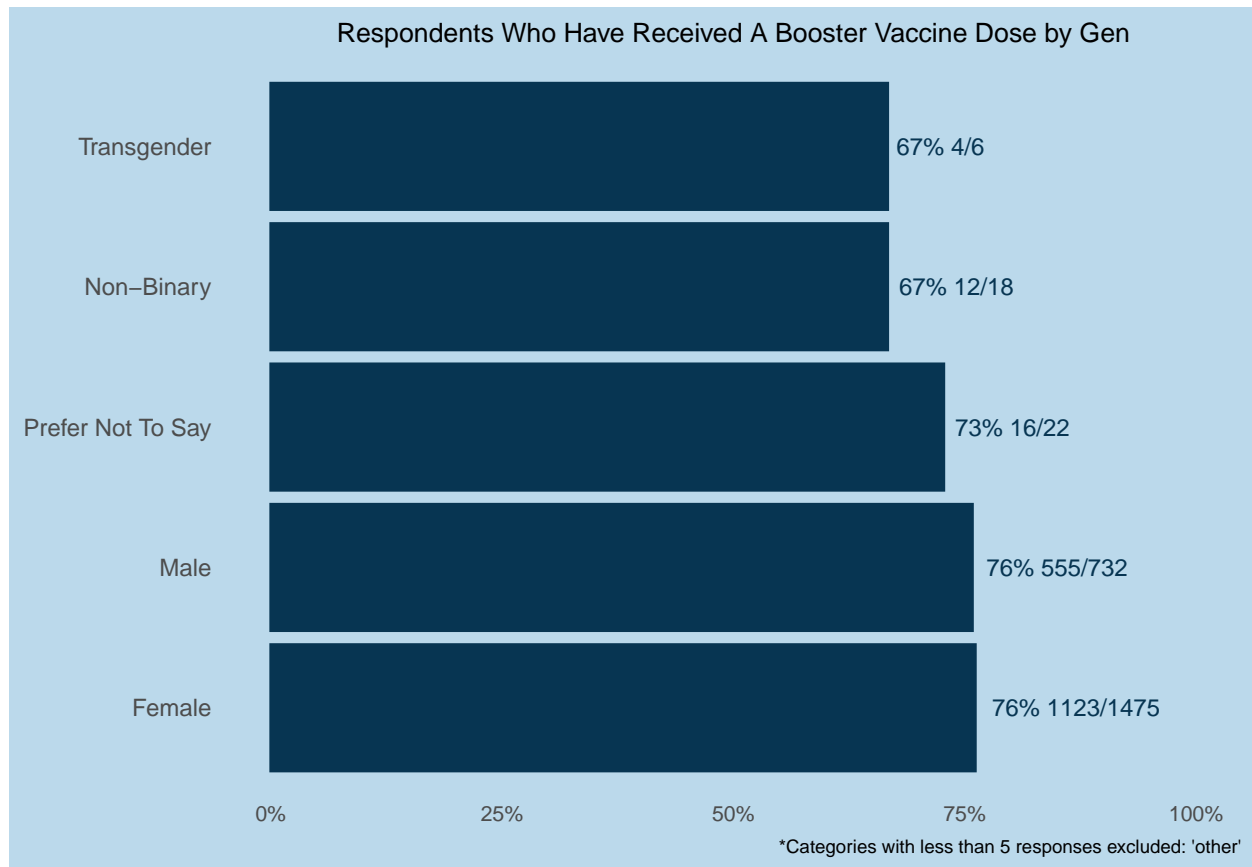
```
make_plots(wrangled, by_vars=demographics, hyp_var="boost_bi", title = "Respondents who have received a
```

```
## $borough  
## $borough$plot
```

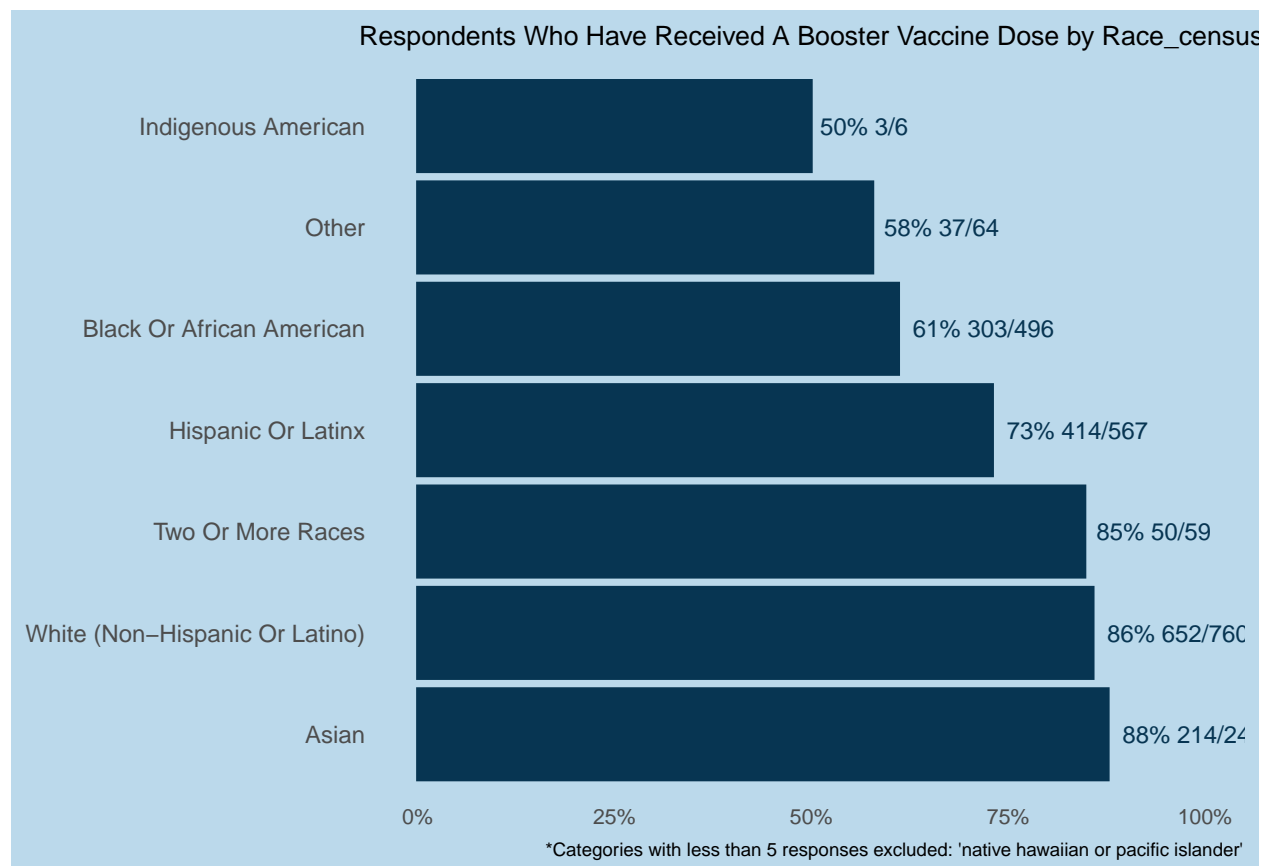


```
##  
## $borough$p.values  
## $borough$p.values$boost_bi  
##          staten island  queens  brooklyn  bronx  manhattan  
## staten island      NA      NA      NA      NA  3.6e-09  
## queens            NA      NA      NA      NA  8.3e-14  
## brooklyn           NA      NA      NA      NA  1.1e-12
```

```
## bronx          NA      NA      NA      NA      1.3e-05
## manhattan    3.6e-09 8.3e-14 1.1e-12 1.3e-05      NA
##
##
##
## $gen
## $gen$plot
```

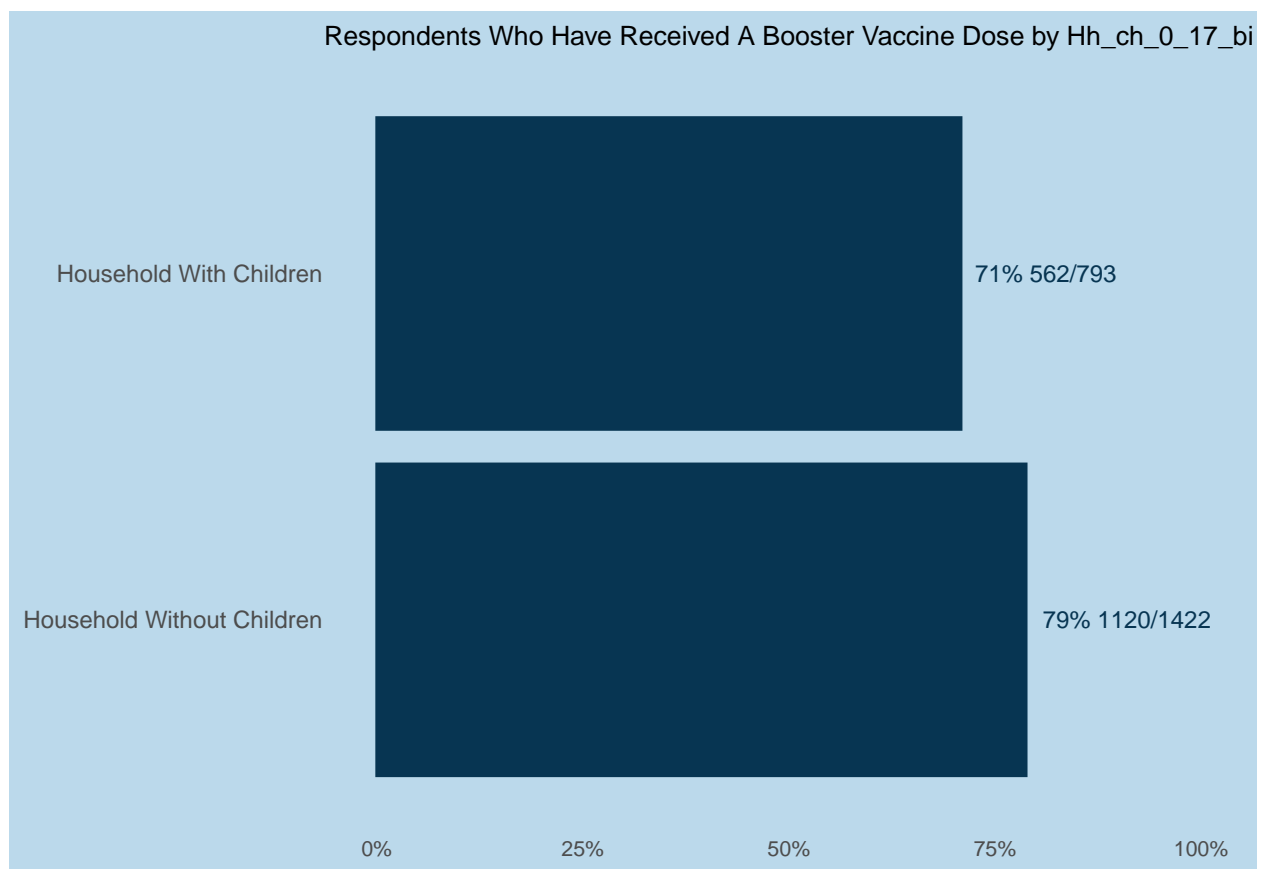


```
##
## $gen$p.values
## $gen$p.values$boost_bi
##          non-binary transgender prefer not to say male female
## non-binary          NA          NA          NA  NA  NA
## transgender          NA          NA          NA  NA  NA
## prefer not to say     NA          NA          NA  NA  NA
## male                  NA          NA          NA  NA  NA
## female                NA          NA          NA  NA  NA
##
##
##
## $race_census
## $race_census$plot
```

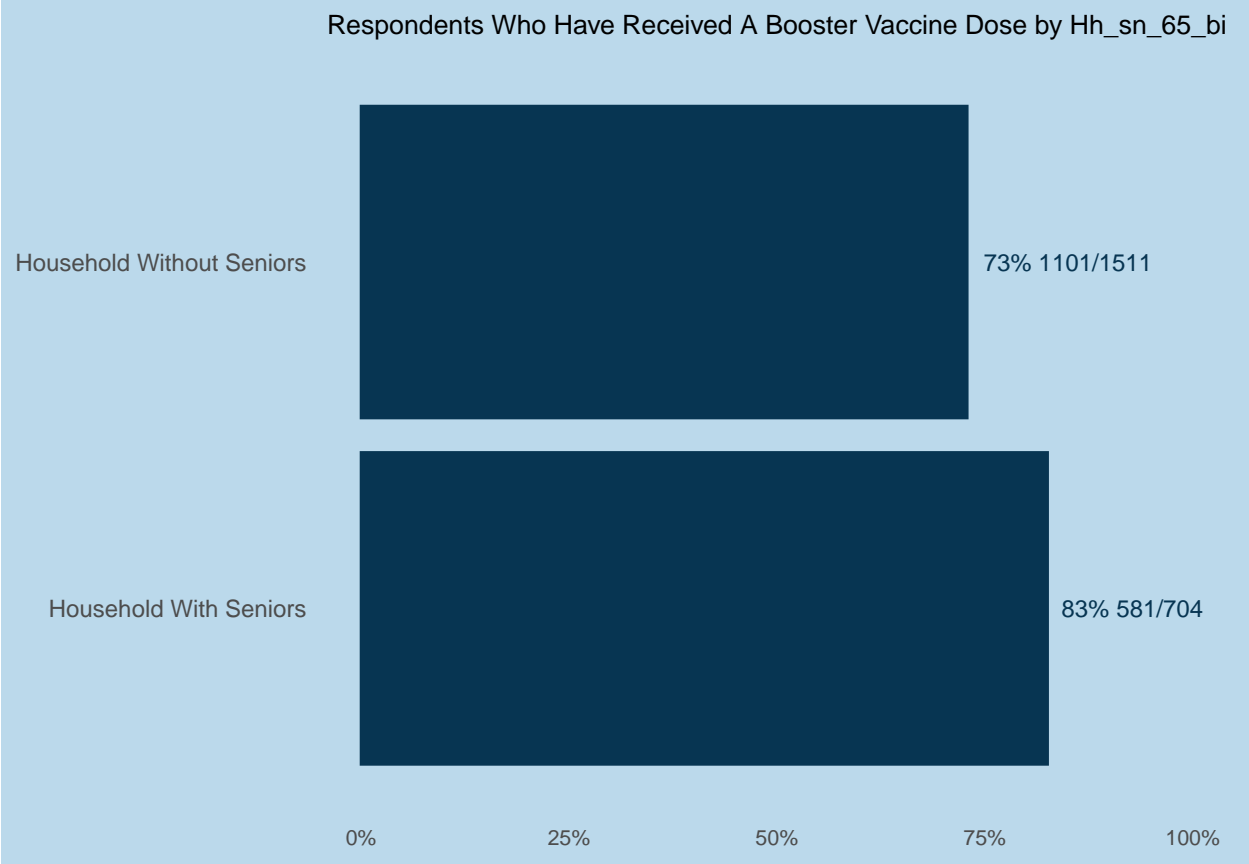


```
##
## $race_census$p.values
## $race_census$p.values$boost_bi
##
## Indigenous American    other
## Indigenous American    NA      NA
## other                  NA      NA
## black or african american    NA      NA
## hispanic or latinx          NA      NA
## two or more races          NA 2.1e-03
## white (non-hispanic or latino)    NA 1.8e-08
## asian                    NA 1.2e-07
##
## black or african american hispanic or latinx
## Indigenous American          NA      NA
## other                        NA      NA
## black or african american          NA 4.6e-05
## hispanic or latinx          4.6e-05      NA
## two or more races          6.1e-04      NA
## white (non-hispanic or latino)    2.4e-23 1.1e-08
## asian                    2.3e-13 6.9e-06
##
## two or more races white (non-hispanic or latino)
## Indigenous American          NA      NA
## other                        0.00210 1.8e-08
## black or african american          0.00061 2.4e-23
## hispanic or latinx          NA      1.1e-08
## two or more races          NA      NA
## white (non-hispanic or latino)    NA      NA
```

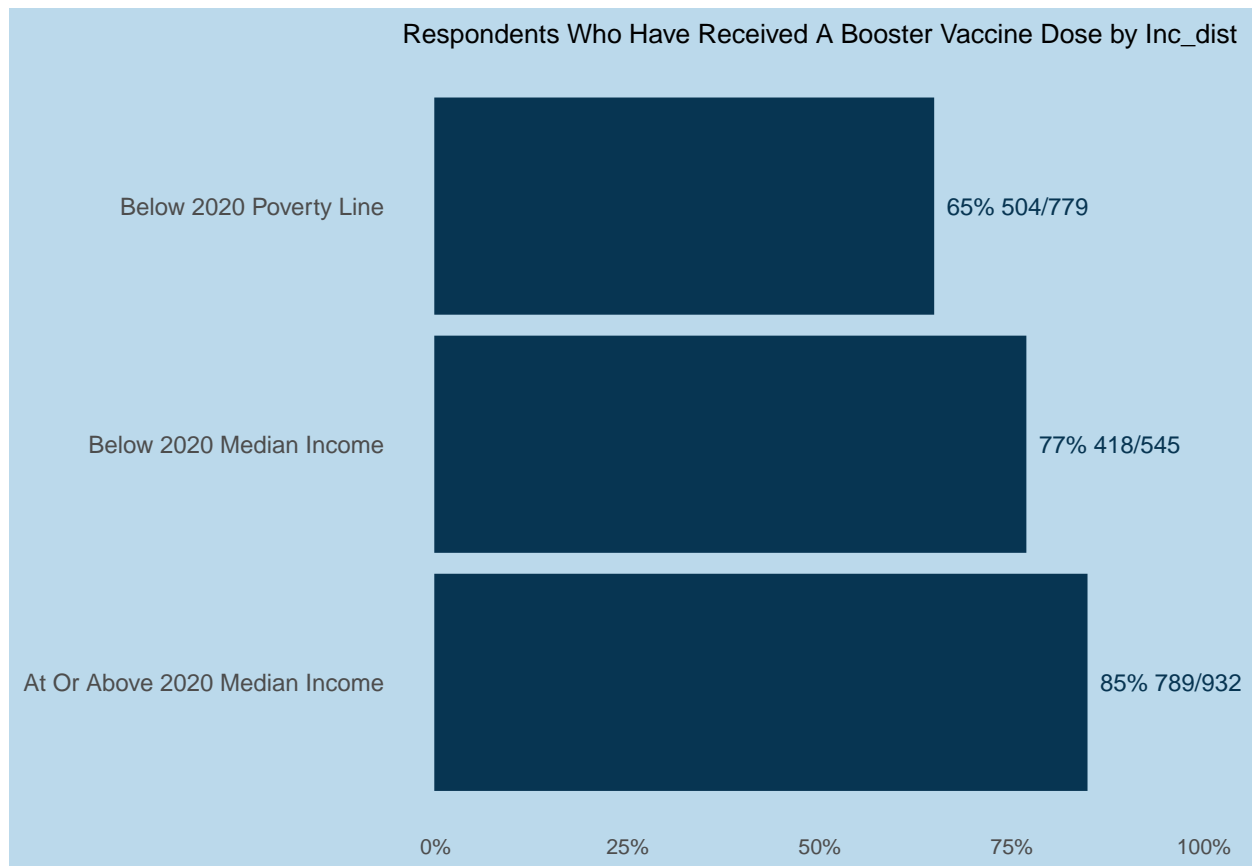
```
## asian
##
## Indigenous American
## other
## black or african american
## hispanic or latinx
## two or more races
## white (non-hispanic or latino)
## asian
##
##
## $hh_ch_0_17_bi
## $hh_ch_0_17_bi$plot
```



```
##
## $hh_ch_0_17_bi$p.values
## $hh_ch_0_17_bi$p.values$boost_bi
## household with children household without children
## household with children NA 3.9e-05
## household without children 3.9e-05 NA
##
##
## $hh_sn_65_bi
## $hh_sn_65_bi$plot
```



```
##
## $hh_sn_65_bi$p.values
## $hh_sn_65_bi$p.values$boost_bi
##           household without seniors household with seniors
## household without seniors              NA              9.6e-07
## household with seniors              9.6e-07              NA
##
##
##
## $inc_dist
## $inc_dist$plot
```



```
##
## $inc_dist$p.values
## $inc_dist$p.values$boost_bi
##
## below 2020 poverty line below 2020 median income
## below 2020 poverty line NA 4.0e-06
## below 2020 median income 4.0e-06 NA
## at or above 2020 median income 1.9e-21 1.8e-04
##
## at or above 2020 median income
## below 2020 poverty line 1.9e-21
## below 2020 median income 1.8e-04
## at or above 2020 median income NA
```

```
cat("Plot for gen is not showing at least one statistically significant result")
```

```
## Plot for gen is not showing at least one statistically significant result
```

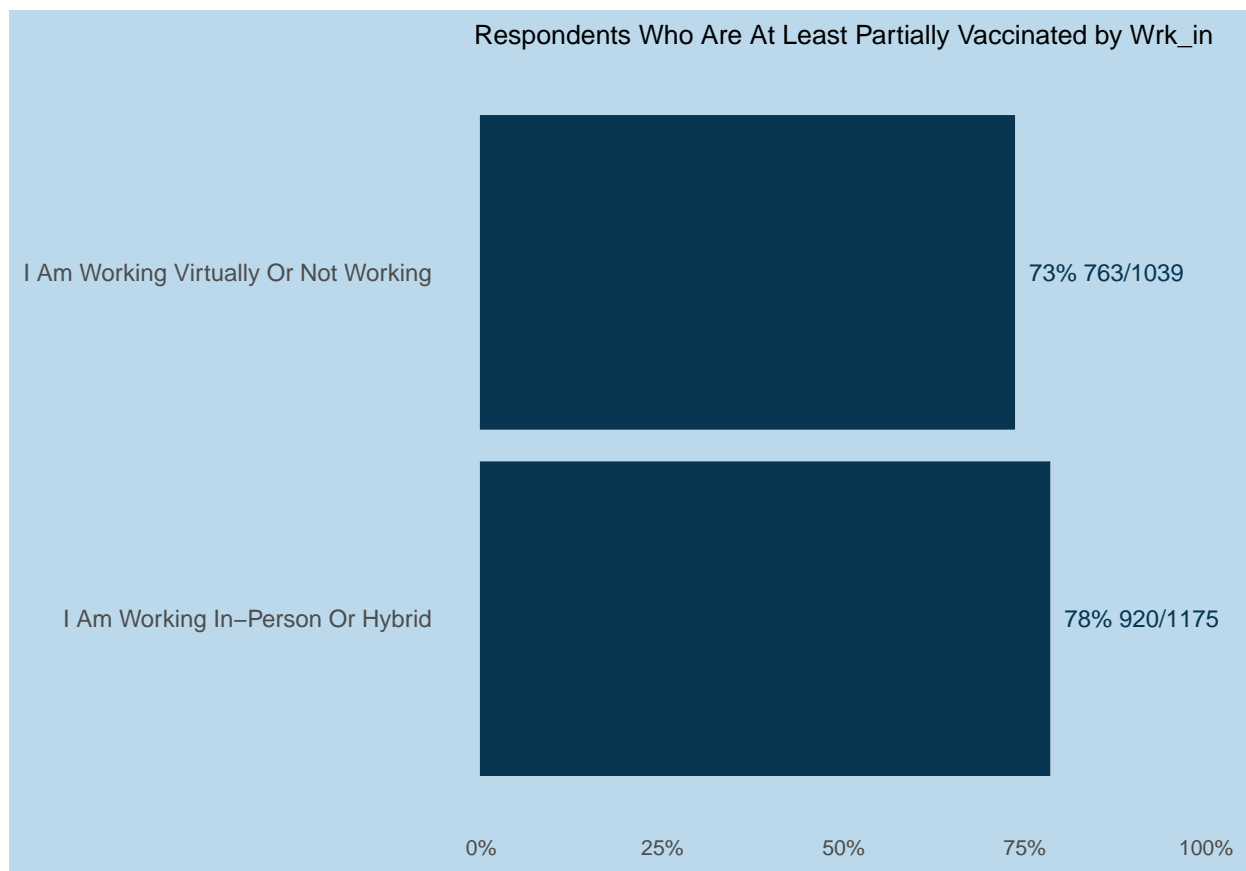
#### 4.4) People who returned to work in person are more likely to be boosted against COVID-19

1. Find respondents who returned to work in person or hybrid [19]
  - a. Find proportion of subset who reported being at least partially vaccinated against COVID-19 [32]pothes

- b. Find proportion not in subset who reported being at least partially vaccinated against COVID-19 and compare (test unequal proportions)

```
make_plots(wrangled, by_vars = "wrk_in", "boost_bi", title = "Respondents who are at least partially vaccinated")
```

```
## $wrk_in
## $wrk_in$plot
```



```
##
## $wrk_in$p.values
## $wrk_in$p.values$boost_bi
##
## I am working virtually or not working NA
## I am working in-person or hybrid 0.0087
##
## I am working in-person or hybrid
## I am working virtually or not working 0.0087
## I am working in-person or hybrid NA
```

## 4.5, 4.14

4.5) People who ranked the government response as average or above average were more likely to be boosted or planning to get boosted against COVID-19

1. Find respondents who rated the government response as average, good, or excellent [31]



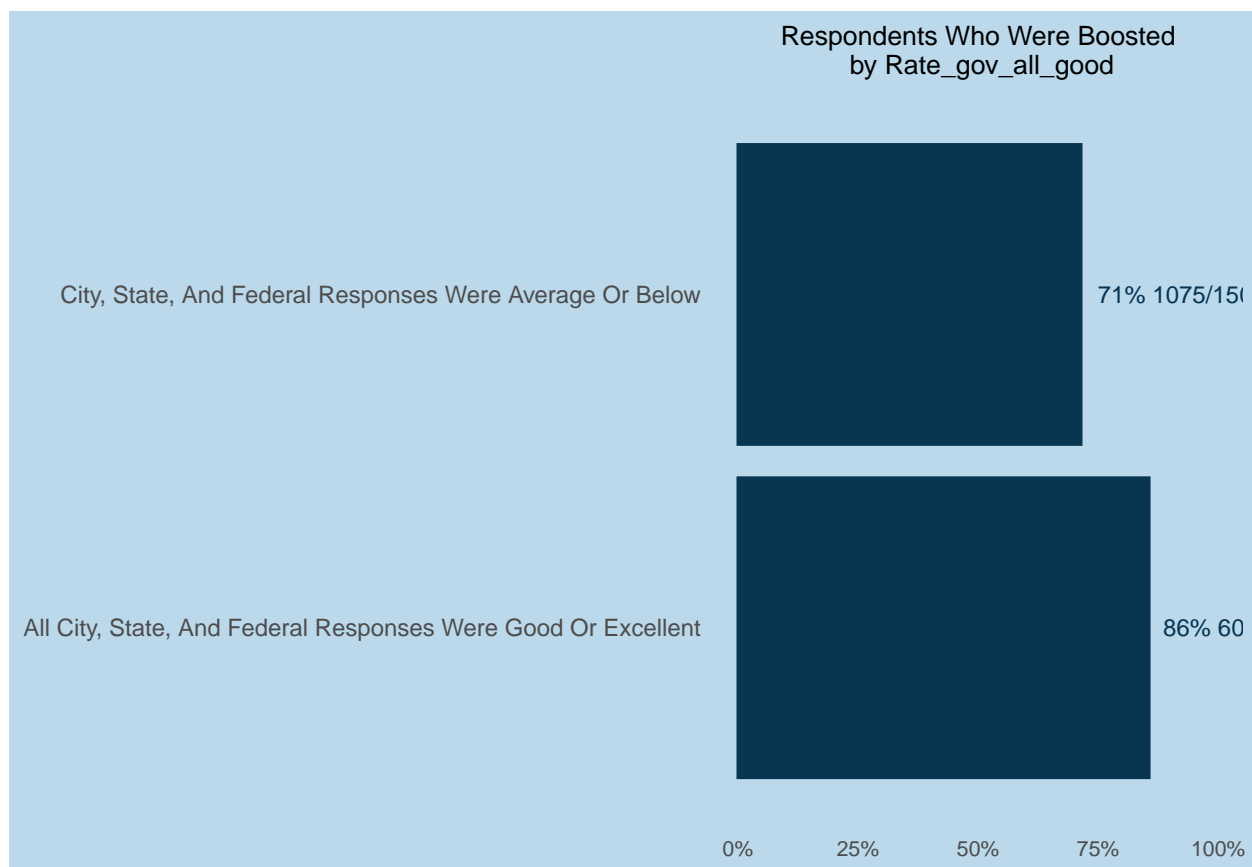
- Find proportion of subset who reported being boosted at least partially planning to get boosted against COVID-19 [37]
- Find proportion not in subset who reported being at boosted or at least partly getting boosted against COVID-19 and compare (test unequal proportions)

#### 4.14) People who rated government services (response of government) above average were more likely to be boosted

- Find proportion of population who rate government services above average (city/state or federal)
- Find the booster status amongst the population
- Compare and contrast with the population who rated the response of the government below average.

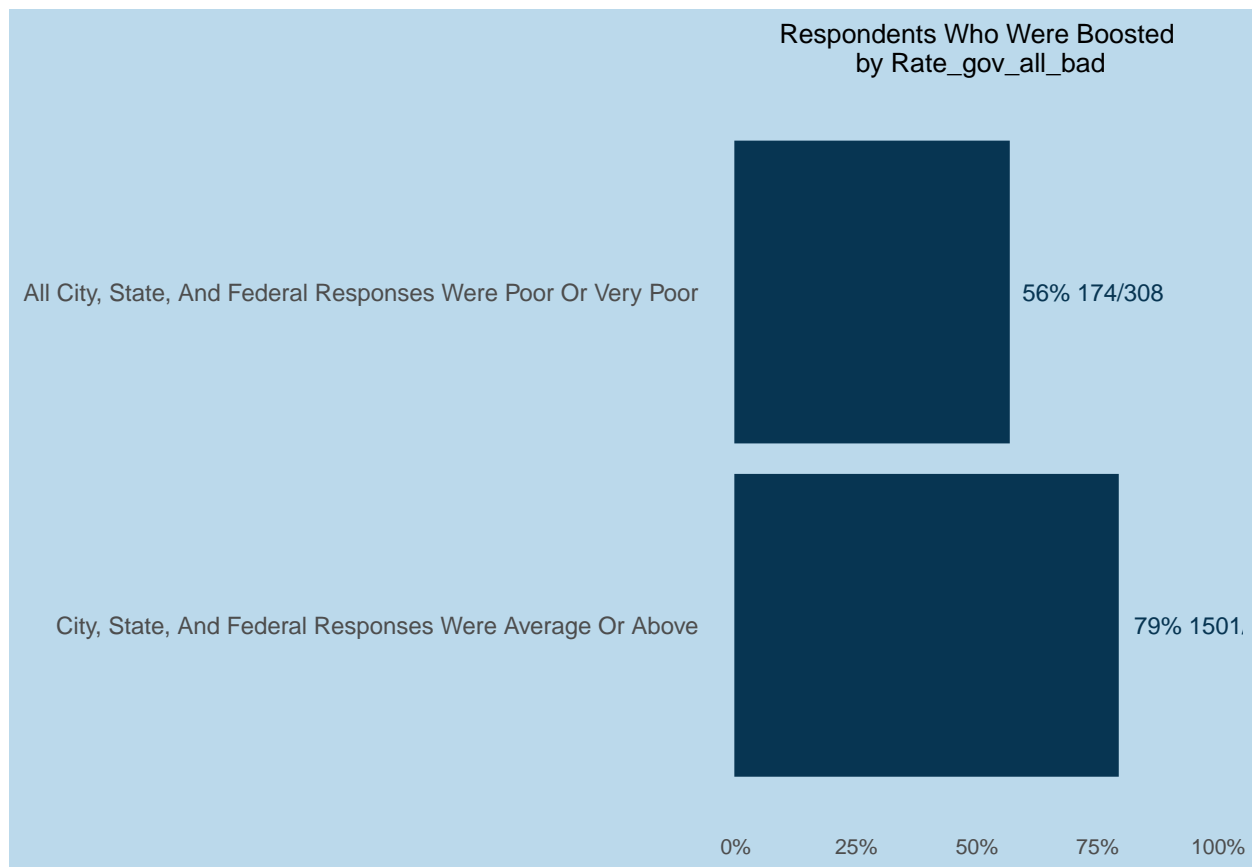
```
make_plots(wrangled, by_var = c("rate_gov_all_good", "rate_gov_all_bad"), hyp_var = "boost_bi",
           title = "Respondents who were boosted\n")
```

```
## $rate_gov_all_good
## $rate_gov_all_good$plot
```



```
##
## $rate_gov_all_good$p.values
## $rate_gov_all_good$p.values$boost_bi
##
## city, state, and federal responses were
## city, state, and federal responses were average or below
```

```
## all city, state, and federal responses were good or excellent
## all city, state, and federal responses
## city, state, and federal responses were average or below
## all city, state, and federal responses were good or excellent
##
##
##
## $rate_gov_all_bad
## $rate_gov_all_bad$plot
```



```
##
## $rate_gov_all_bad$p.values
## $rate_gov_all_bad$p.values$boost_bi
## all city, state, and federal responses
## all city, state, and federal responses were poor or very poor
## city, state, and federal responses were average or above
## city, state, and federal responses were
## all city, state, and federal responses were poor or very poor
## city, state, and federal responses were average or above
```

#### 4.6) People who have been discriminated against or are worried about discrimination due to COVID-19 [35]

1. Run distribution over population

2. Run distribution by sub-demographics

a. Compare and find gaps (test unequal proportions)

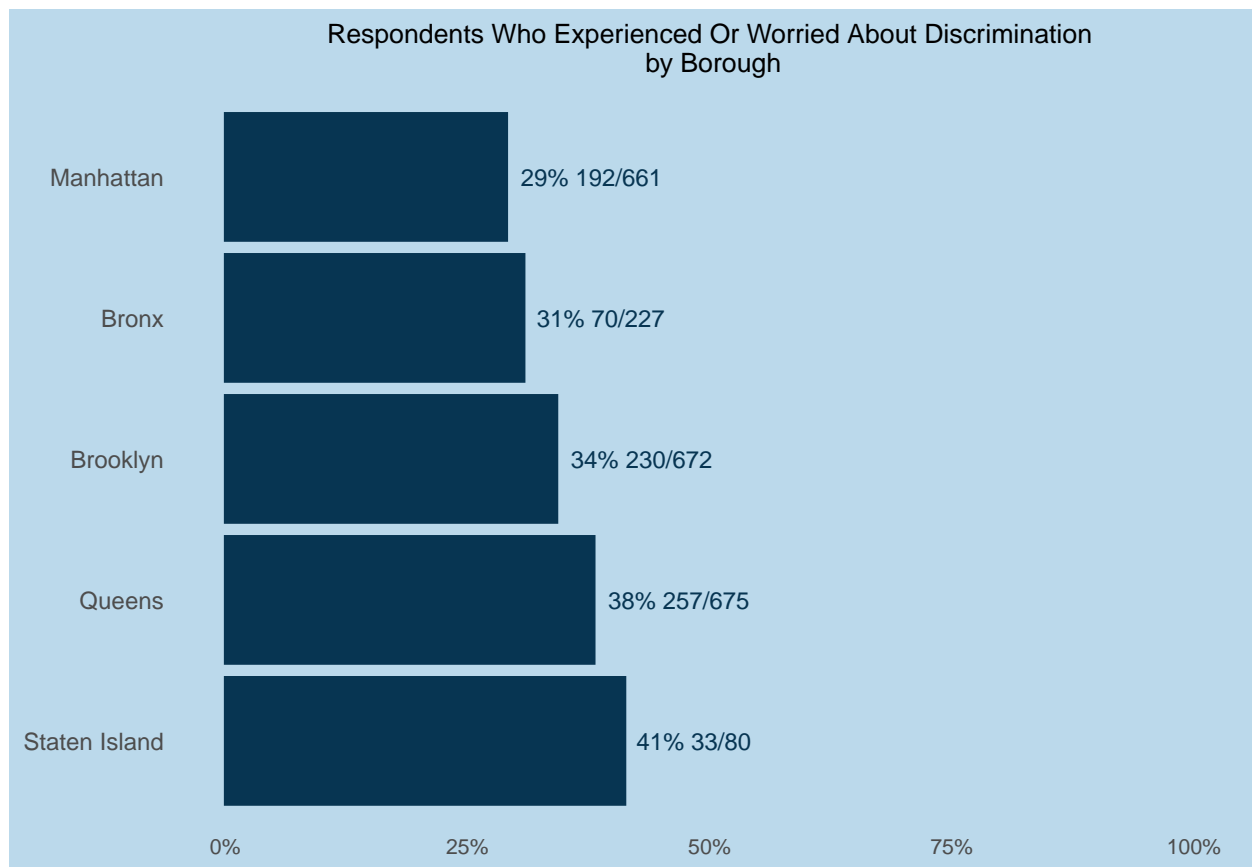
```
mean(wrangled$discrim_bi, na.rm = TRUE)
```

```
## [1] 0.337797
```

```
make_plots(wrangled, demographics, "discrim_bi", title = "Respondents who experienced or worried about discrimination by borough")
```

```
## $borough
```

```
## $borough$plot
```



```
##
```

```
## $borough$p.values
```

```
## $borough$p.values$discrim_bi
```

```
##      manhattan bronx brooklyn  queens staten island
```

```
## manhattan      NA      NA      NA 0.00059          NA
```

```
## bronx          NA      NA      NA      NA          NA
```

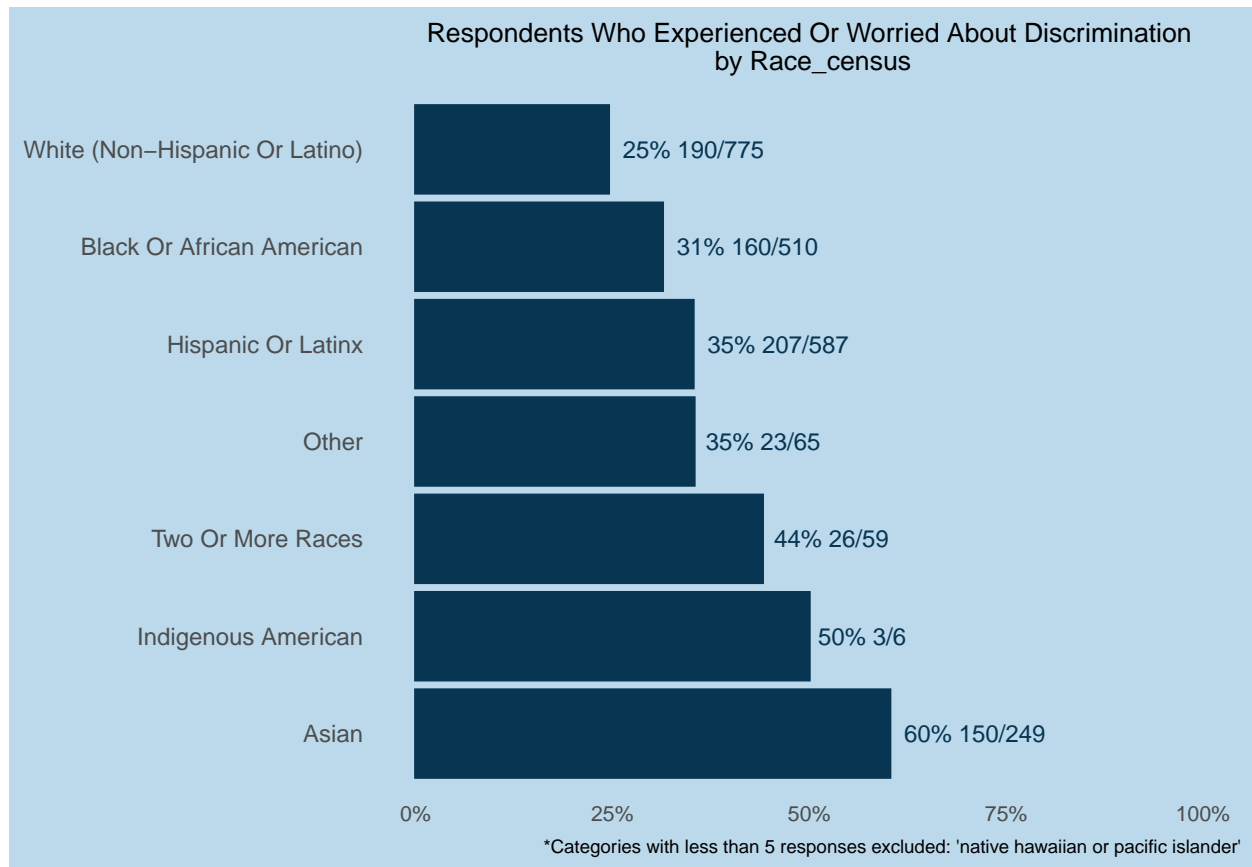
```
## brooklyn       NA      NA      NA      NA          NA
```

```
## queens         0.00059      NA      NA      NA          NA
```

```
## staten island   NA      NA      NA      NA          NA
```

```
##
```

```
##
##
## $gen
## NULL
##
## $race_census
## $race_census$plot
```

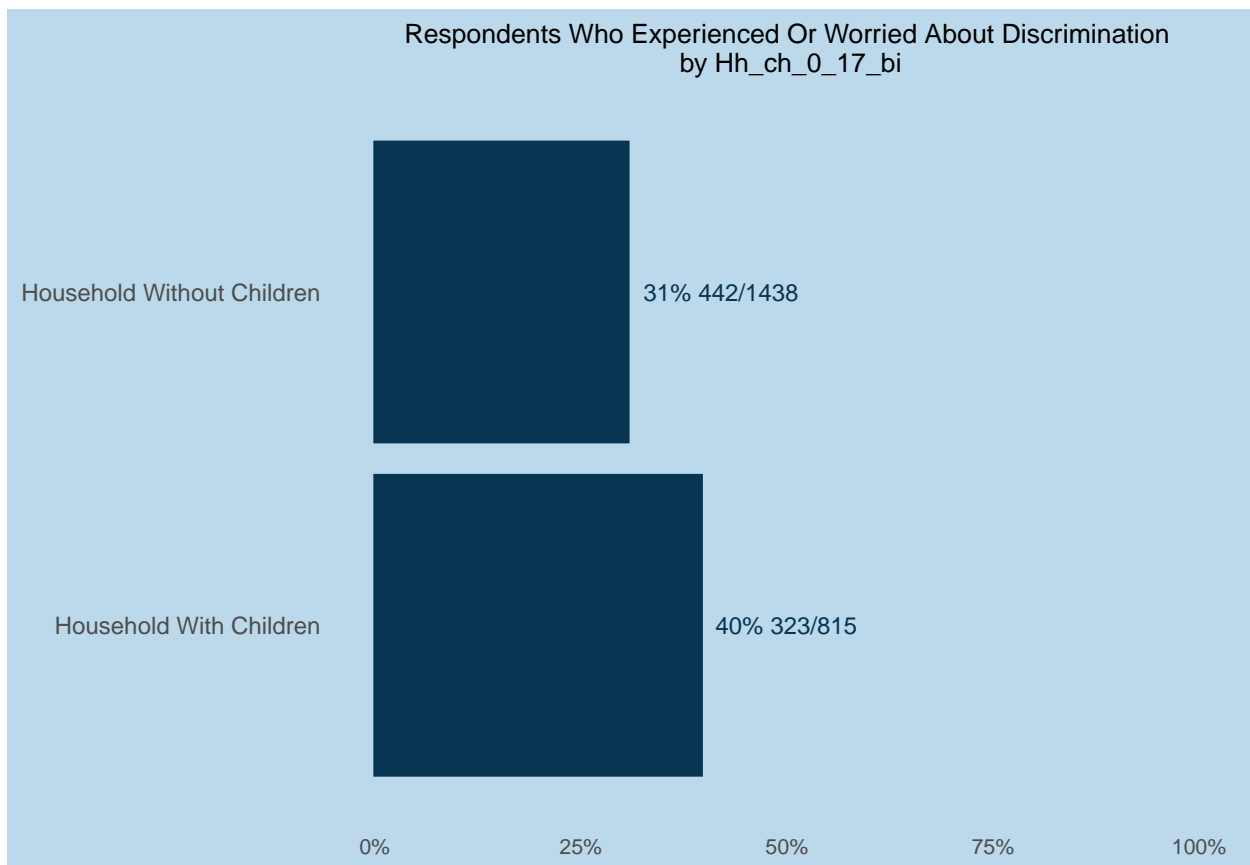


```
##
## $race_census$p.values
## $race_census$p.values$discrim_bi
##
## white (non-hispanic or latino)
## white (non-hispanic or latino) NA
## black or african american 8.4e-03
## hispanic or latinx 2.0e-05
## other NA
## two or more races 1.6e-03
## Indigenous American NA
## asian 4.8e-25
##
## black or african american hispanic or latinx
## white (non-hispanic or latino) 8.4e-03 2.0e-05
## black or african american NA NA
## hispanic or latinx NA NA
## other NA NA
## two or more races NA NA
```

```

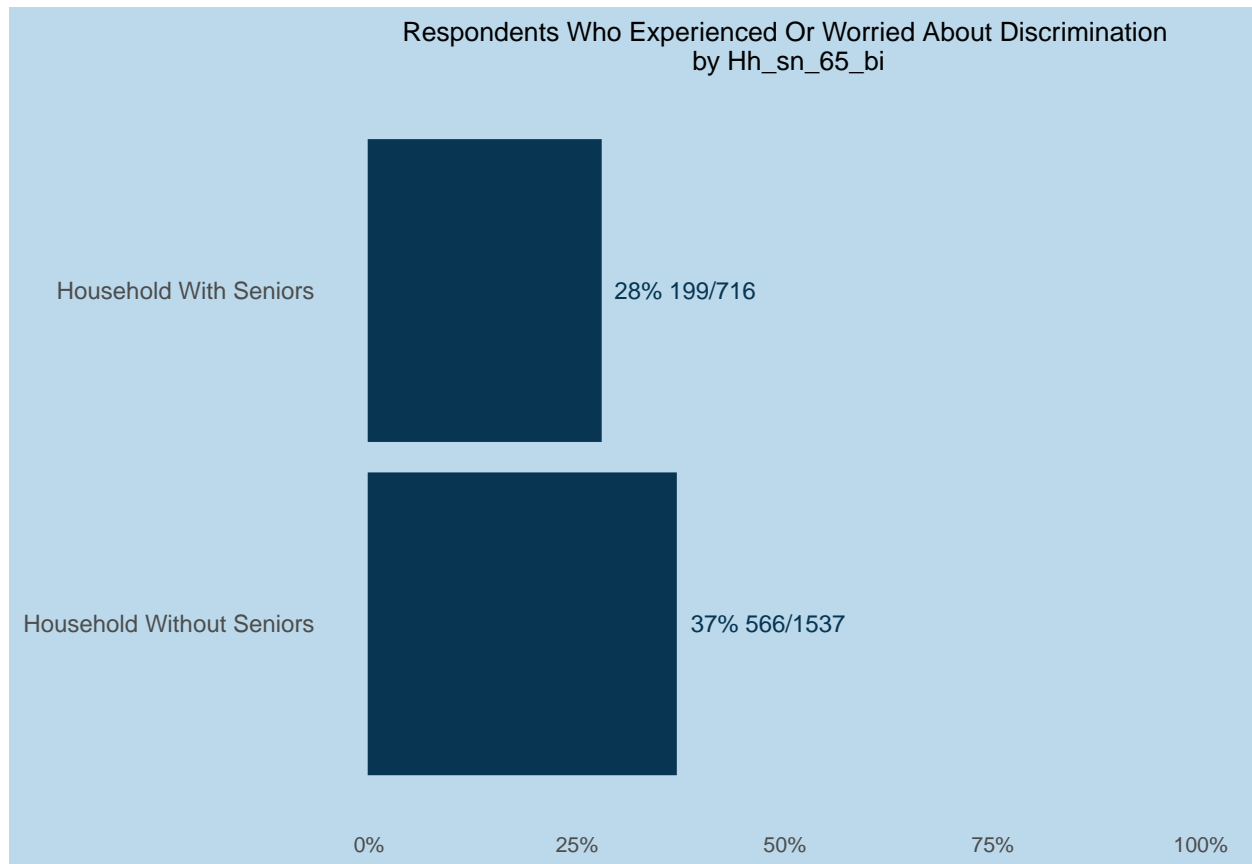
## Indigenous American          NA          NA
## asian                        5.6e-14      4.1e-11
##                               other two or more races Indigenous American
## white (non-hispanic or latino) NA          0.0016          NA
## black or african american     NA          NA          NA
## hispanic or latinx            NA          NA          NA
## other                         NA          NA          NA
## two or more races             NA          NA          NA
## Indigenous American           NA          NA          NA
## asian                        0.00057      NA          NA
##                               asian
## white (non-hispanic or latino) 4.8e-25
## black or african american     5.6e-14
## hispanic or latinx            4.1e-11
## other                         5.7e-04
## two or more races             NA
## Indigenous American           NA
## asian                        NA
##
##
##
## $hh_ch_0_17_bi
## $hh_ch_0_17_bi$plot

```



```
##
```

```
## $hh_ch_0_17_bi$p.values
## $hh_ch_0_17_bi$p.values$discrim_bi
## household without children household with children
## household without children NA 2.3e-05
## household with children 2.3e-05 NA
##
##
##
## $hh_sn_65_bi
## $hh_sn_65_bi$plot
```



```
##
## $hh_sn_65_bi$p.values
## $hh_sn_65_bi$p.values$discrim_bi
## household with seniors household without seniors
## household with seniors NA 3.1e-05
## household without seniors 3.1e-05 NA
##
##
##
## $inc_dist
## NULL
```

```
cat("Plots for gen and inc_dist do not show at least one statistically significant result")
```

```
## Plots for gen and inc_dist do not show at least one statistically significant result
```

## 4.7) People who have experienced abuse or violence due to COVID-19 [34]

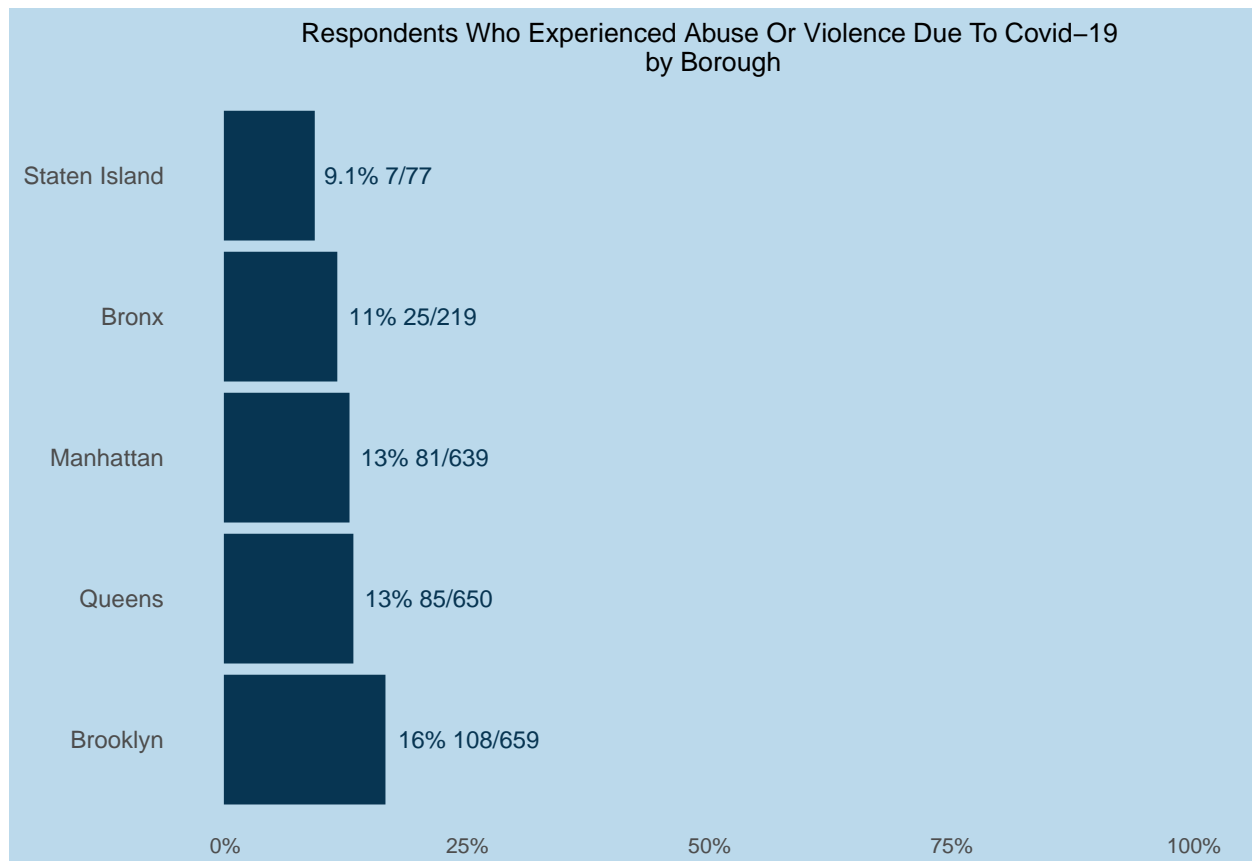
1. Run distribution over population
2. Run distribution by sub-demographics (a-k)
  - a. Compare and find gaps (test unequal proportions)

```
mean(wrangled$exp_ab_or_vi, na.rm = TRUE)
```

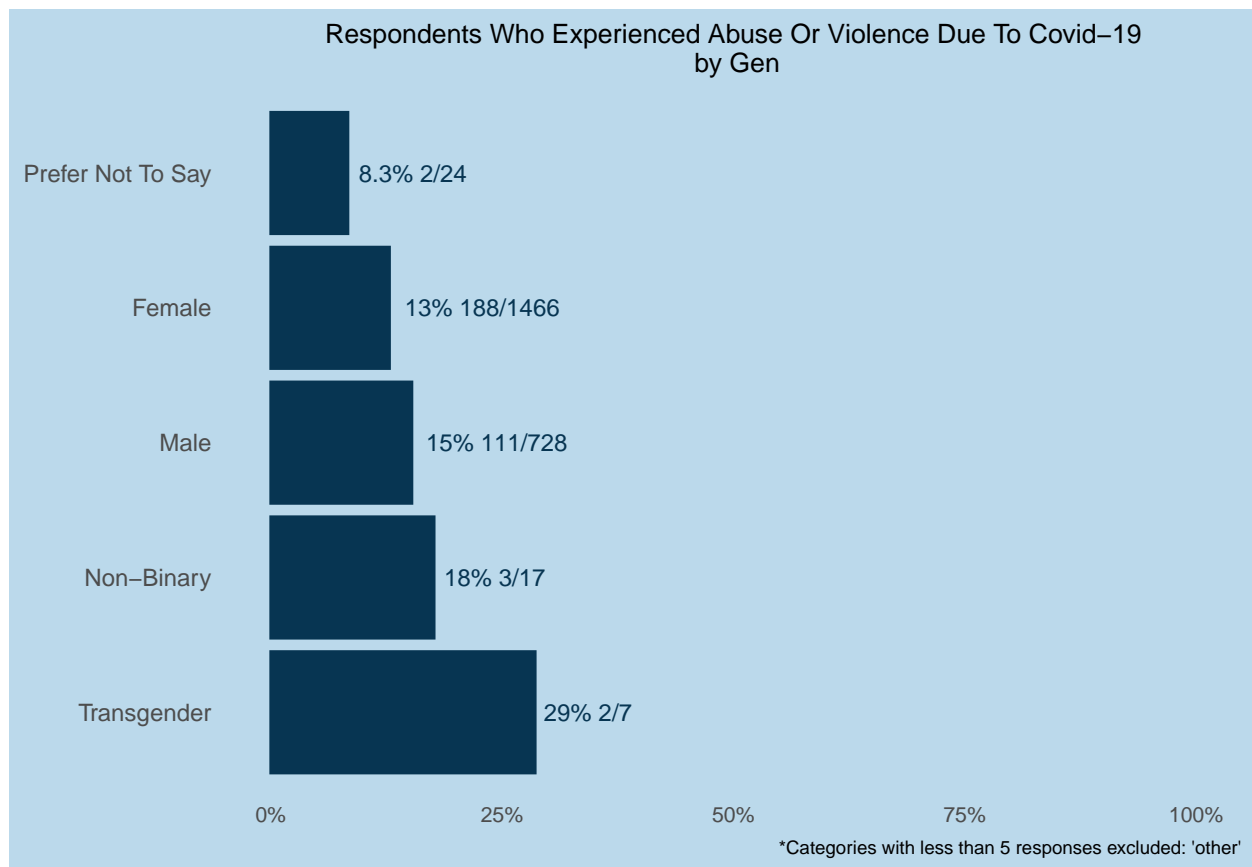
```
## [1] 0.1363636
```

```
#poa %>% mutate(exp_ab_or_vi = exp_ab_or_vi == 1)
make_plots(wrangled, by_vars = demographics, hyp_var = "exp_ab_or_vi",
           title = "Respondents who experienced Abuse or Violence due to COVID-19\n", show = TRUE)
```

```
## $borough
## $borough$plot
```



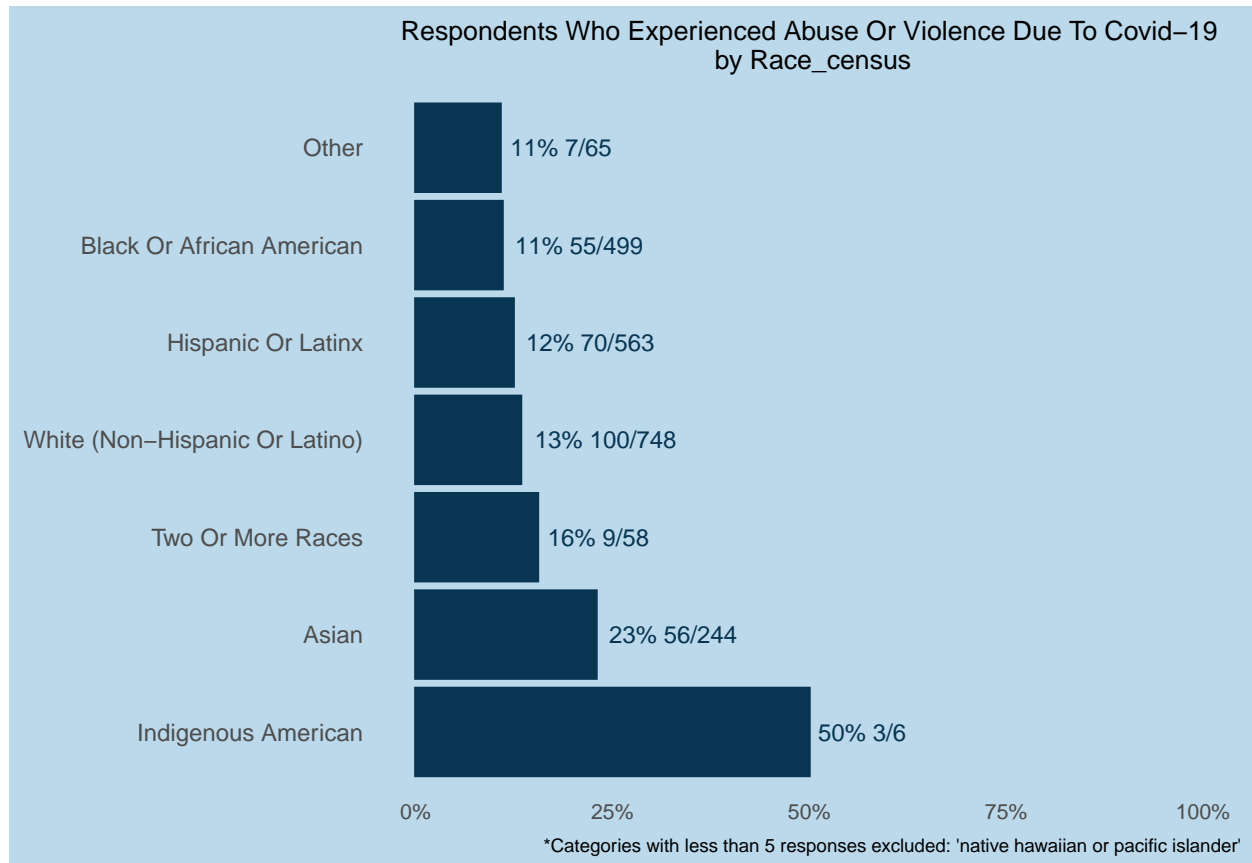
```
##
## $borough$p.values
## $borough$p.values$exp_ab_or_vi
##          staten island bronx manhattan queens brooklyn
## staten island          NA    NA          NA    NA          NA
## bronx                  NA    NA          NA    NA          NA
## manhattan              NA    NA          NA    NA          NA
## queens                 NA    NA          NA    NA          NA
## brooklyn               NA    NA          NA    NA          NA
##
##
##
## $gen
## $gen$plot
```



```
##
## $gen$p.values
## $gen$p.values$exp_ab_or_vi
##          prefer not to say female male non-binary transgender
## prefer not to say          NA    NA    NA          NA          NA
## female                    NA    NA    NA          NA          NA
## male                      NA    NA    NA          NA          NA
## non-binary                NA    NA    NA          NA          NA
## transgender               NA    NA    NA          NA          NA
##
```

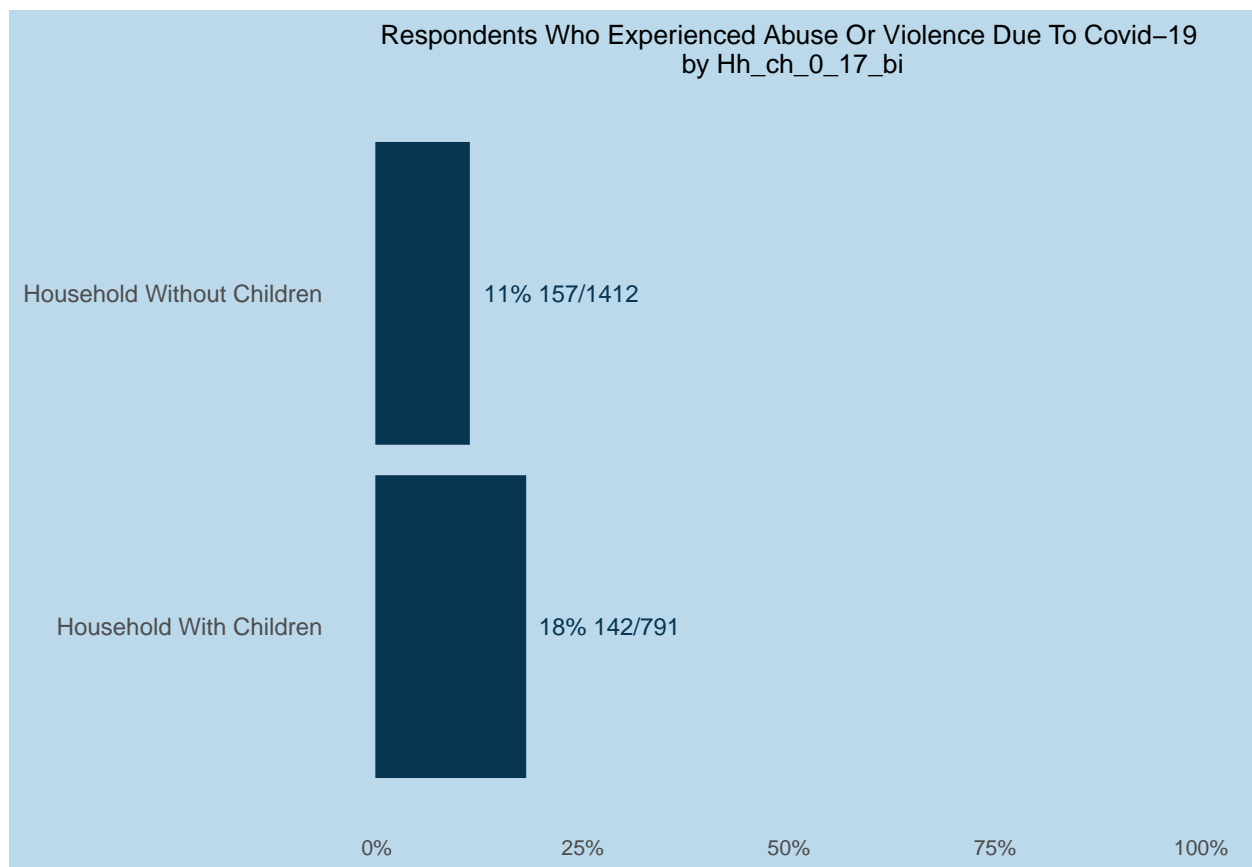


```
##
##
## $race_census
## $race_census$plot
```



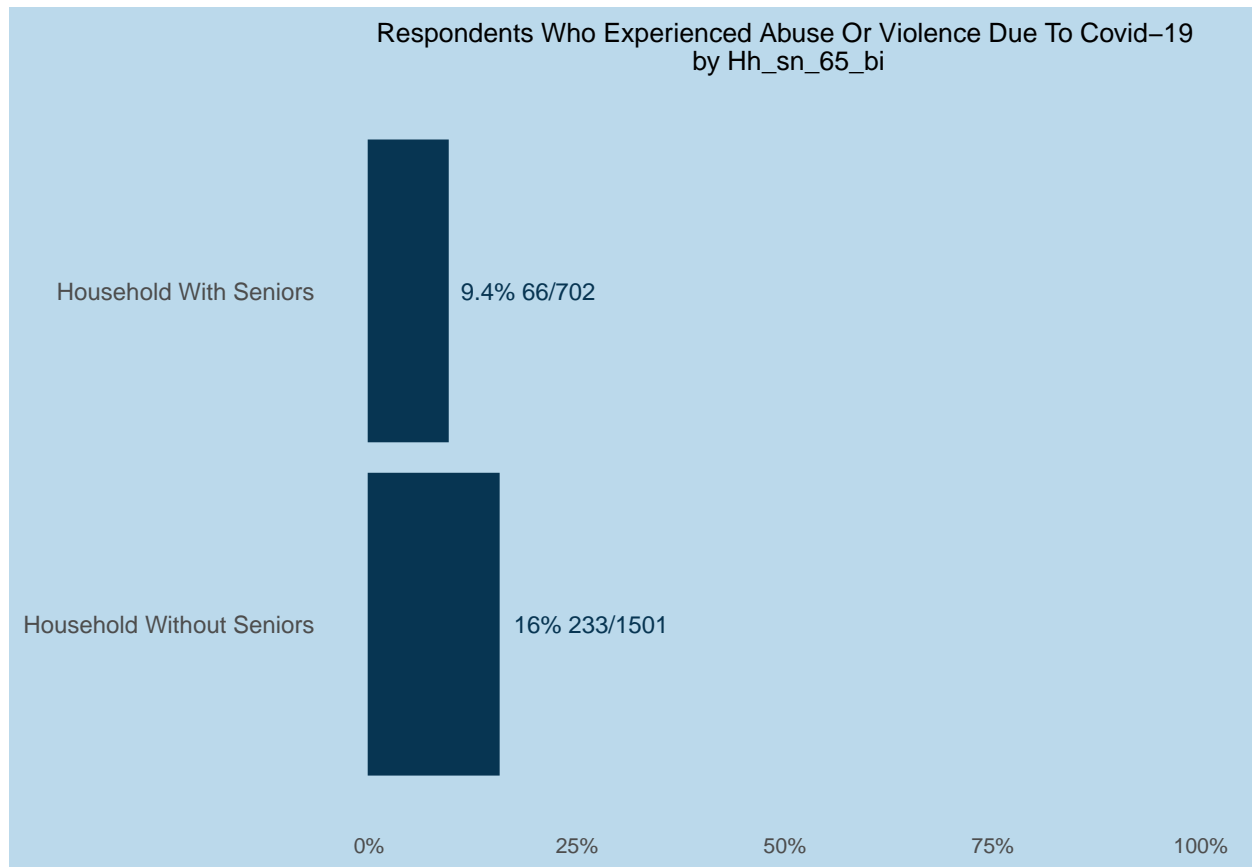
```
##
## $race_census$p.values
## $race_census$p.values$exp_ab_or_vi
##
## other black or african american
## other NA NA
## black or african american NA NA
## hispanic or latinx NA NA
## white (non-hispanic or latino) NA NA
## two or more races NA NA
## asian NA 3e-05
## Indigenous American NA NA
##
## hispanic or latinx
## other NA
## black or african american NA
## hispanic or latinx NA
## white (non-hispanic or latino) NA
## two or more races NA
## asian 0.00024
## Indigenous American NA
##
## white (non-hispanic or latino) two or more races
```

```
## other NA NA
## black or african american NA NA
## hispanic or latinx NA NA
## white (non-hispanic or latino) NA NA
## two or more races NA NA
## asian 0.00052 NA
## Indigenous American NA NA
##
##      asian Indigenous American
## other NA NA
## black or african american 0.00003 NA
## hispanic or latinx 0.00024 NA
## white (non-hispanic or latino) 0.00052 NA
## two or more races NA NA
## asian NA NA
## Indigenous American NA NA
##
##
## $hh_ch_0_17_bi
## $hh_ch_0_17_bi$plot
```

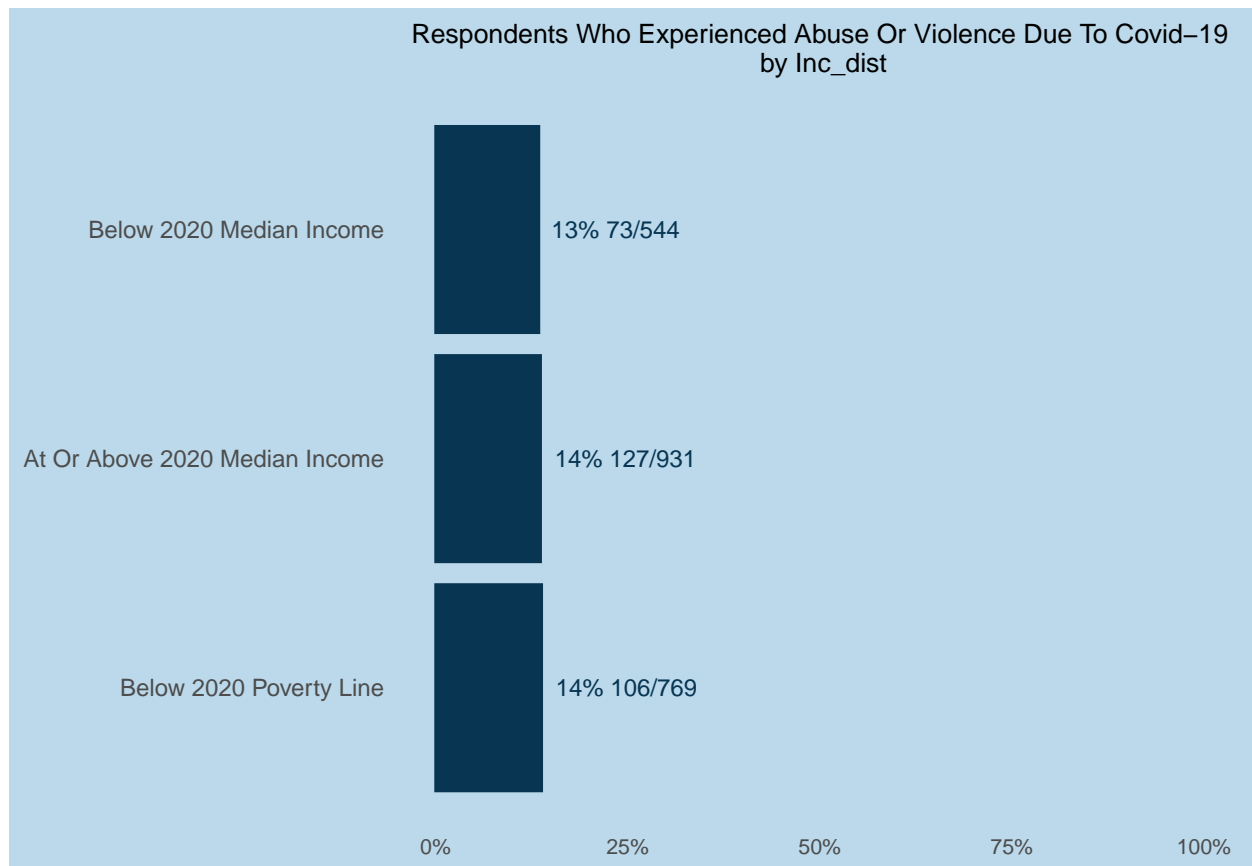


```
##
## $hh_ch_0_17_bi$p.values
## $hh_ch_0_17_bi$p.values$exp_ab_or_vi
##      household without children household with children
```

```
## household without children      NA      9.5e-06
## household with children        9.5e-06      NA
##
##
##
## $hh_sn_65_bi
## $hh_sn_65_bi$plot
```



```
##
## $hh_sn_65_bi$p.values
## $hh_sn_65_bi$p.values$exp_ab_or_vi
## household with seniors household without seniors
## household with seniors      NA      0.00012
## household without seniors    0.00012      NA
##
##
##
## $inc_dist
## $inc_dist$plot
```



```
##
## $inc_dist$p.values
## $inc_dist$p.values$exp_ab_or_vi
##
## below 2020 median income
## below 2020 median income NA
## at or above 2020 median income NA
## below 2020 poverty line NA
##
## at or above 2020 median income
## below 2020 median income NA
## at or above 2020 median income NA
## below 2020 poverty line NA
##
## below 2020 poverty line
## below 2020 median income NA
## at or above 2020 median income NA
## below 2020 poverty line NA

cat("Plots for borough, gen, and inc_dist do not show at least one statistically significant result")
```

```
## Plots for borough, gen, and inc_dist do not show at least one statistically significant result
```

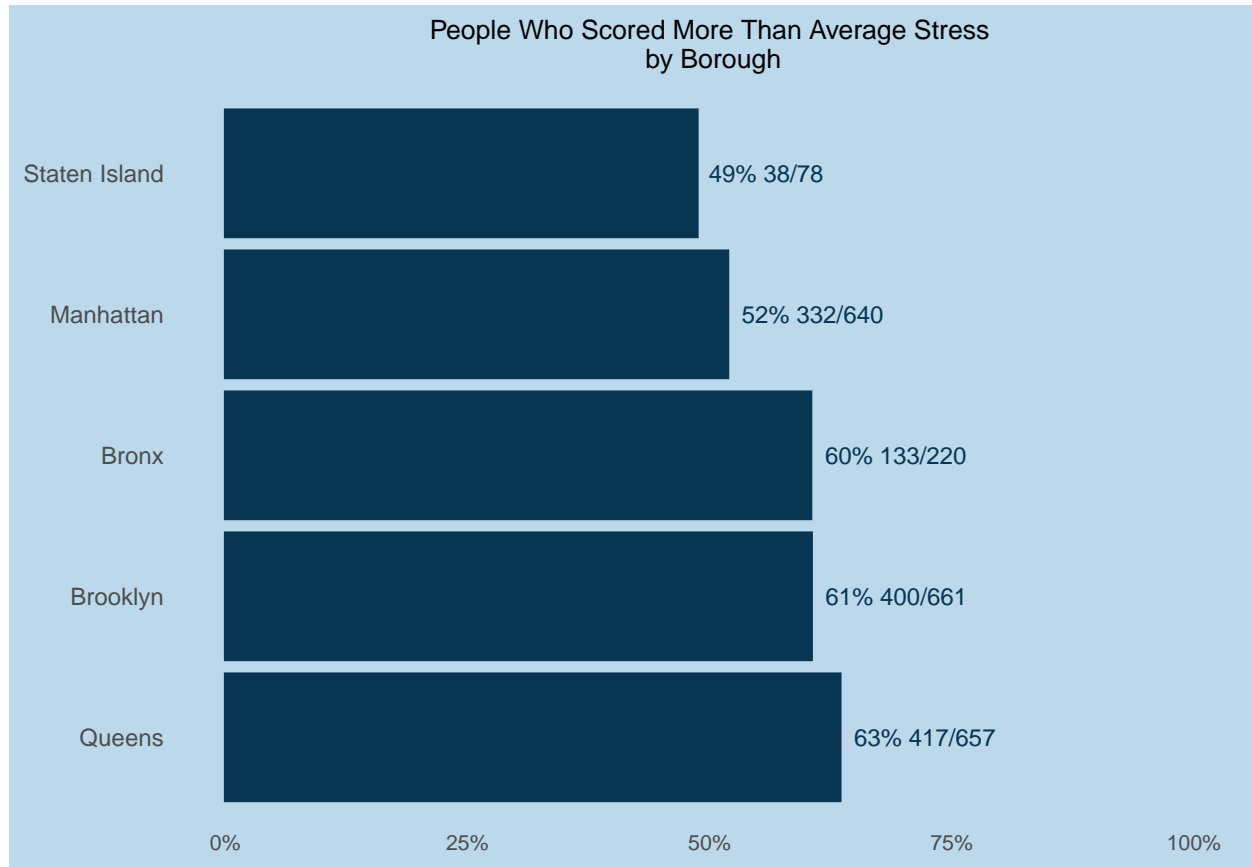
#### 4.8) People who score more than average on the quantified mental health quantified questions [38]

1. Run categorical distribution over population

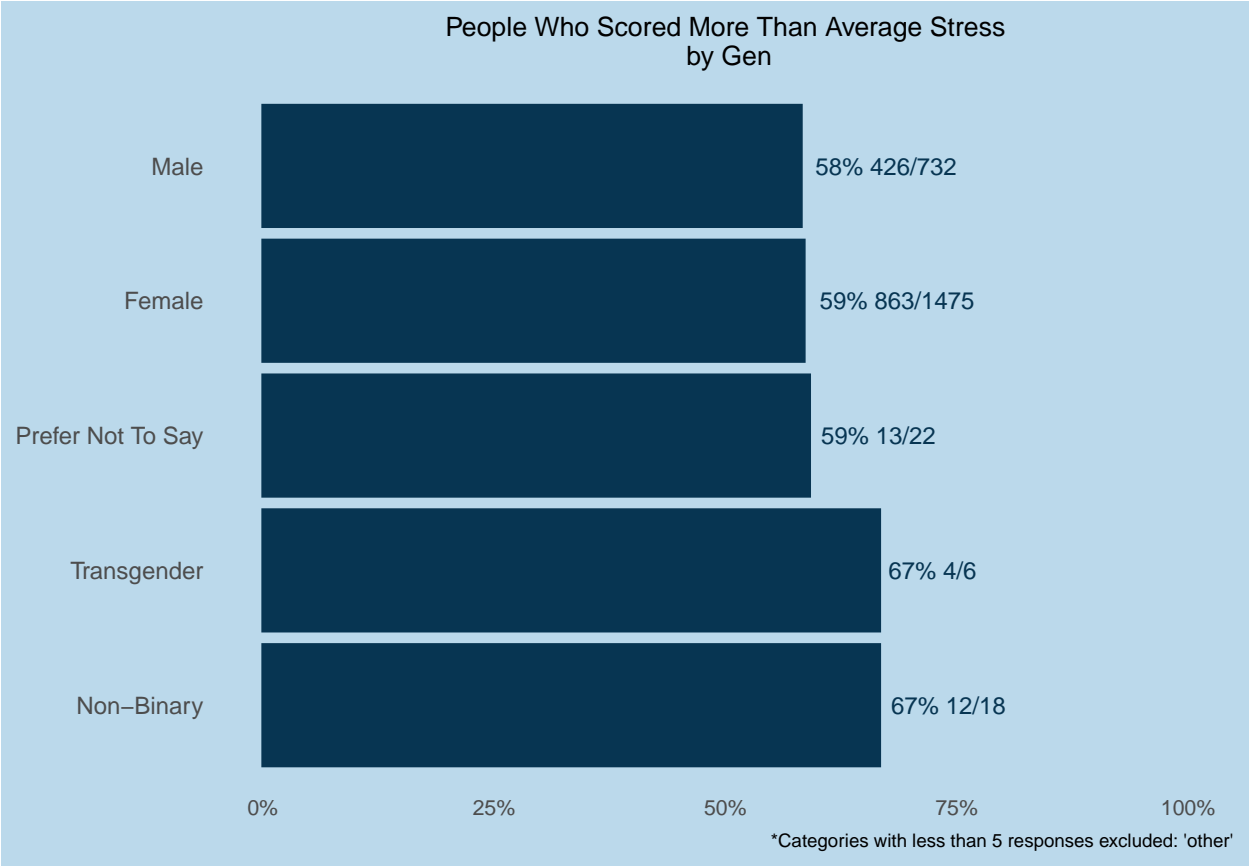
## 2. Run distribution by sub-demographics

```
make_plots(wrangled, demographics, "stress_bi", title = "People who scored more than average stress\n",
```

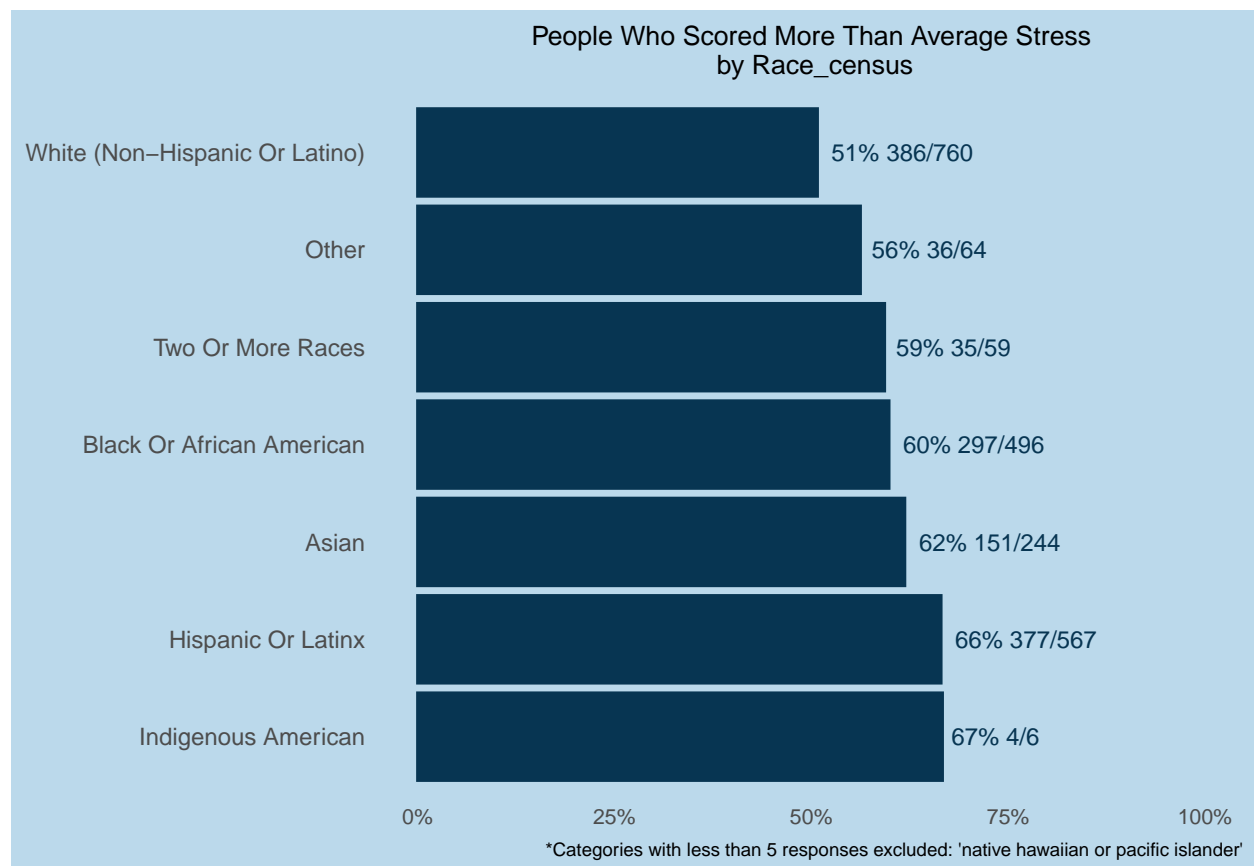
```
## $borough
## $borough$plot
```



```
##
## $borough$p.values
## $borough$p.values$stress_bi
##          staten island manhattan bronx brooklyn queens
## staten island          NA          NA          NA          NA          NA
## manhattan              NA          NA          NA          0.002 3e-05
## bronx                  NA          NA          NA          NA          NA
## brooklyn               NA          2e-03          NA          NA          NA
## queens                 NA          3e-05          NA          NA          NA
##
##
##
## $gen
## $gen$plot
```

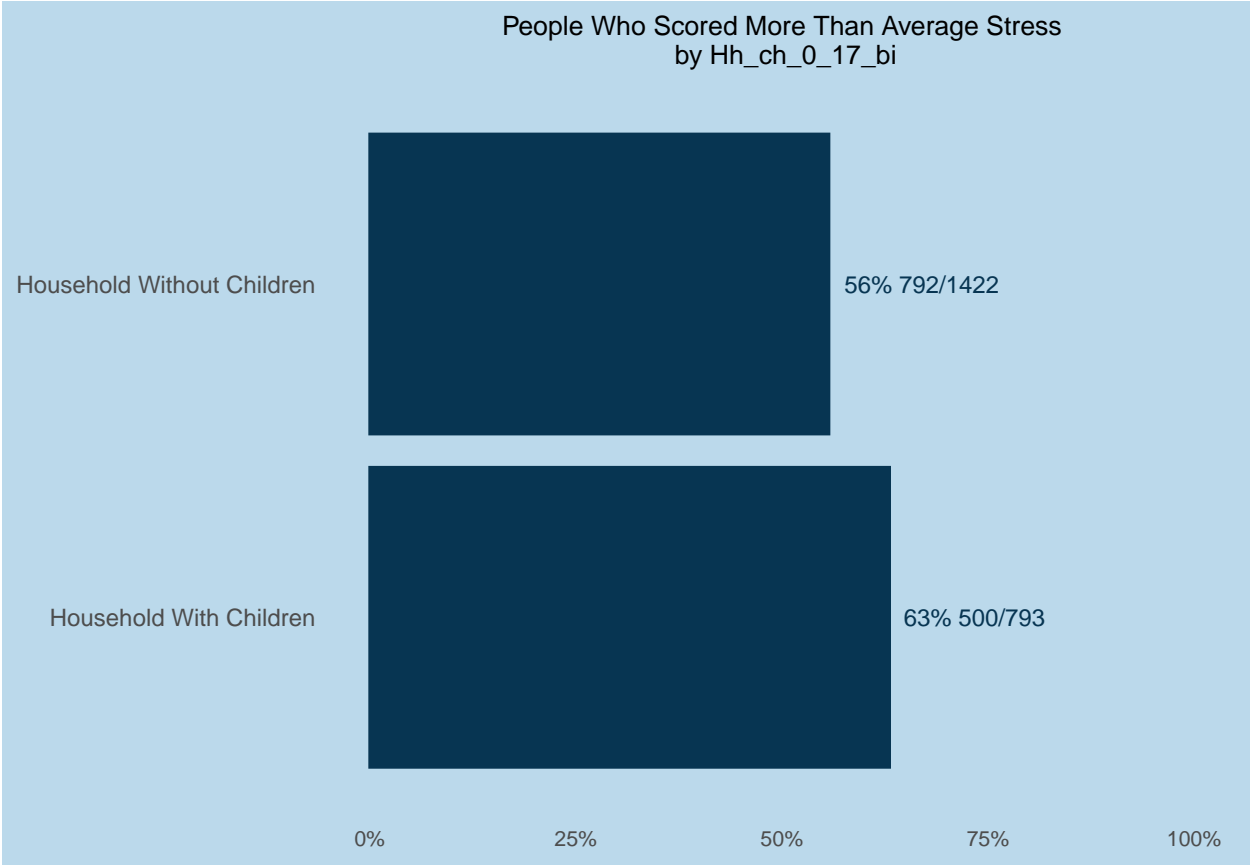


```
##
## $gen$p.values
## $gen$p.values$stress_bi
##           male female prefer not to say non-binary transgender
## male           NA    NA              NA      NA              NA
## female          NA    NA              NA      NA              NA
## prefer not to say NA    NA              NA      NA              NA
## non-binary       NA    NA              NA      NA              NA
## transgender      NA    NA              NA      NA              NA
##
##
##
## $race_census
## $race_census$plot
```



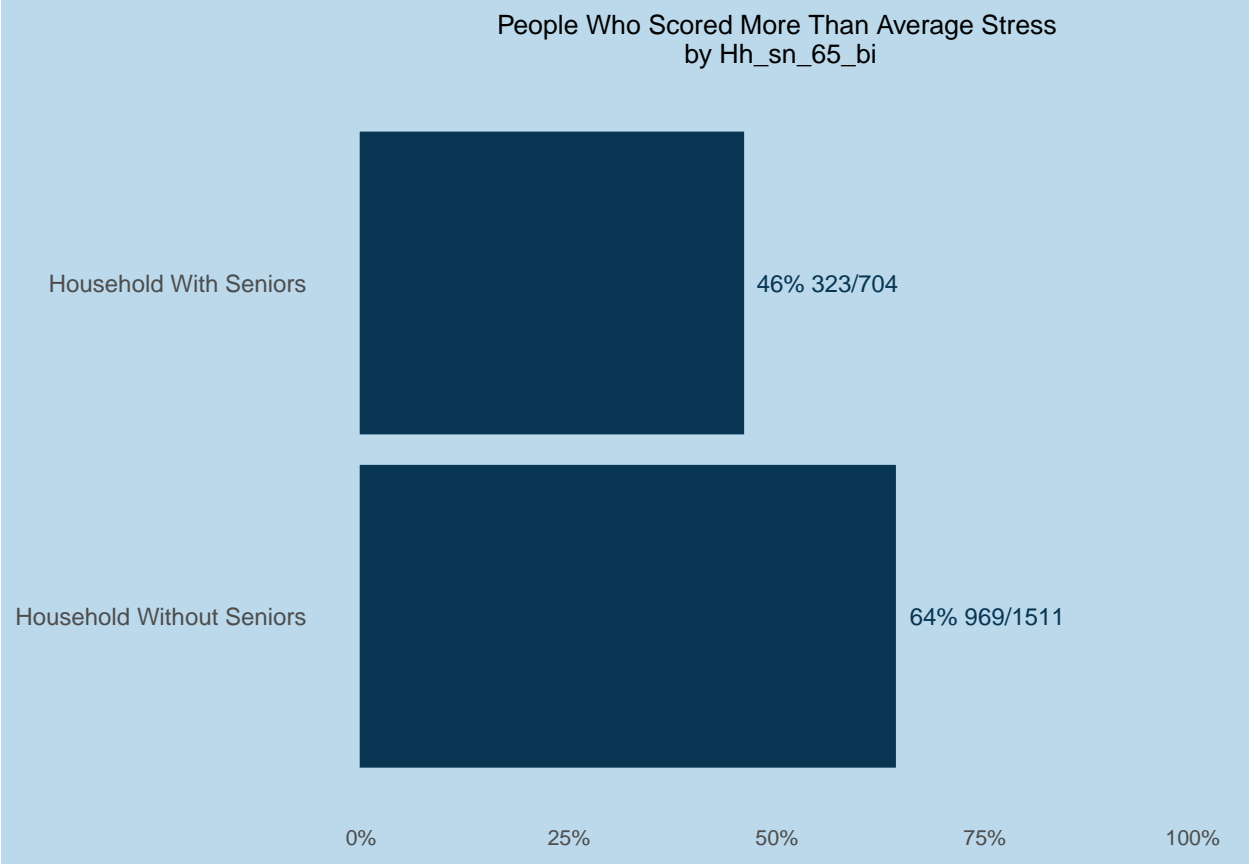
```
##
## $race_census$p.values
## $race_census$p.values$stress_bi
##               white (non-hispanic or latino) other
## white (non-hispanic or latino)              NA  NA
## other                                         NA  NA
## two or more races                          NA  NA
## black or african american                  1.9e-03 NA
## asian                                       3.2e-03 NA
## hispanic or latinx                        1.5e-08 NA
## Indigenous American                       NA  NA
##               two or more races black or african american
## white (non-hispanic or latino)            NA      0.0019
## other                                       NA      NA
## two or more races                         NA      NA
## black or african american                 NA      NA
## asian                                     NA      NA
## hispanic or latinx                       NA      NA
## Indigenous American                      NA      NA
##               asian hispanic or latinx Indigenous American
## white (non-hispanic or latino) 0.0032      1.5e-08      NA
## other                          NA      NA      NA
## two or more races              NA      NA      NA
## black or african american      NA      NA      NA
## asian                          NA      NA      NA
## hispanic or latinx             NA      NA      NA
```

```
## Indigenous American      NA      NA      NA
##
##
##
## $hh_ch_0_17_bi
## $hh_ch_0_17_bi$plot
```

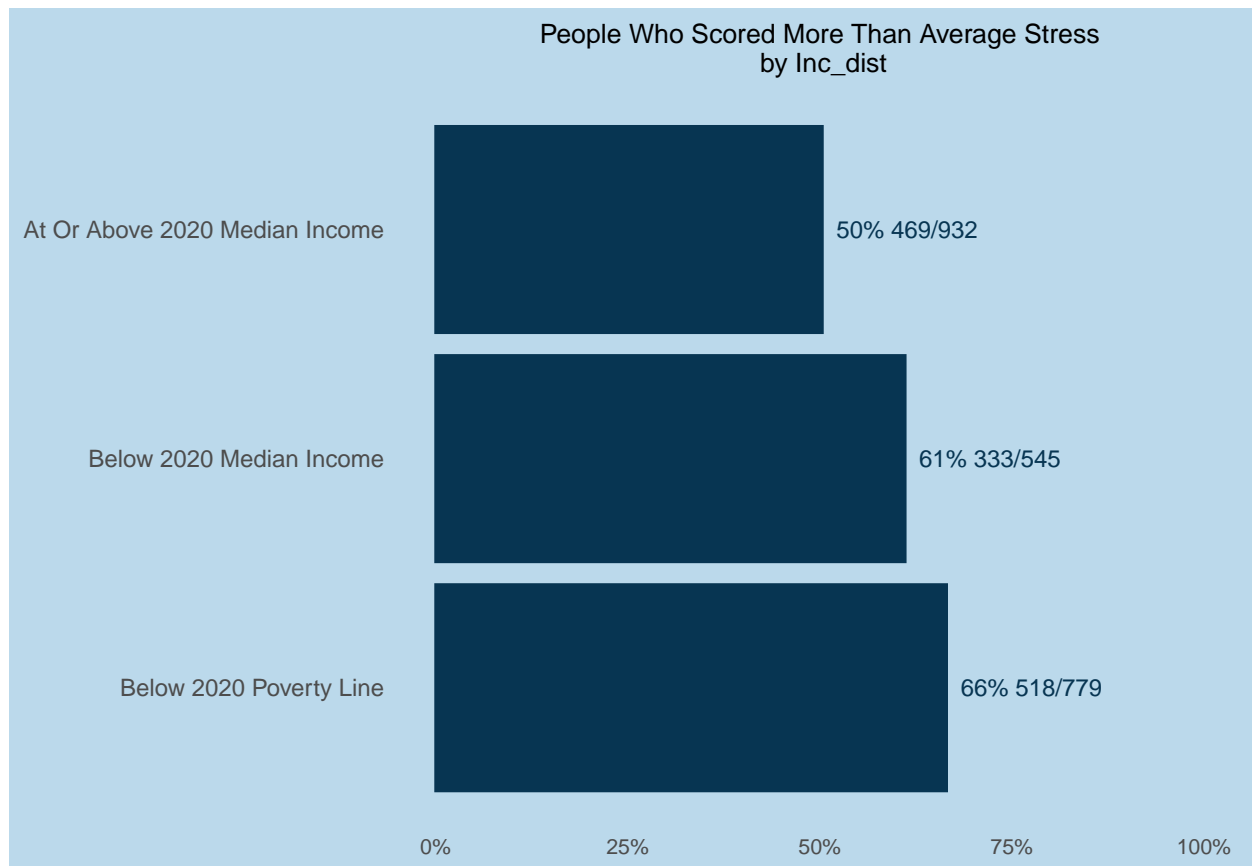


```
##
## $hh_ch_0_17_bi$p.values
## $hh_ch_0_17_bi$p.values$stress_bi
##           household without children household with children
## household without children           NA           9e-04
## household with children           9e-04           NA
##
##
##
## $hh_sn_65_bi
## $hh_sn_65_bi$plot
```





```
##
## $hh_sn_65_bi$p.values
## $hh_sn_65_bi$p.values$stress_bi
##           household with seniors household without seniors
## household with seniors              NA              7.3e-16
## household without seniors          7.3e-16              NA
##
##
##
## $inc_dist
## $inc_dist$plot
```



```
##
## $inc_dist$p.values
## $inc_dist$p.values$stress_bi
##
## at or above 2020 median income
## at or above 2020 median income NA
## below 2020 median income 7.5e-05
## below 2020 poverty line 2.2e-11
##
## below 2020 median income below 2020 poverty line
## at or above 2020 median income 7.5e-05 2.2e-11
## below 2020 median income NA NA
## below 2020 poverty line NA NA
```

```
cat("Plot for gen does not show at least one statistically significant result")
```

```
## Plot for gen does not show at least one statistically significant result
```

## 4.9 - 4.12

**4.9) People who rated government response poorly were more likely to score higher than average on the mental health score.**

1. Find proportion of population who rated government services below average
2. Find mental health score amongst people who rated government services below average as compared to those who rated government services above average.

#### 4.10) People who are currently unemployed are more likely to score higher than average on the mental health score

1. Find respondents who are currently unemployed [15]
  - a. Find proportion of subset that scored higher than average on the mental health score [38]
  - b. Find proportion not in subset and compare (test unequal proportions)

#### 4.11) People who have experienced abuse or violence are more likely to score higher than average on the mental health score

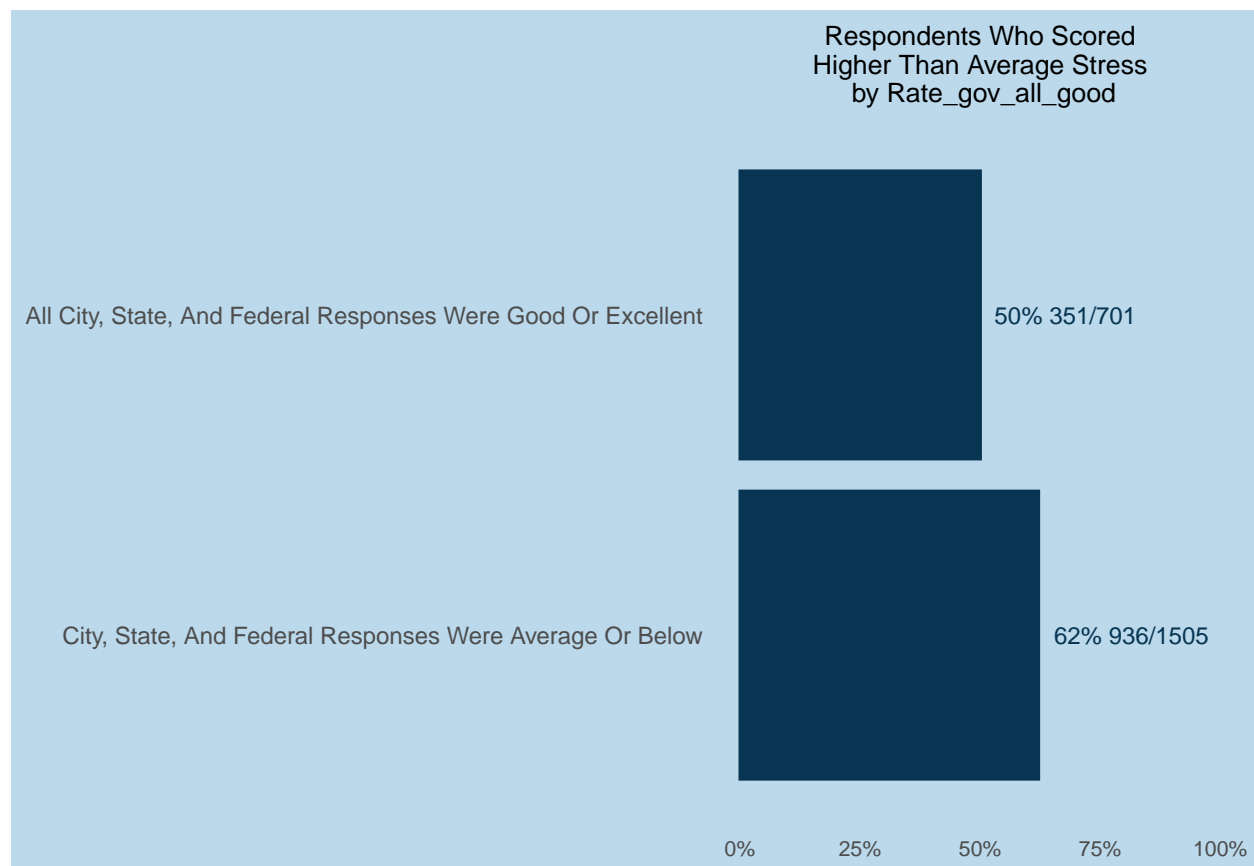
1. Run binary distribution over population [36]
  - a. Yes = experienced verbal abuse or physical abuse
  - b. No = has not experienced any abuse or violence
2. Find respondents who have experienced violence or abuse
  - a. Find proportion of subset that scored higher than average on the mental health score [38]
  - b. Find proportion not in subset and compare (test unequal proportions)

#### 4.12) People who have been discriminated against are more likely to score higher than average on the mental health score

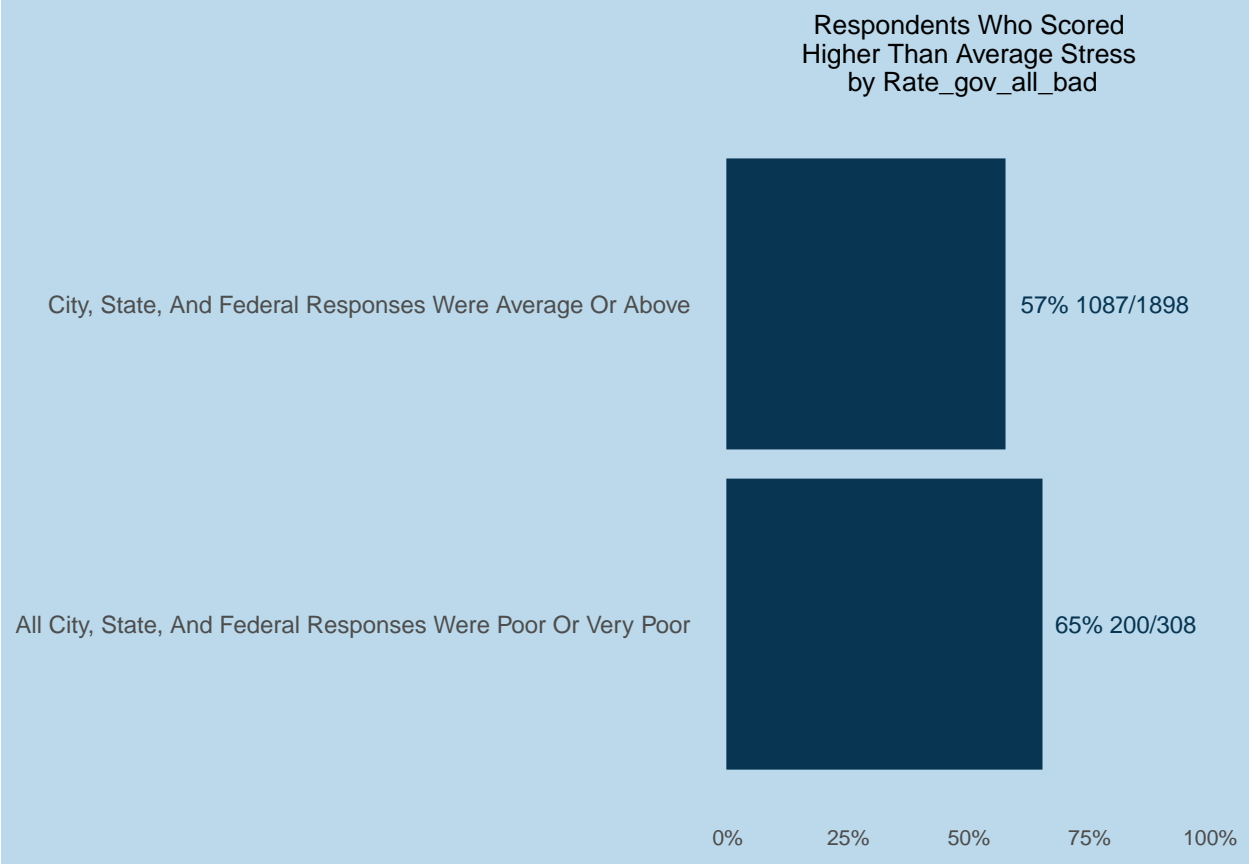
1. Find respondents who have been discriminated against [37]
  - a. Find proportion of subset that scored higher than average on the mental health score [38]
  - b. Find proportion not in subset and compare (test unequal proportions)

```
make_plots(wrangled,
  c(
    # 4.9
    "rate_gov_all_good", "rate_gov_all_bad",
    # 4.10
    "emp_after_un",
    # 4.11
    "exp_ab_or_vi",
    "exp_ab_and_vi",
    # 4.12
    "discrim_bi"),
  "stress_bi", title = "Respondents who scored\nhigher than average stress\n", show = TRUE)

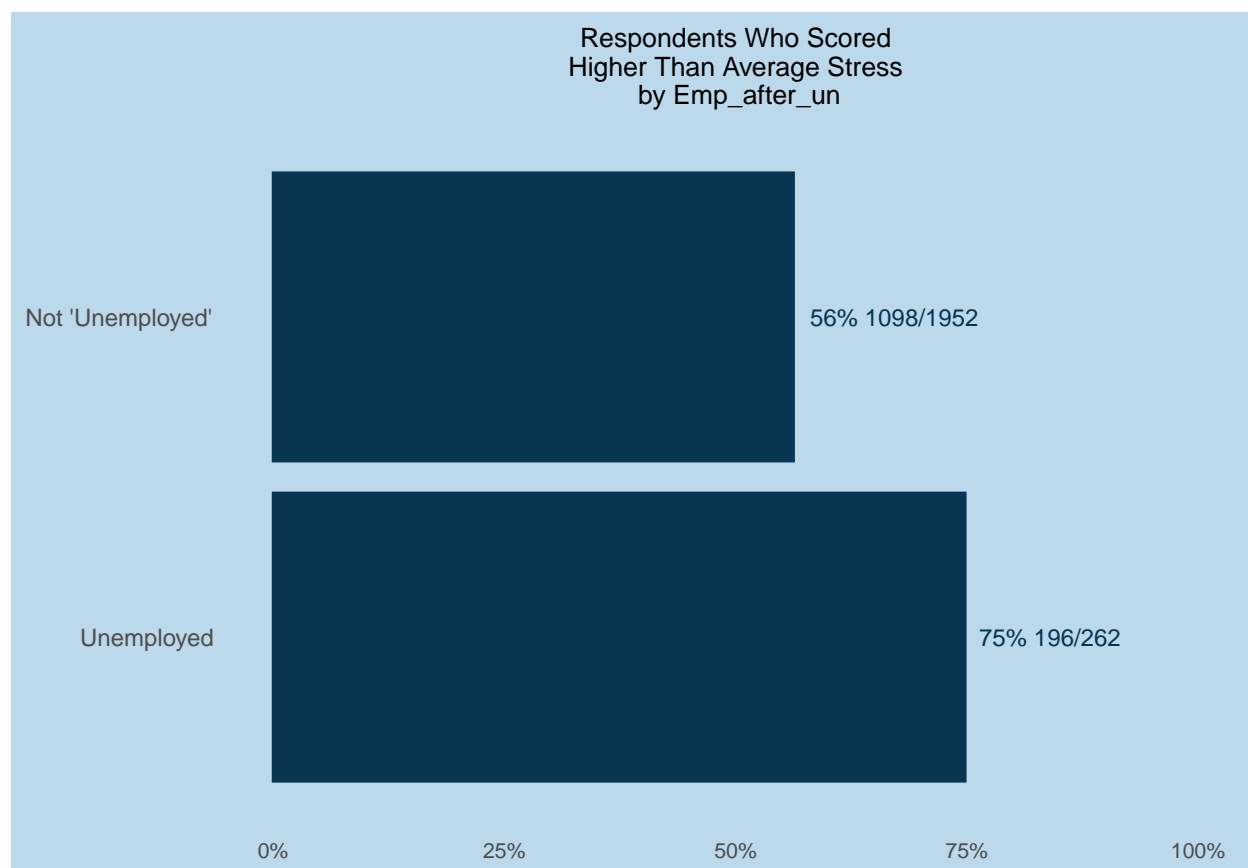
## $rate_gov_all_good
## $rate_gov_all_good$plot
```



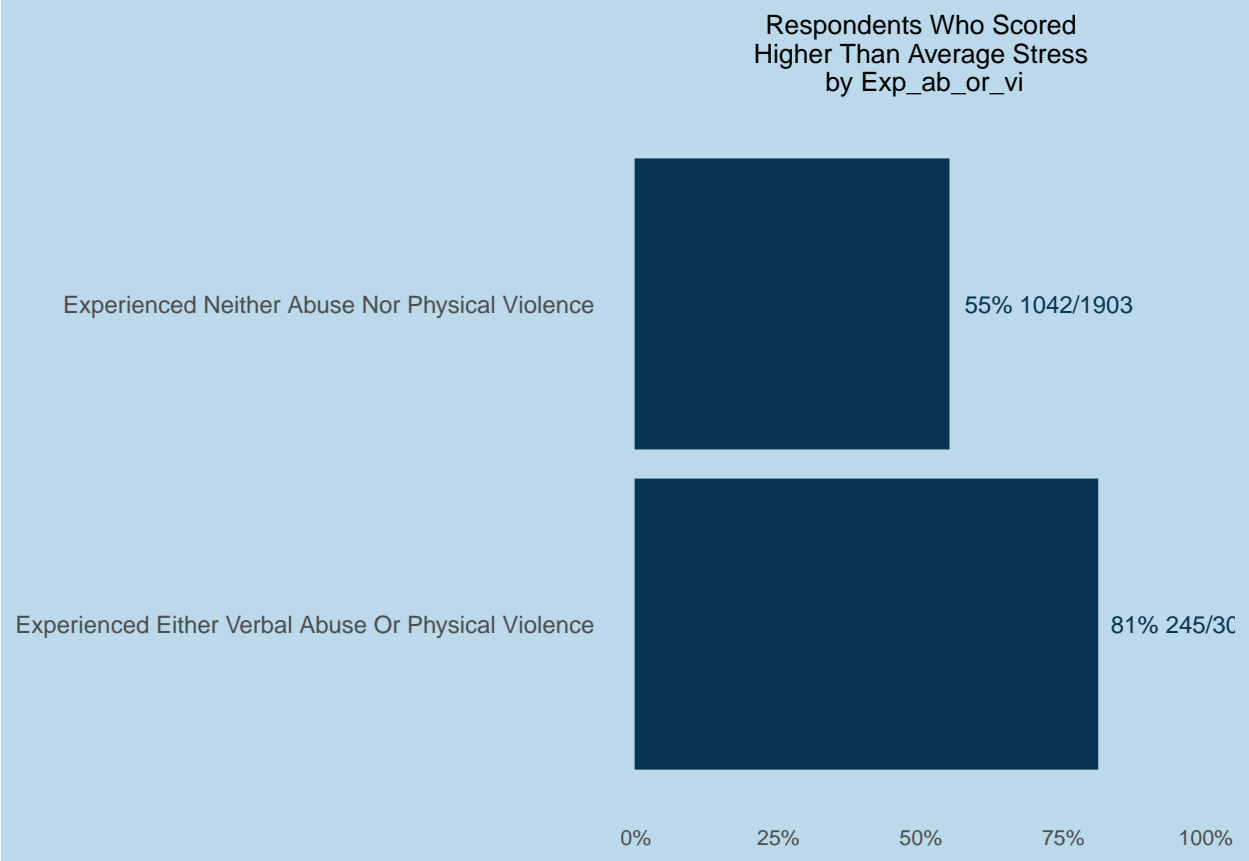
```
##
## $rate_gov_all_good$p.values
## $rate_gov_all_good$p.values$stress_bi
## all city, state, and federal responses
## all city, state, and federal responses were good or excellent
## city, state, and federal responses were average or below
## all city, state, and federal responses were good or excellent
## city, state, and federal responses were average or below
##
##
## $rate_gov_all_bad
## $rate_gov_all_bad$plot
```



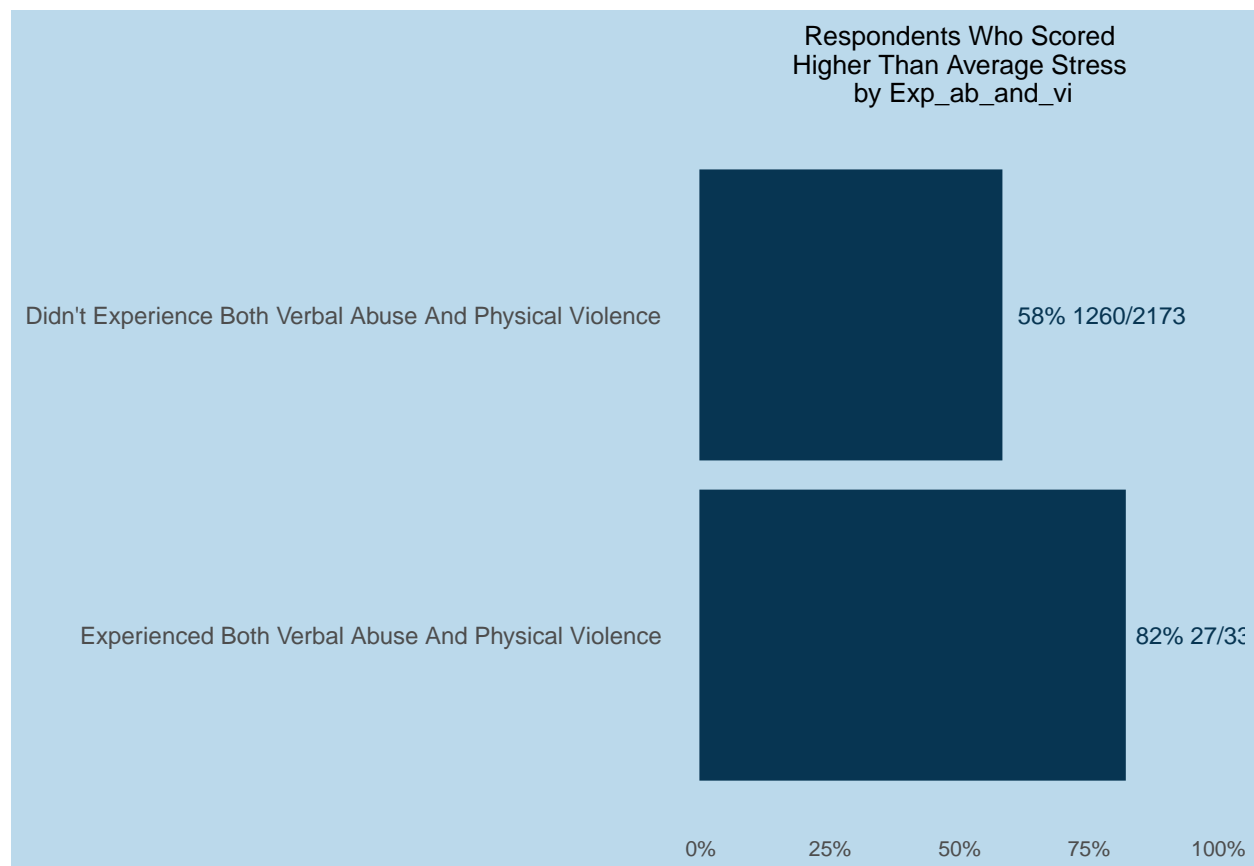
```
##
## $rate_gov_all_bad$p.values
## $rate_gov_all_bad$p.values$stress_bi
##
## city, state, and federal responses were average or above
## all city, state, and federal responses were poor or very poor
##
## city, state, and federal responses were average or above
## all city, state, and federal responses were poor or very poor
##
##
##
## $emp_after_un
## $emp_after_un$plot
```



```
##
## $emp_after_un$p.values
## $emp_after_un$p.values$stress_bi
##           not 'unemployed' unemployed
## not 'unemployed'           NA      1.5e-08
## unemployed           1.5e-08           NA
##
##
##
## $exp_ab_or_vi
## $exp_ab_or_vi$plot
```

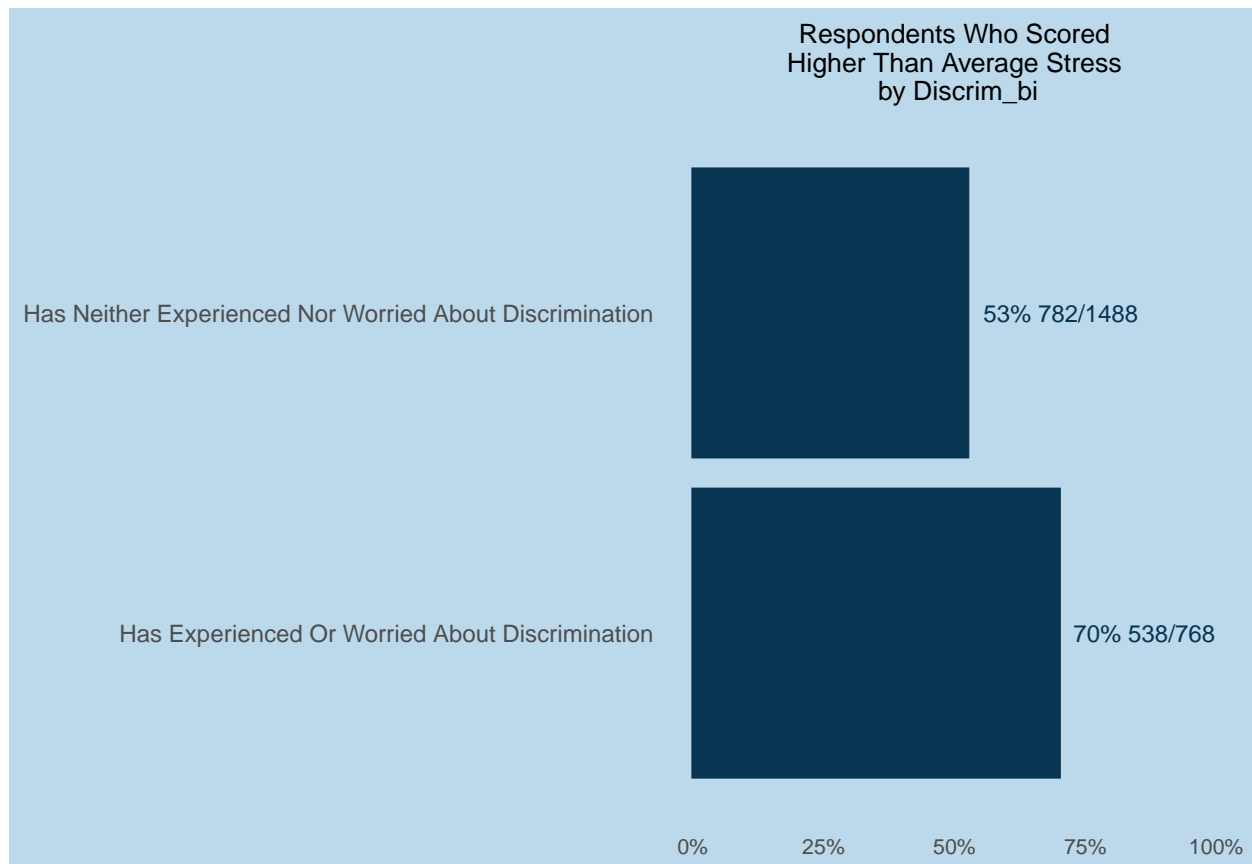


```
##
## $exp_ab_or_vi$p.values
## $exp_ab_or_vi$p.values$stress_bi
##
## experienced neither abuse nor physical violence NA
## experienced either verbal abuse or physical violence 1.9e-17
## experienced neither abuse nor physical violence 1.9e-17
## experienced either verbal abuse or physical violence 1.9e-17
##
##
## $exp_ab_and_vi
## $exp_ab_and_vi$plot
```



```
##
## $exp_ab_and_vi$p.values
## $exp_ab_and_vi$p.values$stress_bi
##
## didn't experience both verbal abuse and physical violence
## experienced both verbal abuse and physical violence
##
## didn't experience both verbal abuse and physical violence
## experienced both verbal abuse and physical violence
##
##
##
## $discrim_bi
## $discrim_bi$plot
```





```
##
## $discrim_bi$p.values
## $discrim_bi$p.values$stress_bi
## has neither experienced nor worried about discrimination
## has experienced or worried about discrimination
## has experienced or worried about discrimination
## has experienced or worried about discrimination
## has experienced or worried about discrimination
## has experienced or worried about discrimination
```

```
cat("Plots for rate_gov_all_bad and exp_ab_and_vi are not showing at least one statistically significant")
```

```
## Plots for rate_gov_all_bad and exp_ab_and_vi are not showing at least one statistically significant
```

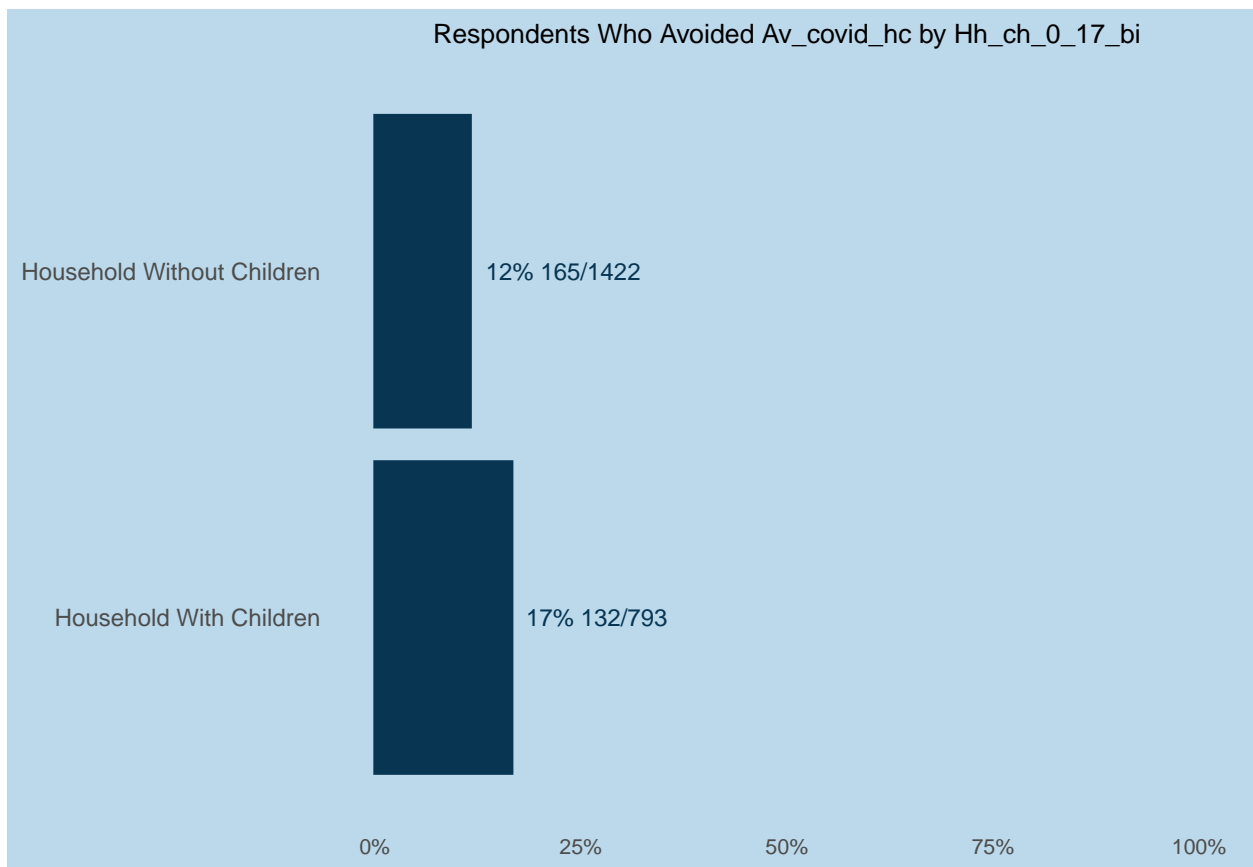
#### 4.13) Run distribution over each activity which has been avoided because of Covid-19

1. Find respondents who avoided any activity because of Covid-19
2. Run sub demographic distribution over each activity

```
av_covid <- wrangled %>% select(starts_with("av_covid_") & !ends_with("text")) %>% colnames
```

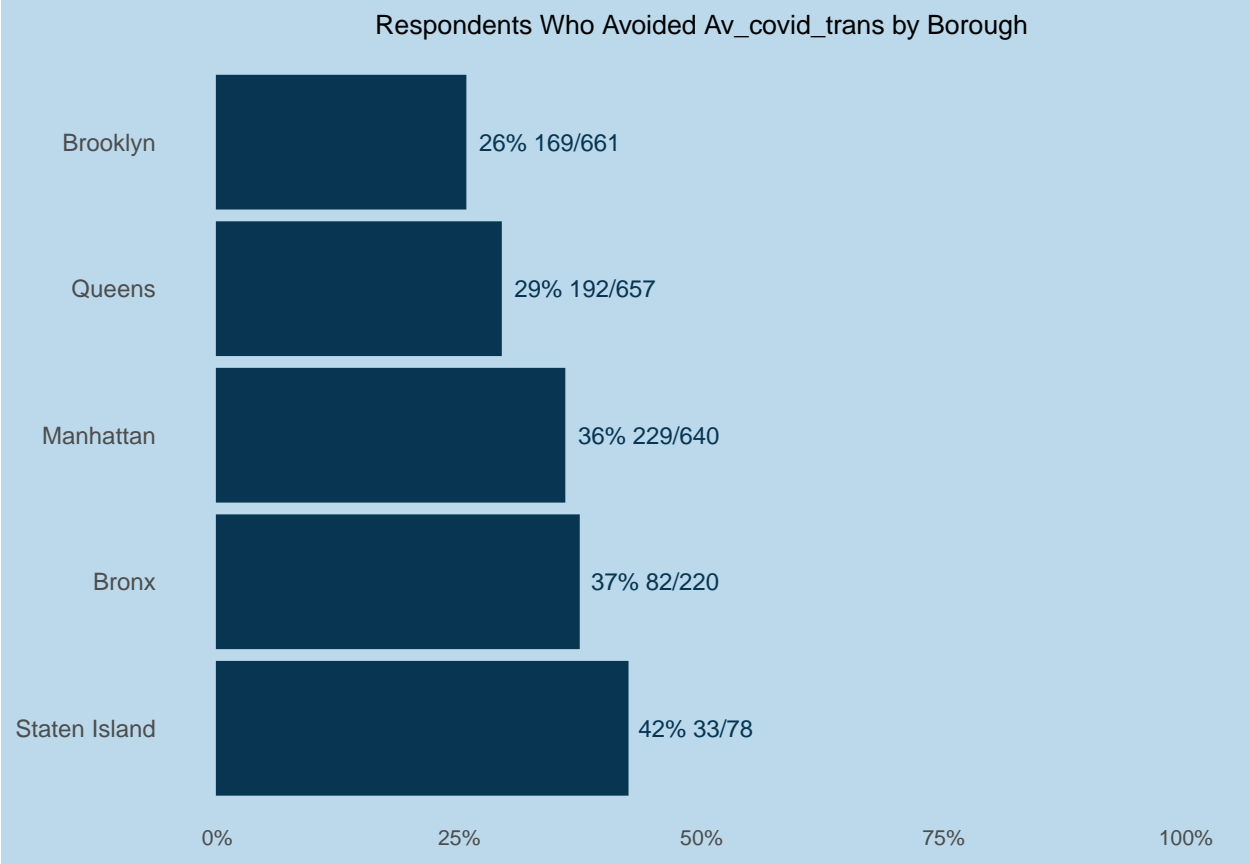
```
lapply(av_covid, function(var) {
  make_plots(wrangled, demographics, var, title = glue::glue("Respondents who avoided {var}"))
})
```

```
## [[1]]
## [[1]]$borough
## NULL
##
## [[1]]$gen
## NULL
##
## [[1]]$race_census
## NULL
##
## [[1]]$hh_ch_0_17_bi
## [[1]]$hh_ch_0_17_bi$plot
```

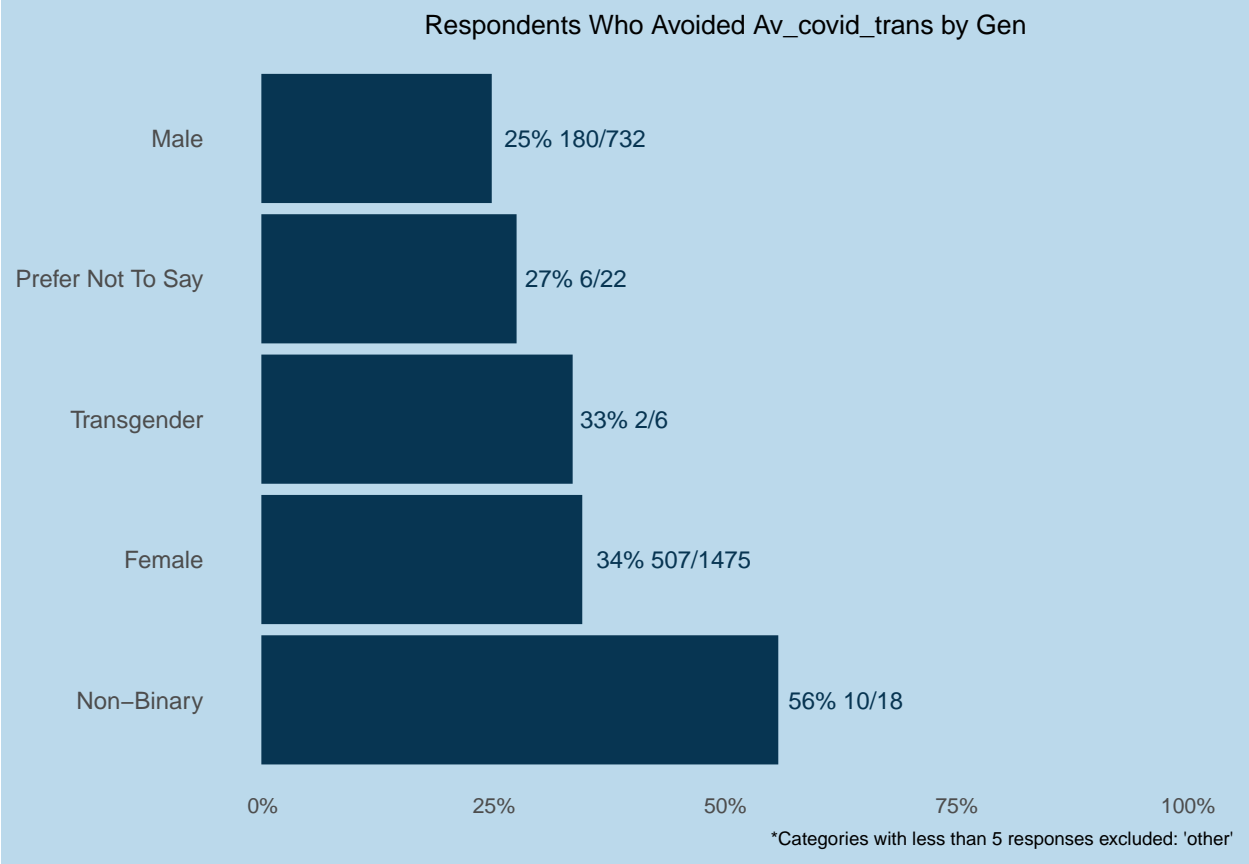


```
##
## [[1]]$hh_ch_0_17_bi$p.values
## [[1]]$hh_ch_0_17_bi$p.values$av_covid_hc
## household without children household with children
## household without children NA 0.0011
## household with children 0.0011 NA
##
```

```
##
##
## [[1]]$hh_sn_65_bi
## NULL
##
## [[1]]$inc_dist
## NULL
##
##
## [[2]]
## [[2]]$borough
## NULL
##
## [[2]]$gen
## NULL
##
## [[2]]$race_census
## NULL
##
## [[2]]$hh_ch_0_17_bi
## NULL
##
## [[2]]$hh_sn_65_bi
## NULL
##
## [[2]]$inc_dist
## NULL
##
##
## [[3]]
## [[3]]$borough
## [[3]]$borough$plot
```

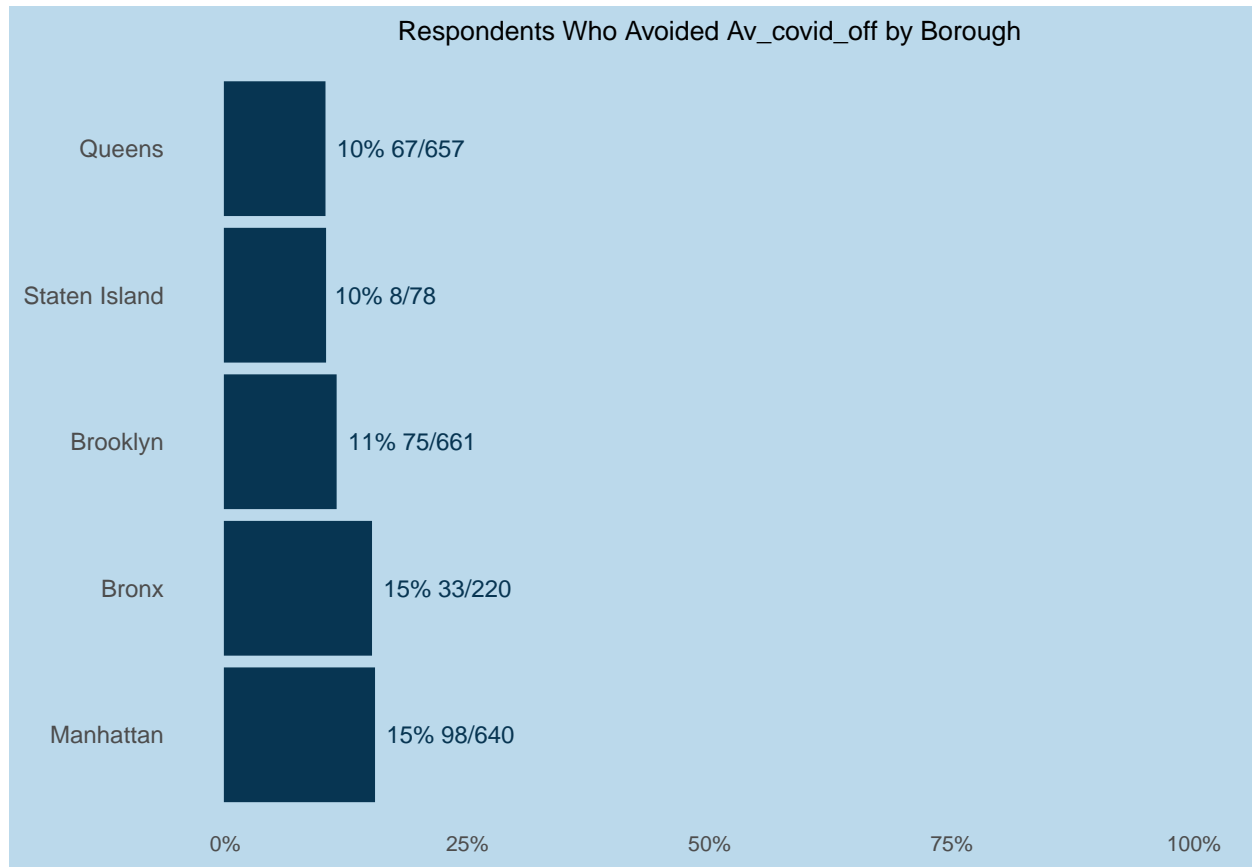


```
##
## [[3]]$borough$p.values
## [[3]]$borough$p.values$av_covid_trans
##      brooklyn queens manhattan  bronx staten island
## brooklyn      NA    NA  8.3e-05 0.0012      0.0027
## queens        NA    NA      NA    NA      NA
## manhattan     8.3e-05  NA      NA    NA      NA
## bronx         1.2e-03  NA      NA    NA      NA
## staten island  2.7e-03  NA      NA    NA      NA
##
##
##
## [[3]]$gen
## [[3]]$gen$plot
```



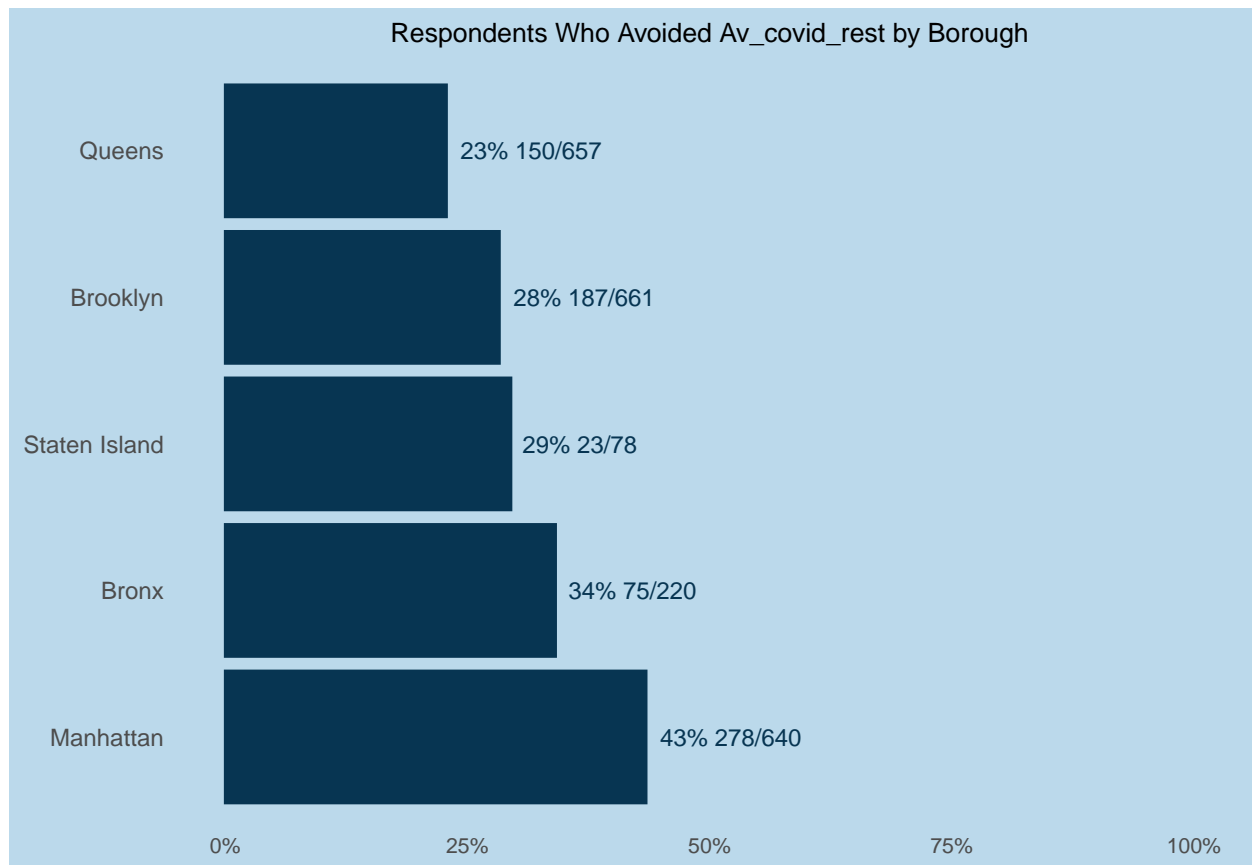
```
##
## [[3]]$gen$p.values
## [[3]]$gen$p.values$av_covid_trans
##           male prefer not to say transgender  female non-binary
## male           NA                NA          NA 3.8e-06        NA
## prefer not to say  NA                NA          NA      NA        NA
## transgender        NA                NA          NA      NA        NA
## female           3.8e-06              NA          NA      NA        NA
## non-binary         NA                NA          NA      NA        NA
##
##
##
## [[3]]$race_census
## NULL
##
## [[3]]$hh_ch_0_17_bi
## NULL
##
## [[3]]$hh_sn_65_bi
## NULL
##
## [[3]]$inc_dist
## NULL
##
##
## [[4]]
```

```
## [[4]]$borough
## [[4]]$borough$plot
```

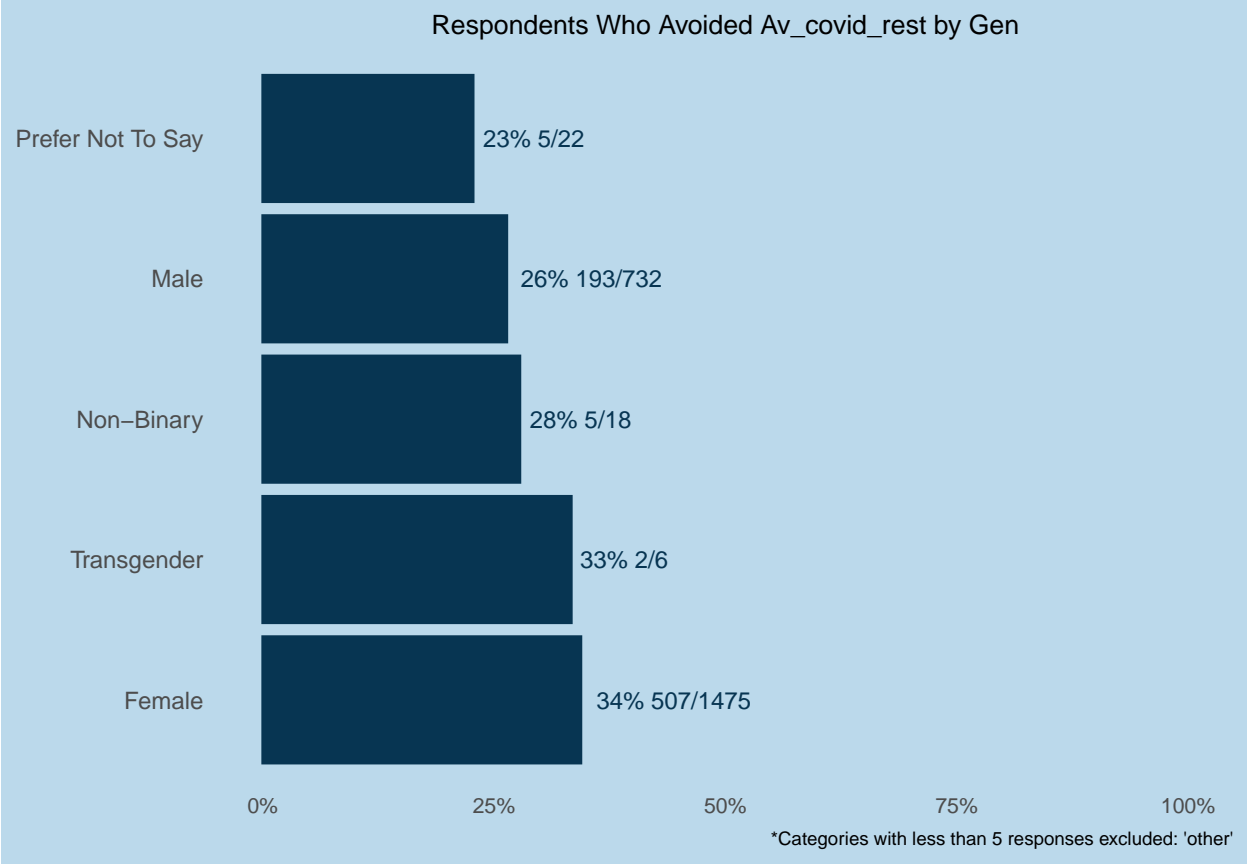


```
##
## [[4]]$borough$p.values
## [[4]]$borough$p.values$av_covid_off
##               queens staten island brooklyn bronx manhattan
## queens              NA              NA      NA      NA      0.0074
## staten island        NA              NA      NA      NA      NA
## brooklyn             NA              NA      NA      NA      NA
## bronx                NA              NA      NA      NA      NA
## manhattan            0.0074          NA      NA      NA      NA
##
##
##
## [[4]]$gen
## NULL
##
## [[4]]$race_census
## NULL
##
## [[4]]$hh_ch_0_17_bi
## NULL
##
## [[4]]$hh_sn_65_bi
```

```
## NULL
##
## [[4]]$inc_dist
## NULL
##
##
## [[5]]
## [[5]]$borough
## [[5]]$borough$plot
```

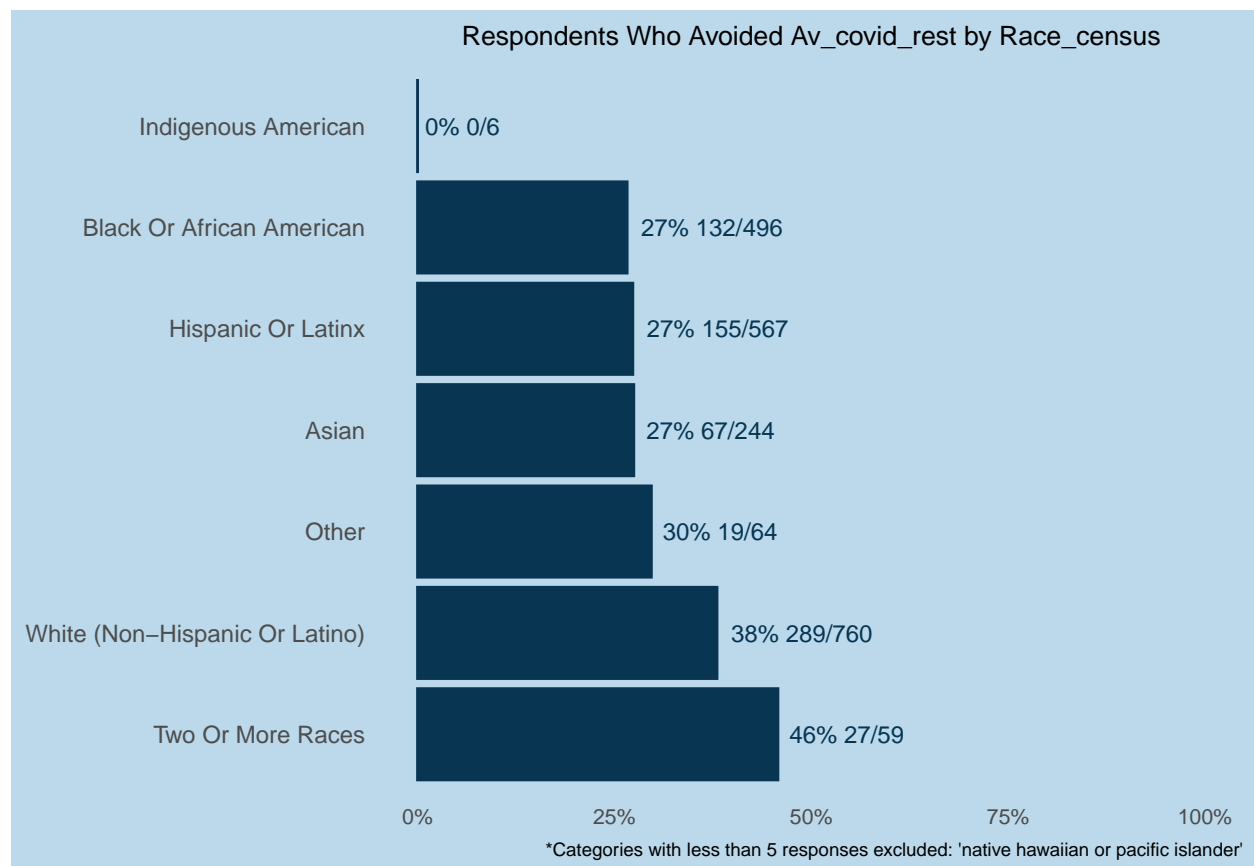


```
##
## [[5]]$borough$p.values
## [[5]]$borough$p.values$av_covid_rest
##
##      queens brooklyn staten island  bronx manhattan
## queens      NA      NA      NA 0.0013  4.8e-15
## brooklyn      NA      NA      NA      NA  1.7e-08
## staten island      NA      NA      NA      NA      NA
## bronx      1.3e-03      NA      NA      NA      NA
## manhattan  4.8e-15  1.7e-08      NA      NA      NA
##
##
##
## [[5]]$gen
## [[5]]$gen$plot
```



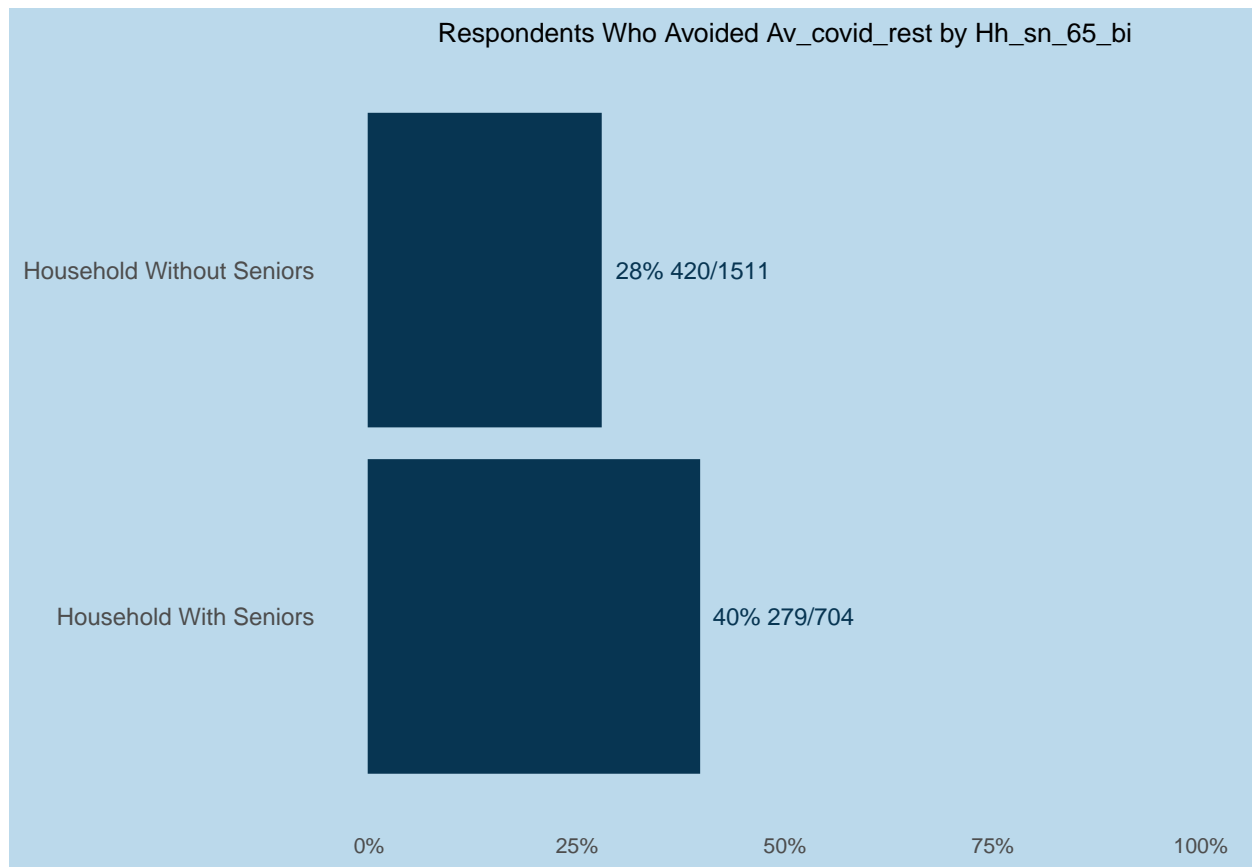
```
##
## [[5]]$gen$p.values
## [[5]]$gen$p.values$av_covid_rest
##               prefer not to say      male non-binary transgender  female
## prefer not to say              NA      NA      NA      NA      NA
## male                          NA      NA      NA      NA 0.00017
## non-binary                    NA      NA      NA      NA      NA
## transgender                    NA      NA      NA      NA      NA
## female                        NA 0.00017      NA      NA      NA
##
##
##
## [[5]]$race_census
## [[5]]$race_census$plot
```



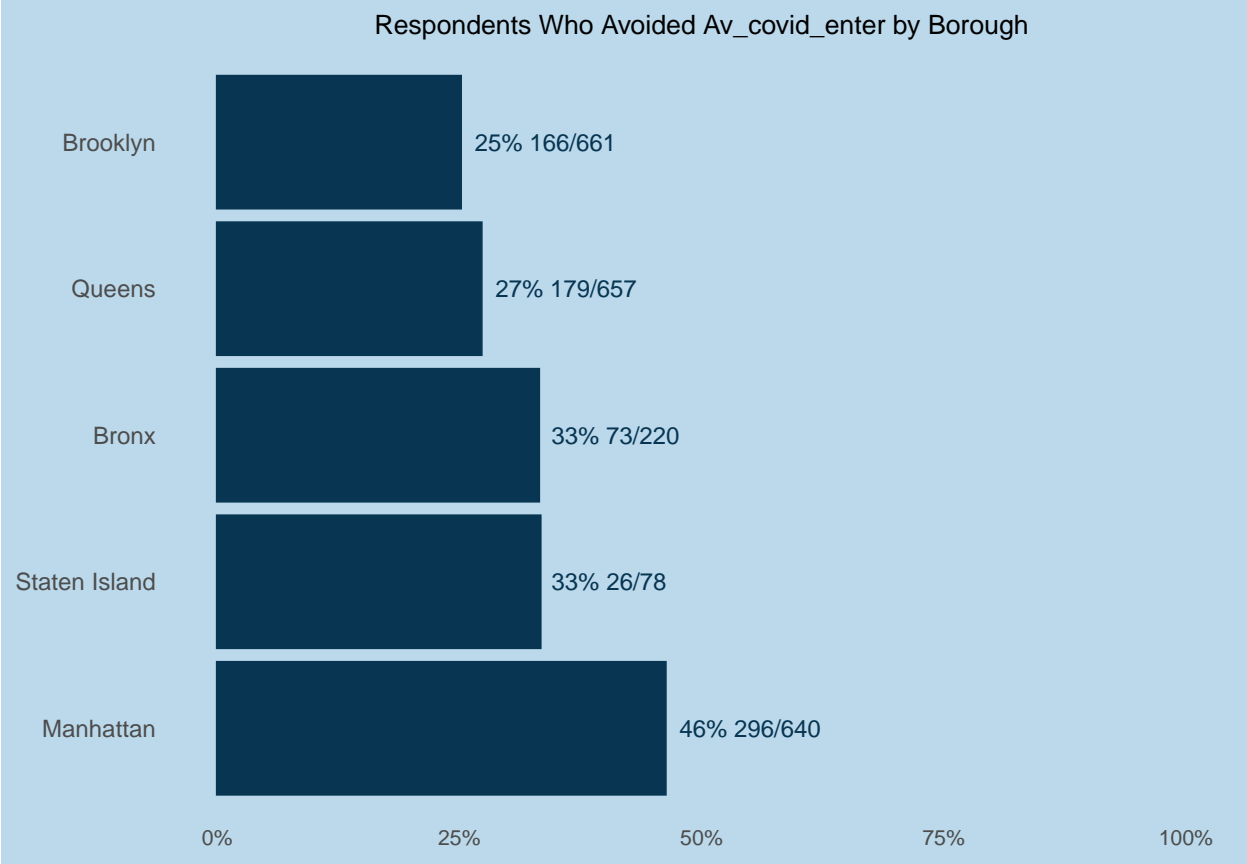


```
##
## [[5]]$race_census$p.values
## [[5]]$race_census$p.values$av_covid_rest
##           Indigenous American black or african american
## Indigenous American           NA                       NA
## black or african american       NA                       NA
## hispanic or latinx              NA                       NA
## asian                          NA                       NA
## other                          NA                       NA
## white (non-hispanic or latino)  NA                      3.7e-05
## two or more races              NA                      3.5e-03
##           hispanic or latinx asian other
## Indigenous American           NA      NA      NA
## black or african american       NA      NA      NA
## hispanic or latinx              NA      NA      NA
## asian                          NA      NA      NA
## other                          NA      NA      NA
## white (non-hispanic or latino)  5.7e-05 0.0034      NA
## two or more races              4.9e-03 0.0100      NA
##           white (non-hispanic or latino) two or more races
## Indigenous American           NA                       NA
## black or african american       3.7e-05                 0.0035
## hispanic or latinx              5.7e-05                 0.0049
## asian                          3.4e-03                 0.0100
## other                          NA                       NA
## white (non-hispanic or latino)  NA                       NA
```

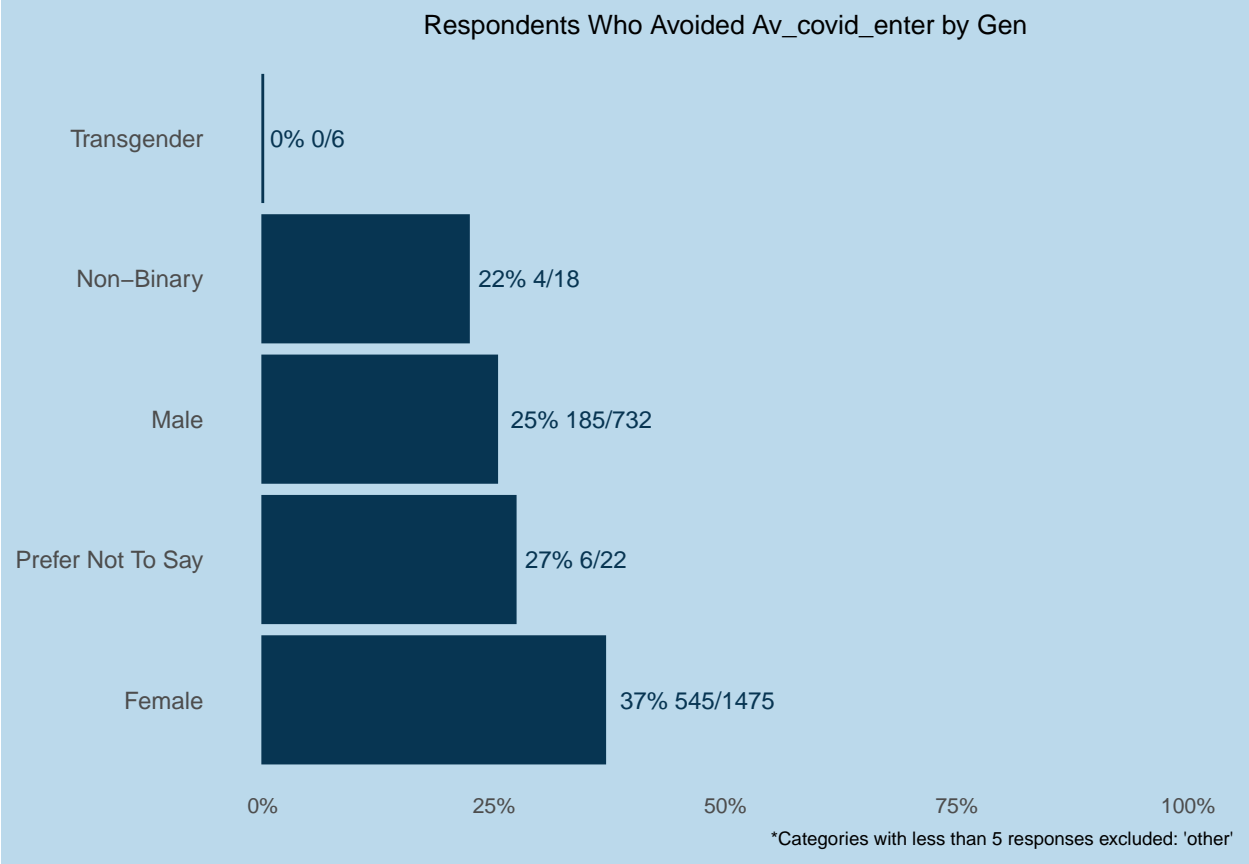
```
## two or more races
##
##
##
## [[5]]$hh_ch_0_17_bi
## NULL
##
## [[5]]$hh_sn_65_bi
## [[5]]$hh_sn_65_bi$plot
```



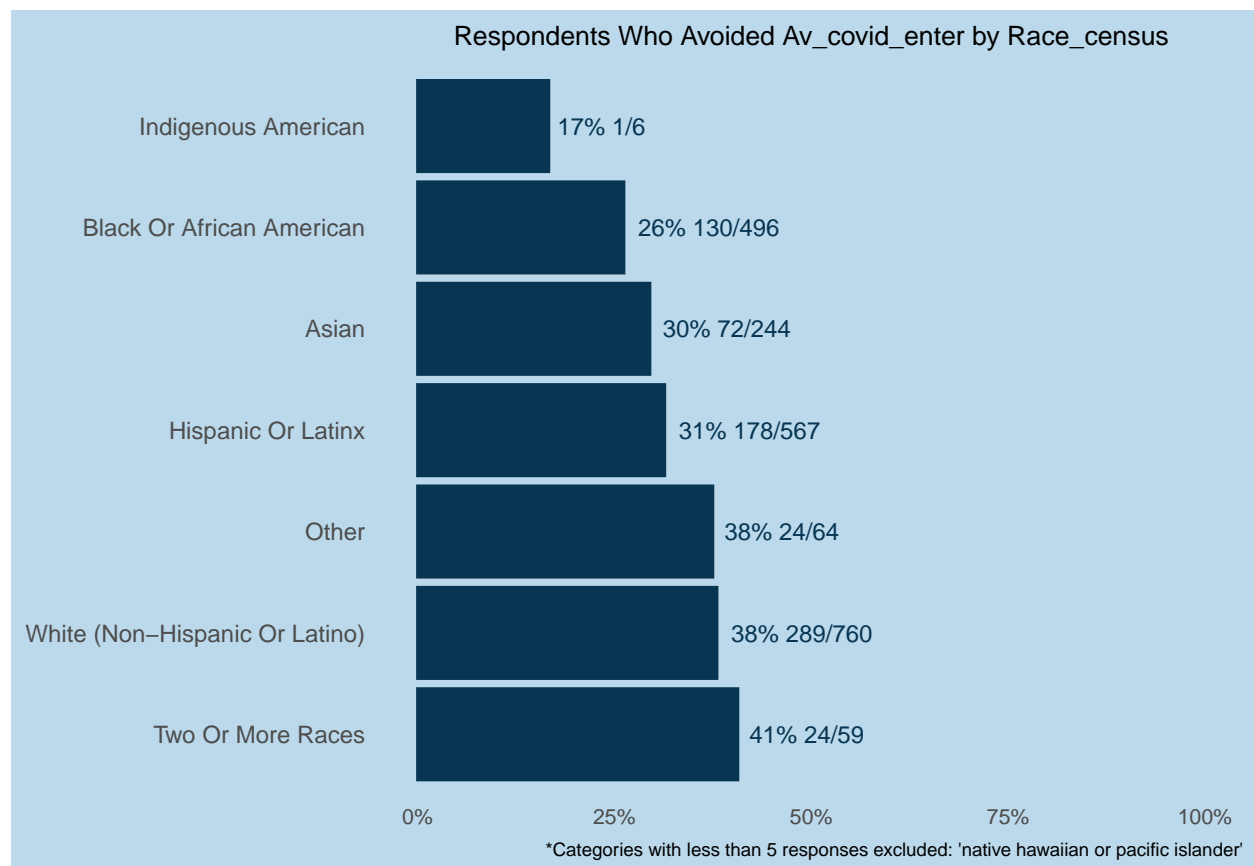
```
##
## [[5]]$hh_sn_65_bi$p.values
## [[5]]$hh_sn_65_bi$p.values$av_covid_rest
## household without seniors household with seniors
## household without seniors NA 3.2e-08
## household with seniors 3.2e-08 NA
##
##
##
## [[5]]$inc_dist
## NULL
##
##
## [[6]]
## [[6]]$borough
## [[6]]$borough$plot
```



```
##
## [[6]]$borough$p.values
## [[6]]$borough$p.values$av_covid_enter
##          brooklyn  queens  bronx  staten  island  manhattan
## brooklyn          NA     NA     NA          NA  2.6e-15
## queens            NA     NA     NA          NA  1.9e-12
## bronx             NA     NA     NA          NA  9.7e-04
## staten island      NA     NA     NA          NA     NA
## manhattan          2.6e-15 1.9e-12 0.00097          NA     NA
##
##
##
## [[6]]$gen
## [[6]]$gen$plot
```

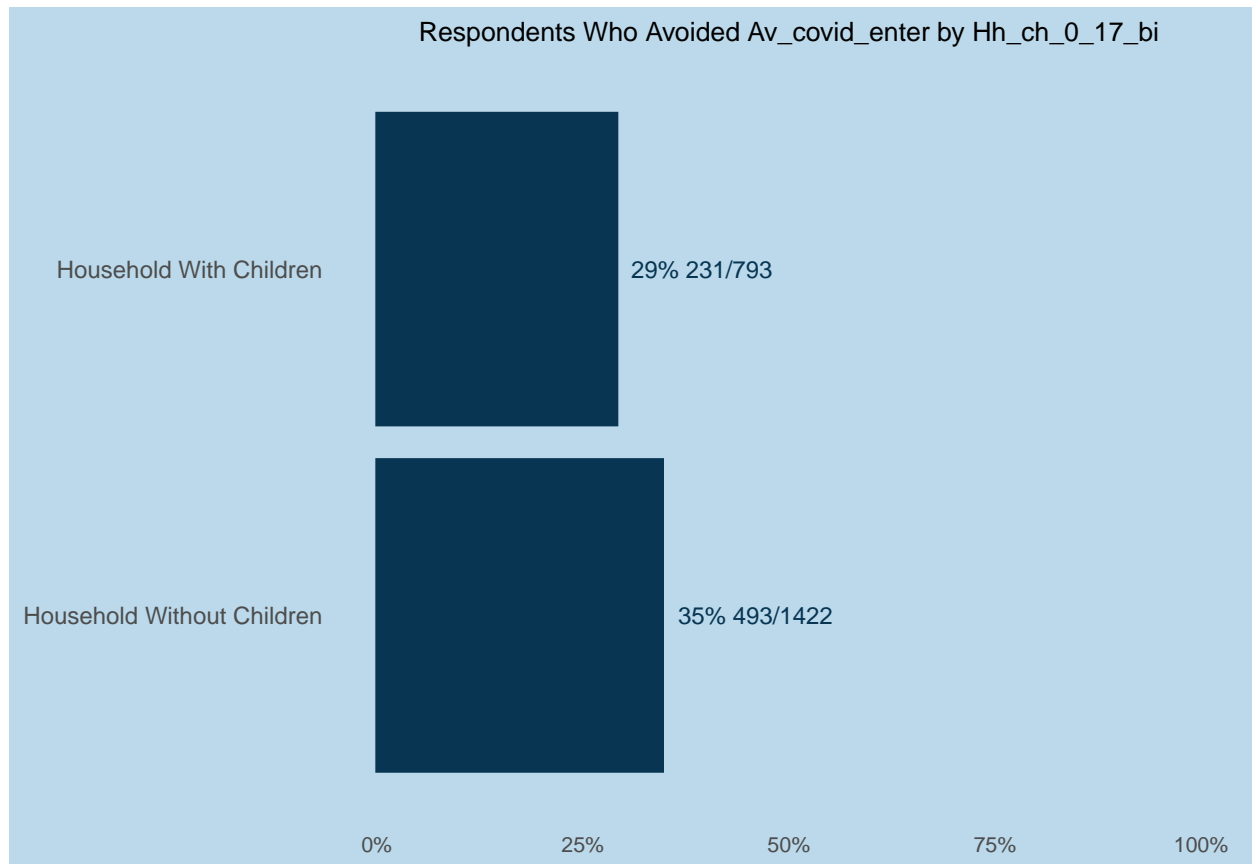


```
##
## [[6]]$gen$p.values
## [[6]]$gen$p.values$av_covid_enter
##               transgender non-binary      male prefer not to say  female
## transgender              NA         NA         NA              NA      NA
## non-binary                NA         NA         NA              NA      NA
## male                      NA         NA         NA              NA  5.3e-08
## prefer not to say         NA         NA         NA              NA      NA
## female                    NA         NA  5.3e-08              NA      NA
##
##
##
## [[6]]$race_census
## [[6]]$race_census$plot
```

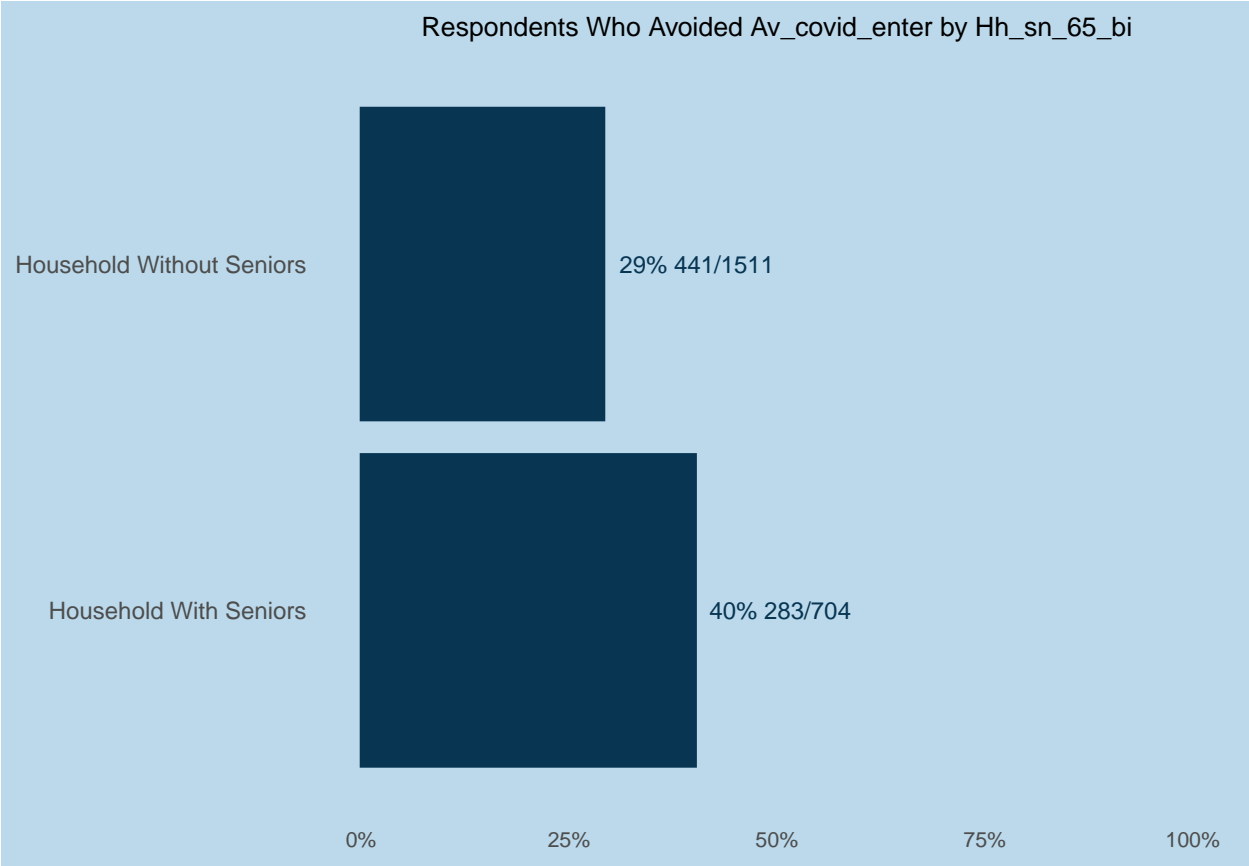


```
##
## [[6]]$race_census$p.values
## [[6]]$race_census$p.values$av_covid_enter
##           Indigenous American black or african american
## Indigenous American                NA                NA
## black or african american          NA                NA
## asian                             NA                NA
## hispanic or latinx                 NA                NA
## other                             NA                NA
## white (non-hispanic or latino)     NA                1.9e-05
## two or more races                  NA                NA
##           asian hispanic or latinx other
## Indigenous American          NA          NA          NA
## black or african american     NA          NA          NA
## asian                         NA          NA          NA
## hispanic or latinx            NA          NA          NA
## other                         NA          NA          NA
## white (non-hispanic or latino) NA          NA          NA
## two or more races             NA          NA          NA
##           white (non-hispanic or latino) two or more races
## Indigenous American                NA                NA
## black or african american          1.9e-05            NA
## asian                             NA                NA
## hispanic or latinx                 NA                NA
## other                             NA                NA
## white (non-hispanic or latino)     NA                NA
```

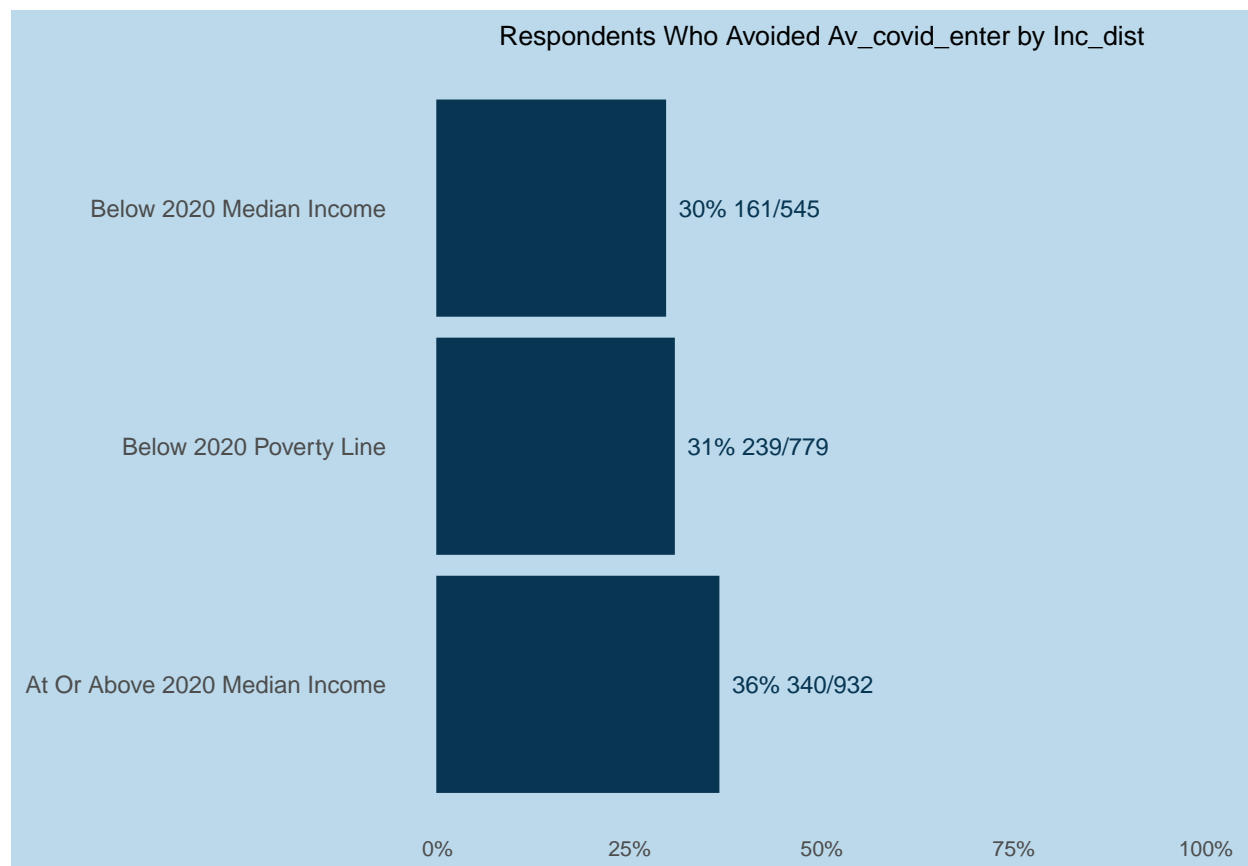
```
## two or more races
##
##
##
## [[6]]$hh_ch_0_17_bi
## [[6]]$hh_ch_0_17_bi$plot
```



```
##
## [[6]]$hh_ch_0_17_bi$p.values
## [[6]]$hh_ch_0_17_bi$p.values$av_covid_enter
##
## household with children household without children
## household with children NA 0.0089
## household without children 0.0089 NA
##
##
##
## [[6]]$hh_sn_65_bi
## [[6]]$hh_sn_65_bi$plot
```

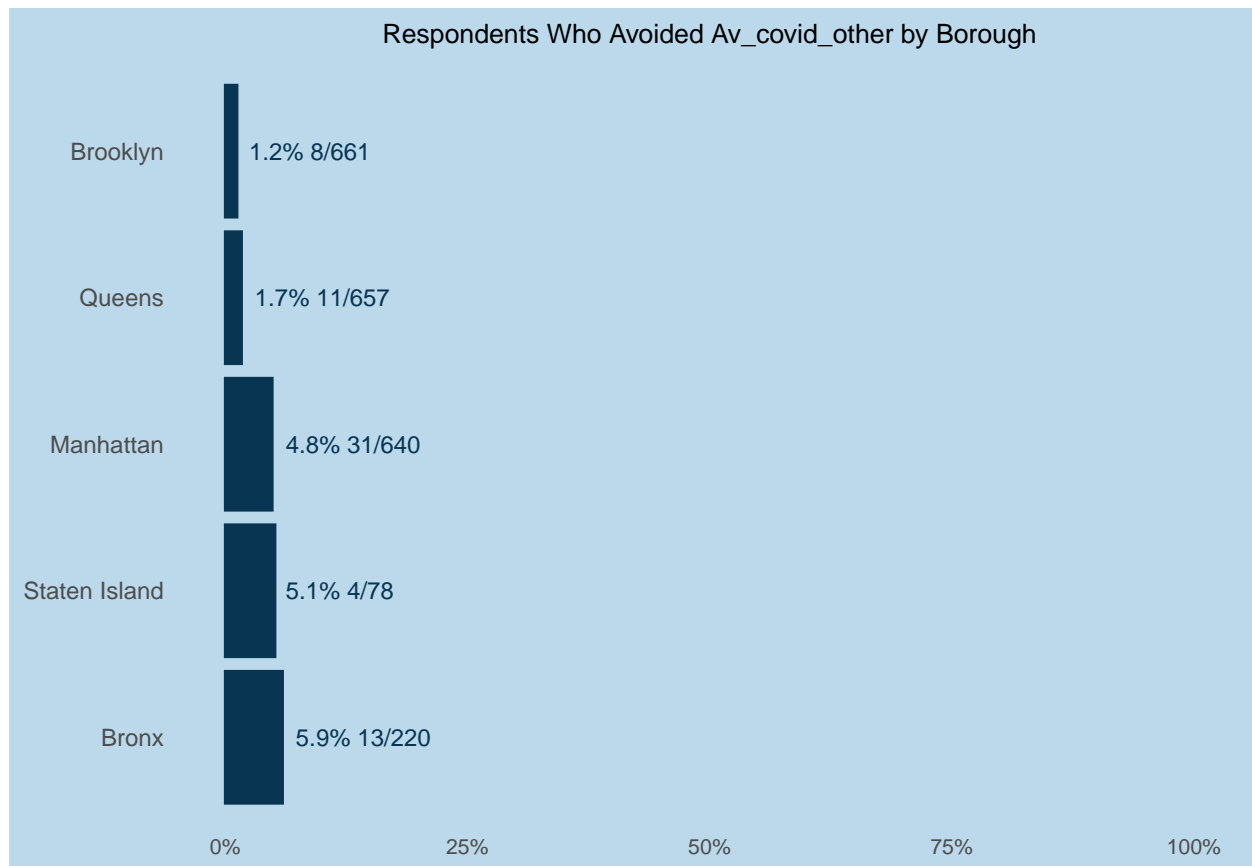


```
##
## [[6]]$hh_sn_65_bi$p.values
## [[6]]$hh_sn_65_bi$p.values$av_covid_enter
##           household without seniors household with seniors
## household without seniors           NA           3.5e-07
## household with seniors           3.5e-07           NA
##
##
##
## [[6]]$inc_dist
## [[6]]$inc_dist$plot
```

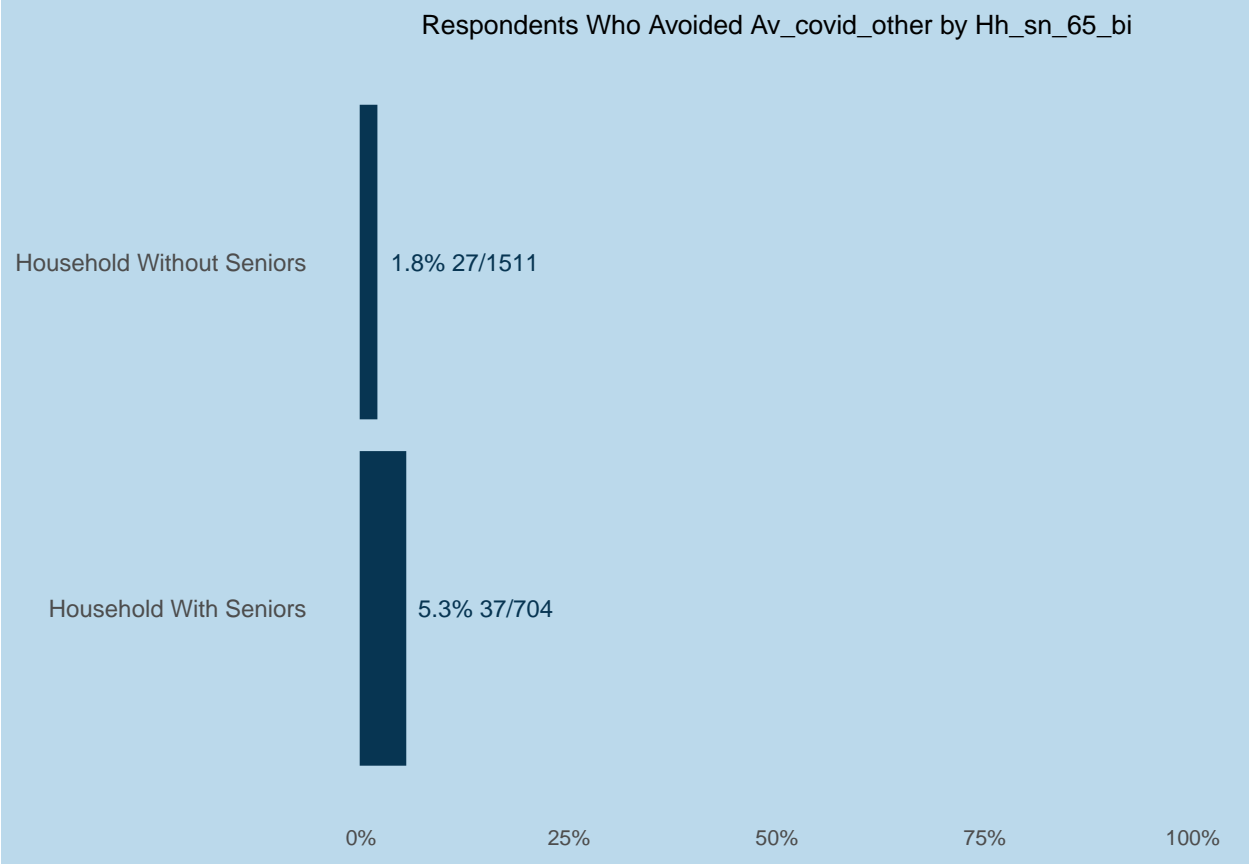


```
##
## [[6]]$inc_dist$p.values
## [[6]]$inc_dist$p.values$av_covid_enter
##          below 2020 median income below 2020 poverty line
## below 2020 median income          NA          NA
## below 2020 poverty line          NA          NA
## at or above 2020 median income    0.0078          NA
##          at or above 2020 median income
## below 2020 median income          0.0078
## below 2020 poverty line          NA
## at or above 2020 median income          NA
##
##
##
## [[7]]
## [[7]]$borough
## [[7]]$borough$plot
```

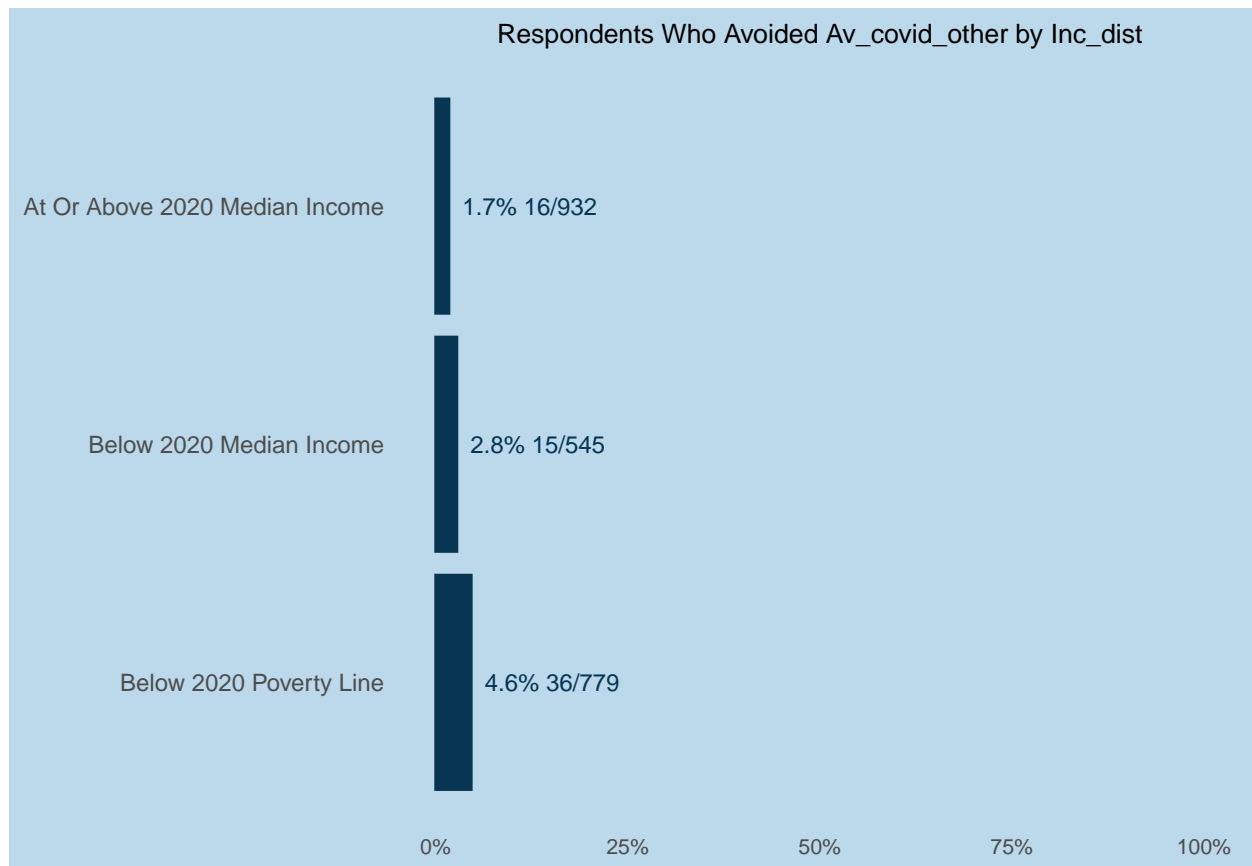




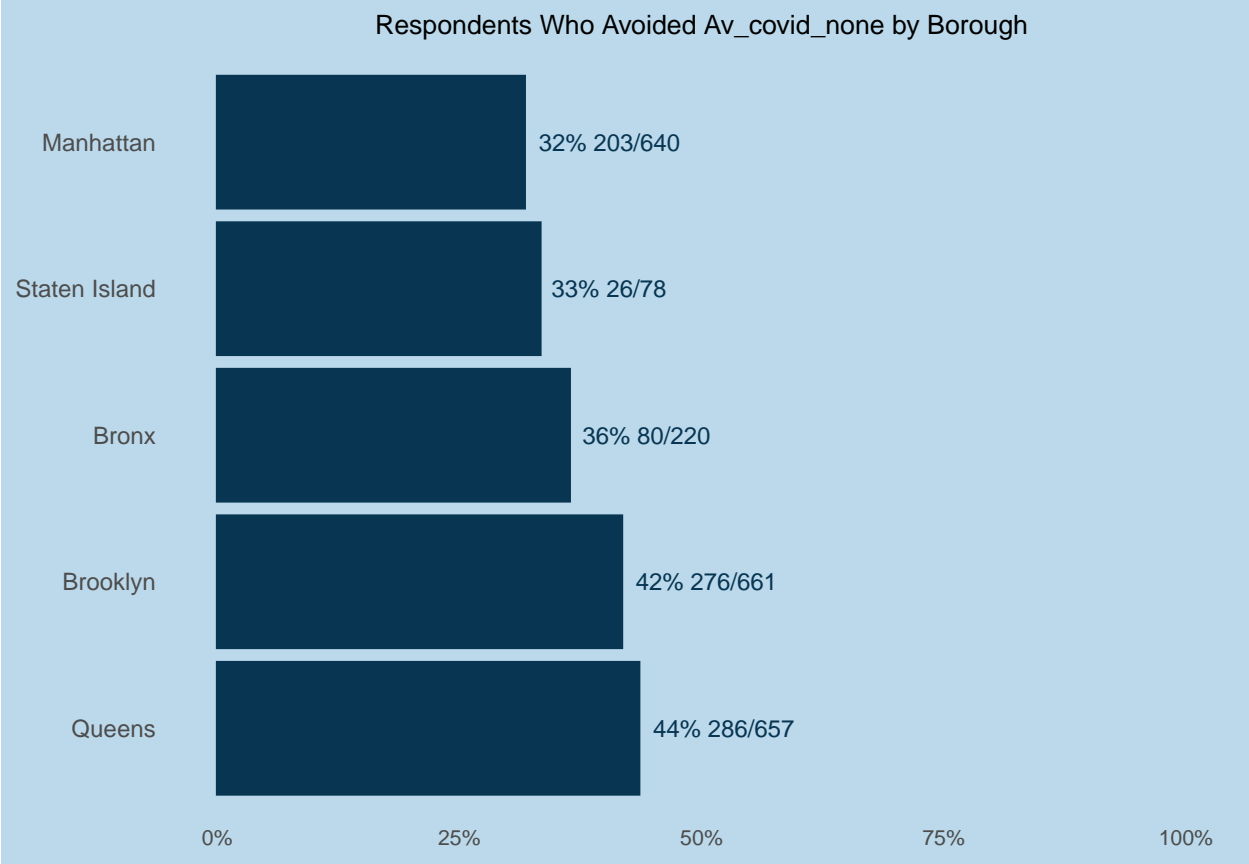
```
##
## [[7]]$borough$p.values
## [[7]]$borough$p.values$av_covid_other
##               brooklyn queens manhattan staten island  bronx
## brooklyn          NA      NA    0.00023              NA 0.00021
## queens            NA      NA    0.00220              NA 0.00200
## manhattan         0.00023 0.0022         NA              NA      NA
## staten island      NA      NA         NA              NA      NA
## bronx              0.00021 0.0020         NA              NA      NA
##
##
##
## [[7]]$gen
## NULL
##
## [[7]]$race_census
## NULL
##
## [[7]]$hh_ch_0_17_bi
## NULL
##
## [[7]]$hh_sn_65_bi
## [[7]]$hh_sn_65_bi$plot
```



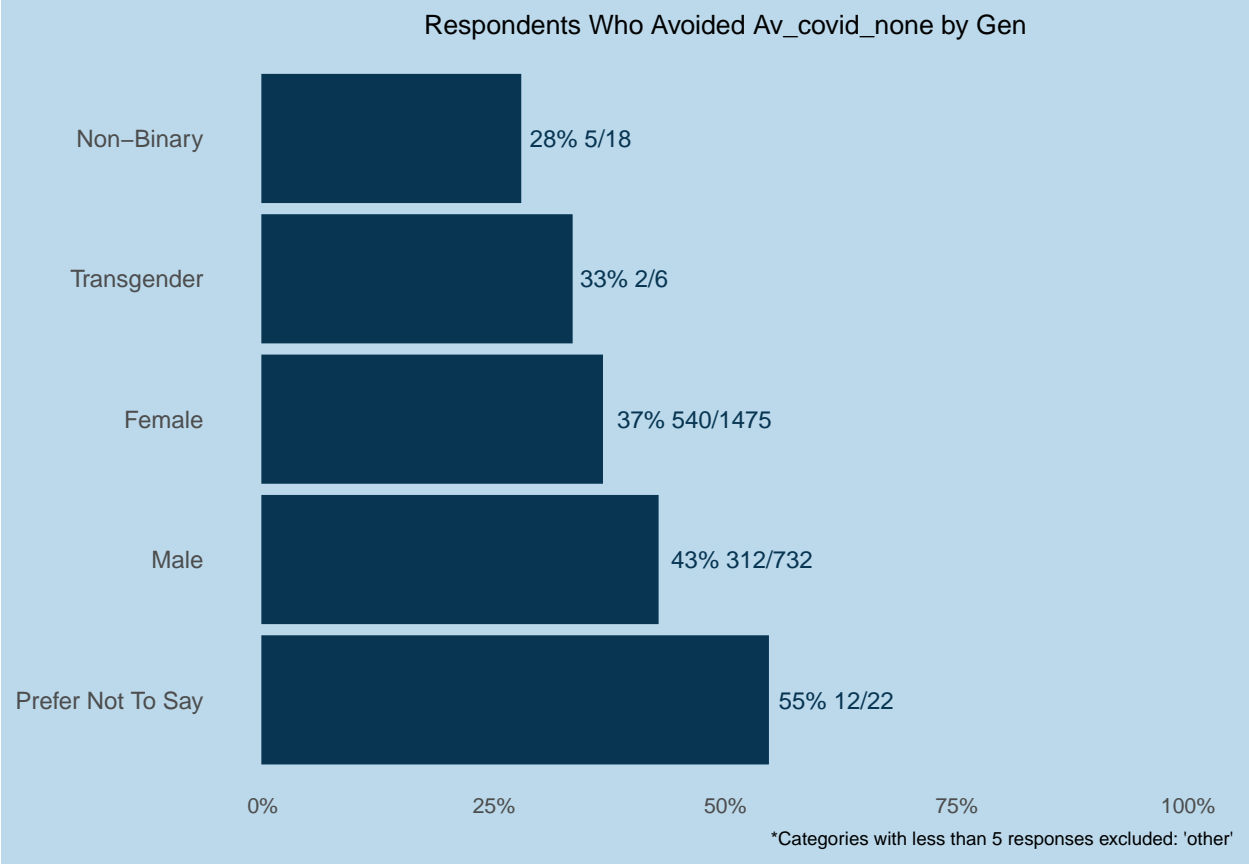
```
##
## [[7]]$hh_sn_65_bi$p.values
## [[7]]$hh_sn_65_bi$p.values$av_covid_other
##           household without seniors household with seniors
## household without seniors           NA           1.1e-05
## household with seniors           1.1e-05           NA
##
##
##
## [[7]]$inc_dist
## [[7]]$inc_dist$plot
```



```
##
## [[7]]$inc_dist$p.values
## [[7]]$inc_dist$p.values$av_covid_other
##               at or above 2020 median income
## at or above 2020 median income                NA
## below 2020 median income                      NA
## below 2020 poverty line                      0.00083
##               below 2020 median income below 2020 poverty line
## at or above 2020 median income                NA                0.00083
## below 2020 median income                      NA                NA
## below 2020 poverty line                      NA                NA
##
##
##
## [[8]]
## [[8]]$borough
## [[8]]$borough$plot
```



```
##
## [[8]]$borough$p.values
## [[8]]$borough$p.values$av_covid_none
##      manhattan staten island bronx brooklyn queens
## manhattan      NA           NA  NA  0.00022 1.5e-05
## staten island    NA           NA  NA      NA      NA
## bronx            NA           NA  NA      NA      NA
## brooklyn         2.2e-04        NA  NA      NA      NA
## queens           1.5e-05        NA  NA      NA      NA
##
##
##
## [[8]]$gen
## [[8]]$gen$plot
```



```
##
## [[8]]$gen$p.values
## [[8]]$gen$p.values$av_covid_none
##               non-binary transgender female   male prefer not to say
## non-binary           NA           NA    NA    NA           NA
## transgender          NA           NA    NA    NA           NA
## female               NA           NA    NA 0.0072          NA
## male                 NA           NA 0.0072    NA           NA
## prefer not to say    NA           NA    NA    NA           NA
##
##
##
## [[8]]$race_census
## NULL
##
## [[8]]$hh_ch_0_17_bi
## NULL
##
## [[8]]$hh_sn_65_bi
## NULL
##
## [[8]]$inc_dist
## NULL
```

```
cat("Plots with no statistically significant result are omitted from printing. See those marked with 'NUL
```

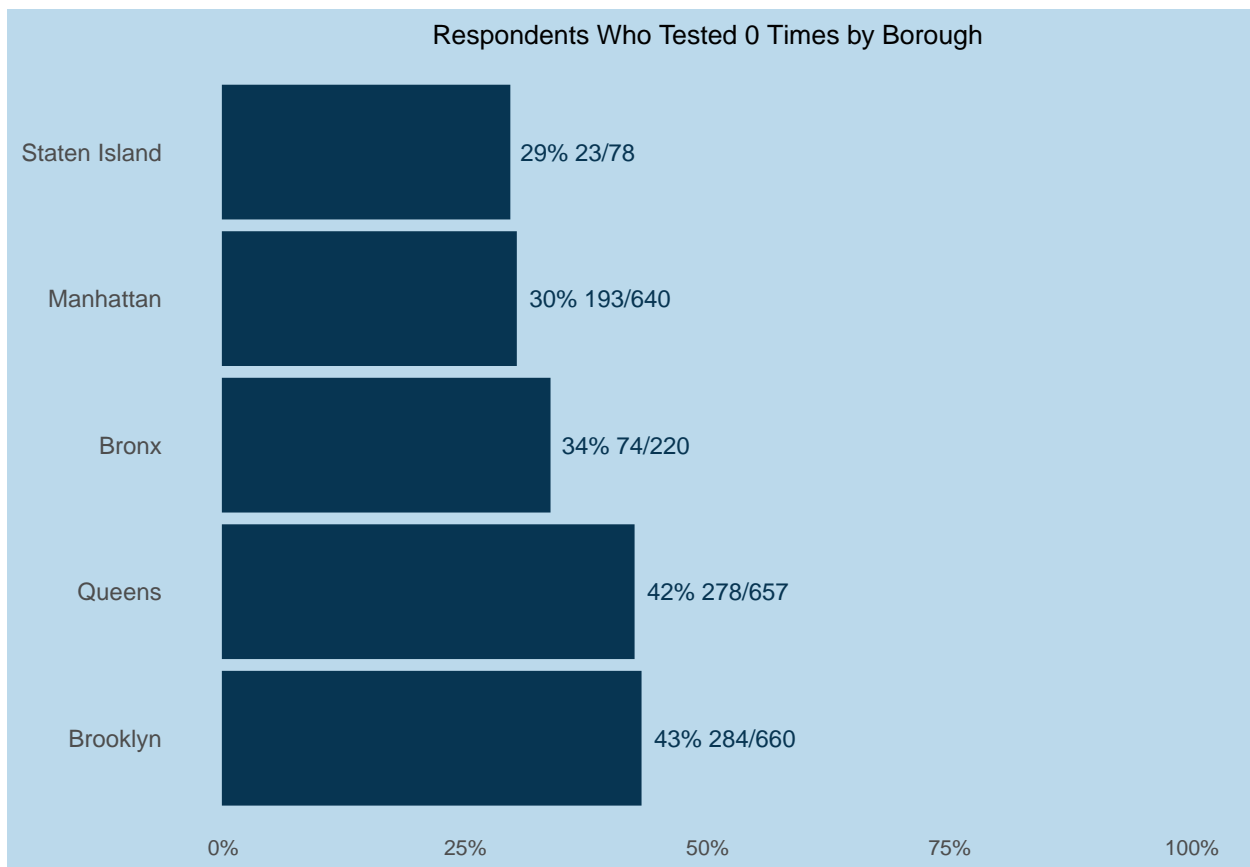
```
## Plots with no statistically significant result are omitted from printing. See those marked with 'NUL
```

## 4.15) Distribution over frequency of testing

1. Run each type of frequency over the sub demographic groups

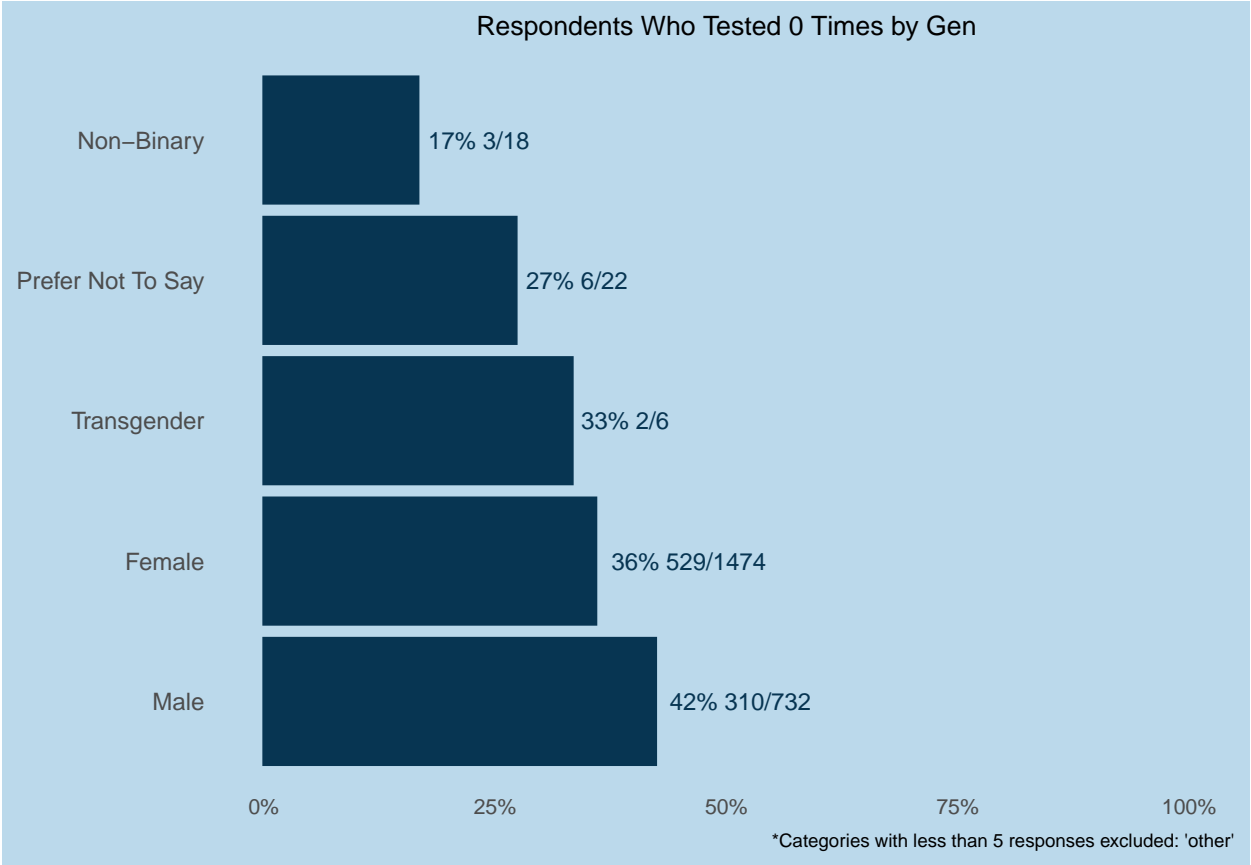
```
lapply(1:4, function(var) {
  freq <- names(attributes(wrangled$test)$labels[var])
  make_plots(wrangled %>% mutate(test_freq = test == var), demographics, "test_freq",
    title = glue::glue("Respondents who tested {freq}"))
})
```

```
## [[1]]
## [[1]]$borough
## [[1]]$borough$plot
```

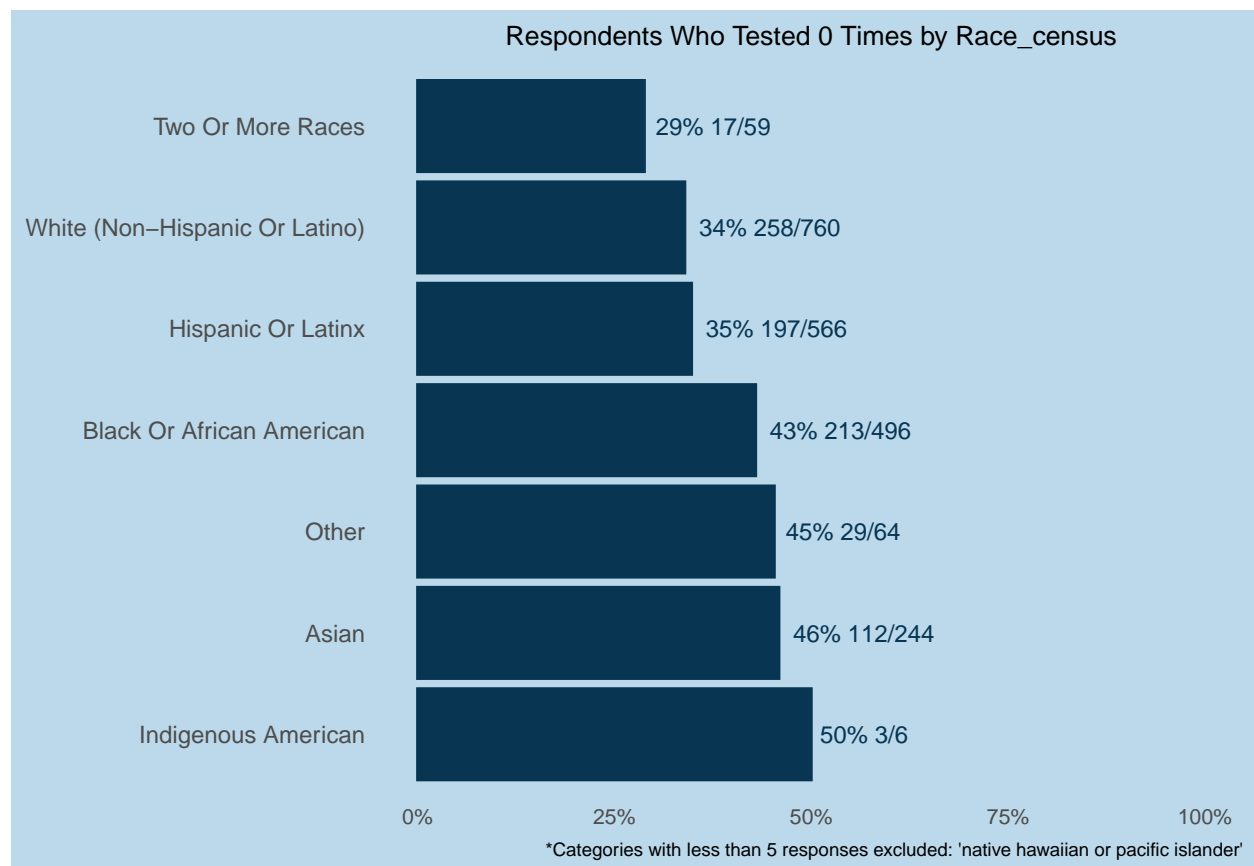


```
##
## [[1]]$borough$p.values
## [[1]]$borough$p.values$test_freq
##          staten island manhattan bronx queens brooklyn
```

```
## staten island      NA      NA      NA      NA      NA
## manhattan         NA      NA      NA 7e-06  2e-06
## bronx             NA      NA      NA      NA      NA
## queens            NA  7e-06      NA      NA      NA
## brooklyn          NA  2e-06      NA      NA      NA
##
##
##
## [[1]]$gen
## [[1]]$gen$plot
```



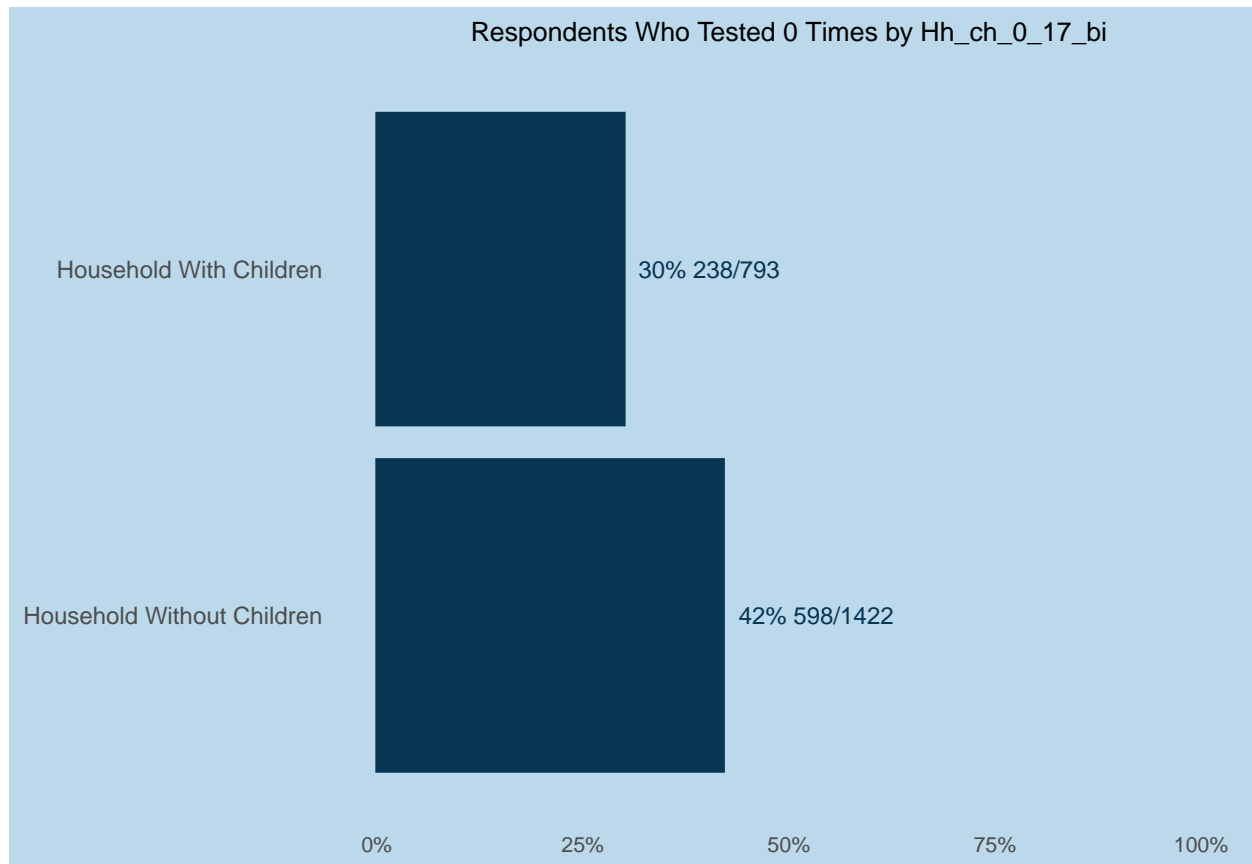
```
##
## [[1]]$gen$p.values
## [[1]]$gen$p.values$test_freq
## non-binary prefer not to say transgender female male
## non-binary      NA      NA      NA      NA      NA
## prefer not to say      NA      NA      NA      NA      NA
## transgender      NA      NA      NA      NA      NA
## female          NA      NA      NA      NA 0.0038
## male           NA      NA      NA 0.0038      NA
##
##
##
## [[1]]$race_census
## [[1]]$race_census$plot
```



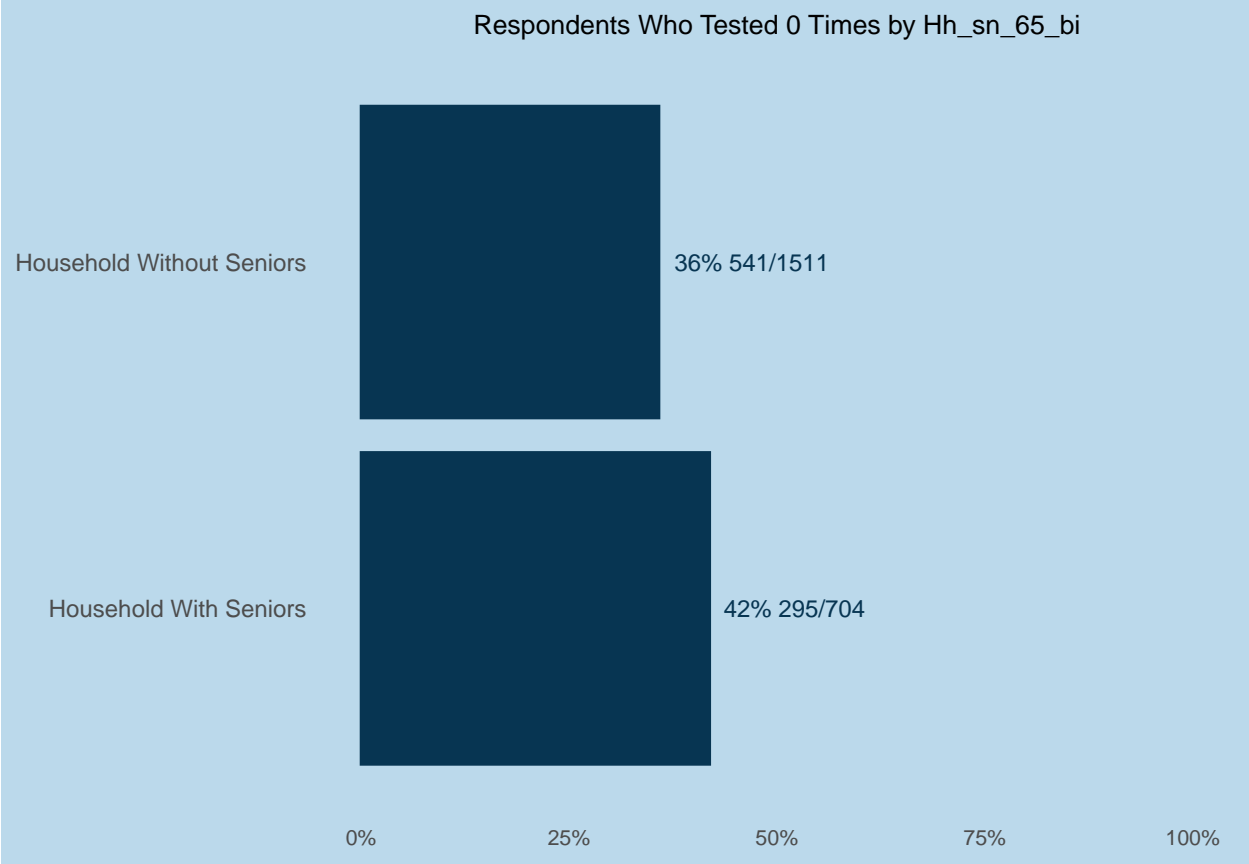
```
##
## [[1]]$race_census$p.values
## [[1]]$race_census$p.values$test_freq
##               two or more races white (non-hispanic or latino)
## two or more races                NA                          NA
## white (non-hispanic or latino)    NA                          NA
## hispanic or latinx                 NA                          NA
## black or african american          NA                        0.0016
## other                             NA                          NA
## asian                             NA                        0.0010
## Indigenous American                NA                          NA
##               hispanic or latinx black or african american
## two or more races                NA                          NA
## white (non-hispanic or latino)    NA                        0.0016
## hispanic or latinx                 NA                        0.0079
## black or african american          NA                        0.0079
## other                             NA                          NA
## asian                             NA                        0.0037
## Indigenous American                NA                          NA
##               other  asian Indigenous American
## two or more races    NA    NA                NA
## white (non-hispanic or latino) NA 0.0010        NA
## hispanic or latinx    NA 0.0037        NA
## black or african american NA    NA        NA
## other                  NA    NA        NA
## asian                  NA    NA        NA
```



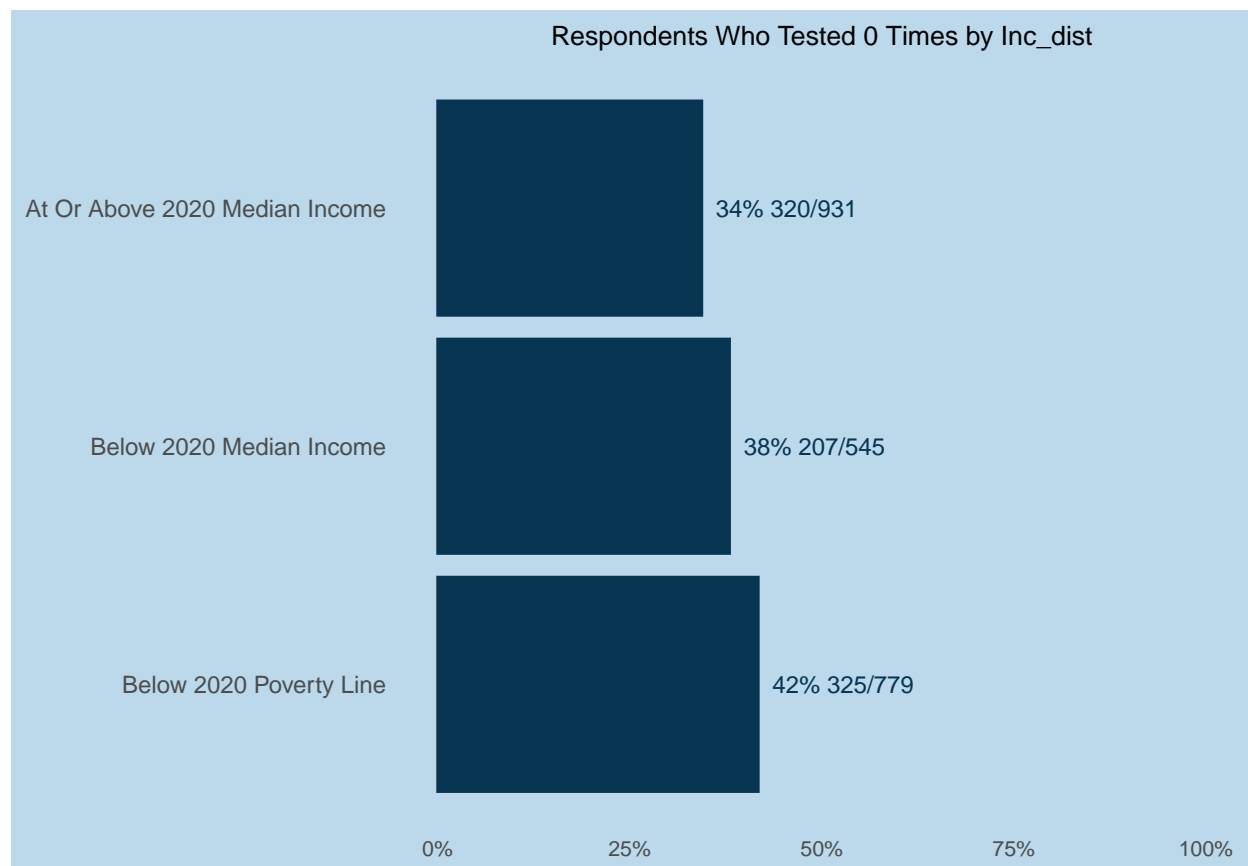
```
## Indigenous American      NA      NA      NA
##
##
##
## [[1]]$hh_ch_0_17_bi
## [[1]]$hh_ch_0_17_bi$plot
```



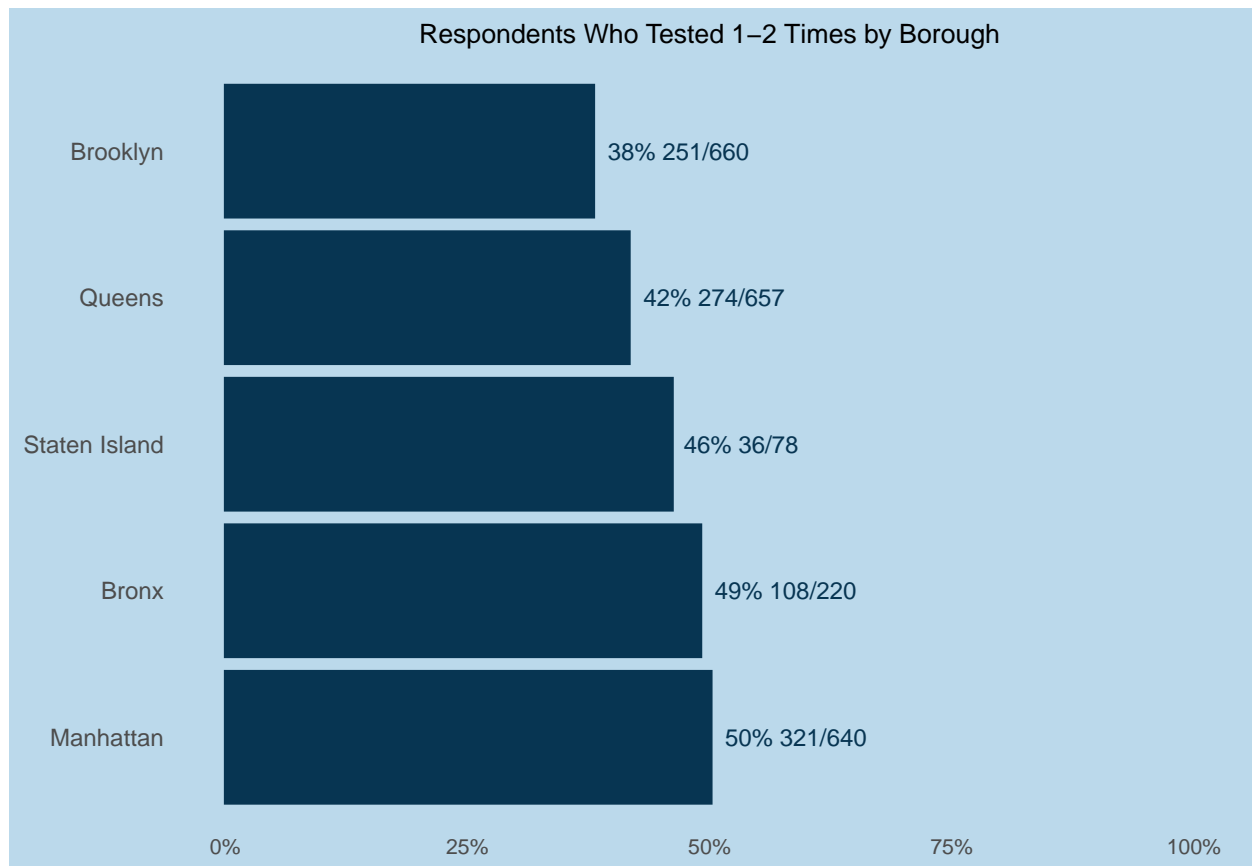
```
##
## [[1]]$hh_ch_0_17_bi$p.values
## [[1]]$hh_ch_0_17_bi$p.values$test_freq
##           household with children household without children
## household with children      NA      2.7e-08
## household without children  2.7e-08      NA
##
##
##
## [[1]]$hh_sn_65_bi
## [[1]]$hh_sn_65_bi$plot
```



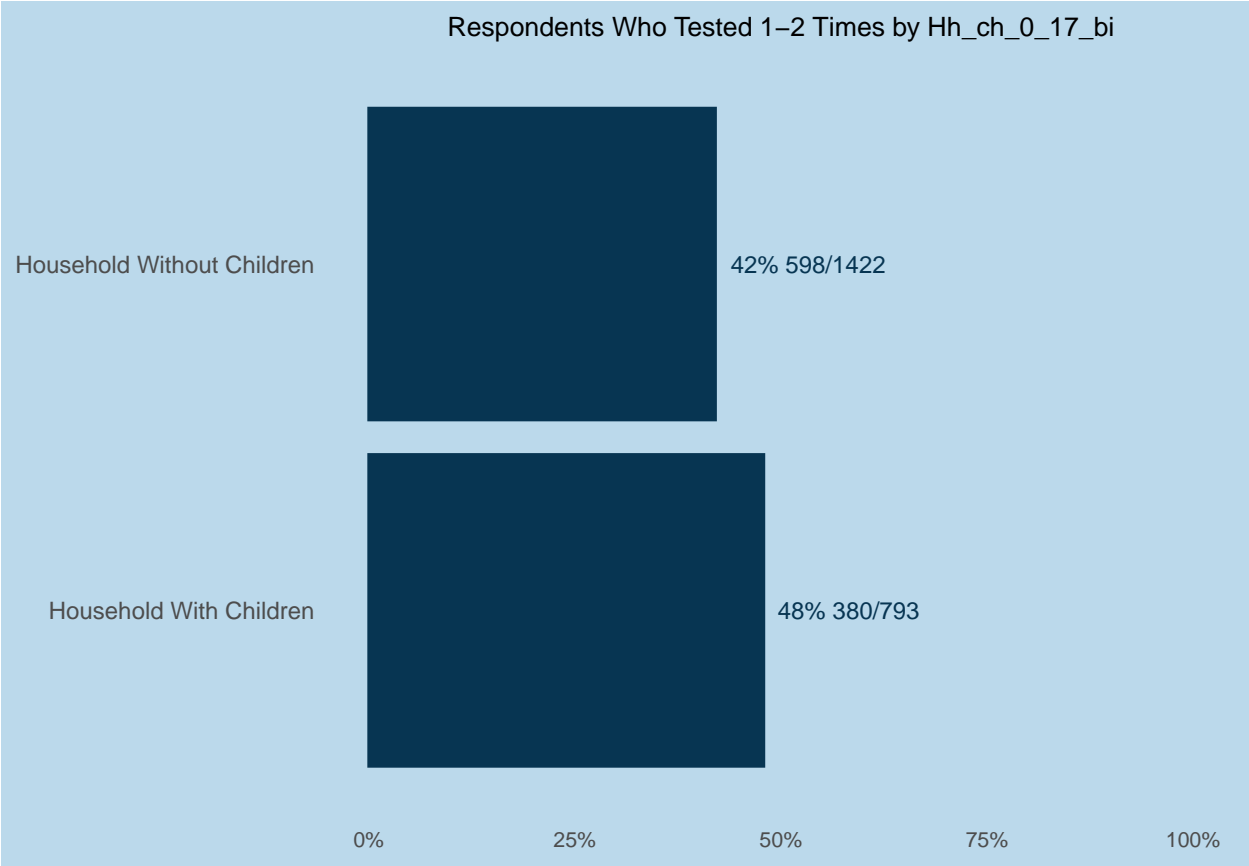
```
##  
## [[1]]$hh_sn_65_bi$p.values  
## [[1]]$hh_sn_65_bi$p.values$test_freq  
##  
## household without seniors household with seniors  
## household without seniors NA 0.0067  
## household with seniors 0.0067 NA  
##  
##  
##  
## [[1]]$inc_dist  
## [[1]]$inc_dist$plot
```



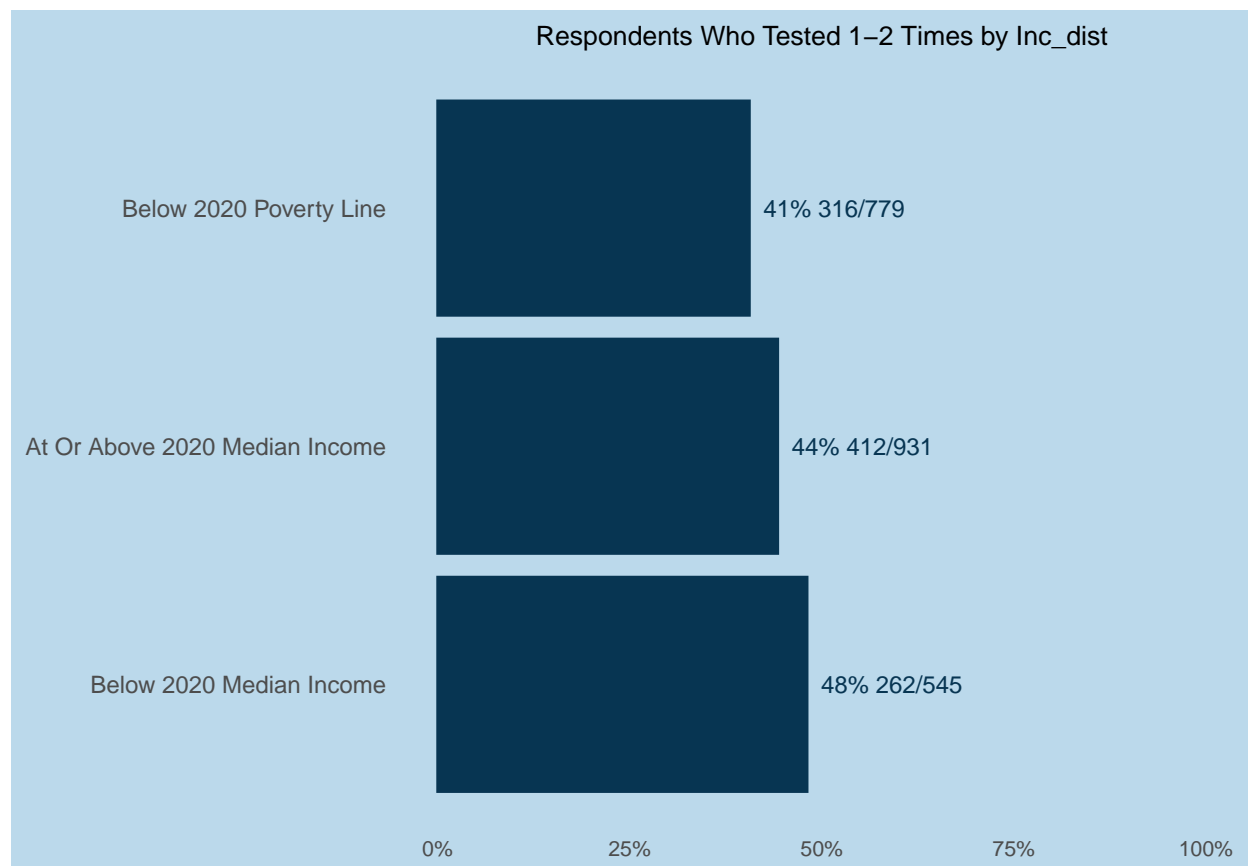
```
##
## [[1]]$inc_dist$p.values
## [[1]]$inc_dist$p.values$test_freq
##                               at or above 2020 median income
## at or above 2020 median income                               NA
## below 2020 median income                                     NA
## below 2020 poverty line                                     0.0021
##                               below 2020 median income below 2020 poverty line
## at or above 2020 median income                               NA                               0.0021
## below 2020 median income                                     NA                               NA
## below 2020 poverty line                                     NA                               NA
##
##
##
## [[2]]
## [[2]]$borough
## [[2]]$borough$plot
```



```
##
## [[2]]$borough$p.values
## [[2]]$borough$p.values$test_freq
##          brooklyn queens staten island  bronx manhattan
## brooklyn          NA      NA          NA 0.0049  1.4e-05
## queens            NA      NA          NA  NA    2.7e-03
## staten island      NA      NA          NA  NA      NA
## bronx              4.9e-03  NA          NA  NA      NA
## manhattan          1.4e-05 0.0027      NA  NA      NA
##
##
##
## [[2]]$gen
## NULL
##
## [[2]]$race_census
## NULL
##
## [[2]]$hh_ch_0_17_bi
## [[2]]$hh_ch_0_17_bi$plot
```

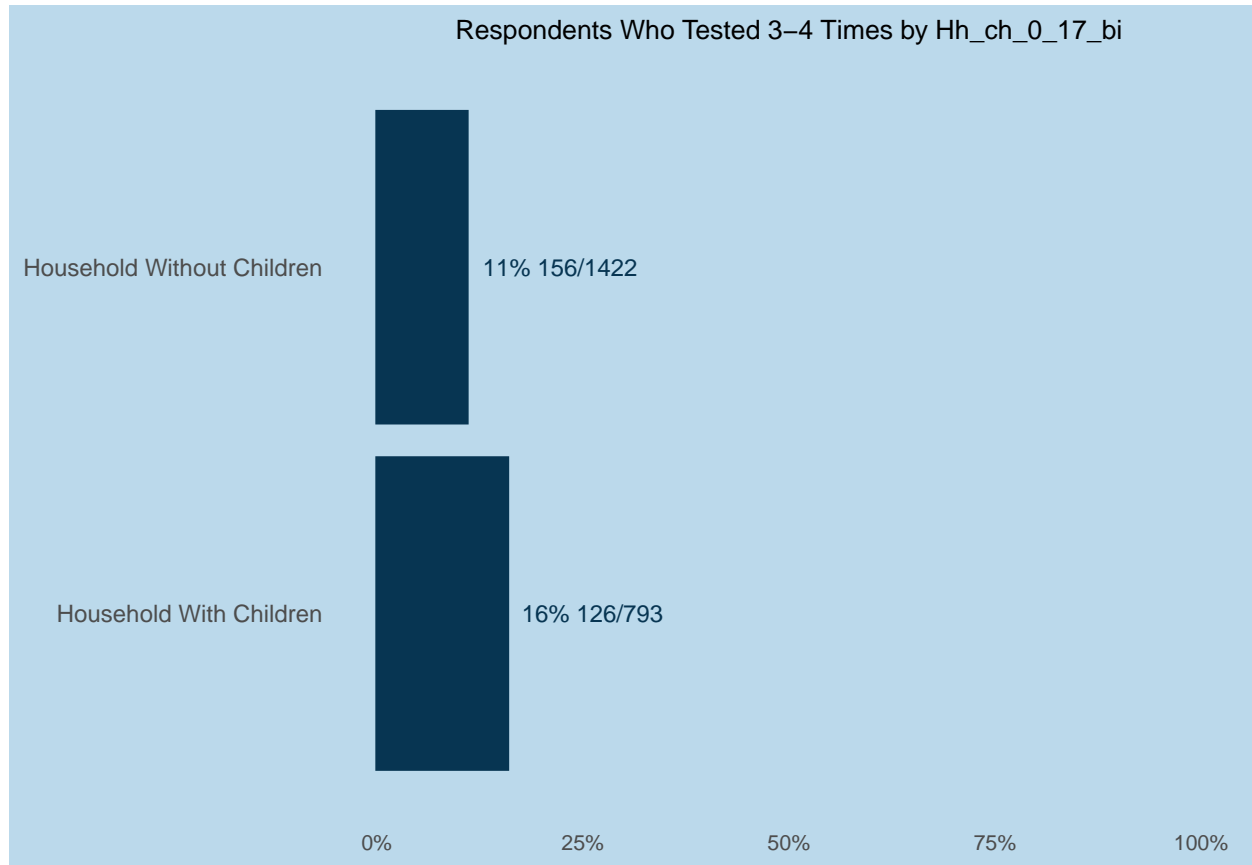


```
##  
## [[2]]$hh_ch_0_17_bi$p.values  
## [[2]]$hh_ch_0_17_bi$p.values$test_freq  
## household without children household with children  
## household without children NA 0.0088  
## household with children 0.0088 NA  
##  
##  
##  
## [[2]]$hh_sn_65_bi  
## NULL  
##  
## [[2]]$inc_dist  
## [[2]]$inc_dist$plot
```



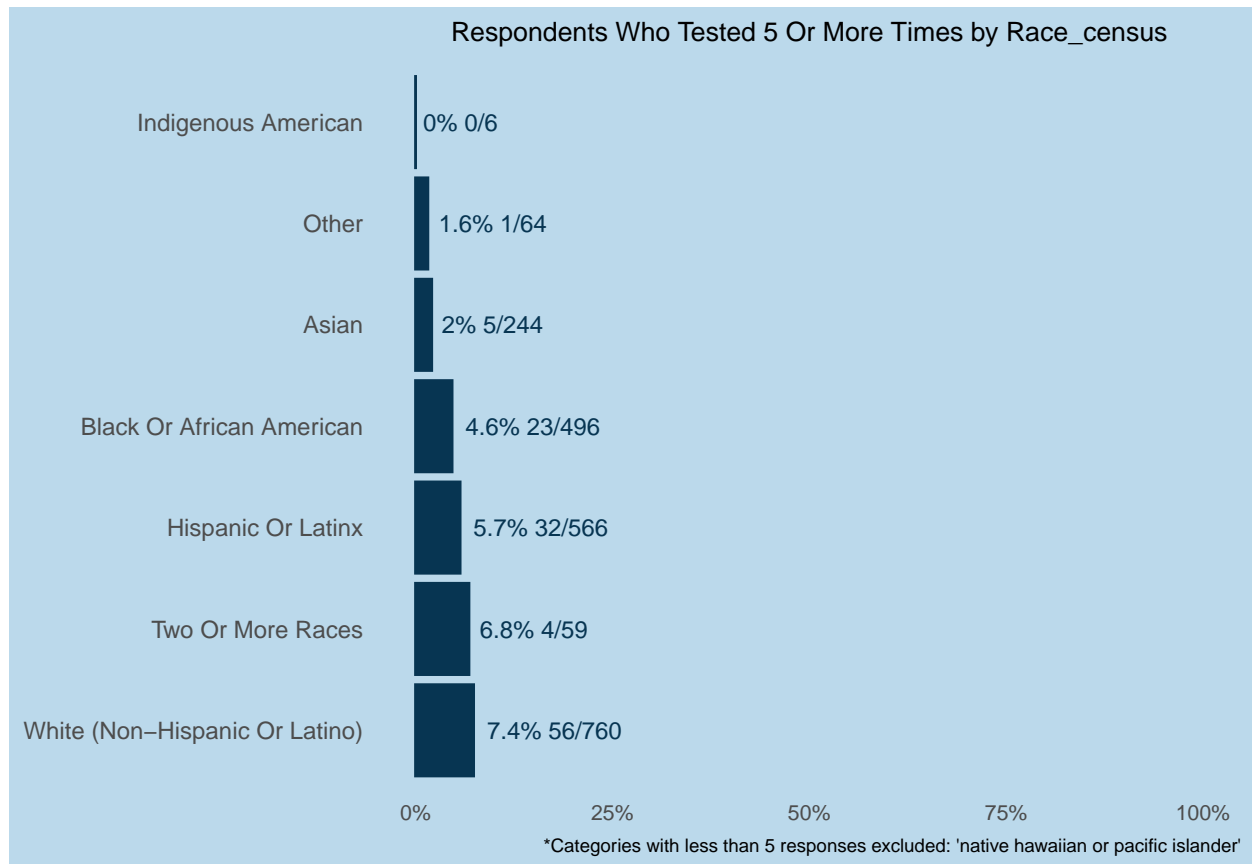
```
##
## [[2]]$inc_dist$p.values
## [[2]]$inc_dist$p.values$test_freq
##          below 2020 poverty line
## below 2020 poverty line          NA
## at or above 2020 median income    NA
## below 2020 median income          0.0079
##          at or above 2020 median income
## below 2020 poverty line          NA
## at or above 2020 median income    NA
## below 2020 median income          NA
##          below 2020 median income
## below 2020 poverty line          0.0079
## at or above 2020 median income    NA
## below 2020 median income          NA
##
##
##
##
## [[3]]
## [[3]]$borough
## NULL
##
## [[3]]$gen
## NULL
##
```

```
## [[3]]$race_census
## NULL
##
## [[3]]$hh_ch_0_17_bi
## [[3]]$hh_ch_0_17_bi$plot
```



```
##
## [[3]]$hh_ch_0_17_bi$p.values
## [[3]]$hh_ch_0_17_bi$p.values$test_freq
## household without children household with children
## household without children NA 0.0011
## household with children 0.0011 NA
##
##
##
## [[3]]$hh_sn_65_bi
## NULL
##
## [[3]]$inc_dist
## NULL
##
##
## [[4]]
## [[4]]$borough
## NULL
```

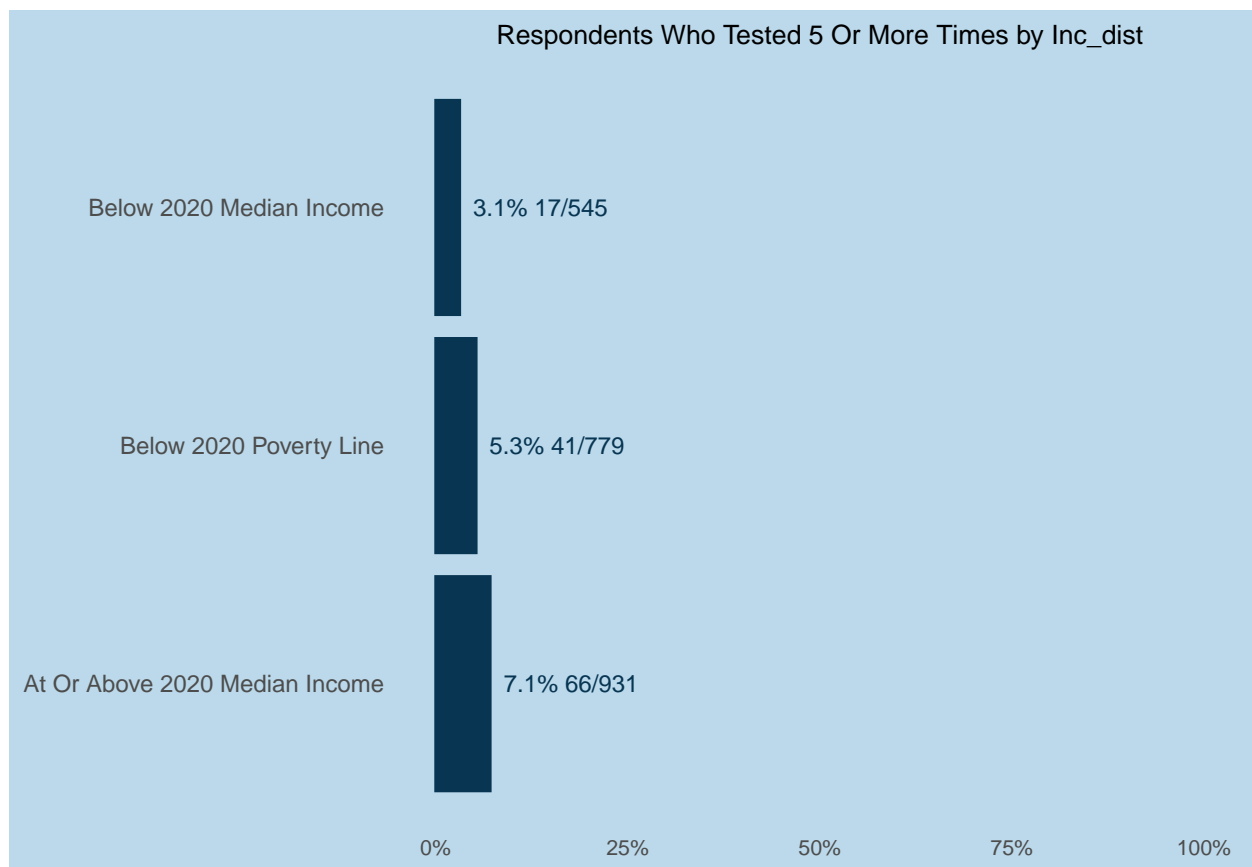
```
##
## [[4]]$gen
## NULL
##
## [[4]]$race_census
## [[4]]$race_census$plot
```



```
##
## [[4]]$race_census$p.values
## [[4]]$race_census$p.values$test_freq
##
## Indigenous American other asian
## Indigenous American NA NA NA
## other NA NA NA
## asian NA NA NA
## black or african american NA NA NA
## hispanic or latinx NA NA NA
## two or more races NA NA NA
## white (non-hispanic or latino) NA NA 0.0041
##
## black or african american hispanic or latinx
## Indigenous American NA NA
## other NA NA
## asian NA NA
## black or african american NA NA
## hispanic or latinx NA NA
## two or more races NA NA
```



```
## white (non-hispanic or latino) NA NA
## two or more races white (non-hispanic or latino)
## Indigenous American NA NA
## other NA NA
## asian NA 0.0041
## black or african american NA NA
## hispanic or latinx NA NA
## two or more races NA NA
## white (non-hispanic or latino) NA NA
##
##
##
## [[4]]$hh_ch_0_17_bi
## NULL
##
## [[4]]$hh_sn_65_bi
## NULL
##
## [[4]]$inc_dist
## [[4]]$inc_dist$plot
```



```
##
## [[4]]$inc_dist$p.values
## [[4]]$inc_dist$p.values$test_freq
## below 2020 median income below 2020 poverty line
```

```
## below 2020 median income          NA          NA
## below 2020 poverty line           NA          NA
## at or above 2020 median income     0.0021      NA
##                                at or above 2020 median income
## below 2020 median income          0.0021
## below 2020 poverty line           NA
## at or above 2020 median income     NA
```

```
cat("Plots with no statistically significant result are omitted from printing. See those marked with 'N")
```

```
## Plots with no statistically significant result are omitted from printing. See those marked with 'NUL
```

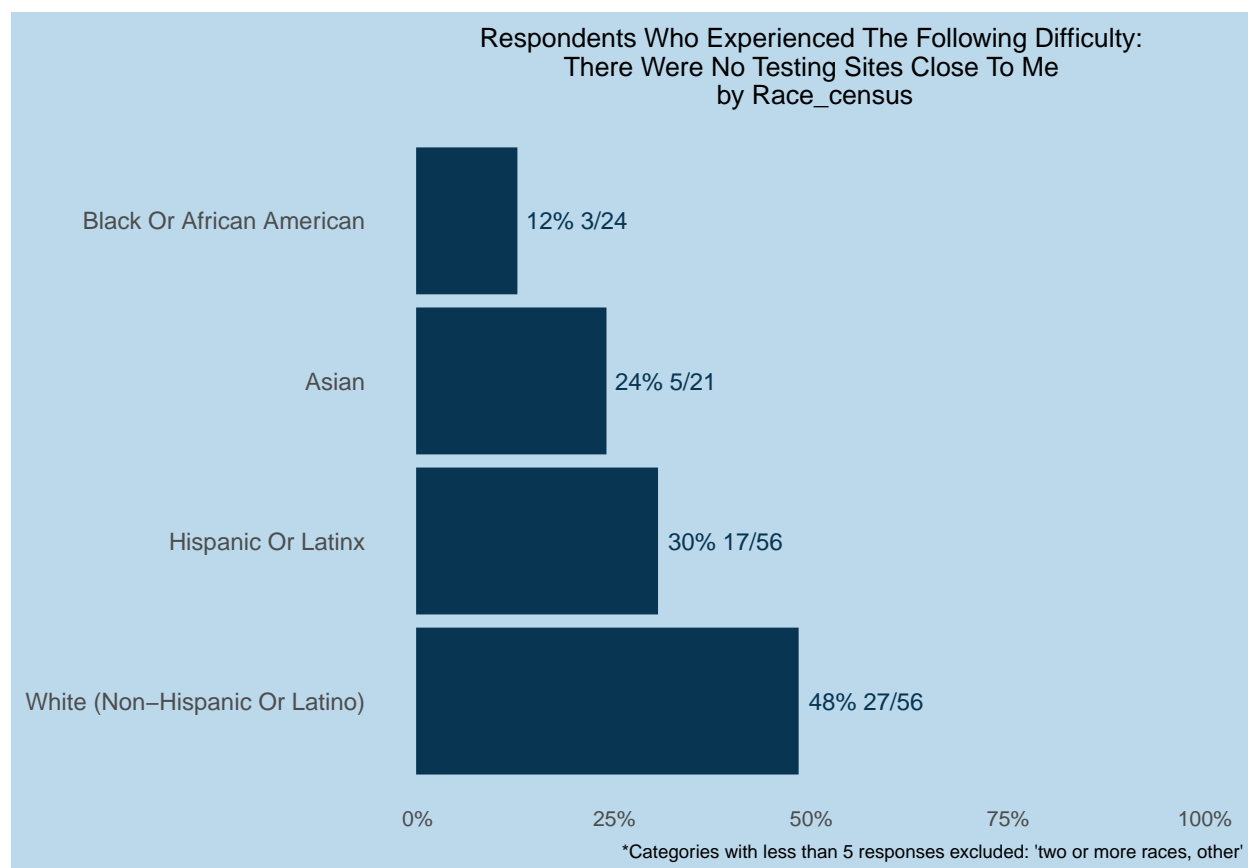
## 4.16) Distribution over each difficulty faced while testing

1. Run each type of difficulty over sub demographic groups

```
test_diffs <- wrangled %>% select(starts_with("test_diffs_") & !ends_with("text")) %>% colnames
lapply(test_diffs, function(test_diff) {
  difficulty <-
    paste0(survey_codebook_labelled$to_name[survey_codebook_labelled$full_name == test_diff], "\n")

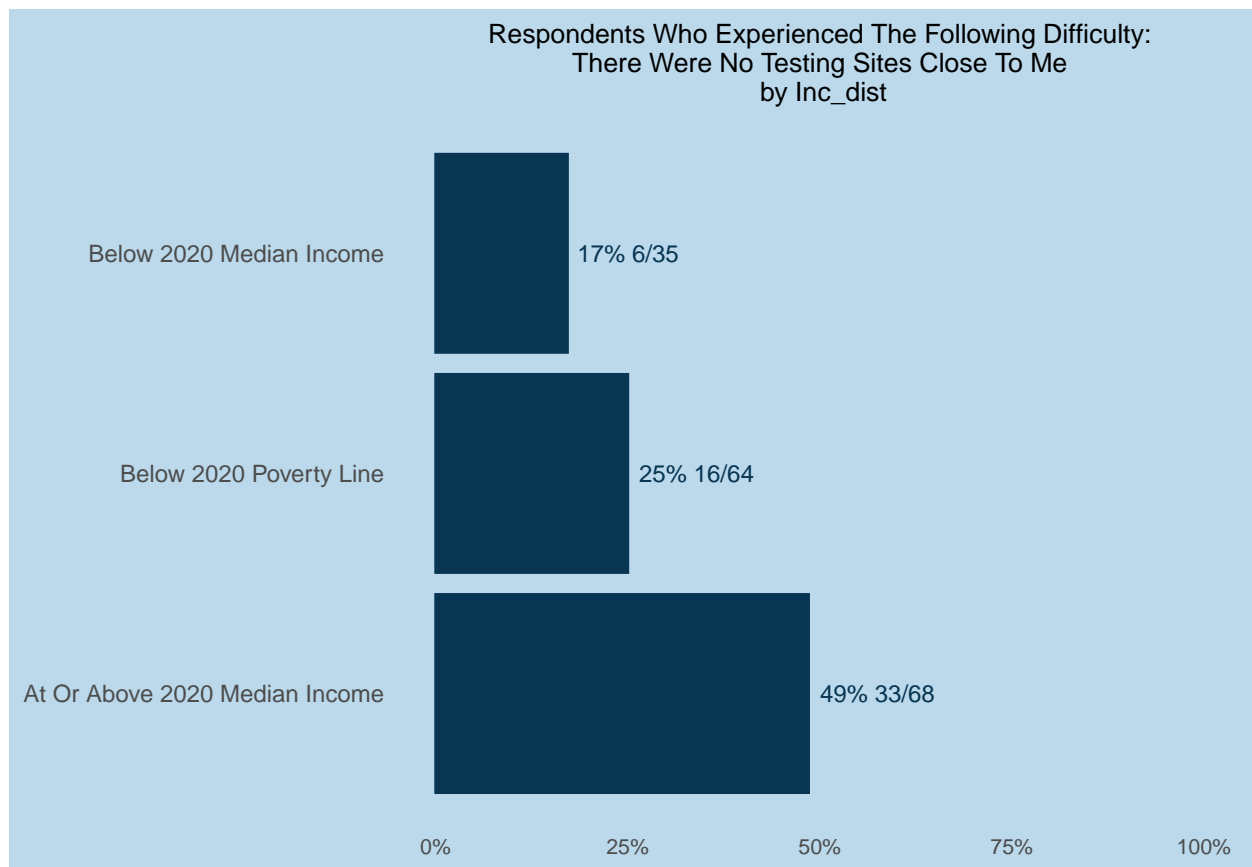
  make_plots(wrangled, demographics, test_diff,
             title = paste("Respondents who experienced the following difficulty:", difficulty, sep = "
  })
```

```
## [[1]]
## [[1]]$borough
## NULL
##
## [[1]]$gen
## NULL
##
## [[1]]$race_census
## [[1]]$race_census$plot
```



```
##
## [[1]]$race_census$p.values
## [[1]]$race_census$p.values$test_diffs_close
##               black or african american asian
## black or african american              NA    NA
## asian                                NA    NA
## hispanic or latinx                     NA    NA
## white (non-hispanic or latino)         0.0056  NA
##               hispanic or latinx
## black or african american              NA
## asian                                NA
## hispanic or latinx                     NA
## white (non-hispanic or latino)         NA
##               white (non-hispanic or latino)
## black or african american              0.0056
## asian                                NA
## hispanic or latinx                     NA
## white (non-hispanic or latino)         NA
##
##
## [[1]]$hh_ch_0_17_bi
## NULL
##
## [[1]]$hh_sn_65_bi
## NULL
```

```
##
## [[1]]$inc_dist
## [[1]]$inc_dist$plot
```



```
##
## [[1]]$inc_dist$p.values
## [[1]]$inc_dist$p.values$test_diffs_close
##
##          below 2020 median income below 2020 poverty line
## below 2020 median income          NA          NA
## below 2020 poverty line          NA          NA
## at or above 2020 median income    0.0038    0.0089
##
##          at or above 2020 median income
## below 2020 median income    0.0038
## below 2020 poverty line    0.0089
## at or above 2020 median income    NA
##
##
##
## [[2]]
## [[2]]$borough
## NULL
##
## [[2]]$gen
## NULL
```

```

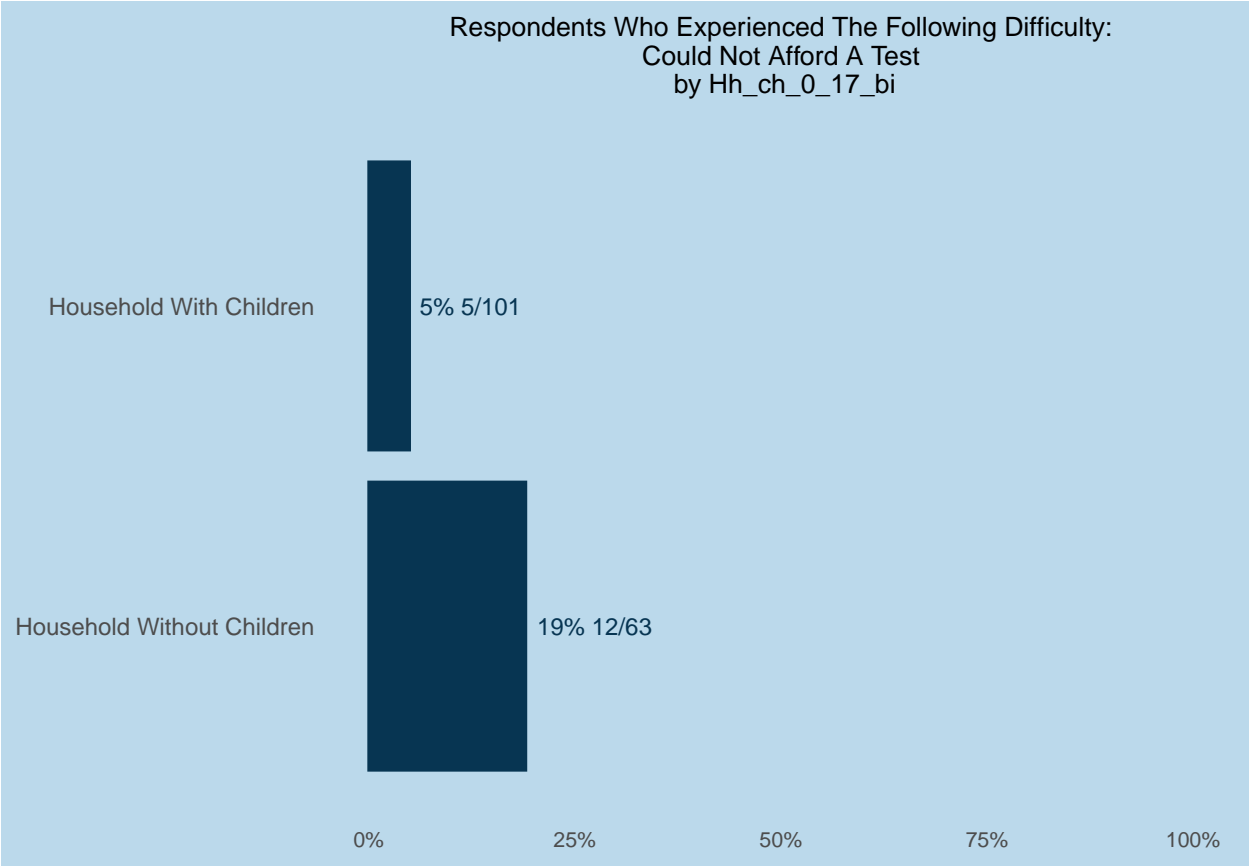
##
## [[2]]$race_census
## NULL
##
## [[2]]$hh_ch_0_17_bi
## NULL
##
## [[2]]$hh_sn_65_bi
## NULL
##
## [[2]]$inc_dist
## NULL
##
##
## [[3]]
## [[3]]$borough
## NULL
##
## [[3]]$gen
## NULL
##
## [[3]]$race_census
## NULL
##
## [[3]]$hh_ch_0_17_bi
## NULL
##
## [[3]]$hh_sn_65_bi
## NULL
##
## [[3]]$inc_dist
## NULL
##
##
## [[4]]
## [[4]]$borough
## NULL
##
## [[4]]$gen
## NULL
##
## [[4]]$race_census
## NULL
##
## [[4]]$hh_ch_0_17_bi
## NULL
##
## [[4]]$hh_sn_65_bi
## NULL
##
## [[4]]$inc_dist
## NULL
##
##

```

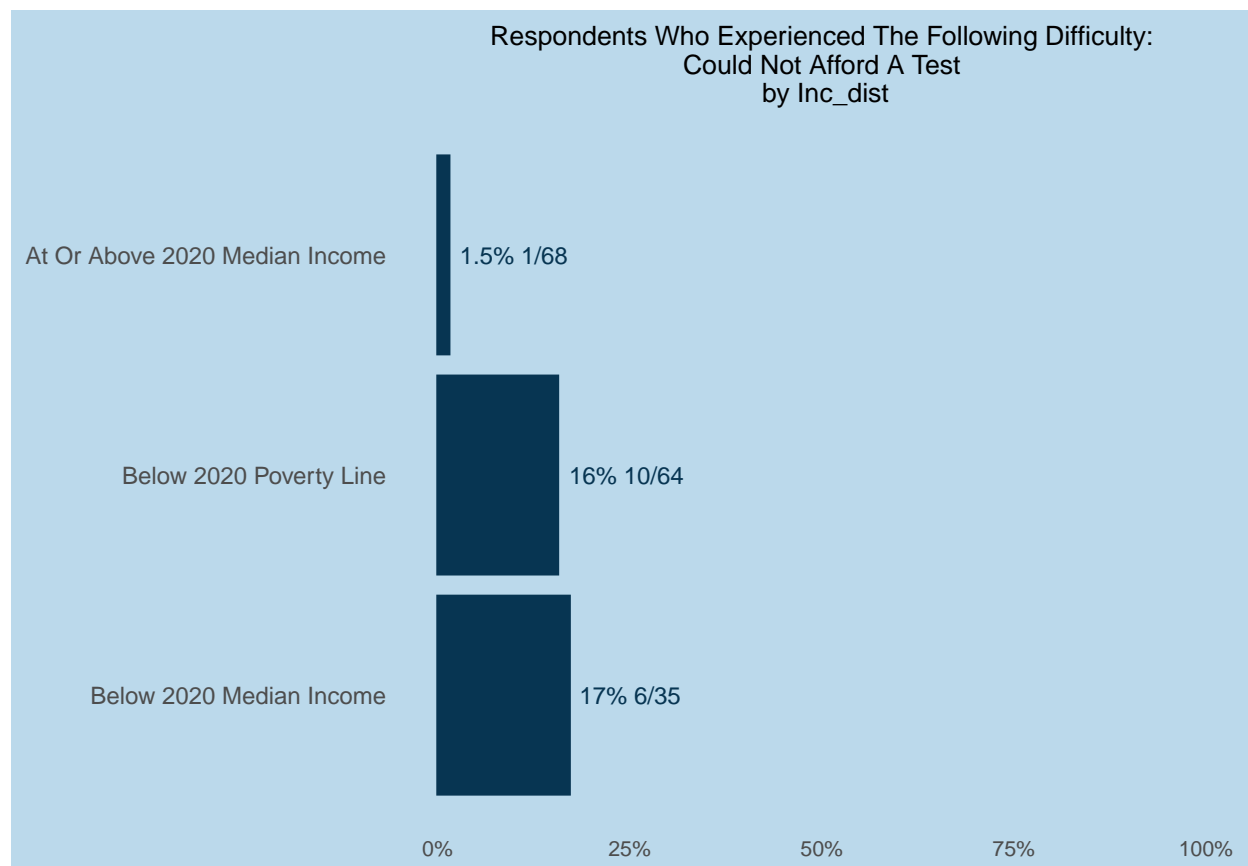
```

## [[5]]
## [[5]]$borough
## NULL
##
## [[5]]$gen
## NULL
##
## [[5]]$race_census
## NULL
##
## [[5]]$hh_ch_0_17_bi
## NULL
##
## [[5]]$hh_sn_65_bi
## NULL
##
## [[5]]$inc_dist
## NULL
##
##
## [[6]]
## [[6]]$borough
## NULL
##
## [[6]]$gen
## NULL
##
## [[6]]$race_census
## NULL
##
## [[6]]$hh_ch_0_17_bi
## [[6]]$hh_ch_0_17_bi$plot

```



```
##
## [[6]]$hh_ch_0_17_bi$p.values
## [[6]]$hh_ch_0_17_bi$p.values$test_diffs_aff
##
##           household with children household without children
## household with children           NA           0.0089
## household without children       0.0089           NA
##
##
##
## [[6]]$hh_sn_65_bi
## NULL
##
## [[6]]$inc_dist
## [[6]]$inc_dist$plot
```



```
##
## [[6]]$inc_dist$p.values
## [[6]]$inc_dist$p.values$test_diffs_aff
##               at or above 2020 median income
## at or above 2020 median income                NA
## below 2020 poverty line                      0.0087
## below 2020 median income                      NA
##               below 2020 poverty line below 2020 median income
## at or above 2020 median income                0.0087                NA
## below 2020 poverty line                      NA                NA
## below 2020 median income                      NA                NA
##
##
##
## [[7]]
## [[7]]$borough
## NULL
##
## [[7]]$gen
## NULL
##
## [[7]]$race_census
## NULL
##
## [[7]]$hh_ch_0_17_bi
```



```
## NULL
##
## [[7]]$hh_sn_65_bi
## NULL
##
## [[7]]$inc_dist
## NULL
```

```
cat("Plots with no statistically significant result are omitted from printing. See those marked with 'N"
```

```
## Plots with no statistically significant result are omitted from printing. See those marked with 'NUL
```