



# Visualizing a categorical variable

Kelly McConville
Assistant Professor of Statistics

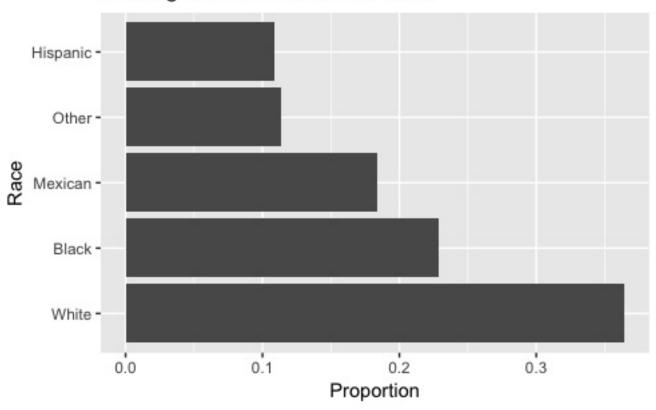


```
library(dplyr)
tab unw <- NHANESraw %>%
   group by(Race1) %>%
   summarize(Freq = n()) %>%
  mutate(Prop = Freq/sum(Freq)) %>%
   arrange(desc(Prop))
tab unw
# A tibble: 5 x 3
    Racel Freq Prop
    <fctr> <int> <dbl>
   White 7393 0.3643128
     Black 4640 0.2286503
   Mexican 3739 0.1842507
     Other 2312 0.1139309
 5 Hispanic 2209 0.1088553
```

### NHANES documentation

```
library(ggplot2)
ggplot(data = tab_unw, mapping = aes(x = Race1, y = Prop)) +
   geom_col() +
   coord_flip() +
   scale_x_discrete(limits = tab_unw$Race1) # Labels layer omitted
```

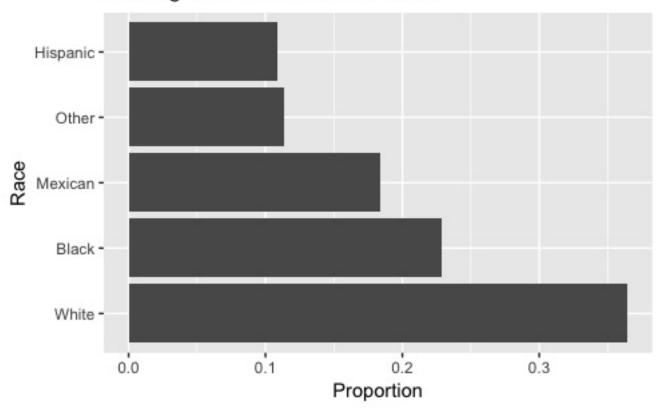
### Unweighted Distribution of Race





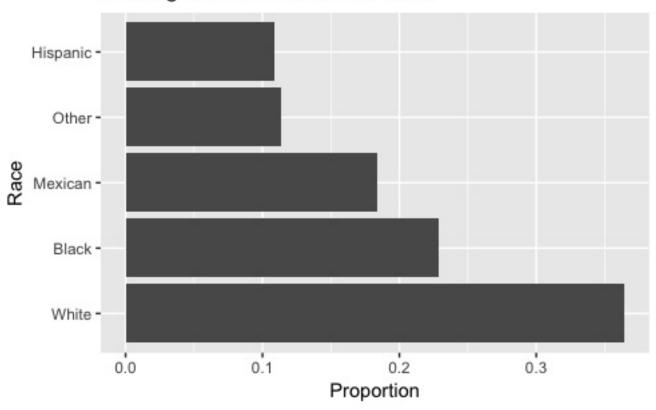
```
library(ggplot2)
ggplot(data = tab_unw, mapping = aes(x = Race1, y = Prop)) +
   geom_bar(stat = "identity") +
   coord_flip() +
   scale_x_discrete(limits = tab_unw$Race1) # Labels layer omitted
```

### Unweighted Distribution of Race



```
library(ggplot2)
ggplot(data = tab_unw, mapping = aes(x = Race1, y = Prop)) +
   geom_col() +
   coord_flip() +
   scale_x_discrete(limits = tab_unw$Race1) # Labels layer omitted
```

### Unweighted Distribution of Race





```
tab_w <- svytable(~Race1, design = NHANES_design) %>%
    as.data.frame() %>%
    mutate(Prop = Freq/sum(Freq)) %>%
    arrange(desc(Prop))

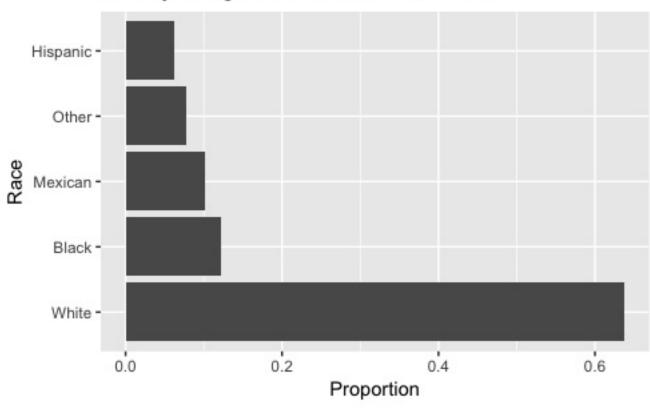
tab_w

Race1    Freq    Prop
    White 193966274 0.63748664
    Black 37241616 0.12239773
    Mexican 30719158 0.10096112
    4    Other 23389002 0.07686994
    5 Hispanic 18951150 0.06228456
```



```
ggplot(data = tab_w, mapping = aes(x = Race1, y = Prop)) +
  geom_col() +
  coord_flip() +
  scale_x_discrete(limits = tab_w$Race1) # Labels layer omitted
```

### Survey-Weighted Distribution of Race







# Let's practice!





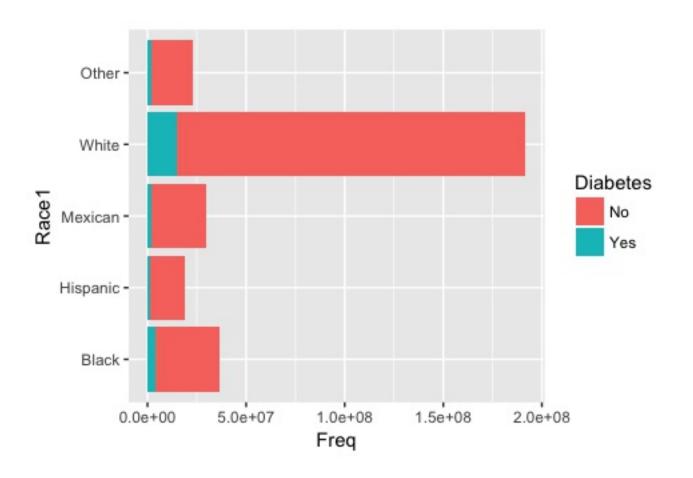
# Exploring two categorical variables

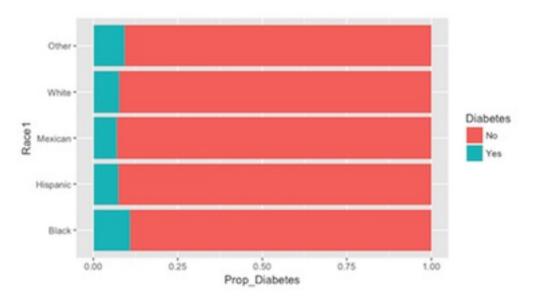
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Assistant Professor of Statistics

```
svytable(~Diabetes,
        design = NHANES design)
Diabetes
      No
           Yes
275814034 24335536
tab w <- svytable(~Race1 + Diabetes,</pre>
                 design = NHANES design)
tab w
         Diabetes
                     Yes
Race1
  Black 32697528
                     4003497
 Hispanic 17258245
                     1370393
 Mexican 27886500
                     2081657
 White 177088354
                    14708094
  Other 20883407
                    2171895
```



```
tab w <- as.data.frame(tab w)</pre>
tab w
      Racel Diabetes
                          Freq
                      32697528
      Black
   Hispanic
                      17258245
   Mexican
                      27886500
     White
                 No 177088354
    0ther
                      20883407
      Black
                      4003497
             Yes
  Hispanic
                 Yes
                      1370393
   Mexican
                       2081657
                 Yes
     White
                 Yes
                      14708094
10
      0ther
                 Yes
                       2171895
ggplot(data = tab w,
       mapping = aes(x = Race1,
                     fill = Diabetes,
                     y = Freq)) +
  geom col() + coord flip()
```









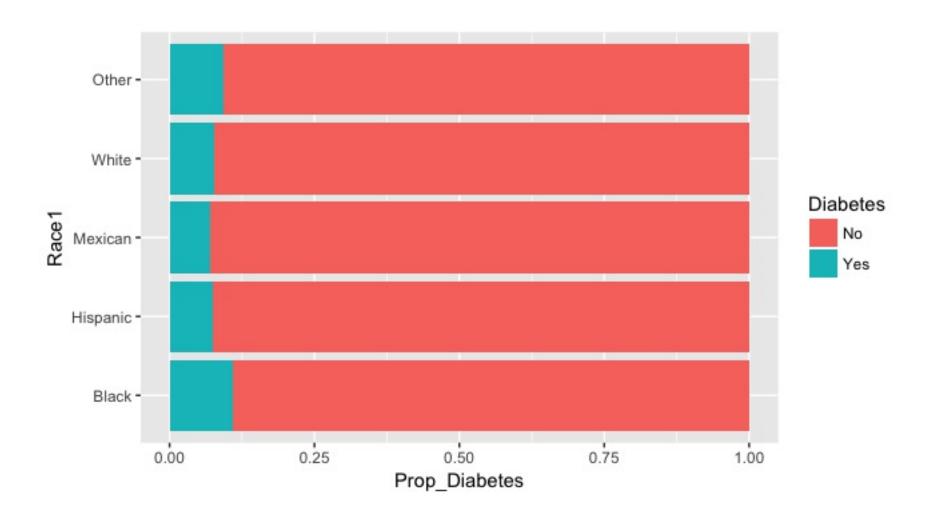
# Let's practice!





# Inference for categorical variables

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# Inference: Chi-square Test

**Null Hypothesis**: Prevalence of diabetes is not associated with race.

**Alternative Hypothesis**: Prevalence of diabetes is associated with race.

```
svychisq(~Race1 + Diabetes, design = NHANES_design, statistic = "Chisq")

Pearsons X^2: Rao & Scott adjustment

data: svychisq(~Race1 + Diabetes, design = NHANES_design, statistic = "Chisq")
    X-squared = 37.708, df = 4, p-value = 0.0001177
```





# Let's practice!