NYPD_Shooting

Edison

2022-06-20

Introduction

Data is on every shooting that occurred in NYC from 2006 to the end of the previous calender year which should mean end of December 2021.

Every row in the data corresponds to a single shooting incident. Location and demographic data are included.

Load Packages

```
library(tidyverse)
## -- Attaching packages ------ tidyverse 1.3.1 --
## v ggplot2 3.3.6
                   v purrr
                            0.3.4
## v tibble 3.1.7
                   v dplyr
                           1.0.9
## v tidyr
          1.2.0
                   v stringr 1.4.0
## v readr
          2.1.2
                   v forcats 0.5.1
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                 masks stats::lag()
library(pander) # only for the single markdown table, can skip
```

Read Data

Let's load in the data from: https://catalog.data.gov/dataset/nypd-shooting-incident-data-historic

```
data_raw <- read_csv("https://data.cityofnewyork.us/api/views/833y-fsy8/rows.csv?accessType=DOWNLOAD")
```

```
## Rows: 25596 Columns: 19
## -- Column specification -------
## Delimiter: ","
## chr (10): OCCUR_DATE, BORO, LOCATION_DESC, PERP_AGE_GROUP, PERP_SEX, PERP_R...
## dbl (7): INCIDENT_KEY, PRECINCT, JURISDICTION_CODE, X_COORD_CD, Y_COORD_CD...
## lgl (1): STATISTICAL_MURDER_FLAG
## time (1): OCCUR_TIME
##
## i Use 'spec()' to retrieve the full column specification for this data.
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
```

Preparing Data

A tibble: 6 x 19

##

##

1

INCIDENT KEY OCCUR DATE OCCUR TIME BORO

24050482 08/27/2006 05:35

<dbl> <chr>

<time>

summary(data raw) INCIDENT_KEY OCCUR_DATE OCCUR_TIME BORO ## ## : 9953245 Length: 25596 Length: 25596 Length: 25596 1st Qu.: 61593633 Class : character Class1:hms Class : character Median: 86437258 Mode : character Class2:difftime Mode : character ## Mean :112382648 Mode :numeric 3rd Qu.:166660833 :238490103 ## Max. ## ## PRECINCT JURISDICTION_CODE LOCATION_DESC STATISTICAL_MURDER_FLAG : 1.00 ## Min. Min. :0.0000 Length:25596 Mode :logical 1st Qu.: 44.00 1st Qu.:0.0000 Class : character FALSE: 20668 Median : 69.00 Median :0.0000 TRUE: 4928 Mode :character ## Mean : 65.87 Mean :0.3316 ## 3rd Qu.: 81.00 3rd Qu.:0.0000 Max. :123.00 ## Max. :2.0000 ## NA's :2 ## PERP_AGE_GROUP PERP_SEX PERP_RACE VIC_AGE_GROUP ## Length: 25596 Length: 25596 Length: 25596 Length: 25596 Class : character Class : character Class : character Class : character ## Mode :character Mode :character Mode :character Mode :character ## ## ## ## ## VIC_SEX VIC RACE X COORD CD Y_COORD_CD ## Length: 25596 Length: 25596 : 914928 :125757 Min. Min. Class : character Class : character 1st Qu.:1000011 1st Qu.:182782 ## Mode :character Mode :character Median :1007715 Median :194038 ## Mean :1009455 Mean :207894 ## 3rd Qu.:1016838 3rd Qu.:239429 ## Max. :1066815 Max. :271128 ## ## Latitude Longitude Lon_Lat :40.51 :-74.25 Length: 25596 1st Qu.:40.67 1st Qu.:-73.94 Class : character Median :40.70 Median :-73.92 Mode : character ## Mean :40.74 Mean :-73.91 3rd Qu.:40.82 3rd Qu.:-73.88 :40.91 :-73.70 ## Max. Max. ## head(data_raw)

<chr>>

BRONX

PRECINCT JURISDICTION_CODE

<dbl>

0

<dbl>

52

```
## 2
       77673979 03/11/2011 12:03
                                   QUEENS
                                                 106
## 3
      226950018 04/14/2021 21:08
                                   BRONX
                                                  42
                                                                  0
                                                                  0
## 4
       237710987 12/10/2021 19:30
                                   BRONX
                                                 52
## 5
       224701998 02/22/2021 00:18
                                                 34
                                                                  0
                                   MANHATTAN
## 6
       225295736 03/07/2021 06:15
                                   BROOKLYN
                                                 75
                                                                  0
## # ... with 13 more variables: LOCATION_DESC <chr>,
      PERP_RACE <chr>, VIC_AGE_GROUP <chr>, VIC_SEX <chr>, VIC_RACE <chr>,
## #
## #
      X_COORD_CD <dbl>, Y_COORD_CD <dbl>, Latitude <dbl>, Longitude <dbl>,
## #
      Lon_Lat <chr>>
```

Let's take a look at the column descriptions:

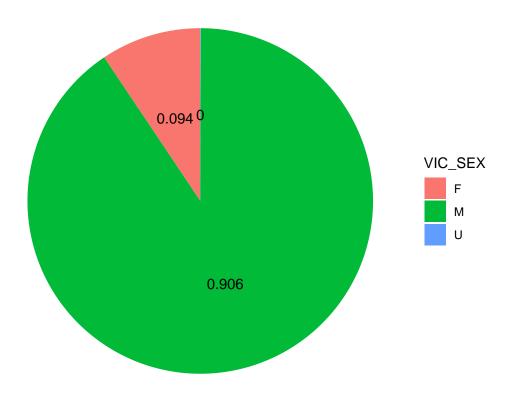
source: https://data.cityofnewyork.us/Public-Safety/NYPD-Shooting-Incident-Data-Historic-/833y-fsy8

Column Name	Description
INCIDENT_KEY	Randomly generated persistent ID for each arrest
OCCUR DATE	Exact date of the shooting incident
OCCUR TIME	Exact time of the shooting incident
BORO	Borough where the shooting incident occurred
PRECINCT	Precinct where the shooting incident occurred
JURISDICTION_CODE	Jurisdiction where the shooting incident occurred.
	Jurisdiction codes 0(Patrol), 1(Transit) and
	2(Housing) represent NYPD whilst codes 3 and
	more represent non NYPD jurisdictions
LOCATION_DESC	Location of the shooting incident
STATISTICAL_MURDER_FLAG	Shooting resulted in the victim's death which would
	be counted as a murder
PERP_AGE_GROUP	Perpetrator's age within a category
PERP_SEX	Perpetrator's sex description
PERP_RACE	Perpetrator's race description
VIC_AGE_GROUP	Victim's age within a category
VIC_SEX	Victim's sex description
VIC_RACE	Victim's race description
X_COORD_CD	Midblock X-coordinate for New York State Plane
	Coordinate System, Long Island Zone, NAD 83,
	units feet (FIPS 3104)
Y_COORD_CD	Midblock Y-coordinate for New York State Plane
	Coordinate System, Long Island Zone, NAD 83,
	units feet (FIPS 3104)
Latitude	Latitude coordinate for Global Coordinate System,
	WGS 1984, decimal degrees (EPSG 4326)
Longitude	Longitude coordinate for Global Coordinate
	System, WGS 1984, decimal degrees (EPSG 4326)
Lon_Lat	Longitude and Latitude Coordinates for mapping

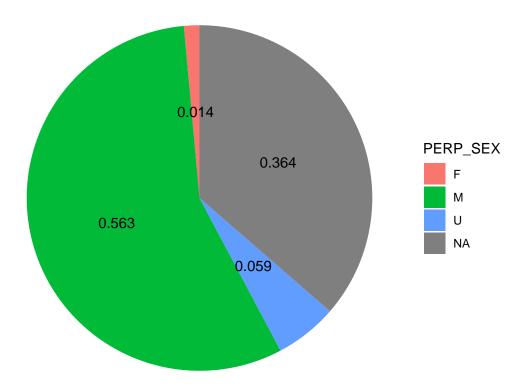
Data Visualization

Let's see how the genders stacked up in the dataset.

First we'll take a look at the victims.



Now let's take a look at the perp's sex distribution.



Males make the majority of victims and perps. Of the known data, women are even less likely to be the perp. The perp data does have a lot of NA and UNKNOWN data.

What else could we explore with this data? Well we could see if a popular statistic is also true for this dataset.

I believe the statistic is that murders tend to be largely within race.

Let's trim this down to relevant columns.

```
data <- data_raw[c("PERP_RACE", "VIC_RACE")]
colSums(is.na(data))

## PERP_RACE VIC_RACE
## 9310 0</pre>
```

The missing data for these these two columns are only in PERP_RACE. It must be when the perp is not caught and is unknown.

Let's remove the rows with NA.

```
print(nrow(data))
```

[1] 25596

```
data_no_na <- na.omit(data)
print(nrow(data_no_na))</pre>
```

[1] 16286

Over a third of the data is dropped which is quite a lot. However, having missing PERP_RACE data would not be helpful for our purposes.

Let's see how the PERP_RACE values look like:

data_no_na %>% count(PERP_RACE)

```
## # A tibble: 7 x 2
##
    PERP_RACE
                                         n
##
     <chr>>
                                      <int>
## 1 AMERICAN INDIAN/ALASKAN NATIVE
                                          2
## 2 ASIAN / PACIFIC ISLANDER
                                        141
## 3 BLACK
                                      10668
## 4 BLACK HISPANIC
                                       1203
## 5 UNKNOWN
                                       1836
## 6 WHITE
                                        272
## 7 WHITE HISPANIC
                                       2164
```

Now let's take a look at the corresponding VIC_RACE values:

data_no_na %>% count(VIC_RACE)

```
## # A tibble: 7 x 2
##
     VIC_RACE
                                         n
     <chr>>
                                      <int>
## 1 AMERICAN INDIAN/ALASKAN NATIVE
## 2 ASIAN / PACIFIC ISLANDER
                                        257
## 3 BLACK
                                     11117
## 4 BLACK HISPANIC
                                       1641
## 5 UNKNOWN
                                         47
## 6 WHITE
                                        515
## 7 WHITE HISPANIC
                                       2702
```

Seems like there is a lot of similarity across the two. The UNKNOWN labels seem like they would be problematic as the number for UNKNOWN is a lot higher than the UNKNOWN for the VIC_RACE.

I will drop any row that contains UNKNOWN as well.

```
data_complete <- data_no_na[rowSums(data_no_na == "UNKNOWN")==0, , drop = FALSE]
data_complete %>% count(VIC_RACE)
```

```
## 3 BLACK 9758
## 4 BLACK HISPANIC 1486
## 5 WHITE 473
## 6 WHITE HISPANIC 2447
```

data_complete %>% count(PERP_RACE)

```
## # A tibble: 6 x 2
##
    PERP_RACE
                                         n
     <chr>>
##
                                     <int>
## 1 AMERICAN INDIAN/ALASKAN NATIVE
                                         2
## 2 ASIAN / PACIFIC ISLANDER
                                       141
## 3 BLACK
                                     10644
## 4 BLACK HISPANIC
                                      1198
## 5 WHITE
                                       271
## 6 WHITE HISPANIC
                                      2153
```

Data distribution in PERP and VIC race seems pretty similar.

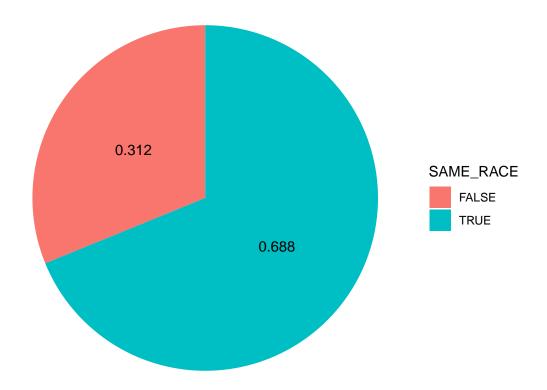
Data went from 25596 rows to 14409 rows so we lost about half our data which is a lot.

Let's make a new column that prints 1 if both VIC_RACE and PERP_RACE is exactly the same.

```
data_complete$SAME_RACE <- as.factor(data_complete$VIC_RACE == data_complete$PERP_RACE)
frequency_table <- data_complete %>% count(SAME_RACE)
frequency_table$n <- frequency_table$n / nrow(data_complete)
frequency_table</pre>
```

```
## # A tibble: 2 x 2
## SAME_RACE n
## <fct> <dbl>
## 1 FALSE 0.312
## 2 TRUE 0.688
```

Let's visualize this really quick



While not high as I expected, in about 69% of all shootings, where both the victim and perp races are known, they are of the same race label.

To get some insight on why it might not be higher, let's take a look at the counts between PERP_VIC of each race type.

data_complete\$PERP_VIC <- paste(data_complete\$PERP_RACE, "-", data_complete\$VIC_RACE)
data_complete %>% count(PERP_VIC)

```
## # A tibble: 27 x 2
      PERP_VIC
##
                                                               n
##
      <chr>
                                                           <int>
##
   1 AMERICAN INDIAN/ALASKAN NATIVE - BLACK
                                                               2
  2 ASIAN / PACIFIC ISLANDER - ASIAN / PACIFIC ISLANDER
                                                              43
   3 ASIAN / PACIFIC ISLANDER - BLACK
                                                              51
  4 ASIAN / PACIFIC ISLANDER - BLACK HISPANIC
##
                                                              13
##
   5 ASIAN / PACIFIC ISLANDER - WHITE
                                                              11
   6 ASIAN / PACIFIC ISLANDER - WHITE HISPANIC
                                                              23
  7 BLACK - AMERICAN INDIAN/ALASKAN NATIVE
                                                               4
   8 BLACK - ASIAN / PACIFIC ISLANDER
                                                             135
## 9 BLACK - BLACK
                                                            8471
## 10 BLACK - BLACK HISPANIC
                                                             749
## # ... with 17 more rows
```

As we can see BLACK HISPANIC to BLACK wouldn't count as same race using our current matching method, so if we wanted to include those, we would need a function more complex. The TRUE rate would be higher if we included these cases.

Let's leave it as is, but let's be aware of this issue.

Model

Let's see if any PERP race can be used to predict the created SAME_RACE factor better than others.

```
model <- glm(SAME_RACE ~ PERP_RACE, data = data_complete, family = "binomial")
summary(model)</pre>
```

```
##
## Call:
  glm(formula = SAME RACE ~ PERP RACE, family = "binomial", data = data complete)
##
## Deviance Residuals:
                      Median
                                    3Q
##
       Min
                 10
                                             Max
           -0.8530
                      0.6758
                                0.6758
                                          1.6249
##
  -1.7826
##
## Coefficients:
                                      Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                                         -11.57
                                                    139.28
                                                            -0.083
                                                                       0.934
## PERP_RACEASIAN / PACIFIC ISLANDER
                                                             0.077
                                                                       0.939
                                          10.74
                                                    139.28
## PERP_RACEBLACK
                                          12.93
                                                    139.28
                                                             0.093
                                                                       0.926
## PERP_RACEBLACK HISPANIC
                                          10.56
                                                    139.28
                                                             0.076
                                                                       0.940
## PERP_RACEWHITE
                                          11.87
                                                    139.28
                                                             0.085
                                                                       0.932
## PERP RACEWHITE HISPANIC
                                          11.29
                                                    139.28
                                                             0.081
                                                                       0.935
##
##
  (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 17881
                                        degrees of freedom
                              on 14408
## Residual deviance: 15650
                              on 14403
                                        degrees of freedom
## AIC: 15662
##
## Number of Fisher Scoring iterations: 10
```

As we can see, none of the variables are significant. It seems like individual PERP race is not a good indicator of whether the VIC RACE will be exactly the same.

Since percentage of SAME_RACE shooting is about 69% and likely over if we include race subcategory Hispanic, it would likely be that any model will predict that SAME_RACE is true regardless of PERP_RACE category.

Conclusion

In this report, we examined our NYPD dataset to see if it fit a popular FBI statistic that murders largely tend to have victims of the same race. We saw that the NYPD does seem to match this statistic where nearly 69% of the shooting cases where both perpetrator and victim races are known, both are of the same race. There are some conflicts with this category with how Hispanic sub categories can be counted, but that will be left for future analysis. We took this SAME_RACE variable and regressed it on PERP_RACE and found that there wasn't enough evidence to conclude that any race is more likely than others to shoot someone who is of the race.

Is there any possible bias in the data source? Well it is reported by the police department and since the perpetrator can sometimes be missing or unknown, it is difficult to say whether a source of bias can be induced. I am not sure how the perpetrator race factors are determined either and could be based off eyewitness testimony. Those could be unreliable as well as people don't have the best of memory and can be biased as well. Victim race seems a lot less likely to be unbiased since the data would be more accurate due to being dead or at the incident report themselves. Further analysis can be done to see if there are any good indicators to predict missing race or age group categories based on location, borough, and victim demographics.