

2007-2017 OASAS Admissions Analytics

```
# Documentation from data.world recommends package installation from GitHub
devtools::install_github("datadotworld/data.world-r", build_vignettes = TRUE)
```

```
## Using GitHub PAT from the git credential store.
```

```
## Skipping install of 'data.world' from a github remote, the SHA1 (a1fd7656) has not changed since last
##   Use 'force = TRUE' to force installation
```

```
# Load the requisite API token obtained from data.world advanced settings
token <- readLines('api')
saved_cfg <- data.world::save_config(token)
data.world::set_config(saved_cfg)

sql_stmt <- data.world::qry_sql("SELECT * FROM
                                chemical_dependence_treatment_program_admissions_beginning_2007_1")
admissions_df <- data.world::query(
  sql_stmt, "https://data.world/data-ny-gov/ngbt-9rwf")
```

```
## Rows: 72463 Columns: 7
```

```
## -- Column specification -----
## Delimiter: ","
## chr (5): county_of_program_location, program_category, service_type, age_group...
## dbl (2): year, admissions
##
## 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.
```

Data Understanding

- Initial look at dataframe, check data types

```
str(admissions_df)
```

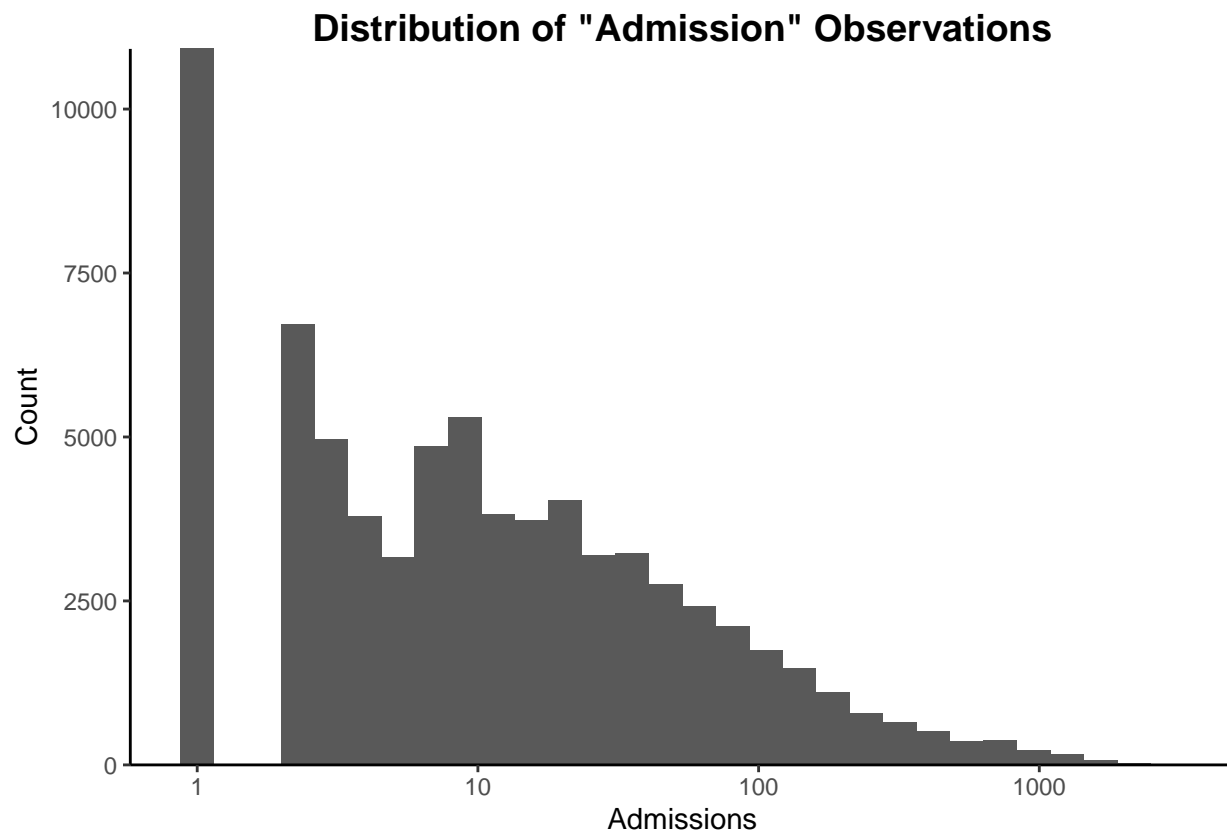
```
## spc_tbl_ [72,463 x 7] (S3: spec_tbl_df/tbl_df/tbl/data.frame)
## $ year                : num [1:72463] 2017 2017 2017 2017 2017 ...
## $ county_of_program_location: chr [1:72463] "Albany" "Albany" "Albany" "Albany" ...
## $ program_category      : chr [1:72463] "Crisis" "Crisis" "Crisis" "Crisis" ...
## $ service_type          : chr [1:72463] "Medically Managed Detoxification" "Medically Managed D...
## $ age_group             : chr [1:72463] "18 thru 24" "18 thru 24" "18 thru 24" "18 thru 24" ...
```

```
## $ primary_substance_group : chr [1:72463] "Alcohol" "All Others" "Cocaine incl Crack" "Heroin" ..
## $ admissions              : num [1:72463] 25 7 1 64 20 140 10 3 1 41 ...
## - attr(*, "spec")=
## .. cols(
## ..   year = col_double(),
## ..   county_of_program_location = col_character(),
## ..   program_category = col_character(),
## ..   service_type = col_character(),
## ..   age_group = col_character(),
## ..   primary_substance_group = col_character(),
## ..   admissions = col_double()
## .. )
## - attr(*, "problems")=<externalptr>
```

- Admissions trends grouped by each variable

```
ggplot(admissions_df, aes(x = admissions)) +
  geom_histogram() +
  labs(title = "Distribution of \"Admission\" Observations",
       x = "Admissions",
       y = "Count") +
  scale_x_log10() +
  scale_y_continuous(expand = c(0,0)) +
  theme_classic() +
  theme(
    plot.title = element_text(face = "bold", size = 14, hjust = 0.5, margin = margin(t = 5))
  )
```

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



```
program_category_summary <- admissions_df %>%
  group_by(program_category) %>%
  summarize(
    min_admissions = min(admissions),
    median_admissions = median(admissions),
    mean_admissions = mean(admissions),
    max_admissions = max(admissions)
  )
print(program_category_summary)
```

```
## # A tibble: 5 x 5
##   program_category    min_admissions median_admissions mean_admissions
##   <chr>              <dbl>             <dbl>             <dbl>
## 1 Crisis              1              15              67.2
## 2 Inpatient           1              15              43.8
## 3 Opioid Treatment Program 1              9              55.4
## 4 Outpatient          1              11              56.1
## 5 Residential         1               3              11.7
## # i 1 more variable: max_admissions <dbl>
```

```
service_type_summary <- admissions_df %>%
  group_by(service_type) %>%
  summarize(
    min_admissions = min(admissions),
    median_admissions = median(admissions),
```

```

    mean_admissions = mean(admissions),
    max_admissions = max(admissions)
  )
print(service_type_summary)

```

```

## # A tibble: 28 x 5
##   service_type min_admissions median_admissions mean_admissions max_admissions
##   <chr>          <dbl>          <dbl>          <dbl>          <dbl>
## 1 Community Re~      1              3           6.70           143
## 2 Inpatient Re~      1             15          43.8          1106
## 3 Intensive Re~      1              7          25.0           516
## 4 Limited Outp~      1              7          15.0           151
## 5 Long Term Re~      1              2           4.79            31
## 6 Med Sup With~      1             14          70.6          2058
## 7 Med Sup With~      1              8          32.1           341
## 8 Medically Ma~      1             19          95.1          2862
## 9 Medically Mo~      1             13          38.9          2516
## 10 Meth to Abst~     1             16          27.4            79
## # i 18 more rows

```

```

age_group_summary <- admissions_df %>%
  group_by(age_group) %>%
  summarize(
    min_admissions = min(admissions),
    median_admissions = median(admissions),
    mean_admissions = mean(admissions),
    max_admissions = max(admissions)
  )
print(age_group_summary)

```

```

## # A tibble: 6 x 5
##   age_group min_admissions median_admissions mean_admissions max_admissions
##   <chr>          <dbl>          <dbl>          <dbl>          <dbl>
## 1 18 thru 24      1              8          31.3          1518
## 2 25 thru 34      1             13          52.9          1876
## 3 35 thru 44      1             10          53.1          2862
## 4 45 thru 54      1              8          58.8          2716
## 5 55 and Older    1              5          30.0          1277
## 6 Under 18        1              4          23.5           661

```

```

primary_substance_group_summary <- admissions_df %>%
  group_by(primary_substance_group) %>%
  summarize(
    min_admissions = min(admissions),
    median_admissions = median(admissions),
    mean_admissions = mean(admissions),
    max_admissions = max(admissions)
  )
print(primary_substance_group_summary)

```

```

## # A tibble: 6 x 5
##   primary_substance_group min_admissions median_admissions mean_admissions

```

```
##   <chr>                                <dbl>                <dbl>                <dbl>
## 1 Alcohol                             1                 21                 91.6
## 2 All Others                           1                  3                  9.92
## 3 Cocaine incl Crack                   1                  7                 29.3
## 4 Heroin                               1                 13                 55.0
## 5 Marijuana incl Hashish               1                  8                 46.0
## 6 Other Opioids                        1                  6                 15.4
## # i 1 more variable: max_admissions <dbl>
```

- Factoring variables and checking for outliers

```
admissions_df <- admissions_df %>%
  mutate(
    program_category = as.factor(program_category),
    service_type = as.factor(service_type),
    age_group = as.factor(age_group),
    primary_substance_group = as.factor(primary_substance_group)
  )

# Manual outlier identification
admissions_outliers <- admissions_df %>%
  mutate(
    mean_admissions = mean(admissions, na.rm = TRUE),
    sd_admissions = sd(admissions, na.rm = TRUE)
  ) %>%
  # Relative to the mean, any values on the lower or
  # upper bounds that are 3 times the standard deviation are filtered
  filter(admissions < mean_admissions - 3 * sd_admissions |
    admissions > mean_admissions + 3 * sd_admissions) %>%
  select(admissions)

admissions_outliers
```

```
## # A tibble: 1,380 x 1
##   admissions
##   <dbl>
## 1     526
## 2     468
## 3     515
## 4     501
## 5     752
## 6     496
## 7     566
## 8     442
## 9     564
## 10    469
## # i 1,370 more rows
```

```
# Remove outliers
rmv_admissions_outliers <- admissions_df %>%
  mutate(
    mean_admissions = mean(admissions, na.rm = TRUE),
    sd_admissions = sd(admissions, na.rm = TRUE)
```

```

) %>%
  filter(!(admissions < mean_admissions - 3 * sd_admissions |
           admissions > mean_admissions + 3 * sd_admissions))

# Note subtracted outliers from new dataframe
str(admissions_df)

## tibble [72,463 x 7] (S3: tbl_df/tbl/data.frame)
## $ year : num [1:72463] 2017 2017 2017 2017 2017 ...
## $ county_of_program_location: chr [1:72463] "Albany" "Albany" "Albany" "Albany" ...
## $ program_category : Factor w/ 5 levels "Crisis","Inpatient",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ service_type : Factor w/ 28 levels "Community Residential",...: 8 8 8 8 8 8 8 8 8 8 .
## $ age_group : Factor w/ 6 levels "18 thru 24","25 thru 34",...: 1 1 1 1 1 2 2 4 4 4
## $ primary_substance_group : Factor w/ 6 levels "Alcohol","All Others",...: 1 2 3 4 6 1 2 2 3 4 ...
## $ admissions : num [1:72463] 25 7 1 64 20 140 10 3 1 41 ...

str(rmv_admissions_outliers$admissions)

```

```
## num [1:71083] 25 7 1 64 20 140 10 3 1 41 ...
```

- It may not be significant that admissions numbers are higher than usual at any given point in time

Data Preparation

- Data is structured relationally prior to analysis

```

# External county code .csv from https://www.dot.ny.gov
county_codes <- read_csv("county_codes.csv")

## Rows: 62 Columns: 2
## -- Column specification -----
## Delimiter: ","
## chr (2): county_of_program_location, county_code
##
## 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.

county_codes

```

```

## # A tibble: 62 x 2
##   county_of_program_location county_code
##   <chr>                  <chr>
## 1 Albany                 AL
## 2 Cattaraugus            CA
## 3 Chenango               CN
## 4 Delaware               DE
## 5 Franklin               FR

```

```
## 6 Hamilton HA
## 7 Lewis LE
## 8 Montgomery MG
## 9 Oneida ON
## 10 Orleans OL
## # i 52 more rows
```

```
county <- admissions_df %>%
  select(county_of_program_location) %>%
  distinct() %>%
  # Join codes with corresponding counties
  left_join(county_codes, by = "county_of_program_location") %>%
  mutate(county_code = case_when(
    # Tagging counties with first two characters and "-NYC"
    county_of_program_location %in% c("Bronx", "Queens", "Kings") ~
      paste(str_to_upper(str_sub(county_of_program_location, 1, 2)),
            "-NYC", sep = ""),
    county_of_program_location == "New York" ~ "NYC",
    # Handle to not treat as NA value, changed code to NS instead
    county_of_program_location == "Nassau" ~ "NS",
    # Handle to not treat as NA value, manually assigned SL
    county_of_program_location == "St Lawrence" ~ "SL",
    TRUE ~ county_code
  )
)
county
```

```
## # A tibble: 61 x 2
##   county_of_program_location county_code
##   <chr>                      <chr>
## 1 Albany                     AL
## 2 Bronx                      BR-NYC
## 3 Broome                     BM
## 4 Dutchess                   DU
## 5 Erie                       ER
## 6 Kings                      KI-NYC
## 7 Monroe                     MO
## 8 Nassau                     NS
## 9 New York                   NYC
## 10 Niagara                   NI
## # i 51 more rows
```

```
# Define abbreviations
program_category_index <- c(
  "Crisis" = "C",
  "Inpatient" = "I",
  "Opioid Treatment Program" = "OTP",
  "Outpatient" = "O",
  "Residential" = "R"
)
# Add program_code column
admissions_df <- admissions_df %>%
  mutate(program_code = recode(program_category,
```

```

    "Crisis" = "C",
    "Inpatient" = "I",
    "Opioid Treatment Program" = "OTP",
    "Outpatient" = "O",
    "Residential" = "R"))

# Add program_category tibble
program_category <- admissions_df %>%
  distinct(program_category, .keep_all = TRUE) %>%
  select(program_code, program_category)

program_category

```

```

## # A tibble: 5 x 2
##   program_code program_category
##   <fct>         <fct>
## 1 C             Crisis
## 2 I             Inpatient
## 3 OTP           Opioid Treatment Program
## 4 O             Outpatient
## 5 R             Residential

```

- program_category contains identifiers assigned to each program category, duplicates

```

# Define index
primary_substance_group_index <- c(
  "Alcohol" = "A",
  "All Others" = "AO",
  "Cocaine incl Crack" = "CC",
  "Heroin" = "H",
  "Marijuana incl Hashish" = "MH",
  "Other Opioids" = "OO"
)

# Add substance_code column
admissions_df <- admissions_df %>%
  mutate(substance_code = recode(primary_substance_group,
    "Alcohol" = "A",
    "All Others" = "AO",
    "Cocaine incl Crack" = "CC",
    "Heroin" = "H",
    "Marijuana incl Hashish" = "MH",
    "Other Opioids" = "OO"))

# Create substance dataframe based on index
primary_substance_group <- admissions_df %>%
  distinct(primary_substance_group, .keep_all = TRUE) %>%
  select(substance_code, primary_substance_group)

primary_substance_group

```

```

## # A tibble: 6 x 2

```



```
## substance_code primary_substance_group
## <fct>          <fct>
## 1 A            Alcohol
## 2 AO           All Others
## 3 CC           Cocaine incl Crack
## 4 H            Heroin
## 5 OO           Other Opioids
## 6 MH           Marijuana incl Hashish
```

```
# Join county_code by county name
admissions_df <- admissions_df %>%
  full_join(county, by = "county_of_program_location")

admissions_df <- admissions_df %>%
  select(
    year,
    county_code,
    program_code,
    service_type,
    age_group,
    primary_substance_group,
    substance_code,
    admissions
  )

str(admissions_df)
```

```
## tibble [72,463 x 8] (S3: tbl_df/tbl/data.frame)
## $ year          : num [1:72463] 2017 2017 2017 2017 2017 ...
## $ county_code   : chr [1:72463] "AL" "AL" "AL" "AL" ...
## $ program_code  : Factor w/ 5 levels "C","I","OTP",...: 1 1 1 1 1 1 1 1 1 ...
## $ service_type  : Factor w/ 28 levels "Community Residential",...: 8 8 8 8 8 8 8 8 8 ...
## $ age_group     : Factor w/ 6 levels "18 thru 24","25 thru 34",...: 1 1 1 1 1 2 2 4 4 ...
## $ primary_substance_group: Factor w/ 6 levels "Alcohol","All Others",...: 1 2 3 4 6 1 2 2 3 4 ...
## $ substance_code : Factor w/ 6 levels "A","AO","CC",...: 1 2 3 4 6 1 2 2 3 4 ...
## $ admissions    : num [1:72463] 25 7 1 64 20 140 10 3 1 41 ...
```

```
# aggregate() to sum the total admissions for every year
annual_admissions <- function() {
  # Store aggregated data in separate tibble
  total_admissions <- aggregate(admissions_df$admissions,
    by = list(year = admissions_df$year),
    sum) %>%
    rename(total = x)
  # Maximum point
  max_point <- total_admissions[which.max(total_admissions$total), ]
  # Line graph
  ggplot(total_admissions, aes(year, total)) +
    geom_line() +
    geom_point() +
    scale_x_continuous(breaks = 2007:2017) +
    scale_y_continuous(limits = c(270000, 320000)) +
    labs(
```

```

    title = "OASAS Annual Admissions",
    x = "Year",
    y = "Total Admissions") +
theme_minimal() +
theme(
  plot.title = element_text(hjust = 0.5, size = 14, face = "bold",
                             margin = margin(t = 5, r = 10, l = 10, b = 10)),
  axis.title.x = element_text(size = 12, margin = margin(t = 10, b = 5)),
  axis.title.y = element_text(size = 12, margin = margin(r = 10, l = 5)),
  margin = margin(r = 10)
) +
  annotate("text",
    x = max_point$year,
    y = max_point$total,
    label = paste("Top Admissions:", max_point$total, " in ", max_point$year),
    vjust = -1)
}

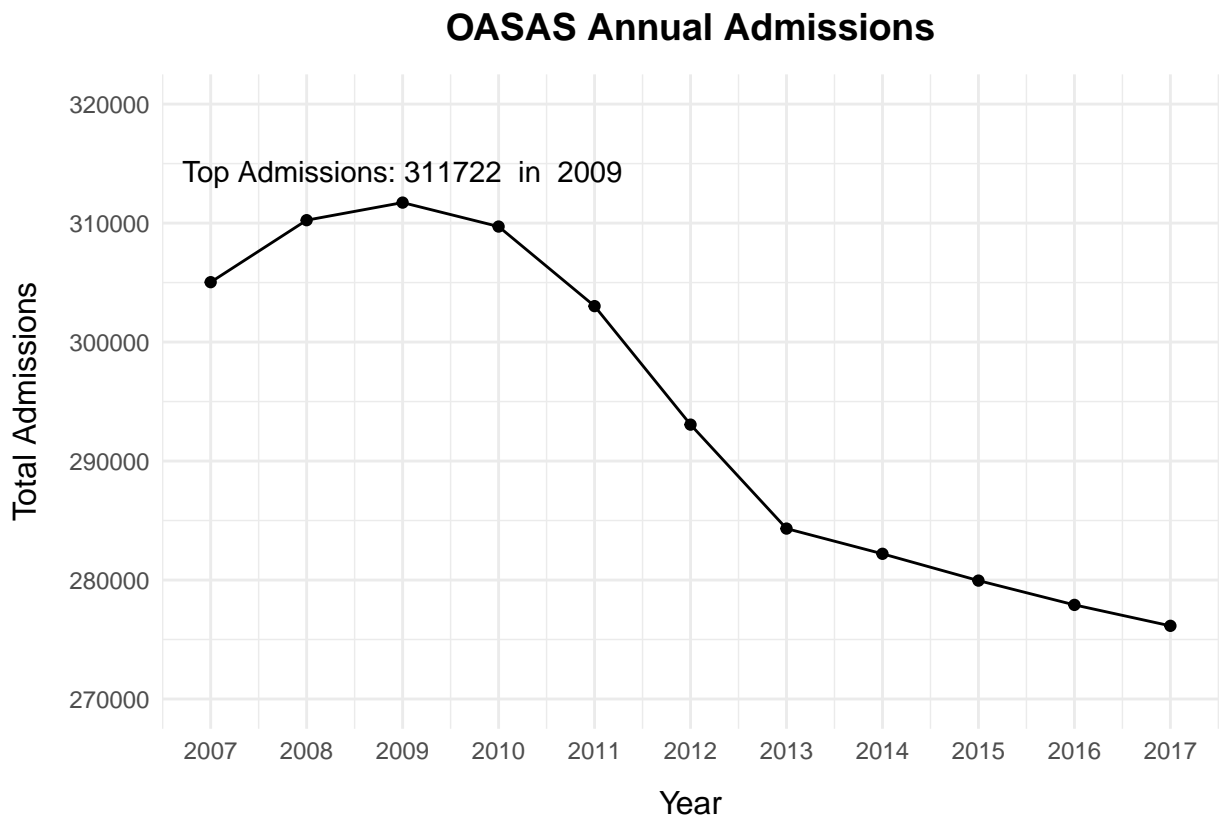
annual_admissions()

```

```

## Warning in plot_theme(plot): The 'margin' theme element is not defined in the
## element hierarchy.

```



- Admissions peak in 2009 with 311,722 patients

- A significant downward trend is observed from 2010 to 2013
- Admissions continue to gradually decline from 2013 onward

Data Evaluation

Admissions by County

```
# Total number of admission in NYS
total_admissions = sum(admissions_df$admissions)

# Calculate percentage of admissions in each county
admissions_by_county <- admissions_df %>%
  select(county_code, admissions) %>%
  group_by(county_code) %>%
  summarize(percentage = ((sum(admissions) / total_admissions * 100))) %>%
  arrange(desc(percentage))

admissions_by_county
```

```
## # A tibble: 61 x 2
##   county_code percentage
##   <chr>          <dbl>
## 1 NYC           15.3
## 2 QU-NYC        8.42
## 3 BR-NYC        8.03
## 4 KI-NYC        7.64
## 5 SU            7.57
## 6 ER            5.45
## 7 WE            5.08
## 8 MO            4.53
## 9 NS            3.66
## 10 OD           3.51
## # i 51 more rows
```

```
top_admissions <- admissions_by_county %>%
  slice_max(n = 5, percentage)

top_admissions
```

```
## # A tibble: 5 x 2
##   county_code percentage
##   <chr>          <dbl>
## 1 NYC           15.3
## 2 QU-NYC        8.42
## 3 BR-NYC        8.03
## 4 KI-NYC        7.64
## 5 SU            7.57
```

Rehab Facility Identification

```
rehab_df <- admissions_df %>%  
  # Only show rehabilitation services  
  filter(str_detect(service_type, regex("Rehab|Rehabilitation", ignore_case = TRUE))) %>%  
  select(service_type, age_group, primary_substance_group, admissions)  
rehab_df
```

```
## # A tibble: 17,319 x 4  
##   service_type      age_group primary_substance_group admissions  
##   <fct>          <fct>      <fct>                <dbl>  
## 1 Inpatient Rehabilitation 18 thru 24 Alcohol                11  
## 2 Inpatient Rehabilitation 18 thru 24 All Others                2  
## 3 Inpatient Rehabilitation 18 thru 24 Cocaine incl Crack        4  
## 4 Inpatient Rehabilitation 18 thru 24 Heroin                 21  
## 5 Inpatient Rehabilitation 18 thru 24 Marijuana incl Hashish    6  
## 6 Inpatient Rehabilitation 18 thru 24 Other Opioids            5  
## 7 Inpatient Rehabilitation 25 thru 34 Alcohol                49  
## 8 Inpatient Rehabilitation 25 thru 34 All Others                7  
## 9 Inpatient Rehabilitation 25 thru 34 Cocaine incl Crack       31  
## 10 Inpatient Rehabilitation 25 thru 34 Heroin                 101  
## # i 17,309 more rows
```

```
top_substance_df <- rehab_df %>%  
  # Only interested in these combinations  
  group_by(service_type, age_group, primary_substance_group) %>%  
  # Take count to show how many admissions exist for each substance in each age group  
  summarize(substance_count = sum(admissions)) %>%  
  # Limit to age group  
  group_by(age_group) %>%  
  # Filter for the substances with the highest count  
  filter(substance_count == max(substance_count)) %>%  
  # Show relevant columns  
  select(service_type, age_group, primary_substance_group, substance_count)
```

```
## 'summarise()' has grouped output by 'service_type', 'age_group'. You can  
## override using the '.groups' argument.
```

```
top_substance_df
```

```
## # A tibble: 6 x 4  
## # Groups:   age_group [6]  
##   service_type      age_group primary_substance_group substance_count  
##   <fct>          <fct>      <fct>                <dbl>  
## 1 Inpatient Rehabilitation 18 thru 24 Heroin                22705  
## 2 Inpatient Rehabilitation 25 thru 34 Heroin                37753  
## 3 Inpatient Rehabilitation 35 thru 44 Alcohol                50698  
## 4 Inpatient Rehabilitation 45 thru 54 Alcohol                69590  
## 5 Inpatient Rehabilitation 55 and Older Alcohol                30051  
## 6 Res Rehab for Youth      Under 18      Marijuana incl Hashish    10643
```

- To identify the most prominent substance used in each age group, we first define a regular expression in a new dataframe that filters all services containing “Rehab” or “Rehabilitation” in the name
- top_substance_df is created to find the top substance per age group
 - The data is grouped by age_group and primary_substance_group because we are only interested in analyses in the context of these variables paired together

Admissions by Substance and Age Group

```
all_substance_df <- rehab_df %>%
  group_by(service_type, age_group, primary_substance_group) %>%
  # Take count to show how many admissions exist for each substance in each age group
  summarize(substance_count = sum(admissions)) %>%
  # Limit to age group
  group_by(age_group) %>%
  # Filter for the substances with the highest count
  mutate(substance_count == max(substance_count)) %>%
  # Show relevant columns
  select(service_type, age_group, primary_substance_group, substance_count)
```

‘summarise()’ has grouped output by ‘service_type’, ‘age_group’. You can
override using the ‘.groups’ argument.

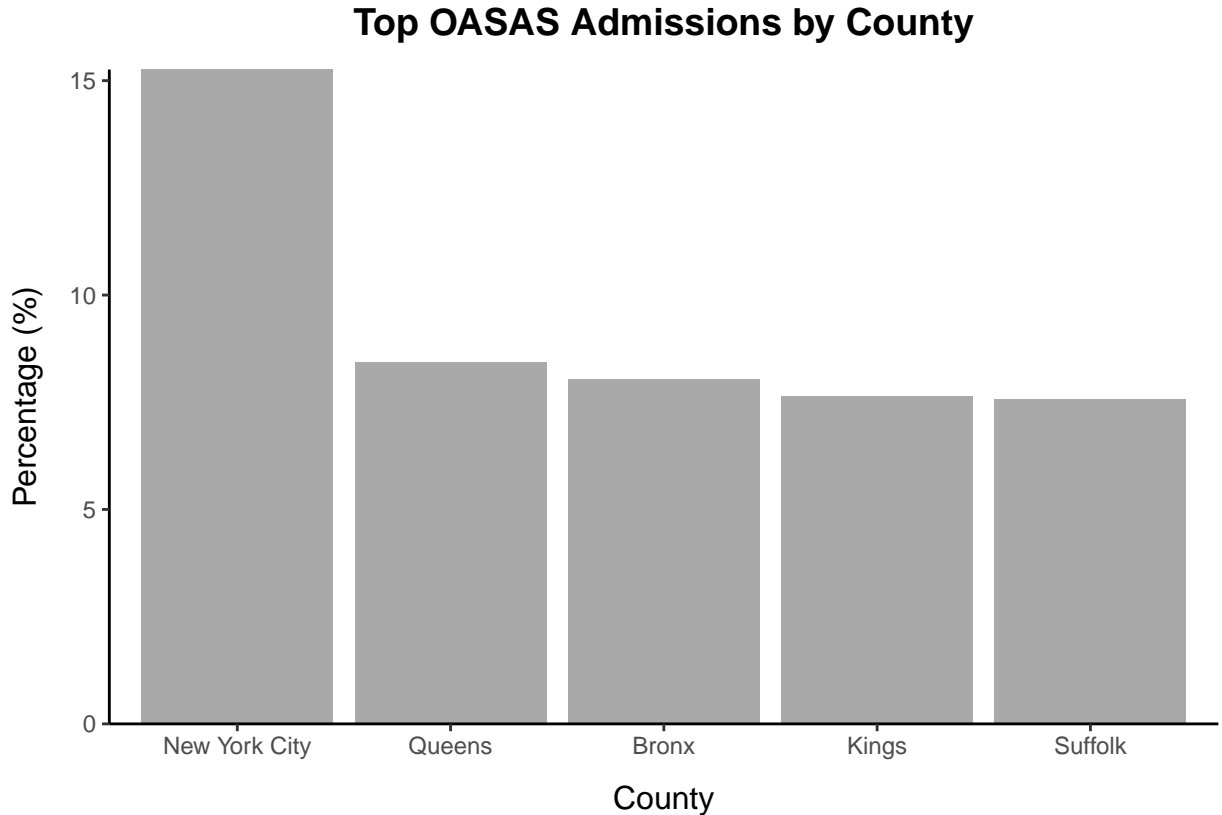
```
all_substance_df
```

```
## # A tibble: 236 x 4
## # Groups:   age_group [6]
##   service_type      age_group primary_substance_group substance_count
##   <fct>          <fct>      <fct>                  <dbl>
## 1 Inpatient Rehabilitation 18 thru 24 Alcohol                10949
## 2 Inpatient Rehabilitation 18 thru 24 All Others                3234
## 3 Inpatient Rehabilitation 18 thru 24 Cocaine incl Crack        4583
## 4 Inpatient Rehabilitation 18 thru 24 Heroin                 22705
## 5 Inpatient Rehabilitation 18 thru 24 Marijuana incl Hashish    10209
## 6 Inpatient Rehabilitation 18 thru 24 Other Opioids             8718
## 7 Inpatient Rehabilitation 25 thru 34 Alcohol                32121
## 8 Inpatient Rehabilitation 25 thru 34 All Others                5640
## 9 Inpatient Rehabilitation 25 thru 34 Cocaine incl Crack       14559
## 10 Inpatient Rehabilitation 25 thru 34 Heroin                 37753
## # i 226 more rows
```

Data Visualization

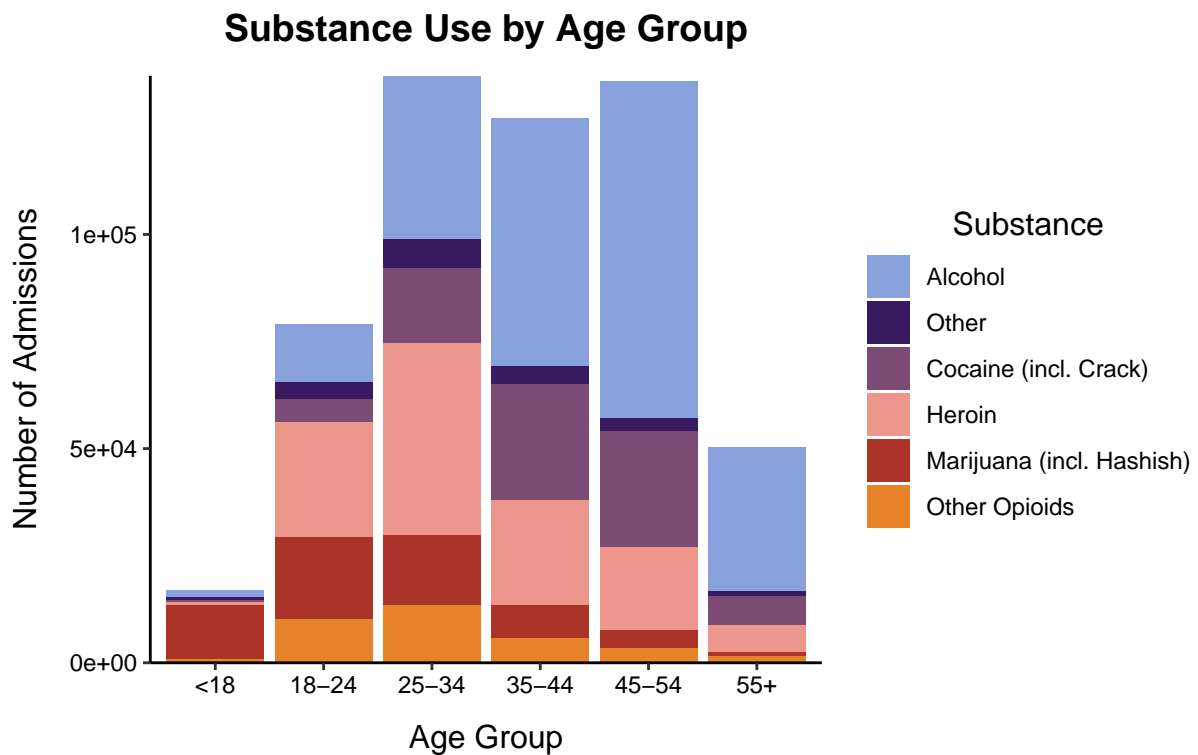
Top Admissions by County

```
ggplot(top_admissions, aes(reorder(county_code, -percentage), percentage)) +  
  geom_col(fill = "darkgray") +  
  labs(x = "County",  
       y = "Percentage (%)",  
       title = "Top OASAS Admissions by County") +  
  scale_y_continuous(expand = c(0,0)) +  
  scale_x_discrete(labels = c("NYC" = "New York City",  
                              "QU-NYC" = "Queens",  
                              "BR-NYC" = "Bronx",  
                              "KI-NYC" = "Kings",  
                              "SU" = "Suffolk")) +  
  
  theme_classic() +  
  theme(  
    plot.title = element_text(size = 14, face = "bold", hjust = 0.5,  
                              margin = margin(t = 5, b = 10)),  
    axis.title.x = element_text(size = 12, color = "black",  
                                margin = margin(t = 10, b = 5)),  
    axis.title.y = element_text(size = 12, color = "black",  
                                margin = margin(r = 10, l = 5))  
  )
```



Substance Use by Age Group

```
ggplot(all_substance_df, aes(factor(age_group,
                                   levels = c("Under 18", "18 thru 24",
                                               "25 thru 34", "35 thru 44",
                                               "45 thru 54", "55 and Older")),
                             substance_count,
                             fill = primary_substance_group)) +
  geom_bar(stat = "identity") +
  labs(
    x = "Age Group",
    y = "Number of Admissions",
    title = "Substance Use by Age Group",
    caption = "Stacked bar showing substance use by age group from 2007-2017.",
    legend = "Substance"
  ) +
  scale_y_continuous(expand = c(0,0)) +
  scale_fill_paletteer_d("MetBrewer::Archambault",
    name = "Substance",
    labels = c(
      "All Others" = "Other",
      "Cocaine incl Crack" = "Cocaine (incl. Crack)",
      "Marijuana incl Hashish" = "Marijuana (incl. Hashish)"
    )
  ) +
  scale_x_discrete(
    labels = c(
      "Under 18" = "<18",
      "18 thru 24" = "18-24",
      "25 thru 34" = "25-34",
      "35 thru 44" = "35-44",
      "45 thru 54" = "45-54",
      "55 and Older" = "55+"
    )
  ) +
  theme_classic() +
  theme(
    axis.title.x = element_text(size = 12, color = "black",
                                margin = margin(t = 10, b = 5)),
    axis.title.y = element_text(size = 12, color = "black",
                                margin = margin(r = 10, l = 10)),
    axis.text = element_text(color = "black"),
    legend.title = element_text(size = 12, color = "black", hjust = 0.5),
    plot.title = element_text(size = 14, face = "bold",
                              margin = margin(b = 10, t = 10), hjust = 0.5),
    plot.caption = element_text(size = 11, hjust = 0.5,
                                margin = margin(b = 5, t = 5))
  )
```



Stacked bar showing substance use by age group from 2007–2017.

Service Use by Age Group

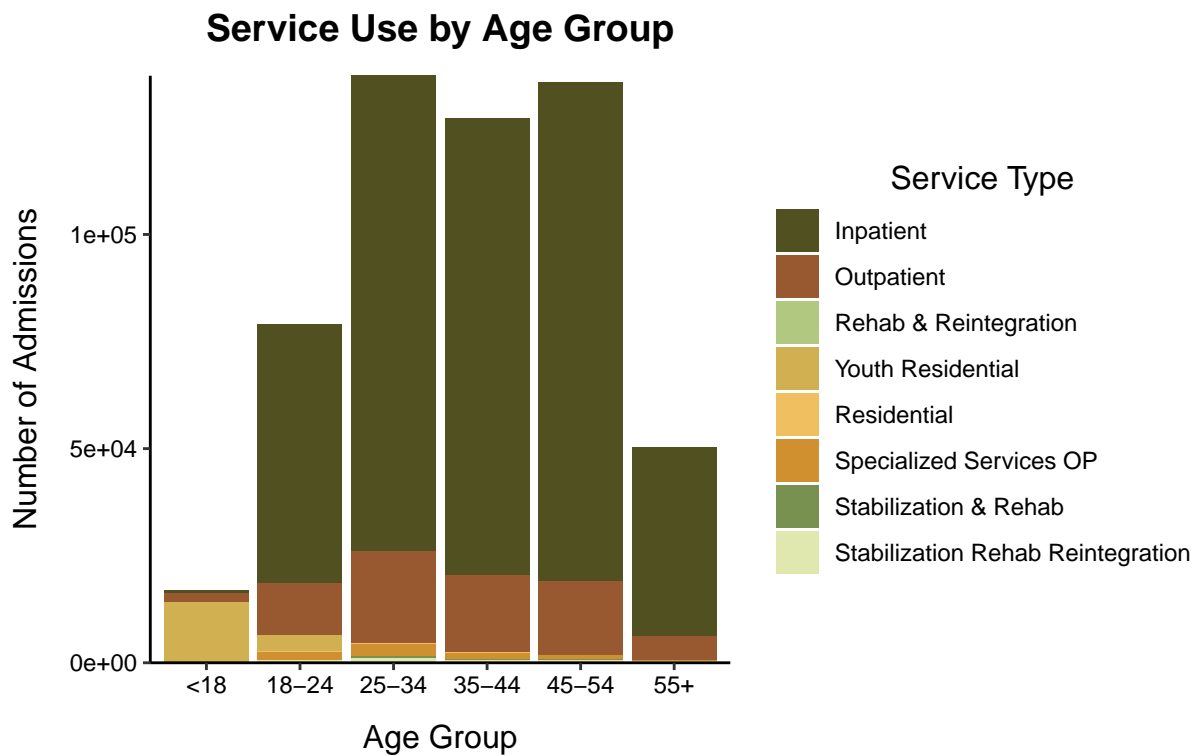
```
ggplot(all_substance_df, aes(factor(age_group, levels = c("Under 18", "18 thru 24",
                                                         "25 thru 34", "35 thru 44",
                                                         "45 thru 54", "55 and Older")),
                             substance_count,
                             fill = service_type)) +
  geom_bar(stat = "identity") +
  scale_y_continuous(expand = c(0, 0)) +
  labs(
    x = "Age Group",
    y = "Number of Admissions",
    title = "Service Use by Age Group",
    caption = "Stacked bar showing service use by age group from 2007–2017."
  ) +
  scale_fill_paletteer_d("palettetown::seedot",
    name = "Service Type",
    labels = c(
      "Inpatient Rehabilitation" = "Inpatient",
      "Outpatient Rehabilitation" = "Outpatient",
      "Rehab and Reintegration" = "Rehab & Reintegration",
      "Res Rehab for Youth" = "Youth Residential",
      "Residential Rehabilitation" = "Residential",
      "Specialized Services OP Rehab" = "Specialized Services OP",
    )
  )
```



```

    "Stabilization and Rehab" = "Stabilization & Rehab"
  )
) +
scale_x_discrete(
  labels = c(
    "18 thru 24" = "18-24",
    "25 thru 34" = "25-34",
    "35 thru 44" = "35-44",
    "45 thru 54" = "45-54",
    "55 and Older" = "55+",
    "Under 18" = "<18"
  )
) +
theme_classic() +
theme(
  axis.title.x = element_text(size = 12, color = "black",
                              margin = margin(t = 10, b = 5)),
  axis.title.y = element_text(size = 12, color = "black",
                              margin = margin(r = 10, l = 10)),
  axis.text = element_text(color = "black"),
  legend.title = element_text(size = 12, color = "black", hjust = 0.5),
  plot.title = element_text(size = 14, face = "bold",
                             margin = margin(b = 10, t = 10), hjust = 0.5),
  plot.caption = element_text(size = 11, hjust = 0.5,
                              margin = margin(b = 5, t = 5))
)

```



Stacked bar showing service use by age group from 2007–2017.

Top Substance Use by Age Group

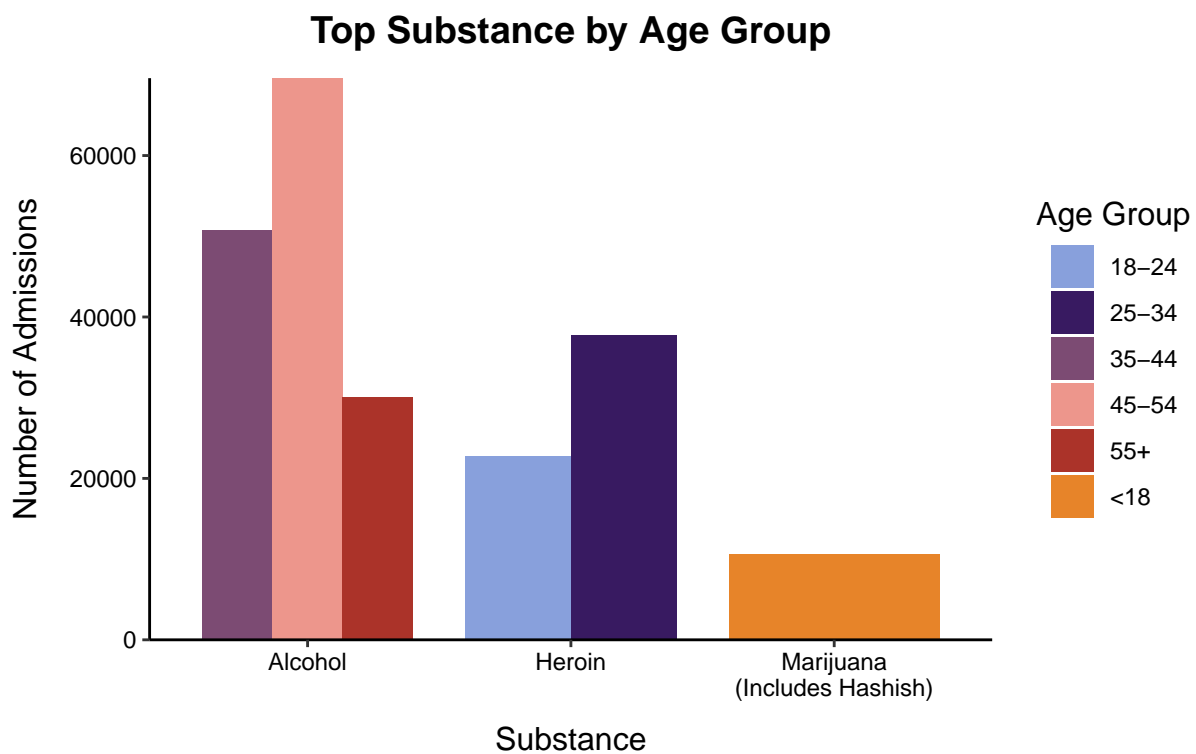
```
ggplot(top_substance_df, aes(primary_substance_group, substance_count,
                             fill = age_group)) +
  geom_bar(stat = "identity", position = "dodge", width = 0.8) +
  scale_y_continuous(expand = c(0,0)) +
  labs(
    y = "Number of Admissions",
    title = "Top Substance by Age Group",
    caption = "Stacked bar showing the top substance used per age group from 2007–2017."
  ) +
  scale_x_discrete(
    name = "Substance",
    labels = c(
      "All Others" = "Other",
      "Cocaine incl Crack" = "Cocaine (Includes Crack)",
      "Marijuana incl Hashish" = "Marijuana\n(Includes Hashish)"
    )
  ) +
  scale_fill_paletteer_d("MetBrewer::Archambault",
    name = "Age Group",
    labels = c(
      "18 thru 24" = "18-24",
      "25 thru 34" = "25-34",

```

```

    "35 thru 44" = "35-44",
    "45 thru 54" = "45-54",
    "55 and Older" = "55+",
    "Under 18" = "<18"
  )
) +
theme_classic() +
theme(
  axis.title.x = element_text(size = 12, color = "black",
                              margin = margin(t = 10, b = 5)),
  axis.title.y = element_text(size = 12, color = "black",
                              margin = margin(r = 10, l = 10)),
  axis.text = element_text(color = "black"),
  legend.title = element_text(size = 12, color = "black", hjust = 0.5),
  plot.title = element_text(size = 14, face = "bold",
                             margin = margin(b = 10, t = 10), hjust = 0.5),
  plot.caption = element_text(size = 11, hjust = 0.5,
                              margin = margin(b = 5, t = 5))
)

```



Stacked bar showing the top substance used per age group from 2007–2017.