

Final Report

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```
install.packages("taRifx")
install.packages("fastDummies")
library(tidyverse)
library(dplyr)
library(taRifx)
library(fastDummies)
library(infer)
library(parsnip)
```

```
drug <- readr::read_csv("Drug_Consumption.csv")
```

Abstract:

Background and Significance:

Methods: a) Data Collection and Variables

b) Exploratory Data Analysis

```
drug1 <- drug %>%
  mutate(across(Alcohol:VSA,destring))
```

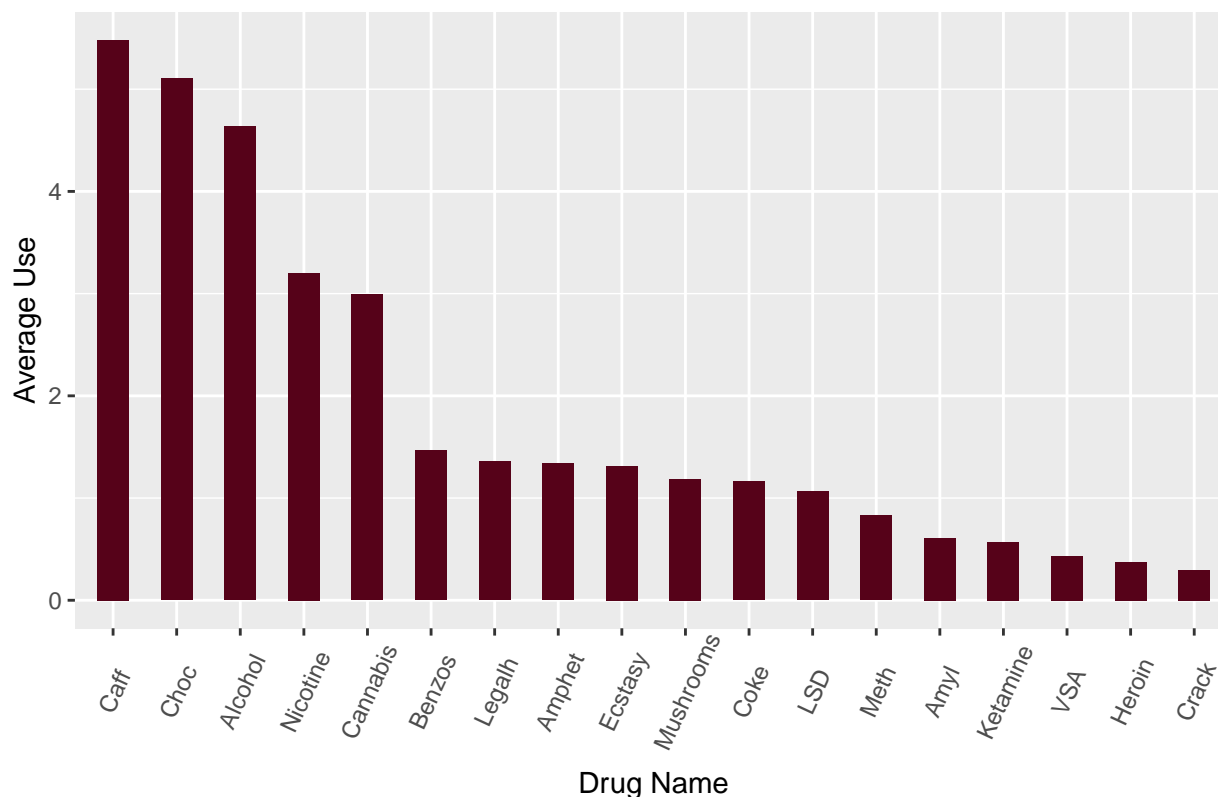
```
drug1[,14:32] <- sapply(drug1[,14:32], as.numeric)
drugmeans <- colMeans(drug1[, 14:32])
```

```
drug_name <- c('Alcohol', 'Amphet', 'Amyl', 'Benzos', 'Caff', 'Cannabis', 'Choc', 'Coke', 'Crack', 'Ecs')
average_use <- c(4.63481953, 1.34023355, 0.60721868, 1.46496815, 5.48354565, 2.99097665, 5.10668790, 1.34023355,
```

```
drug_averages <- data.frame(drug_name, average_use)
```

```
drug_averages$drug_name <- factor(drug_averages$drug_name,
                                  levels = drug_averages$drug_name[order(drug_averages$average_use, decreasing = TRUE)])
ggplot(drug_averages, aes(x=drug_name, y=average_use)) +
  geom_bar(stat="identity", width=.5, fill="#560219") +
  labs(title="Average Drug Use by Drug",
       x = "Drug Name",
       y = "Average Use") +
  theme(axis.text.x = element_text(angle=65, vjust=0.6))
```

Average Drug Use by Drug



```
numdrug <- drug1 %>%
  mutate(Age = replace(Age, Age == "18-24", 0), Age = replace(Age, Age == "25-34", 1), Age = replace(Age, Age == "35-44", 2), Age = replace(Age, Age == "45-54", 3), Age = replace(Age, Age == "55-64", 4), Age = replace(Age, Age == "65+", 5))

numdrug <- mutate_all(numdrug, function(x) as.numeric(as.character(x)))

numdrug2 = select(numdrug, -Alcohol, -Amphet, -Amyl, -Benzos, -Caff, -Cannabis, -Choc, -Coke, -Crack, -Ecstasy, -Heroin, -Ketamine, -LSD, -Meth, -Mushrooms, -VSA)

head(numdrug2)
```

```
## # A tibble: 6 x 14
##   ID   Age Gender Education Country Ethnicity Nscore Escore Oscore AScore
##   <dbl> <dbl> <dbl>   <dbl>   <dbl>   <dbl>   <dbl> <dbl>   <dbl> <dbl>
## 1     2     1     1       8       5       6 -0.678  1.94   1.44  0.761
## 2     3     2     1       5       5       6 -0.467  0.805 -0.847 -1.62
## 3     4     0     0       7       5       6 -0.149 -0.806 -0.0193 0.590
## 4     5     2     0       8       5       6  0.735 -1.63  -0.452 -0.302
## 5     6     5     0       3       1       6 -0.678 -0.300 -1.56   2.04
## 6     7     3     1       7       6       6 -0.467 -1.09  -0.452 -0.302
## # ... with 4 more variables: Cscore <dbl>, Impulsive <dbl>, SS <dbl>,
## #   Mushrooms <dbl>
```

```
correlation_matrix <- round(cor(numdrug2),2)
head(correlation_matrix)
```

```
##           ID   Age Gender Education Country Ethnicity Nscore Escore Oscore
## ID       1.00 -0.27  0.02   -0.01   0.10     0.01   0.02  -0.05   0.17
## Age      -0.27  1.00 -0.10    0.10  -0.06    0.04  -0.14  -0.03  -0.22
```

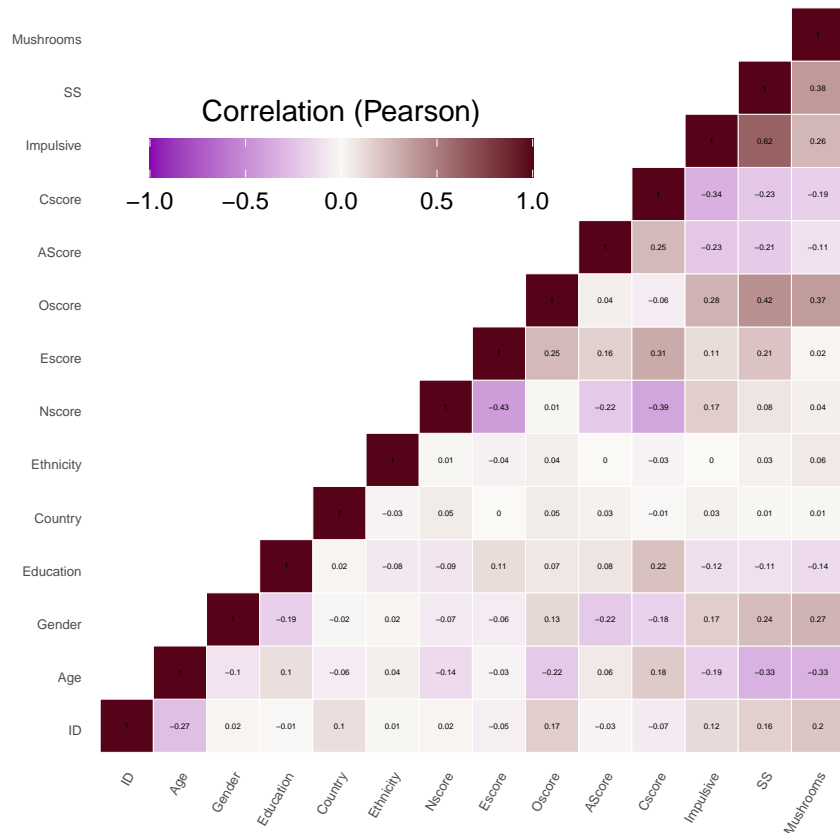
```
## Gender      0.02 -0.10  1.00      -0.19  -0.02      0.02 -0.07 -0.06  0.13
## Education -0.01  0.10 -0.19      1.00   0.02     -0.08 -0.09  0.11  0.07
## Country    0.10 -0.06 -0.02      0.02   1.00     -0.03  0.05  0.00  0.05
## Ethnicity  0.01  0.04  0.02     -0.08  -0.03      1.00  0.01 -0.04  0.04
##           AScore Cscore Impulsive  SS Mushrooms
## ID          -0.03 -0.07      0.12  0.16      0.20
## Age          0.06  0.18     -0.19 -0.33     -0.33
## Gender      -0.22 -0.18      0.17  0.24      0.27
## Education    0.08  0.22     -0.12 -0.11     -0.14
## Country      0.03 -0.01      0.03  0.01      0.01
## Ethnicity    0.00 -0.03      0.00  0.03      0.06
```

```
get_upper_tri<-function(correlation_matrix){
  correlation_matrix[lower.tri(correlation_matrix)] <- NA
  return(correlation_matrix)
}
upper_tri <- get_upper_tri(correlation_matrix)
library(reshape2)
```

```
##
## Attaching package: 'reshape2'

## The following object is masked from 'package:tidyr':
##
## smiths
```

```
melted_cormat <- melt(upper_tri, na.rm = TRUE)
library(ggplot2)
ggplot(data = melted_cormat, aes(Var2, Var1, fill = value))+
  geom_tile(color = "white")+
  scale_fill_gradient2(low = "#8a02b2", high = "#560219", mid = "#FAF9F6",
    midpoint = 0, limit = c(-1,1), space = "Lab",
    name="Correlation (Pearson)") +
  theme_minimal()+
  theme(axis.text.x = element_text(angle = 60, vjust = 1,
    size = 5, hjust = 1), axis.text.y = element_text(vjust = 1, size = 5, hjust = 1))+
  coord_fixed() +
  geom_text(aes(Var2, Var1, label = value), color = "black", size = 1) +
  theme(
    axis.title.x = element_blank(),
    axis.title.y = element_blank(),
    panel.grid.major = element_blank(),
    panel.border = element_blank(),
    panel.background = element_blank(),
    axis.ticks = element_blank(),
    legend.justification = c(1, 0),
    legend.position = c(0.6, 0.7),
    legend.direction = "horizontal",
    legend.key.size = unit(0.5, 'cm'))+
  guides(fill = guide_colorbar(barwidth = 10, barheight = 1,
    title.position = "top", title.hjust = 0.5))
```



```
drug_clean <- numdrug %>%
  mutate(Alcohol_User = as.factor(ifelse(Alcohol > 1, "yes", "no")),
         Amphetamine_User = as.factor(ifelse(Amphet > 1, "yes", "no")),
         AmylNitrite_User = as.factor(ifelse(Amyl > 1, "yes", "no")),
         Benzos_User = as.factor(ifelse(Benzos > 1, "yes", "no")),
         Caffeine_User = as.factor(ifelse(Caff > 1, "yes", "no")),
         Cannabis_User = as.factor(ifelse(Cannabis > 1, "yes", "no")),
         Chocolate_User = as.factor(ifelse(Choc > 1, "yes", "no")),
         Cocaine_User = as.factor(ifelse(Coke > 1, "yes", "no")),
         Crack_User = as.factor(ifelse(Crack > 1, "yes", "no")),
         Ecstasy_User = as.factor(ifelse(Ecstasy > 1, "yes", "no")),
         Heroin_User = as.factor(ifelse(Heroin > 1, "yes", "no")),
         Ketamine_User = as.factor(ifelse(Ketamine > 1, "yes", "no")),
         LegalHighs_User = as.factor(ifelse(Legalh > 1, "yes", "no")),
         LSD_User = as.factor(ifelse(LSD > 1, "yes", "no")),
         Meth_User = as.factor(ifelse(Meth > 1, "yes", "no")),
         Mushrooms_User = as.factor(ifelse(Mushrooms > 1, "yes", "no")),
         Nicotine_User = as.factor(ifelse(Nicotine > 1, "yes", "no")),
         Semeron_User = as.factor(ifelse(Semer > 1, "yes", "no")),
         VSA_User = as.factor(ifelse(VSA > 1, "yes", "no")))

drug_clean_2 <- drug1 %>%
  mutate(Alcohol_User = as.factor(ifelse(Alcohol > 1, "yes", "no")),
         Amphetamine_User = as.factor(ifelse(Amphet > 1, "yes", "no")),
         AmylNitrite_User = as.factor(ifelse(Amyl > 1, "yes", "no")),
         Benzos_User = as.factor(ifelse(Benzos > 1, "yes", "no")),
```

```

Caffeine_User = as.factor(ifelse(Caff > 1, "yes", "no")),
Cannabis_User = as.factor(ifelse(Cannabis > 1, "yes", "no")),
Chocolate_User = as.factor(ifelse(Choc > 1, "yes", "no")),
Cocaine_User = as.factor(ifelse(Coke > 1, "yes", "no")),
Crack_User = as.factor(ifelse(Crack > 1, "yes", "no")),
Ecstasy_User = as.factor(ifelse(Ecstasy > 1, "yes", "no")),
Heroin_User = as.factor(ifelse(Heroin > 1, "yes", "no")),
Ketamine_User = as.factor(ifelse(Ketamine > 1, "yes", "no")),
LegalHighs_User = as.factor(ifelse(Legalh > 1, "yes", "no")),
LSD_User = as.factor(ifelse(LSD > 1, "yes", "no")),
Meth_User = as.factor(ifelse(Meth > 1, "yes", "no")),
Mushrooms_User = as.factor(ifelse(Mushrooms > 1, "yes", "no")),
Nicotine_User = as.factor(ifelse(Nicotine > 1, "yes", "no")),
Semeron_User = as.factor(ifelse(Semer > 1, "yes", "no")),
VSA_User = as.factor(ifelse(VSA > 1, "yes", "no"))

```

```

drug_byuse <- numdrug %>%
  mutate(Alcohol_User = ifelse(Alcohol > 1, 1, 0),
    Amphetamine_User = ifelse(Amphet > 1, 1, 0),
    AmylNitrite_User = ifelse(Amyl > 1, 1, 0),
    Benzos_User = ifelse(Benzos > 1, 1, 0),
    Caffeine_User = ifelse(Caff > 1, 1, 0),
    Cannabis_User = ifelse(Cannabis > 1, 1, 0),
    Chocolate_User = ifelse(Choc > 1, 1, 0),
    Cocaine_User = ifelse(Coke > 1, 1, 0),
    Crack_User = ifelse(Crack > 1, 1, 0),
    Ecstasy_User = ifelse(Ecstasy > 1, 1, 0),
    Heroin_User = ifelse(Heroin > 1, 1, 0),
    Ketamine_User = ifelse(Ketamine > 1, 1, 0),
    LegalHighs_User = ifelse(Legalh > 1, 1, 0),
    LSD_User = ifelse(LSD > 1, 1, 0),
    Meth_User = ifelse(Meth > 1, 1, 0),
    Mushrooms_User = ifelse(Mushrooms > 1, 1, 0),
    Nicotine_User = ifelse(Nicotine > 1, 1, 0),
    Semeron_User = ifelse(Semer > 1, 1, 0),
    VSA_User = ifelse(VSA > 1, 1, 0)) %>%
  dplyr::select(Alcohol_User, Amphetamine_User, AmylNitrite_User, Benzos_User,
    Caffeine_User, Cannabis_User, Chocolate_User, Cocaine_User,
    Crack_User, Ecstasy_User, Heroin_User, Ketamine_User,
    LegalHighs_User, LSD_User, Meth_User, Mushrooms_User,
    Nicotine_User, Semeron_User, VSA_User)

data.frame(yes = colSums(drug_byuse), no = 1885 - colSums(drug_byuse),
  drug = colnames(drug_byuse)) %>%
  summarise(diff = abs(yes - no), drug = drug) %>%
  arrange(diff) %>%
  head()

```

```

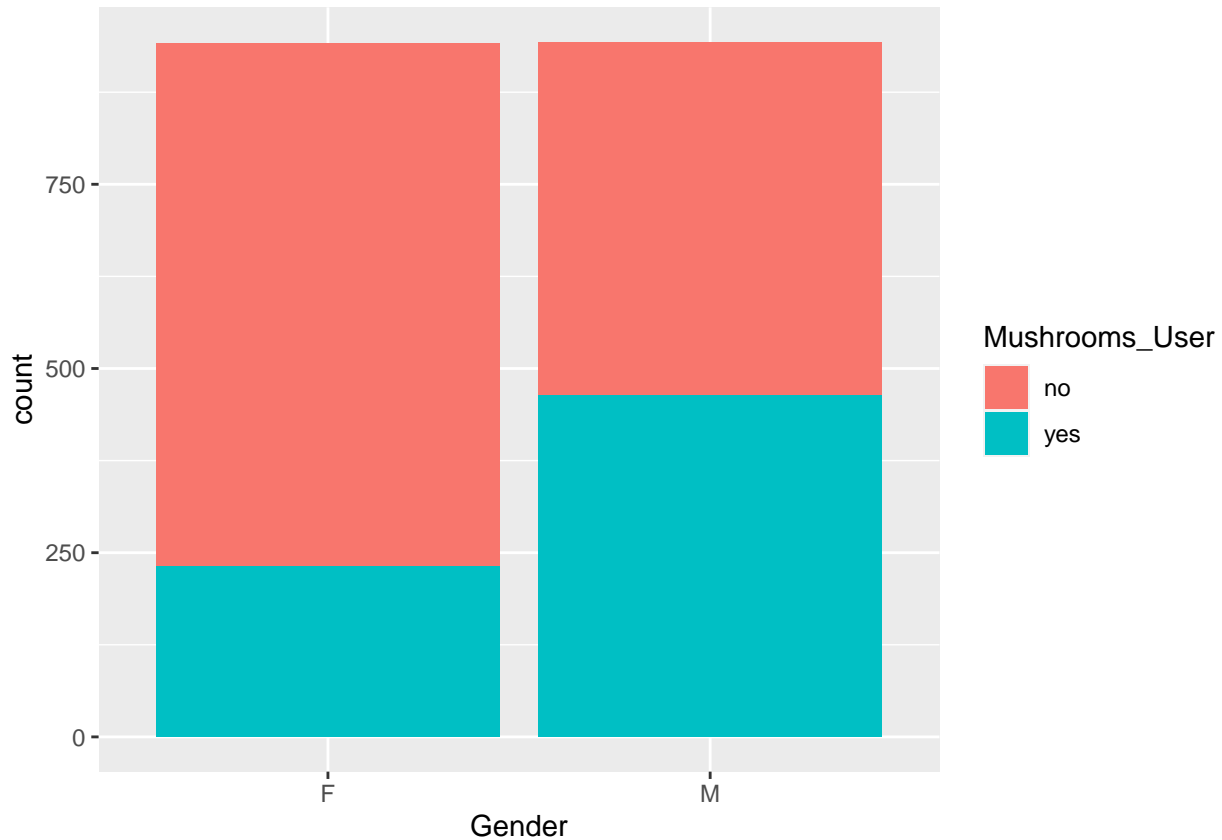
##   diff      drug
## 1  349  Benzos_User
## 2  361 LegalHighs_User
## 3  383  Ecstasy_User
## 4  497  Mushrooms_User

```

```
## 5  511    Cocaine_User
## 6  529 Amphetamine_User
```

c) Analytical Methods

```
drug_clean_2 %>%
  ggplot(aes(x = Gender,
             fill = Mushrooms_User)) +
  geom_bar()
```



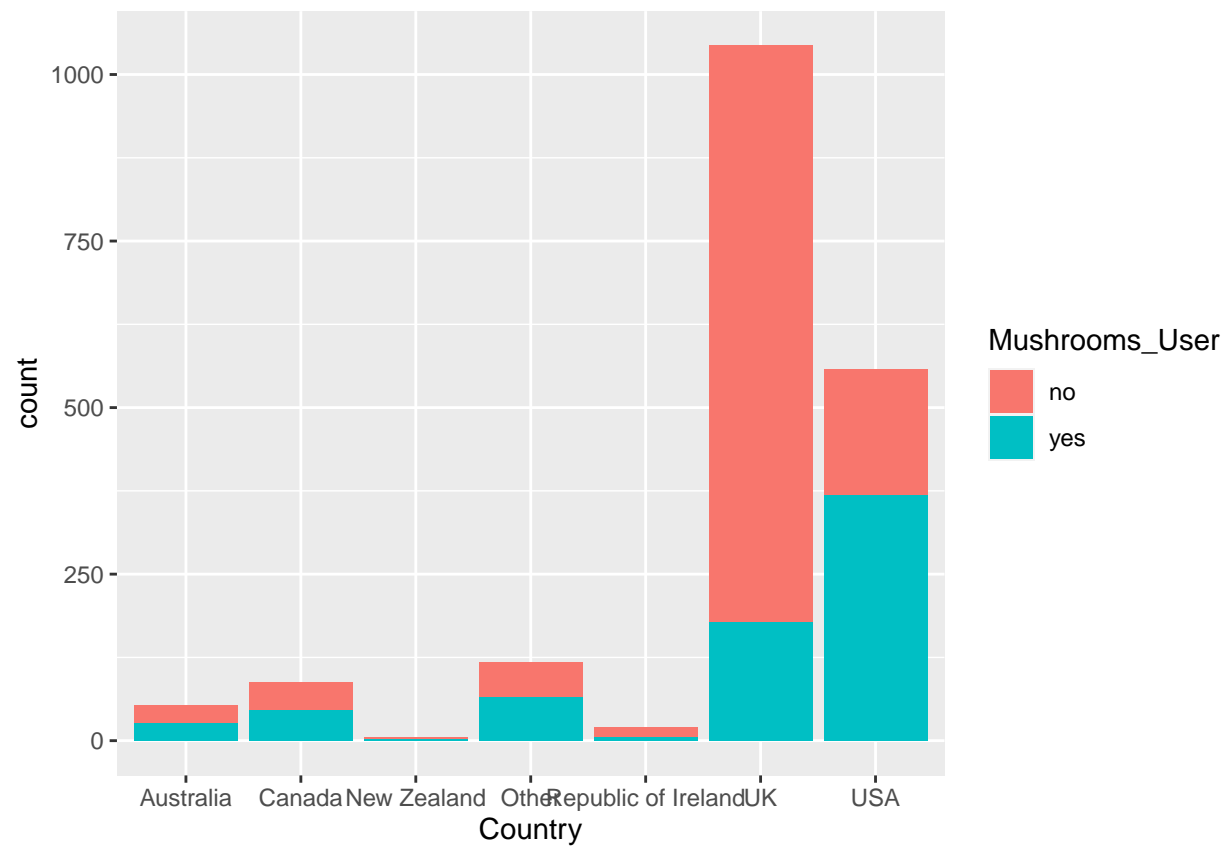
```
fisher.test(drug_clean_2$Gender, drug_clean_2$Mushrooms_User)
```

```
##
## Fisher's Exact Test for Count Data
##
## data: drug_clean_2$Gender and drug_clean_2$Mushrooms_User
## p-value < 2.2e-16
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
##  2.426426 3.624594
## sample estimates:
## odds ratio
##  2.962888
```

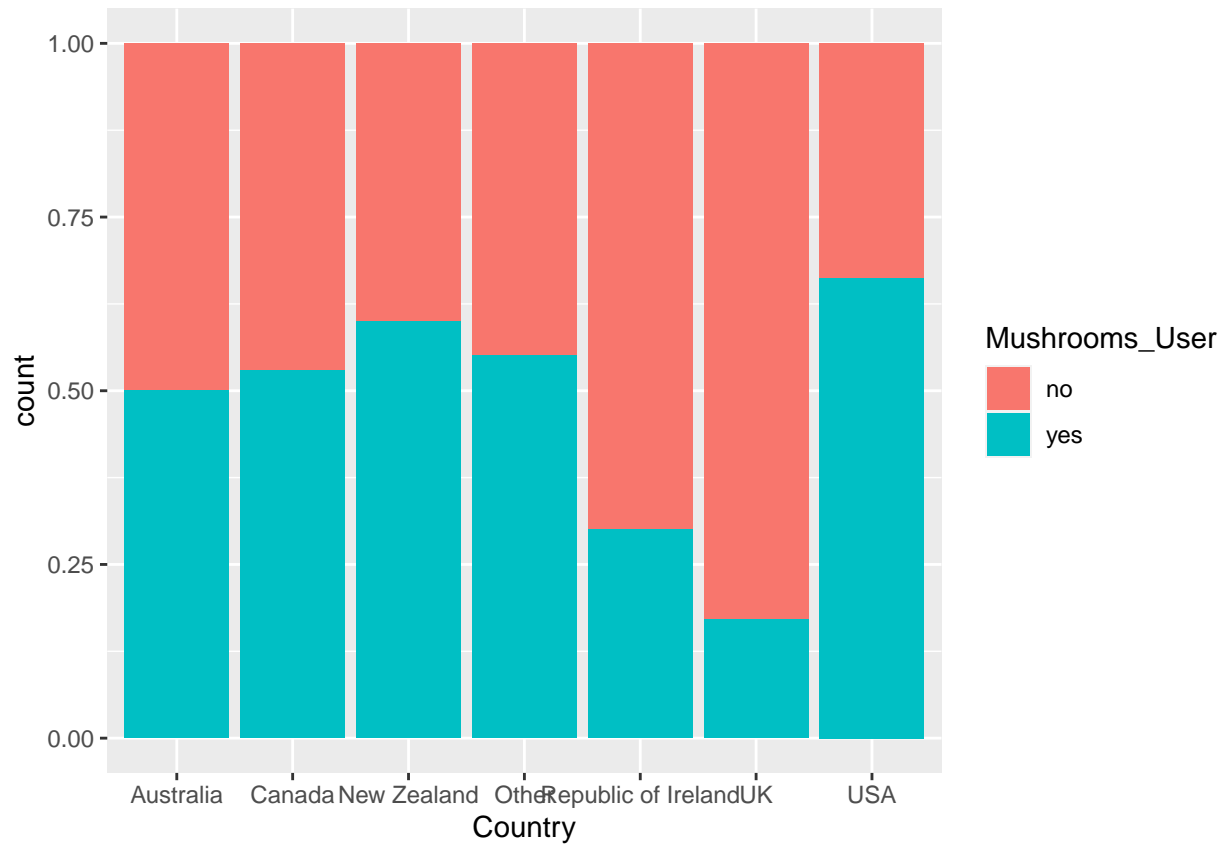
-Since the p-value is less than the significance level, gender is statistically significant.

```
drug_clean_2 %>%
  ggplot(aes(x = Country,
             fill = Mushrooms_User)) +
```

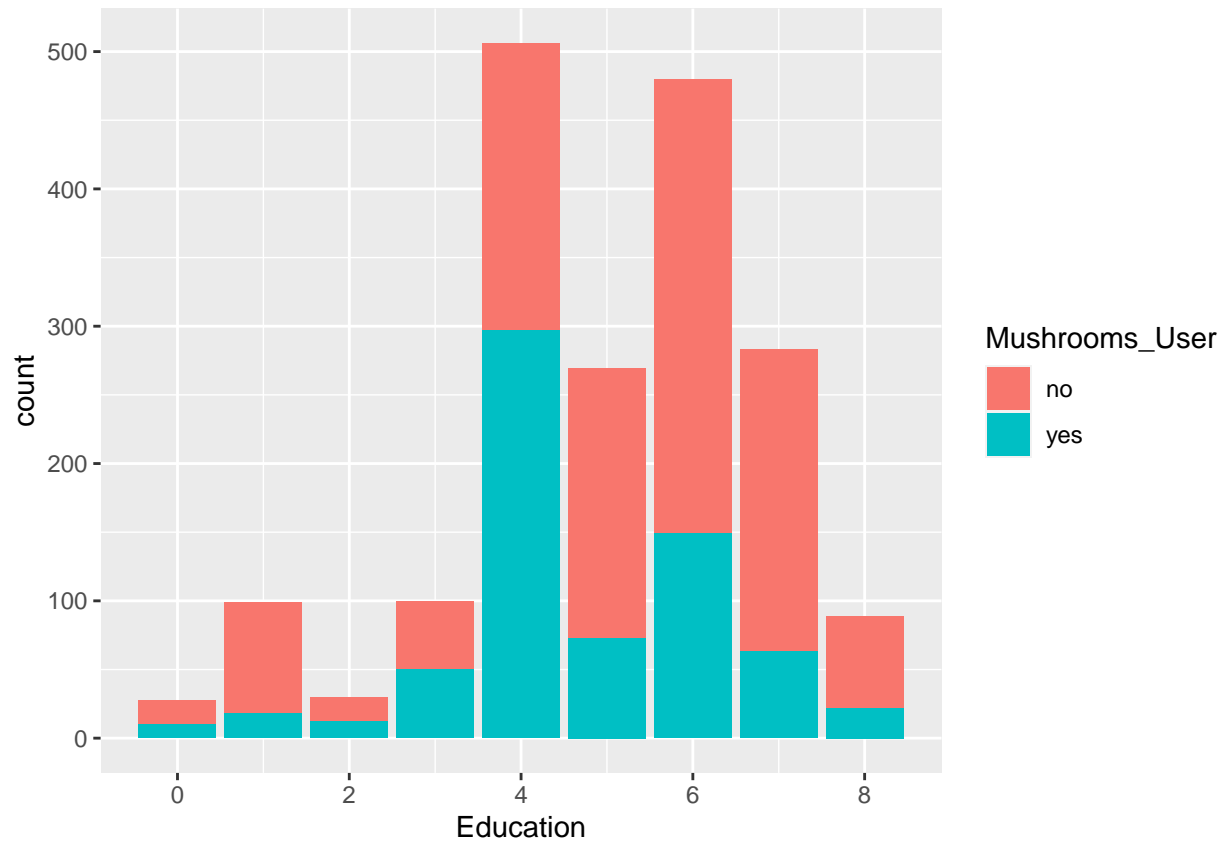
```
geom_bar()
```



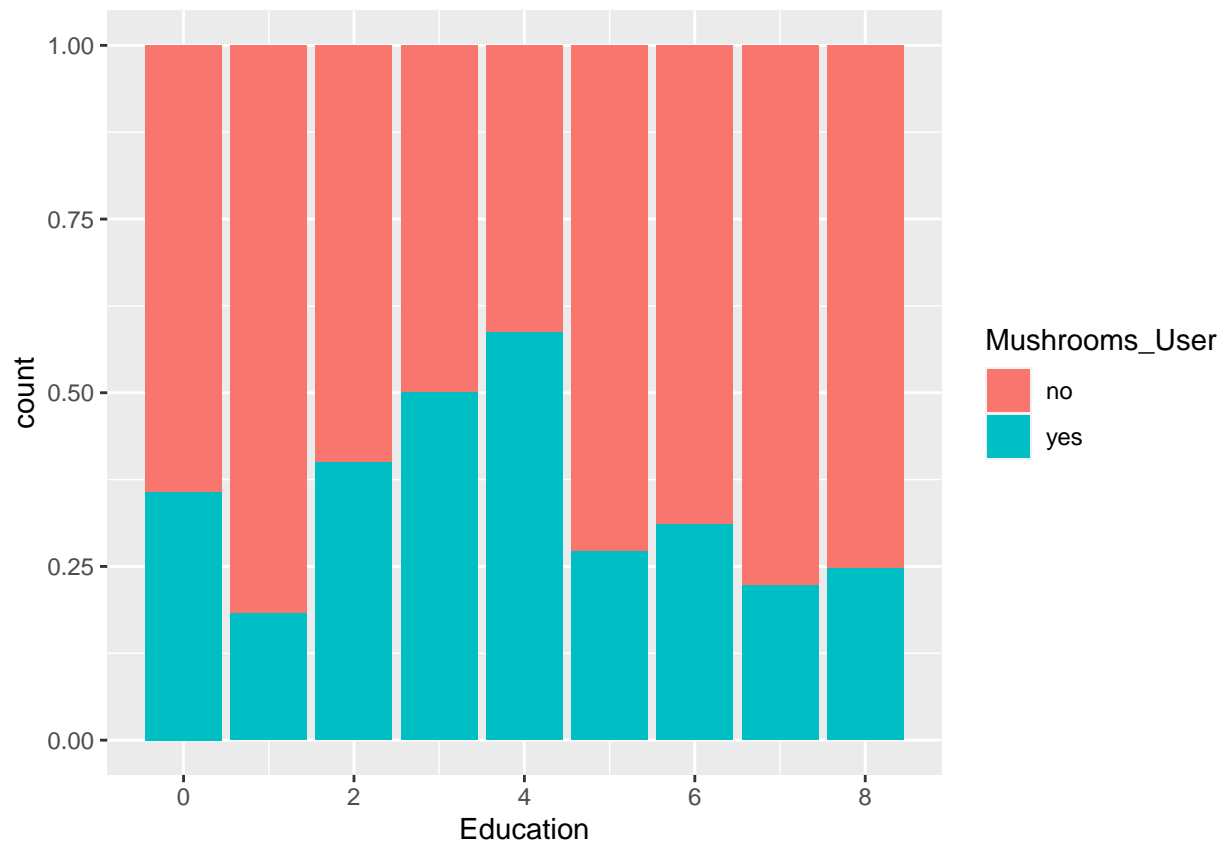
```
drug_clean_2 %>%  
  ggplot(aes(x = Country,  
             fill = Mushrooms_User)) +  
  geom_bar(position = "fill")
```



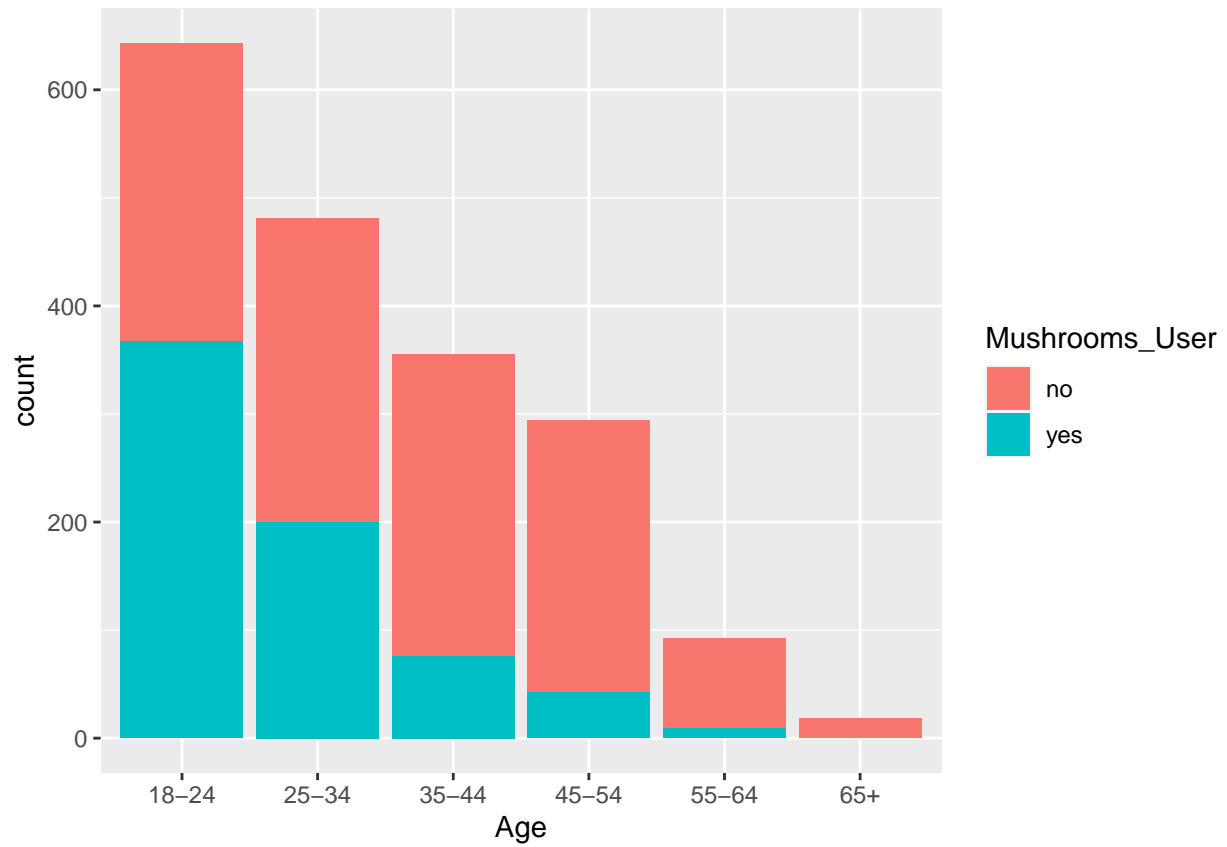
```
drug_clean %>%
  ggplot(aes(x = Education,
             fill = Mushrooms_User)) +
  geom_bar()
```

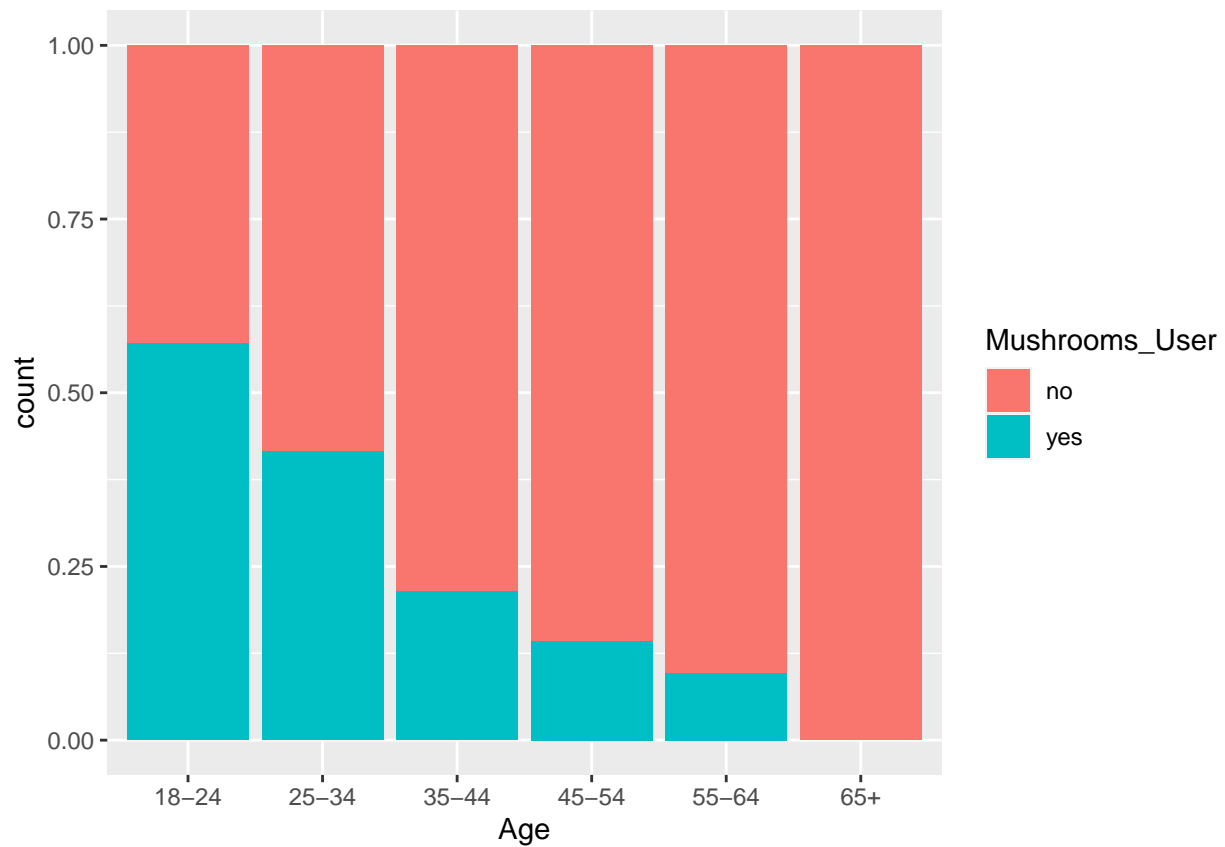
```
drug_clean %>%  
  ggplot(aes(x = Education,  
             fill = Mushrooms_User)) +  
  geom_bar(position = "fill")
```



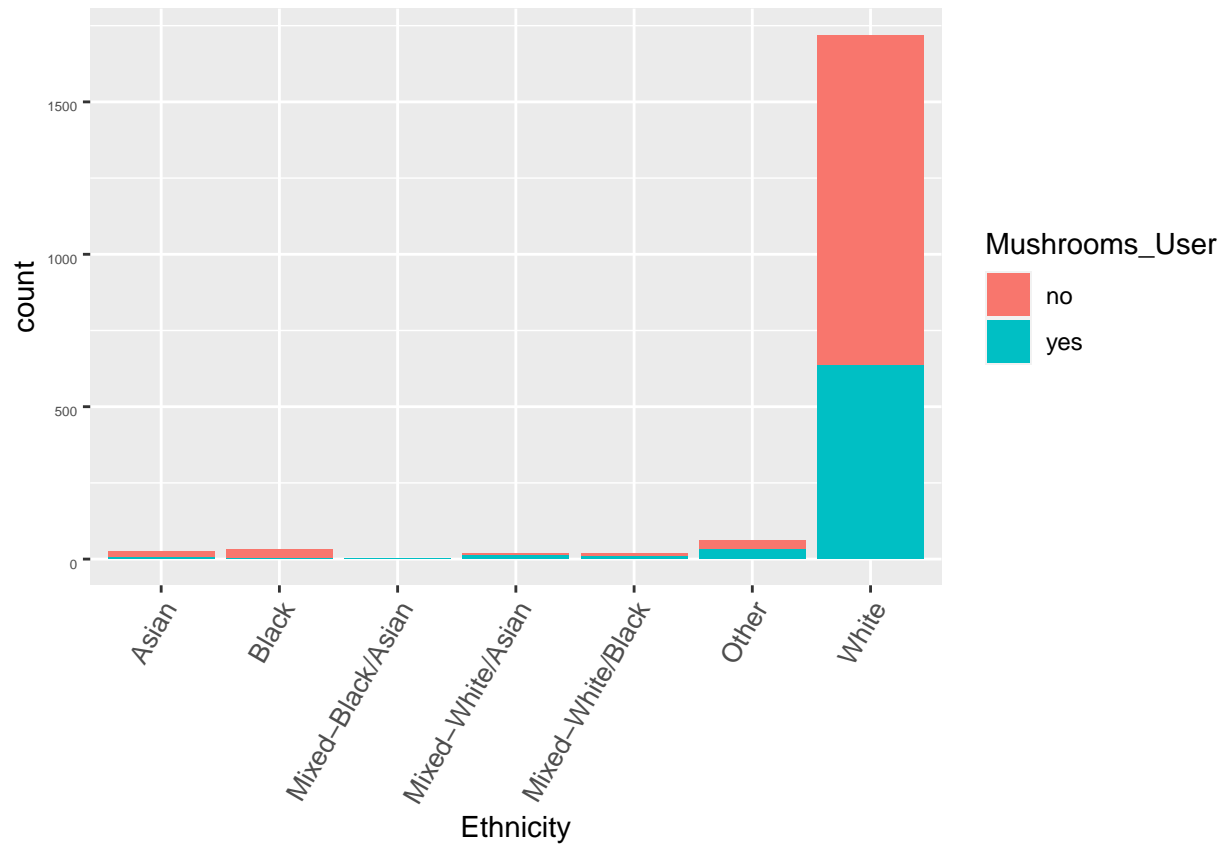
```
drug_clean_2 %>%  
  ggplot(aes(x = Age,  
             fill = Mushrooms_User)) +  
  geom_bar()
```



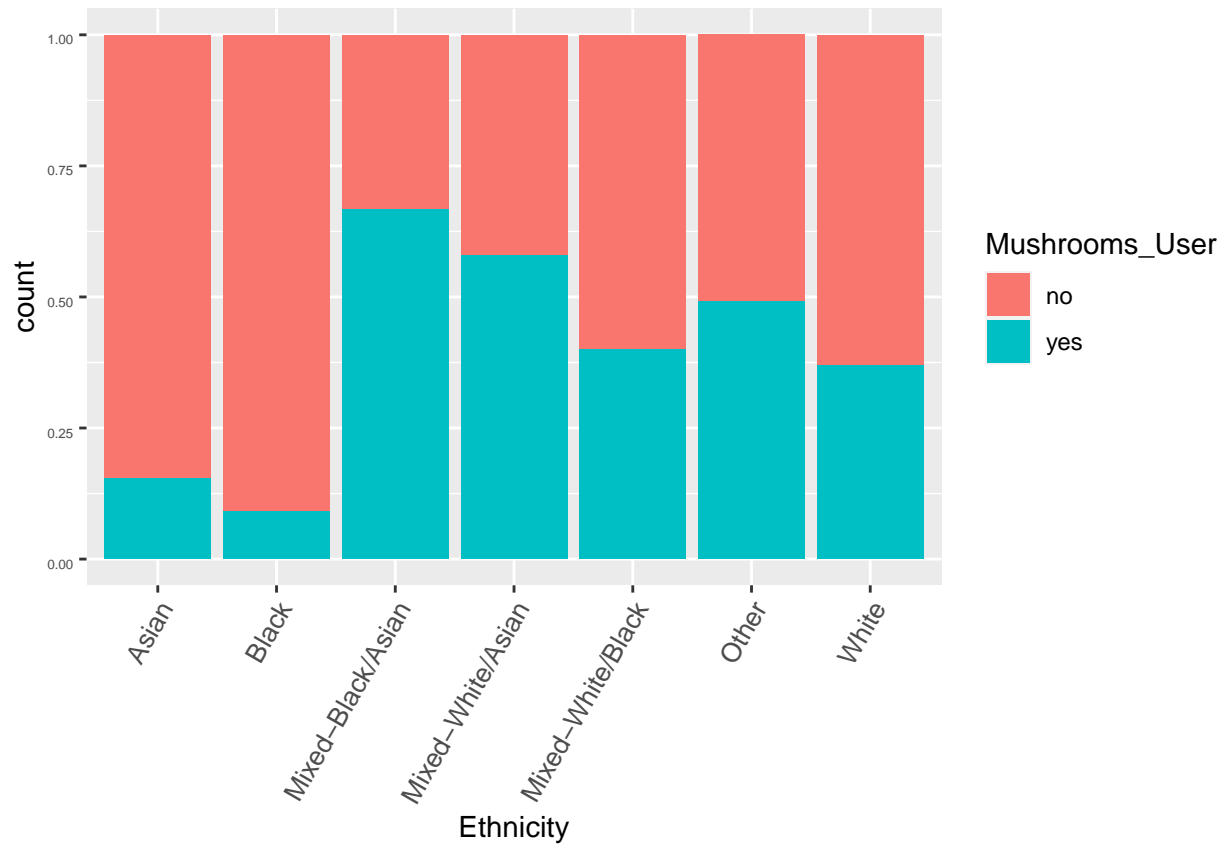
```
drug_clean_2 %>%  
  ggplot(aes(x = Age,  
             fill = Mushrooms_User)) +  
  geom_bar(position = "fill")
```



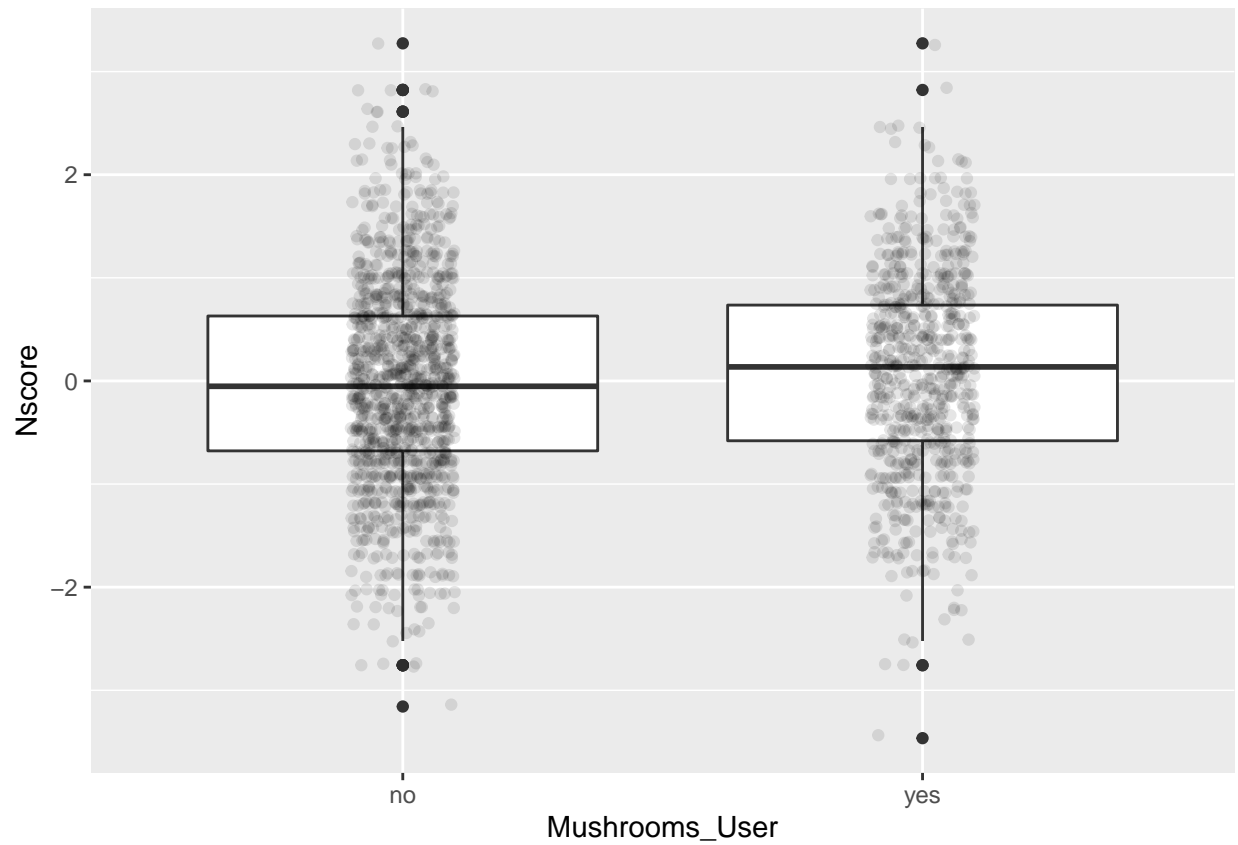
```
drug_clean_2 %>%
  ggplot(aes(x = Ethnicity,
             fill = Mushrooms_User)) +
  geom_bar() +
  theme(axis.text.x = element_text(angle = 60, vjust = 1,
                                     size = 10, hjust = 1), axis.text.y = element_text(vjust = 1, size = 5, hjust = 1))
```



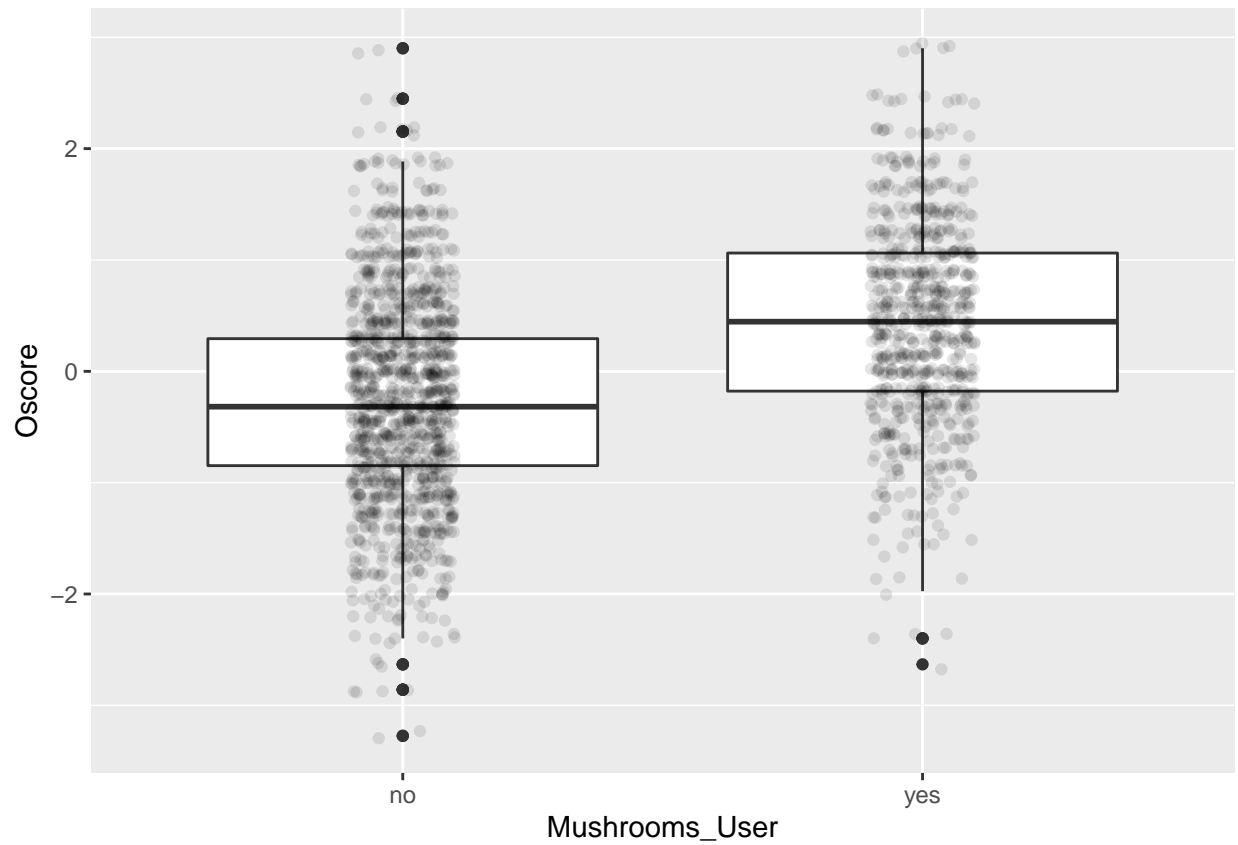
```
drug_clean_2 %>%
  ggplot(aes(x = Ethnicity,
             fill = Mushrooms_User)) +
  geom_bar(position = "fill") +
  theme(axis.text.x = element_text(angle = 60, vjust = 1,
                                    size = 10, hjust = 1), axis.text.y = element_text(vjust = 1, size = 5, hjust = 1))
```



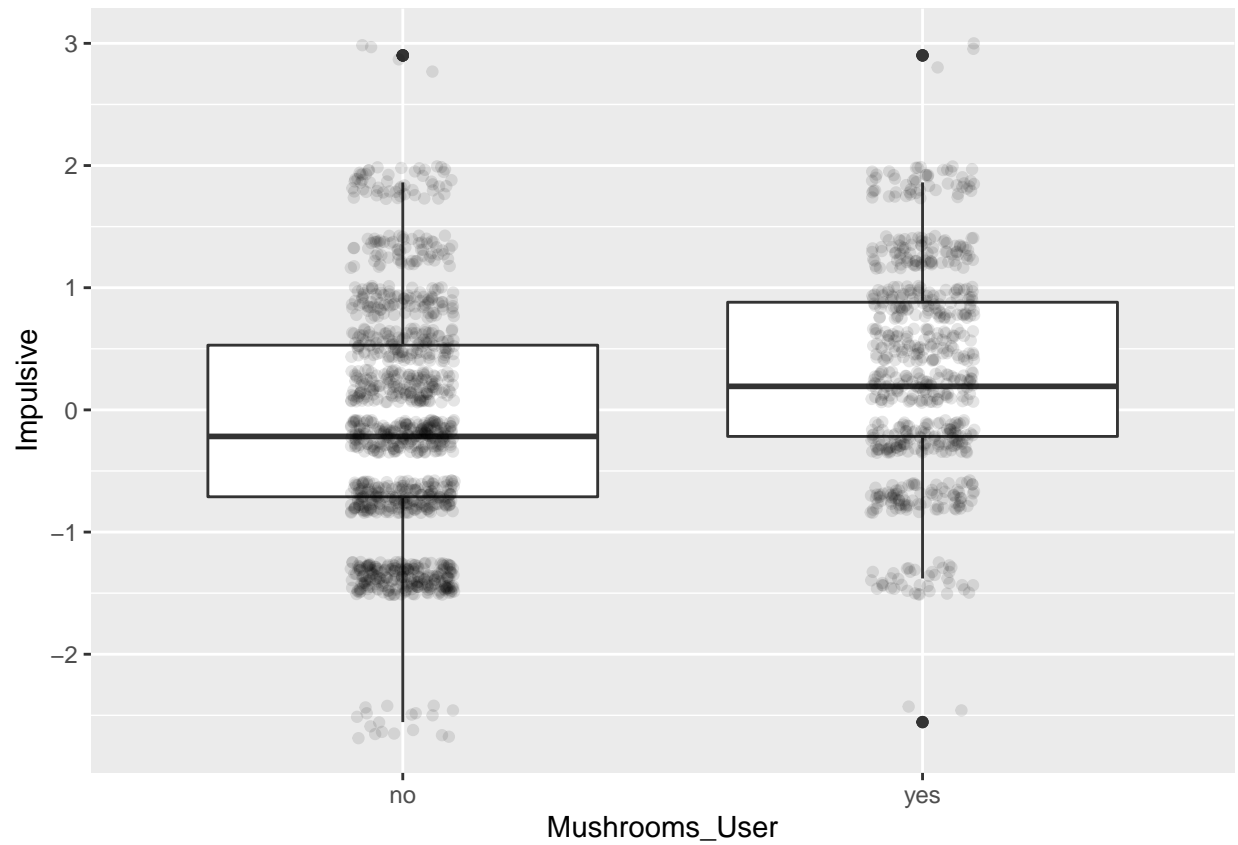
```
drug_clean %>%
  ggplot(aes(Mushrooms_User, Nscore)) +
  geom_boxplot() +
  geom_jitter(width = 0.1, alpha = 0.1)
```



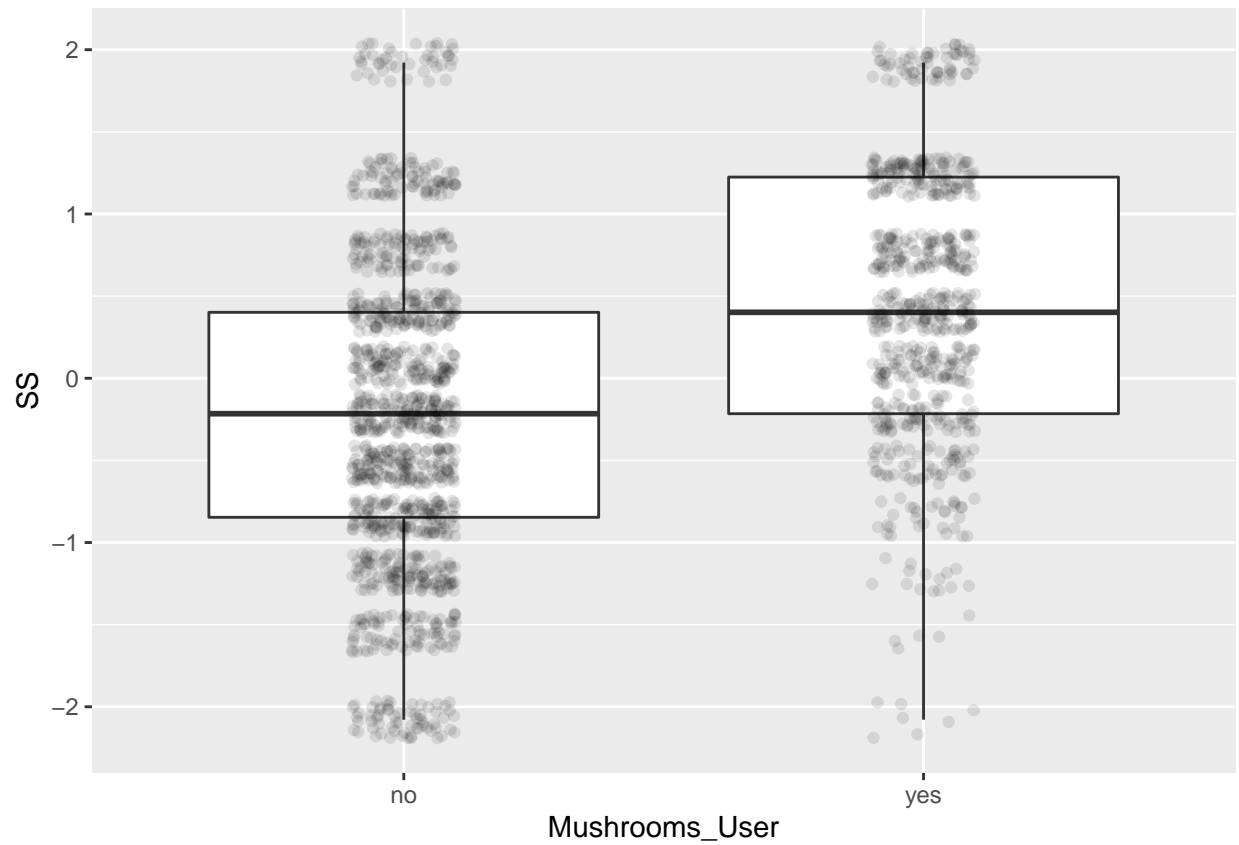
```
drug_clean %>%  
  ggplot(aes(Mushrooms_User, Nscore)) +  
  geom_boxplot() +  
  geom_jitter(width = 0.1, alpha = 0.1)
```



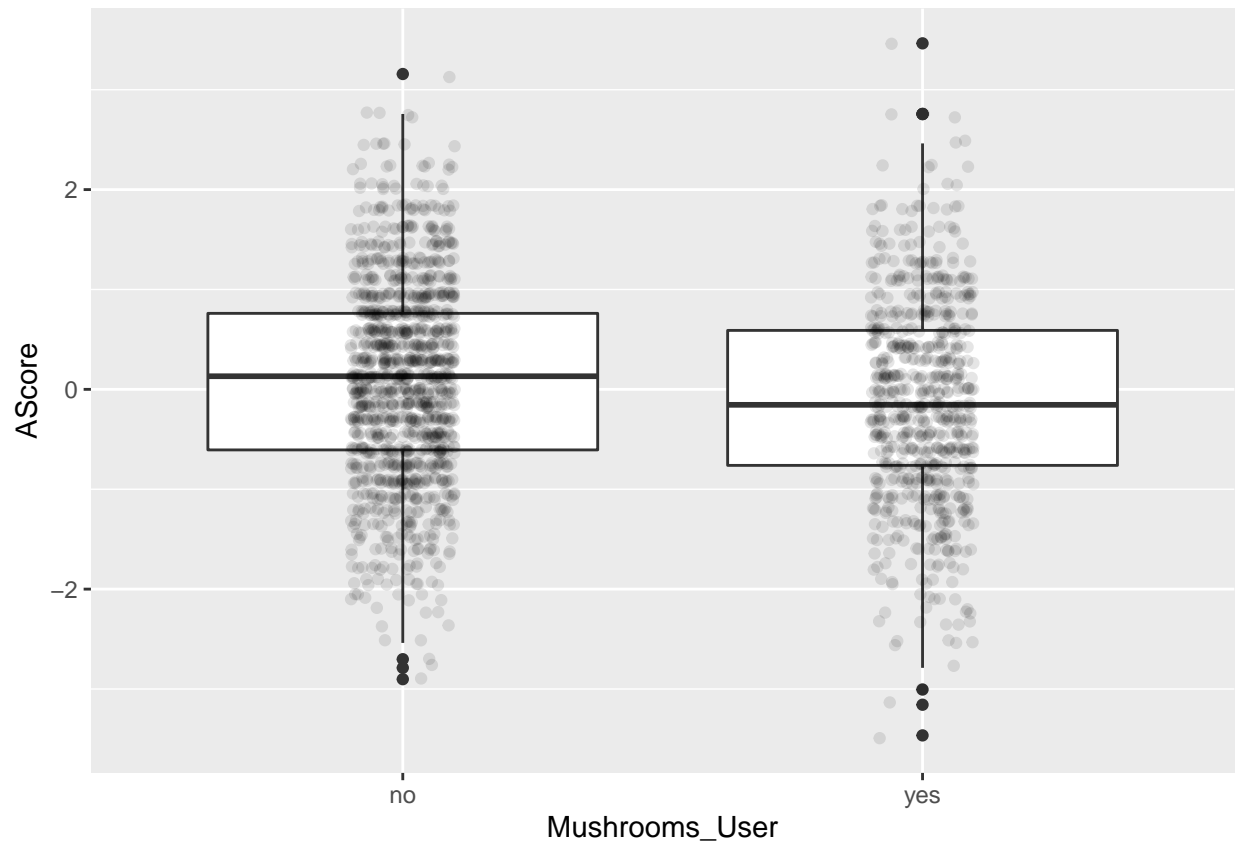
```
drug_clean %>%  
  ggplot(aes(Mushrooms_User, Impulsive)) +  
  geom_boxplot() +  
  geom_jitter(width = 0.1, alpha = 0.1)
```

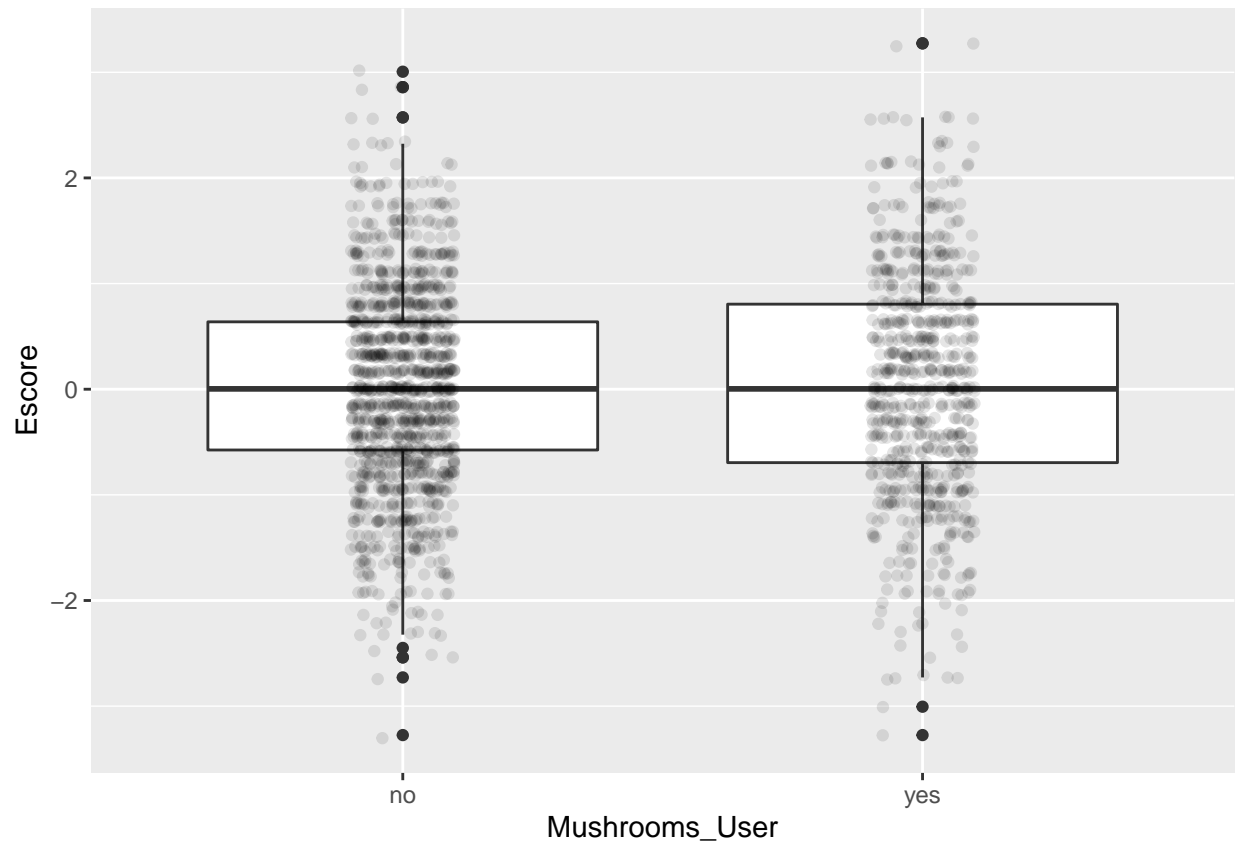
```
drug_clean %>%  
  ggplot(aes(Mushrooms_User, SS)) +  
  geom_boxplot() +  
  geom_jitter(width = 0.1, alpha = 0.1)
```



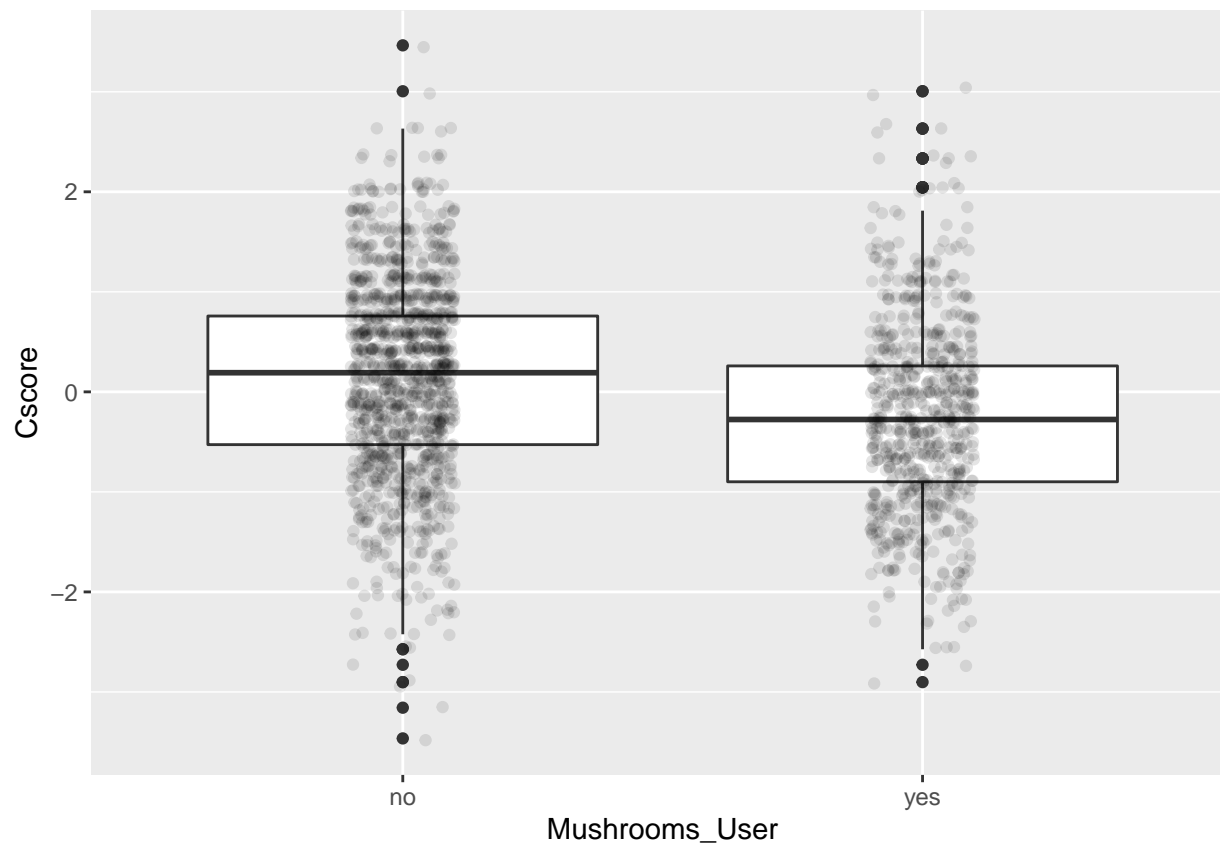
```
drug_clean %>%  
  ggplot(aes(Mushrooms_User, AScore)) +  
  geom_boxplot() +  
  geom_jitter(width = 0.1, alpha = 0.1)
```



```
drug_clean %>%  
  ggplot(aes(Mushrooms_User, Escore)) +  
  geom_boxplot() +  
  geom_jitter(width = 0.1, alpha = 0.1)
```



```
drug_clean %>%  
  ggplot(aes(Mushrooms_User, Cscore)) +  
  geom_boxplot() +  
  geom_jitter(width = 0.1, alpha = 0.1)
```



```
fit_multi <- logistic_reg() %>%
  set_engine("glm") %>%
  fit(Benzos_User ~ Gender + Age + Education + Nscore + Oscore + Impulsive + SS + Cscore + AScore + Esc
result<-tidy(fit_multi, conf.int=TRUE, exponentiate=TRUE)
print(result, n=20)
```

```
## # A tibble: 22 x 7
##   term                                estimate std.error statistic  p.value conf.low conf.high
##   <chr>                                <dbl>     <dbl>     <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)                        0.498     0.271     -2.58  1.00e- 2    0.290    0.842
## 2 GenderM                            1.35      0.112      2.66  7.84e- 3    1.08     1.68
## 3 Age25-34                           1.46      0.145      2.62  8.83e- 3    1.10     1.94
## 4 Age35-44                           1.13      0.162      0.758 4.48e- 1    0.822    1.56
## 5 Age45-54                           1.18      0.175      0.924 3.56e- 1    0.834    1.66
## 6 Age55-64                           0.674     0.278     -1.42 1.56e- 1    0.384    1.15
## 7 Age65+                             0.702     0.633     -0.560 5.76e- 1    0.177    2.23
## 8 EducationLeft schoo~               1.19      0.332      0.528 5.98e- 1    0.622    2.29
## 9 EducationLeft schoo~               1.18      0.467      0.348 7.28e- 1    0.470    2.95
## 10 EducationLeft schoo~              1.25      0.334      0.671 5.02e- 1    0.652    2.42
## 11 EducationLeft schoo~              3.06      0.495      2.26 2.40e- 2    1.17     8.22
## 12 EducationMasters de~              0.703     0.273     -1.29 1.97e- 1    0.413    1.21
## 13 EducationProfession~              0.888     0.278     -0.427 6.69e- 1    0.517    1.54
## 14 EducationSome colle~              1.23      0.269      0.755 4.50e- 1    0.726    2.09
## 15 EducationUniversity~              0.859     0.259     -0.588 5.57e- 1    0.519    1.44
## 16 Nscore                            1.46      0.0634     5.97 2.35e- 9    1.29     1.66
## 17 Oscore                            1.49      0.0611     6.49 8.48e-11    1.32     1.68
```

## 18 Impulsive	1.07	0.0714	0.989	3.22e- 1	0.933	1.23
## 19 SS	1.30	0.0761	3.49	4.89e- 4	1.12	1.51
## 20 Cscore	0.929	0.0621	-1.19	2.35e- 1	0.822	1.05
## # ... with 2 more rows						

Results:

Discussion:

References: