Untitled

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```
library('tidyverse')
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr
              1.1.4
                         v readr
                                     2.1.5
## v forcats 1.0.0
                         v stringr
                                     1.5.1
## v ggplot2 3.5.1
                                     3.2.1
                         v tibble
## v lubridate 1.9.3
                         v tidyr
                                     1.3.1
## v purrr
               1.0.2
## -- Conflicts -----
                                                ----- tidyverse_conflicts() --
## x dplyr::filter()
                           masks stats::filter()
## x dplyr::lag()
                           masks stats::lag()
## x ggplot2::resolution() masks ggvis::resolution()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
library('ggplot2')
library('stats')
library('dplyr')
library('corrplot')
## corrplot 0.94 loaded
# read the file
fast_food= read.csv("C:\\Users\\Vrinda\\Downloads\\FastFoodNutritionScraped20241011V2.csv", header=T, n
```

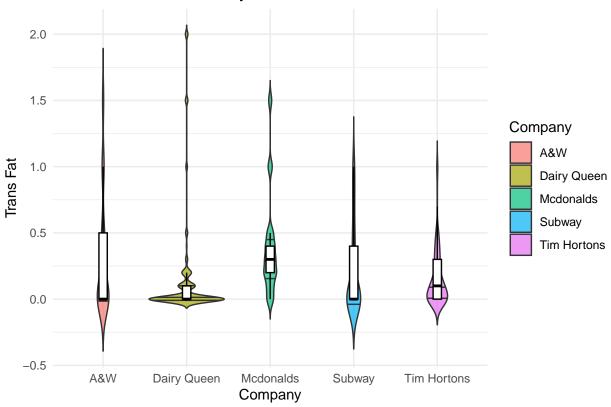
State the hypothesis

library('ggvis')

- 1. Null Hypothesis(Ho) Mean Trans Fat of the two companies is equal. i.e. $\mu 1 = \mu 2$
- 2. Alternative Hypothesis(H1) Mean Trans fat of the two companies are unequal i.e. $\mu 1 \neq \mu 2$

```
#visualisation to see how trans fat is different based on compnay
violin_plot <- ggplot(fast_food, aes(x = Company, y = Trans.Fat..g., fill = Company)) +
geom_violin(trim = FALSE, draw_quantiles = c(0.25, 0.5, 0.75), alpha = 0.7) +
geom_boxplot(width = 0.1, fill = "white", color = "black", outlier.shape = NA) +
labs(title = "Violin Plot of Trans Fat by Restaurant",
x = "Company",
y = "Trans Fat",
fill = "Company") +
theme_minimal()
print(violin_plot)</pre>
```

Violin Plot of Trans Fat by Restaurant



```
#splitting the data based on company wise
company1_data <- fast_food$Trans.Fat..g.[fast_food$Company == 'A&W']
company2_data <- fast_food$Trans.Fat..g.[fast_food$Company == 'Dairy Queen']
company3_data <- fast_food$Trans.Fat..g.[fast_food$Company == 'Mcdonalds']
company4_data <- fast_food$Trans.Fat..g.[fast_food$Company == 'Subway']
company5_data <- fast_food$Trans.Fat..g.[fast_food$Company == 'Tim Hortons']

#check the t- test
# for A&W and Dairy Queen
t.test(company1_data, company2_data, alternative = "two.sided")</pre>
```

```
##
## Welch Two Sample t-test
##
## data: company1_data and company2_data
## t = 3.1064, df = 198.83, p-value = 0.002171
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 0.05414848 0.24240408
## sample estimates:
## mean of x mean of y
## 0.2873874 0.1391111
```

As p-value is less than significance level typically 0.05, We are rejecting the null hypothesis Means there is

a significant difference in Trans Fat between A&W and Dairy Queen.