

# Protein Prediction Based on Nutrition Attributes

ADY201m

Group 5:

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### 1. Understand the Problem

### 1.1Main Objective

- Explore and compare the nutritional composition of fast-food items across companies (e.g., McDonald's, KFC, Burger King, Wendy's).
- Answer practical questions such as:
  - Compare the average overall energy (calorie) levels across products from different fast-food companies.
  - **Examine the correlations** between key nutritional components (e.g., fat, protein, carbohydrates, sodium, etc.).
  - Evaluate which brand may pose a higher cardiovascular risk, based on nutrients related to heart health (e.g., saturated fat, trans fat, cholesterol, sodium).
  - Assess the "quality" of energy intake, with the assumption that higher protein relative to total calories indicates better nutritional quality.
  - **Identify the "best" menu item** according to balanced and healthy nutritional criteria.

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• (Extended goal) Build a predictive model to estimate **Protein** from other nutritional attributes.

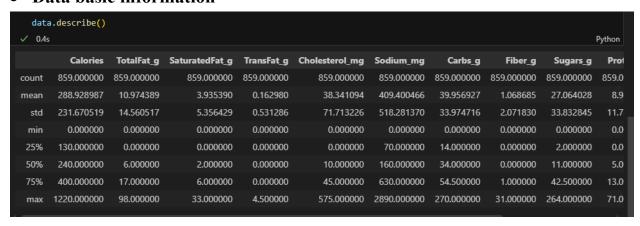
# 2. Data Understanding

### 2.1Dataset Overview

• Size: 859 rows × 13 columns.



### Data basic information



### Columns:

**Company** – The name of the fast food chain or restaurant that offers the menu item (e.g., McDonald's, Burger King, KFC). This attribute allows brand-level comparisons and grouping in analysis.

**Item** – The specific name or description of the menu item. This is a categorical identifier used to distinguish different food or beverage

products.

Calories – The total energy provided by the food item, measured in kilocalories (kcal). This represents the amount of energy a consumer gains from eating the item.

**TotalFat\_g** – The total fat content in grams. Fat contributes to energy intake and plays a significant role in determining the overall caloric value of the item.

**SaturatedFat\_g** – The portion of fat that is saturated, measured in grams. High levels of saturated fat are associated with increased cholesterol and cardiovascular risk.

**TransFat\_g** – The amount of trans fat in grams. Trans fats are artificially produced fats that negatively affect heart health and are often restricted in modern food standards.

**Cholesterol\_mg** – The amount of cholesterol contained in the item, measured in milligrams. Cholesterol levels are important for assessing potential cardiovascular health impacts.

**Sodium\_mg** – The sodium (salt) content in milligrams. Excessive sodium intake is linked to high blood pressure and heart disease, making this an essential nutritional indicator.

**Carbs\_g** – The total carbohydrate content in grams, including starches and sugars. Carbohydrates are the main source of energy in most diets.

**Fiber\_g** – The amount of dietary fiber in grams. Fiber aids digestion, promotes satiety, and helps regulate blood sugar and cholesterol levels.

**Sugars\_g** – The total sugar content in grams, including both natural and added sugars. High sugar levels can contribute to obesity, diabetes, and other metabolic issues.

**Protein\_g** – The protein content in grams. Protein is a critical nutrient for muscle repair, growth, and metabolic processes. This is the **target variable** for prediction in this project.

**WeightWatchers\_Points** – A point-based nutritional scoring system used by the Weight Watchers program to assess the overall dietary impact of a food item. It summarizes calories, fat, and fiber into a single score to guide healthier eating decisions.

- Missing data: none
- **Top companies by item count:** McDonald's (~325), KFC (~200), Burger King (~180), Wendy's (~150).

# 2.2 Data Quality Observations

Some items have Calories = 0, usually drinks like water or diet soda.

# 2.3 Preprocessing (While importing code)

Remove every row containing null values.

**Scaling (for regression later):** Standardize numerical columns if using linear or ridge regression.

# 3. Analysis with SQL

```
--Average calorie content for each company.
select Company, avg(Calories) as AvgCalo
\color{red} \textbf{from} \ \ \textbf{FastFoodNutrition}
group by Company
-- The highest-calorie item from each company.
select Company, Item, Calories
from (select Company, Item, Calories, ROW NUMBER() over (partition by Company order by Calories desc) rn
        from FastFoodNutrition) t
where rn <= 1
--Average amount of saturated fat (SaturatedFat_g) by company.
select Company, avg(SaturatedFat_g) as AvgSaturated
from FastFoodNutrition
group by Company
-- Top 10 items with the highest sodium content (Sodium_mg).
select top 10
    Item,
    Sodium_mg
\color{red} \textbf{from} \ \ \textbf{FastFoodNutrition}
order by Sodium_mg desc
-- Average Calories grouped by WeightWatchers Points.
select WeightWatchers_Points, avg(Calories) AvgCalo
from FastFoodNutrition
group by WeightWatchers_Points
-- Average ratio of Protein to Calories for each company.
select Company, avg(cast(Protein g*1.0 / Calories as decimal(18,4))) as ProPerCalo
from FastFoodNutrition
where Calories > 0
group by Company
-- Top 5 items with the highest protein content but the lowest calories.
select top 5
    Item,
    Protein g,
    Calories,
    cast(Protein_g*1.0 / Calories as decimal(18,4)) as sth
from FastFoodNutrition
where Calories > 0
order by sth desc
```

# 4. Analysis with Python

### **Expected analysis scope**

• Import dataset to SSMS via python code and clean data also.

```
import pandas as pd
     import pyodbc
    data = pd.read_csv('C:\LaLaLa\Data_Storage\FastFoodNutritionMenuV2.csv')
    sever = 'KHANHTUONGDEPTR\SQLEXPRESS'
    datatabase = 'FASTFOOD'
11 v numeric_cols = ['Calories','TotalFat_g','SaturatedFat_g','TransFat_g',
                     'Cholesterol_mg','Sodium_mg','Carbs_g','Fiber_g','Sugars_g',
                     'Protein_g','WeightWatchers_Points']
14 v def to_number(val):
            return float(str(val).replace('g','').replace('mg','').replace('Pnts','').strip())
            return None
20 ∨ for col in numeric_cols:
        data[col] = data[col].apply(to_number)
    data = data.dropna(subset=numeric_cols)
    cnxn = pyodbc.connect('DRIVER={ODBC Driver 11 for SQL Server}; SERVER='+sever+'; DATABASE='+datatabase+'; Tru
    cursor = cnxn.cursor()
29 	imes insert_query = '''INSERT INTO FastFoodNutrition (Company,Item, Calories, TotalFat_g,SaturatedFat_g,TransFa
33 v for row in data.itertuples(index=False):
```

```
cursor = cnxn.cursor()
insert_query = '''INSERT INTO FastFoodNutrition (Company,Item, Calories, TotalFat_g,SaturatedFat_g,TransFat
for row in data.itertuples(index=False):
   values = (
       row.Company,
       float(row.Calories) if pd.notnull(row.Calories) else None,
       float(row.TotalFat_g) if pd.notnull(row.TotalFat_g) else None,
       float(row.SaturatedFat_g) if pd.notnull(row.SaturatedFat_g) else None,
        float(row.TransFat_g) if pd.notnull(row.TransFat_g) else None,
       float(row.Cholesterol_mg) if pd.notnull(row.Cholesterol_mg) else None,
       float(row.Sodium_mg) if pd.notnull(row.Sodium_mg) else None,
       float(row.Carbs_g) if pd.notnull(row.Carbs_g) else None,
       float(row.Fiber_g) if pd.notnull(row.Fiber_g) else None,
       float(row.Sugars_g) if pd.notnull(row.Sugars_g) else None,
        float(row.Protein_g) if pd.notnull(row.Protein_g) else None,
       float(row.WeightWatchers_Points) if pd.notnull(row.WeightWatchers_Points) else None
   cursor.execute(insert_query, values)
cnxn.commit()
cursor.execute ('SELECT * FROM FastFoodNutrition')
```

- Compute mean and correlation between nutrients.
  - -Average calorie content for each company

```
req1 = data.groupby('Company')['Calories'].mean()
```

```
data = pd.read_csv('C:\LaLaLa\ADY201m\Data\FastFoodNutritionMenuV2.csv')
Company
Burger King 359.189944
KFC 210.049751
McDonald's 283.107692
Wendy's 322.500000
Name: Calories, dtype: float64
```

-Relationship between Calories and TotalFat g

```
req3= data[['Calories','TotalFat_g']].corr()

Calories TotalFat_g

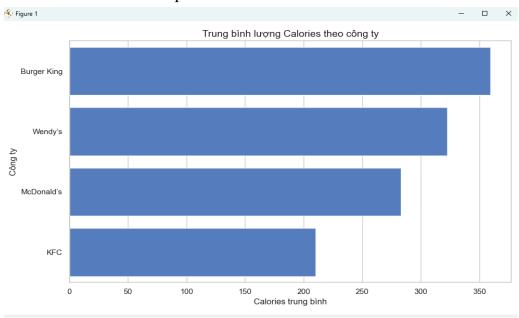
Calories 1.000000 0.824249

TotalFat_g 0.824249 1.000000
```

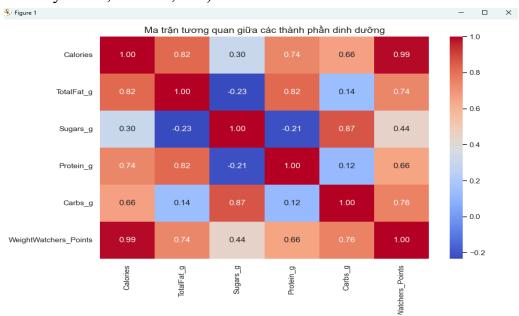
### 5. Visualization

- Use seaborn and matplotlib.pyplot to display queries:

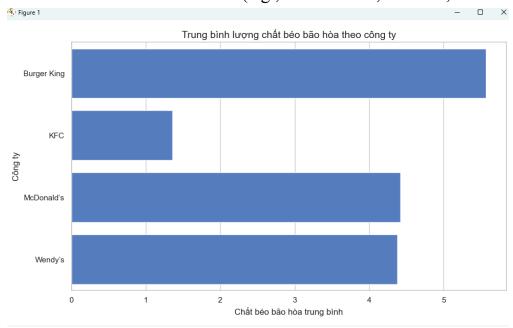
Compare the average overall energy (calorie) levels across products from different fast-food companies.



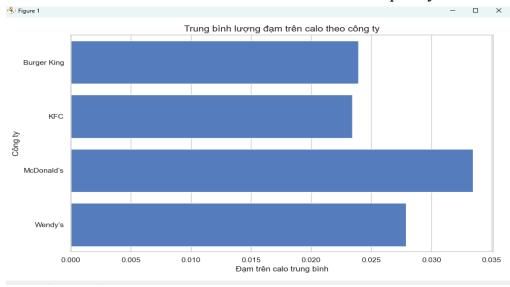
**Examine the correlations** between key nutritional components (e.g., fat, protein, carbohydrates, sodium, etc.).



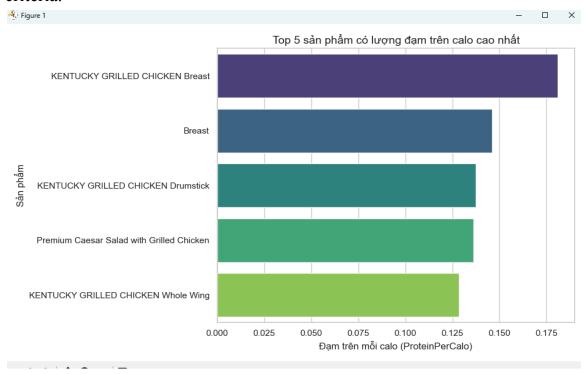
**Evaluate which brand may pose a higher cardiovascular risk**, based on nutrients related to heart health (e.g., saturated fat, trans fat, cholesterol, sodium).



Assess the "quality" of energy intake, with the assumption that higher protein relative to total calories indicates better nutritional quality.



**Identify the "best" menu item** according to balanced and healthy nutritional criteria.



# 6. Regression

Use scikit-learn library to predict "Protein\_g" via other nutrition attributes.

```
import pandas as pd
from sklearn.linear model import LinearRegression
df = pd.read csv("C:\LaLaLa\ADY201m\Data\FastFoodNutritionMenuV2.csv")
numeric cols = df.select dtypes(include=['int64', 'float64']).columns
X = df[numeric cols].drop(columns=['Protein g'])
y = df['Protein_g']
model = LinearRegression()
model.fit(X, y)
print("\nNhâp thông tin món ăn để dư đoán Protein g:")
Calories = float(input("Calories: "))
TotalFat_g = float(input("TotalFat_g: "))
SaturatedFat g = float(input("SaturatedFat g: "))
TransFat g = float(input("TransFat g: "))
Cholesterol mg = float(input("Cholesterol mg: "))
Sodium_mg = float(input("Sodium_mg: "))
Carbs g = float(input("Carbs_g: "))
Fiber_g = float(input("Fiber_g: "))
Sugars_g = float(input("Sugars_g: "))
WeightWatchers Points = float(input("WeightWatchers Points: "))
```

```
sample data = pd.DataFrame([{
         'Calories': Calories,
         'TotalFat g': TotalFat g,
         'SaturatedFat_g': SaturatedFat_g,
         'TransFat g': TransFat g,
         'Cholesterol_mg': Cholesterol_mg,
         'Sodium_mg': Sodium_mg,
         'Carbs_g': Carbs_g,
         'Fiber_g': Fiber_g,
38
         'Sugars_g': Sugars_g,
         'WeightWatchers Points': WeightWatchers Points
     }])
42
     predicted protein = model.predict(sample data)[0]
     print(f"\nDv doán Protein_g cho món mẫu: {predicted_protein:.2f} ")
```

### 7. Analysis tool

-R Studio was employed to validate the regression model and visualize relationships between dependent and independent variables. The *ggplot2* package was used to plot residuals and regression lines, confirming the linear relationship assumption. Additionally, summary statistics from R supported the findings obtained through Python, reinforcing the consistency of the analysis.

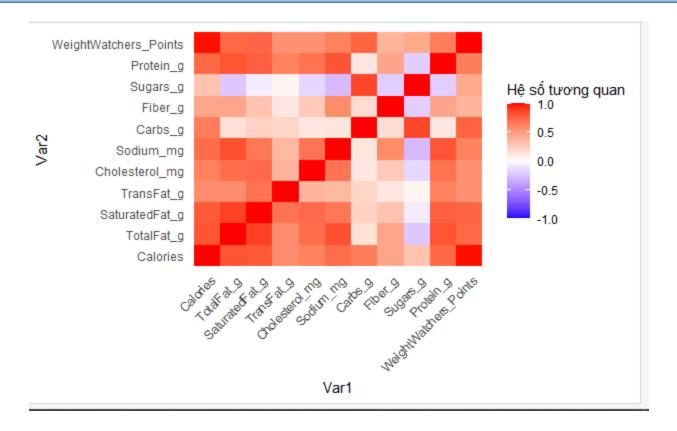
### **ANALYSIS CODE:**

```
# --- Phân tích thống kê mô tả (Descriptive Statistics) ---
COPYDATA <- read.csv("C:/LaLaLa/ADY201m/Data/FastFoodNutritionMenuV2.csv")
# 1. Tổng quan dữ liệu
summary(COPYDATA)
# 2. Trung bình, độ lệch chuẩn, phương sai cho các cột số
mean(COPYDATA$Calories, na.rm = TRUE)
sd(COPYDATA$Calories, na.rm = TRUE)
var(COPYDATA$Calories, na.rm = TRUE)
# Neu muön tinh nhiều bien môt lúc:
num_cols <- sapply(COPYDATA, is.numeric)</pre>
sapply(COPYDATA[, num_cols], mean, na.rm = TRUE)
sapply(COPYDATA[, num_cols], var, na.rm = TRUE)
# Ma trận tương quan giữa các biến số
cor(COPYDATA[, num_cols], use = "complete.obs")
# Vẽ heatmap tương quan
library(ggplot2)
library(reshape2)
num_cols <- sapply(COPYDATA, is.numeric)</pre>
corr_matrix <- cor(COPYDATA[, num_cols], use = "complete.obs")</pre>
melted_corr <- melt(corr_matrix)</pre>
ggplot(data = melted_corr, aes(x=Var1, y=Var2, fill=value)) +
  geom_tile() +
  scale_fill_gradient2(low="blue", high="red", mid="White",
                       midpoint=0, limit=c(-1,1), space="Lab",
                       name="Hê số tương quan") +
  theme_minimal() +
  theme(axis.text.x = element_text(angle=45, vjust=1, hjust=1))
```

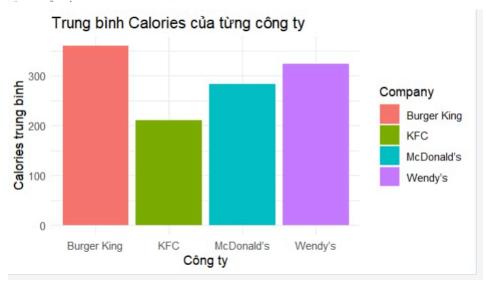
#### **OUTPUT:**

```
> summary(COPYDATA)
                               Calories TotalFat_g
  Company
                  Item
Length:859
               Length:859
                              Min. : 0.0 Min. : 0.00
               Class :character
Class :character
                              1st Qu.: 130.0 1st Qu.: 0.00
                              Median: 240.0 Median: 6.00
Mode :character Mode :character
                              Mean : 288.9 Mean :10.97
                               3rd Qu.: 400.0 3rd Qu.:17.00
                              Max. :1220.0 Max. :98.00
              TransFat_g
SaturatedFat_q
                          Cholesterol_mq Sodium_mq
Min. : 0.000
            Min. :0.000
                          Min. : 0.00 Min. : 0.0
1st Qu.: 0.000 1st Qu.:0.000
                          1st Qu.: 0.00 1st Qu.: 70.0
Median: 2.000 Median: 0.000 Median: 10.00 Median: 160.0
Mean : 3.935 Mean : 0.163 Mean : 38.34 Mean : 409.4
3rd Qu.: 6.000 3rd Qu.:0.000
                          3rd Qu.: 45.00 3rd Qu.: 630.0
Max. :33.000 Max. :4.500 Max. :575.00 Max. :2890.0
                                        Protein_g
  Carbs_g
             Fiber_g
                          Sugars_g
Min. : 0.00 Min. : 0.000 Min. : 0.00 Min. : 0.000
Median: 34.00 Median: 0.000 Median: 11.00 Median: 5.000
Mean : 39.96 Mean : 1.069 Mean : 27.06 Mean : 8.987
3rd Qu.: 54.50 3rd Qu.: 1.000 3rd Qu.: 42.50 3rd Qu.:13.000
Max. :270.00 Max. :31.000 Max. :264.00
                                         Max. :71.000
WeightWatchers_Points
Min. : 0.0
1st Qu.: 142.5
Median : 272.0
Mean : 310.9
3rd Qu.: 430.0
Max. :1317.0
>
```

```
> # 2. Trung bình, độ lệch chuẩn, phương sai cho các cột số
> mean(COPYDATA$Calories, na.rm = TRUE)
[1] 288.929
> sd(COPYDATA$Calories, na.rm = TRUE)
[1] 231.6705
> var(COPYDATA$Calories, na.rm = TRUE)
[1] 53671.23
> |
```



### **VISUALIZATION CODE:**



```
cor(data$Calories, data$TotalFat_g, use = "complete.obs")

ggplot(data, aes(x = TotalFat_g, y = Calories)) +

geom_point(color = "blue") + # Vē các điểm dữ liệu

geom_smooth(method = "lm", se = TRUE, color = "red") + # Thêm đường hồi quy tuyến tính

theme_minimal() +

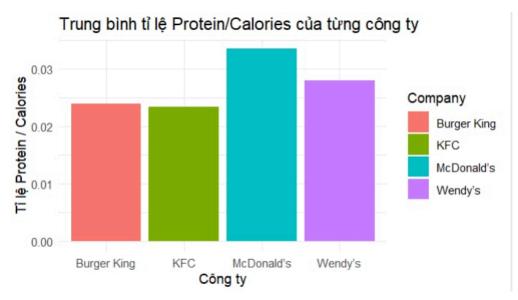
labs(title = "Mối quan hệ giữa Calories và Total Fat",

x = "Total Fat (g)",

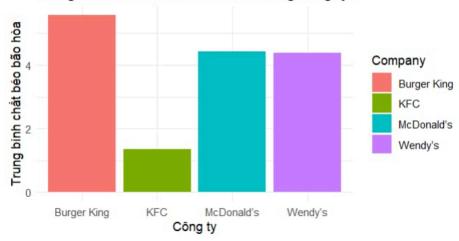
y = "Calories")
```



```
data$Protein_Ratio <- data$Protein_g / data$Calories
avg_ratio <- aggregate(Protein_Ratio ~ Company, data = data, FUN = mean)
ggplot(avg_ratio, aes(x = Company, y = Protein_Ratio, fill = Company)) +
geom_col() +
theme_minimal() +
labs(title = "Trung bình ti lệ Protein/Calories của từng công ty",
        x = "Công ty",
        y = "Ti lệ Protein / Calories")</pre>
```



### Trung bình chất béo bão hòa của từng công ty



#### Top 5 món ăn có tỉ lệ Protein/Calories cao nhất theo công ty

