## **Introduction**

The **goal** of this project is to build predictive models to estimate the probability of individuals receiving the **H1N1 vaccine** and the **seasonal flu vaccine** based on demographic, behavioral, and health-related factors. The dataset used for this analysis comes from the **2009 National H1N1 Flu Survey**, which includes responses from thousands of individuals regarding their attitudes towards vaccination, medical history, and risk perception.

Understanding vaccine uptake patterns is **crucial** for public health authorities to design targeted awareness campaigns and vaccination programs, especially during pandemic situations. By leveraging **machine learning models**, we aim to improve vaccine adoption strategies.

The models developed include:

- **✓ Logistic Regression** A simple baseline model.
- **▼ Random Forest** A tree-based model capturing non-linear relationships.
- **▼ XGBoost** An advanced boosting algorithm.
- ✓ LightGBM A fast gradient boosting model that performed the best.

This report presents an exploratory **data visualization**, data preprocessing steps, and a comparison of **model performances using ROC-AUC scores** 

# **Dataset Description**

The dataset used in this analysis is the 2009 National H1N1 Flu Survey dataset, which contains responses from individuals regarding their vaccination status and various demographic, health-related, and behavioral factors.

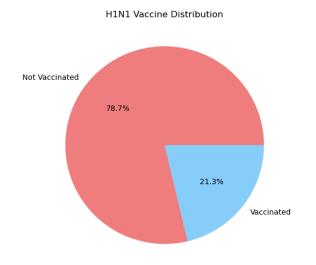
### Dataset Overview:

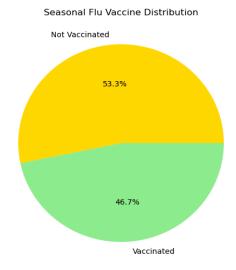
- Total Rows: 26708
- Total Columns: 36
- Target Variables:
  - h1n1\_vaccine: 1 if the person took the H1N1 vaccine, 0 otherwise.
  - seasonal\_vaccine: 1 if the person took the seasonal flu vaccine, 0 otherwise.
- Key Features:
  - Demographics: Age, gender, education level, income level, etc.
  - Health Conditions: Chronic diseases, doctor visits, general health conditions, etc.
  - Vaccination Attitudes: Concern about vaccine safety, effectiveness, trust in government, etc.
  - Behavioral Factors: Frequency of doctor visits, social distancing habits, etc.

A total of 26,708 individuals are included in this dataset, providing a diverse sample for vaccine adoption modeling.

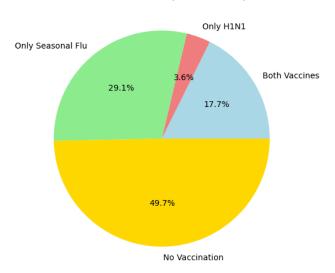
# **DATA VISUALIZATION**

### • PEOPLE VACCINATED STATS

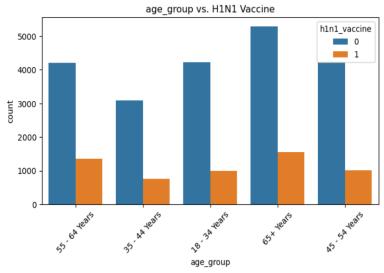


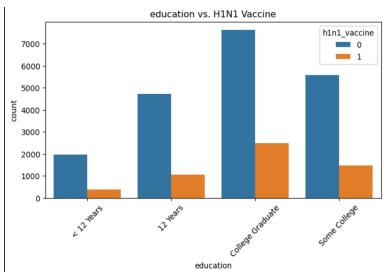


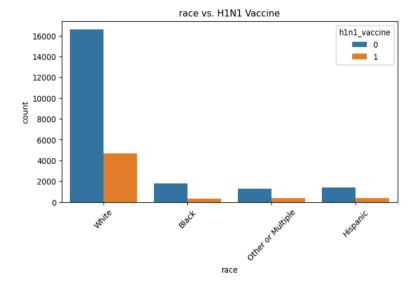
Vaccination Status (Both Vaccines)

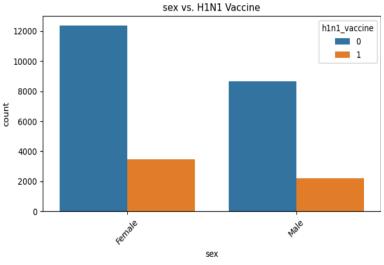


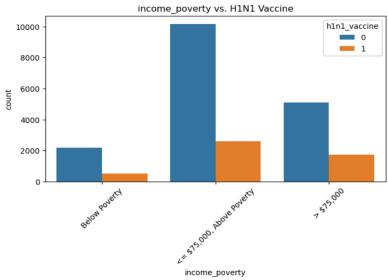
## • FACTORS AFFECTING VACCINATION RATE

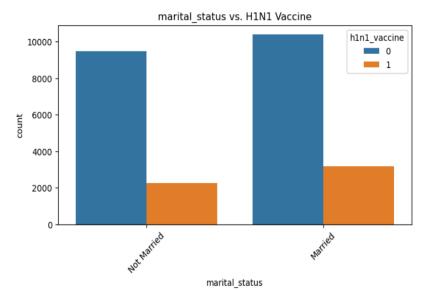


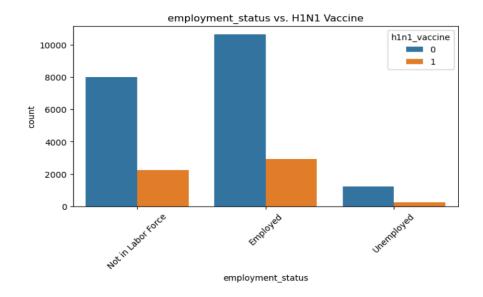


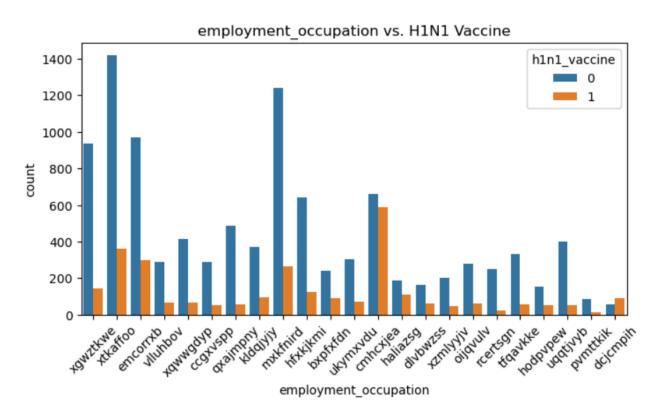


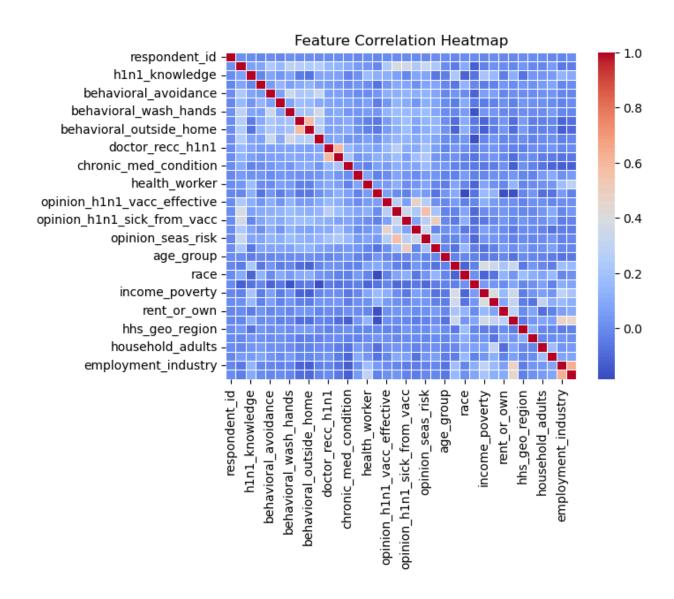












## **Data Visualization Summary and Conclusion**

After conducting exploratory data analysis (EDA) and visualizing the dataset, the following key insights were observed:

### 1. Age Group Distribution:

 The dataset contains individuals from different age groups, with a higher proportion of middle-aged and older individuals.  The vaccination rates vary significantly across different age groups.

### 2. Employment Status & Vaccination Rates:

 A noticeable trend was observed where unemployed individuals had different vaccination patterns compared to employed individuals.

#### 3. Education Level and Awareness:

 Higher education levels were generally correlated with higher vaccination rates, suggesting a possible link between awareness and vaccine acceptance.

### 4. H1N1 and Seasonal Flu Vaccine Uptake:

- The proportion of people receiving the H1N1 vaccine
  was lower than those receiving the seasonal flu vaccine.
- A significant number of individuals who took the seasonal flu vaccine also took the H1N1 vaccine, but the reverse was not as strong.

#### **Conclusion from Data Visualization:**

- **Demographic Factors:** Age, employment status, and education level play a crucial role in vaccine uptake.
- **Behavioral Trends:** Individuals who take the seasonal flu vaccine are more likely to take the H1N1 vaccine.
- **Healthcare Awareness:** People with higher awareness of flurisks show a higher vaccination rate.

# **Steps Followed in Building the Model**

## 1. Data Preprocessing

- **Handling Missing Values:** Used **SimpleImputer** to fill missing values with the most frequent values.
- **Encoding Categorical Variables:** Converted categorical variables into numerical form using one-hot encoding.
- **Feature Selection:** Removed redundant or highly correlated features to improve model efficiency.
- Checked for outliers using boxplots and removed them without overfitting the model

#### 2. Model Selection

Several models were trained and evaluated using **ROC-AUC** score:

- Baseline Model:
  - Logistic Regression achieved an ROC-AUC of 0.8314 (H1N1) and 0.8555 (Seasonal Flu).
- Tree-Based Models:
  - Random Forest: Moderate improvement, but prone to overfitting.
  - XGBoost & LightGBM: Showed strong results due to their ability to handle categorical features and missing values.

### 3. Hyperparameter Tuning

- **LightGBM** was selected as the final model due to its high accuracy and efficiency.
- Optimized parameters:
  - o num\_leaves=20
  - o n\_estimators=200
  - o max\_depth=5

- o learning\_rate=0.05
- colsample\_bytree=0.6
- Final **ROC-AUC Score for LightGBM**:
  - H1N1 Vaccine: 0.865
  - Seasonal Flu Vaccine: 0.877

## The CSV file generated contained three columns:

- respondent\_id The unique identifier for each individual in the test dataset.
- 2. **h1n1\_vaccine** The predicted probability that the individual received the H1N1 vaccine.
- 3. **seasonal\_vaccine** The predicted probability that the individual received the seasonal flu vaccine.

The **submission.csv** file provides key insights about the **likelihood of individuals receiving the H1N1 and seasonal flu vaccines** based on their demographic and behavioral features. Since the values in the h1n1\_vaccine and seasonal\_vaccine columns are probabilities (ranging from 0 to 1), they represent the model's confidence in whether each individual **received the vaccine**.

## **Key Insights from the CSV File**

- A probability close to **1** means the model is very confident that the individual **took the vaccine**.
- A probability close to **0** means the model is confident that the individual **did not take the vaccine**.
- A probability around **0.5** means the model is uncertain and considers the individual as having a **50-50 chance** of taking the vaccine.