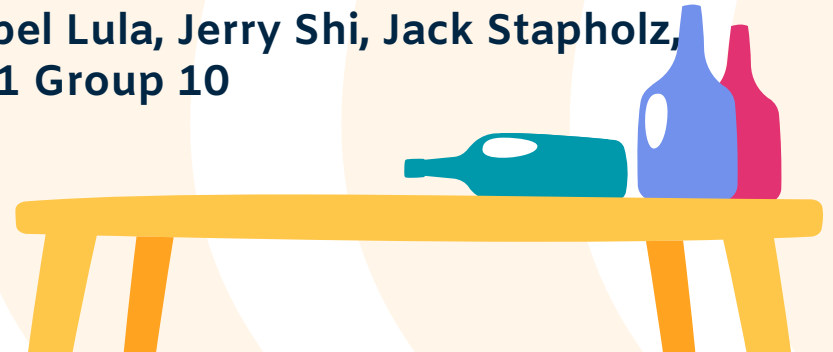


# Predicting People's Alcoholic Status

By: Nathan Kim, Abel Lula, Jerry Shi, Jack Stapholz,  
Rachel Stokol Lec 1 Group 10

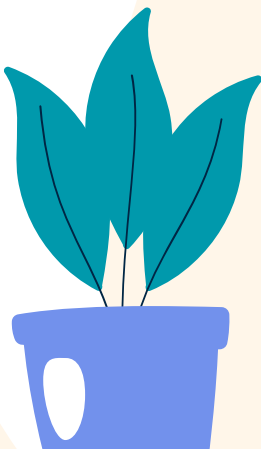


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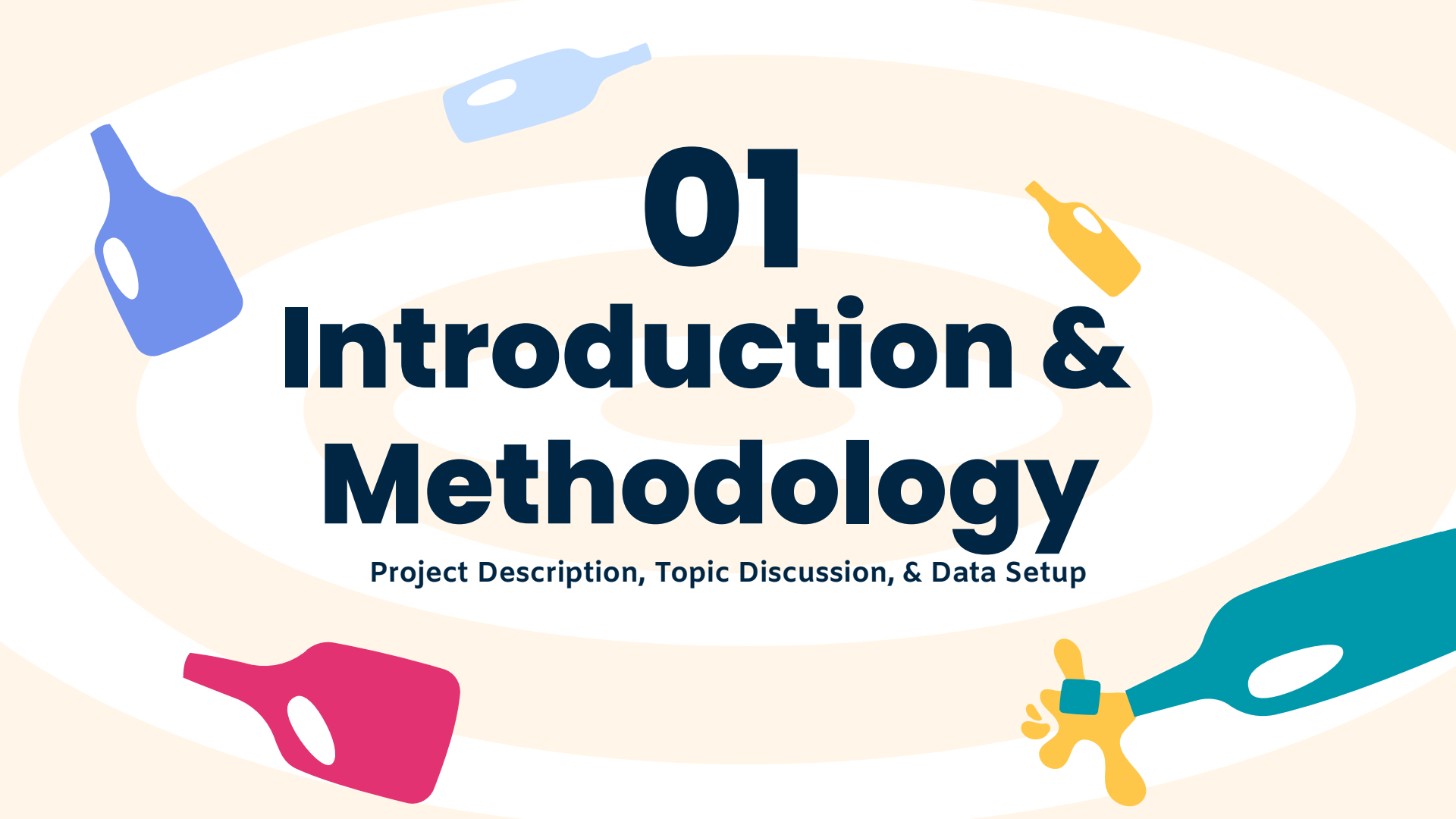
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# 01 Introduction & Methodology

Project Description, Topic Discussion, & Data Setup

# Alcoholic Status Prediction

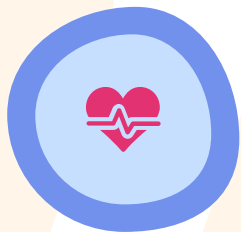


The data discusses a person's alcoholic status based off vitals such as:

- Height
- Weight
- LDL\_Chole

The goal for this project is to utilize the `predict()` function in R and predict the testing data response variable, given the training data, with a modeling method or a combination of methods that we learned in class to predict future alcoholic statuses.

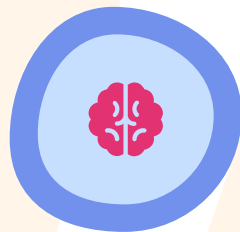
# Testing and Training Dataset



## Observations

Training Data: 70,000 Observations

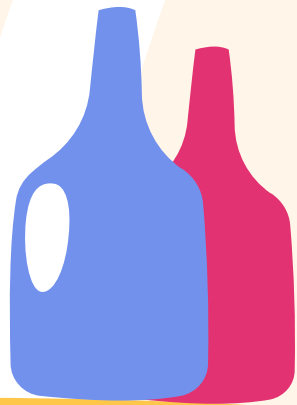
Testing Data: 30,000 Observations



## Dimensions

Training Data: 70,000 x 27

Testing Data: 30,000 x 26



# Process

## Clean Up Data



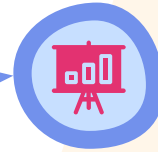
Address missing values and set up categorical variables as factors

## Fit Models



Create multiple models of varying type and number of predictors

## Test & Compare Models



Fit models, identify their accuracy rates, and select the best model according to accuracy



# Identifying Missing Values

**86%**

of observations in the complete dataset had at least one missing value

All 26 predictors had missing values with frequencies between

**6-12%**

We employed three imputation methods to replace the missing values:

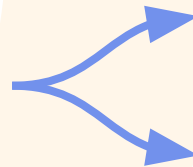
1. Mean, median, and mode
2. MICE
3. Hmisc



# Imputing Missing Values



**Average**

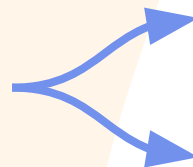


Impute NAs in numerical predictors with the mean or median

Impute NAs in categorical predictors with the mode



**MICE**

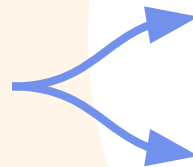


Use mice package in R to create a data frame of imputed values

Merge imputed values and original data to replace NAs



**Hmisc**



Use aregImpute() from Hmisc package in R to impute values (n.impute = 5)

Merge imputed values from the 5th iteration of aregImpute() with original data to replace NAs



The background features a series of concentric, hand-drawn style ovals in shades of light orange and cream. Scattered around these ovals are several stylized, colorful shapes that resemble bottles or containers. These include a light blue bottle at the top, a medium blue bottle on the left, a yellow bottle on the right, a pink bottle at the bottom left, and a teal bottle with a yellow base at the bottom right. The overall aesthetic is clean and modern.

**02**

# **Model Selection**

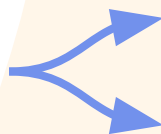
**Model Creation Process**

# Models Tested

Models tested include: logistic regression, random forest, LDA, QDA, GLM CV, KNN, XGBoost, and GAM.



## Logistic Regression

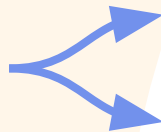


GLM with 6 predictors: sex, age, weight, height, hemoglobin, smoking status

Kaggle accuracy: 0.70513



## Random Forest

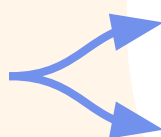


Random forest model with all predictors, 1000 trees, and 7 variables per tree

Kaggle accuracy: 0.72906

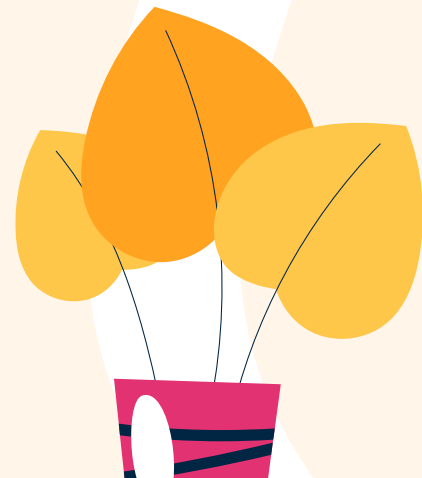


## GAM



GAM with degree 6

Kaggle accuracy: 0.73013



# Predictor Selection

## Step 1

Run backwards BIC stepwise regression to reduce predictors.

## Step 3

Run ANOVA between different degree GAMs with reduced & all predictors and analyze accuracy predicting training response variable.

## Step 2

Fit Generalized Additive Models to training data with degrees 3 & above.

## Final Model

GAM model with all predictors and degree 6 natural splines on numerical predictors.



# Pros and Cons of GAMs



## Interpretability

Easy to interpret



## Flexibility

Flexible functions can learn the distributions of predictors



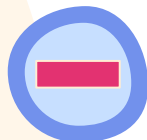
## Accuracy

Generalized Additive Models beat out many other methods with this data set



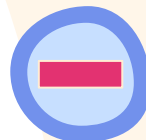
## Variability

The increase in flexibility of a GAM model also increases variability (especially at extremes)



## Numerical

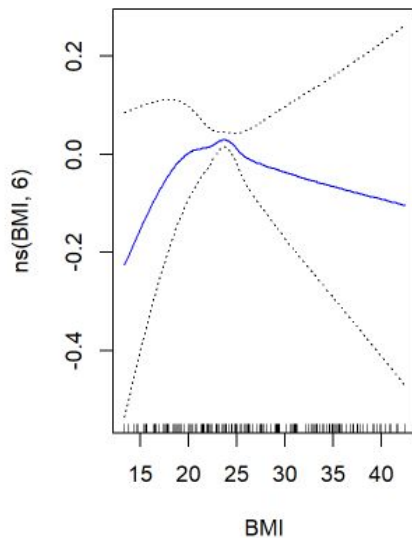
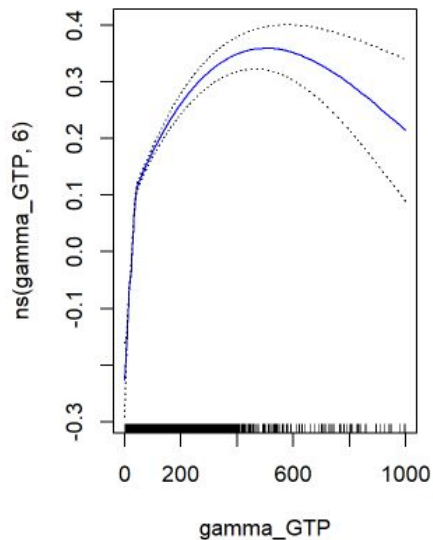
Splines only work with numerical predictors, categorical variables remain unchanged



## Choice

There are a number of different degrees and spline functions to choose from

# Predictor Scatterplots



Generalized Additive Models learn the non-linear distributions of the numerical predictors



**20  
predictors**

Number of numerical  
predictors



**19 predictors**

Number of statistically  
significant predictors

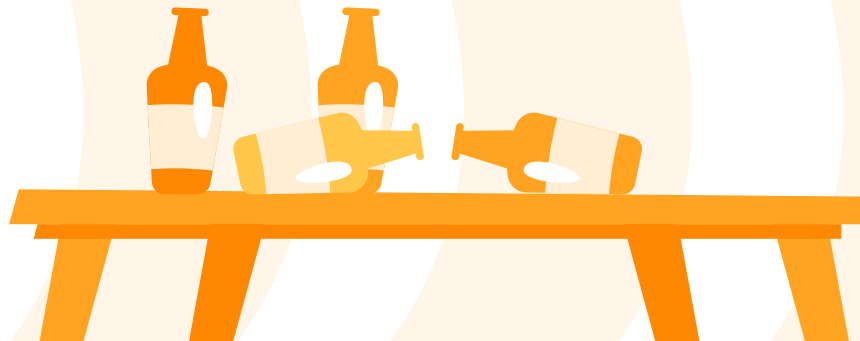
**70,000**

Total observations in  
training data

# ANOVA of GAM with Reduced & All Predictors

Model	Resid DF	Resid Dev	DF	Deviance	P(>Chi)
Red	69927	12649			
Full	69873	12589	54	59.441	<2.2 e-16

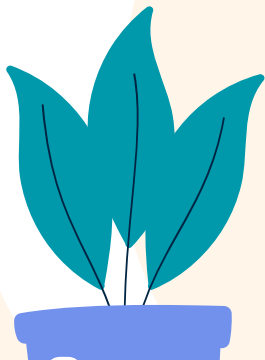
**ANOVA favors the GAM with all predictors with degree 6 over over a reduced degree 6 GAM**



# ANOVA of Degree 3 vs 6

ANOVA favors the higher degree GAM with degree 6 over the GAM with degree 3 splines

Model	Resid DF	Resid Dev	DF	Deviance	P(>Chi)
Deg 3	69930	12702			
Deg 6	69873	12589	57	112.69	<2.2e-16



# Accuracy Predicting Response in Training Data



**72.74%**

**GAM degree 3**

GAM model with all  
predictors



**72.77%**

**GAM degree 6**

GAM model with predictors  
reduced by stepwise  
regression (BIC)



**72.97%**

**GAM degree 6**

GAM model with all  
predictors



# Confusion Matrix of Accuracy Predictions Training Set

y

Accuracy of  
predictions  
compared to  
training data

72.97%

**Sensitivity**

Sensitivity of the  
model

73.23  
%

Predicted

	Yes	No
Yes	25,364	9,399
No	9,523	25,714



03

# Results

Discussion of Results and Model Description

# Final Model

GAM (with Degree 6)

# Predictors

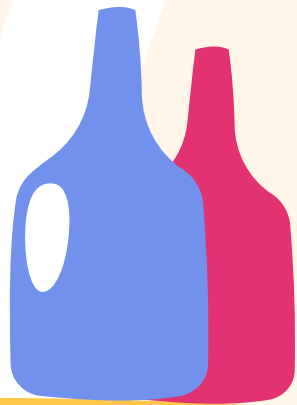
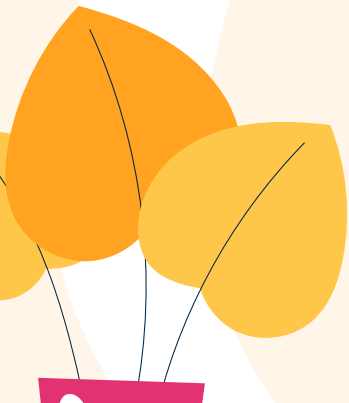
26 Predictors

# Observations

30,000 Alcoholic Statuses  
30x1

# MCR/Rank

MCR: 0.26987  
Rank: 30th



# Important Predictors

## Characteristics

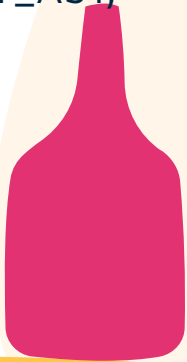
Sex, age, height, weight,  
wasteline

## Blood-Related

SBP, DBP, BLDS, tote\_chole, HDL\_chole,  
LDL\_chole, triglyceride, hemoglobin,  
serum\_creatinine, SGOT\_ALT, SGOT\_AST,  
gamma\_GTP, BMI

## Smoking

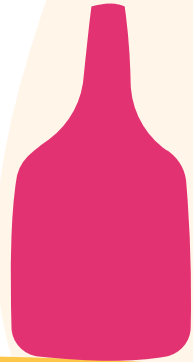
Smoking Status




# Most Important Predictors

Predictors: Sex, Smoking.Status, Hemoglobin, gamma\_GTP, height, age, triglyceride, HDL\_chole, and waistline

From our model, we found that these predictors were most essential at predicting the alcoholic status. These predictors contributed the most to accurately predicting a person's alcoholic status.



The background features a series of concentric, hand-drawn style circles in shades of light orange and cream. Scattered around these circles are several stylized, colorful bottle or flask shapes. There are three blue bottles, one yellow bottle, one pink bottle, and one teal bottle with a yellow base. The overall aesthetic is clean and modern.

**04**

# **Conclusion**

**Discussion/Limitations**

# Conclusion

Our most successful model was a GAM model using Hmisc imputations that had **73.01**% accuracy in predicting the alcoholic status for the test data.

We believe that if additional improvements were made to the imputation method for missing values, this model would be stronger and produce a higher accuracy rate.



# Limitations

## Model Complexity

- High Degree of 6
- Use of ALL Predictors
- Prone to Overfitting

## Data (NA and Features)

- Missing Values in Data
- Lack of Health Domain

Knowledge

## GAM Model Assumptions

- Linearity
- Independence
- Smoothness
- Homoscedasticity



# Better Data, Better Model

## Using Original Full Dataset

- Same Random Forest model on 70,000 vs 1.6 mil rows
- 72% Accuracy vs 89% Accuracy
- Data and Features are as Important as Model Selection

# Future Work

## Suggestions

- Regularization Techniques (PCA, L1/L2, CV) -> Simpler Model
- Feature Engineering with Domain Knowledge
- Better Imputation Methods: KNN
- Using Ensemble or Combination of Methods

The background features a series of concentric, hand-drawn style ovals in shades of light orange and cream. Scattered around the central text are five stylized bottle icons in various colors: light blue, orange, yellow, teal, and pink. The pink bottle on the right is tilted and has a yellow splash at its base.

# Thanks!

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