Interpretable Modelling of Credit Risk

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As detailed in Cynthia Rudin's excellent commentary on interpretability (ArXiV version here), there are a plethora of reasons to avoid the use of black box models when models are being used to make high stakes decisions to may have life-altering effects on real people. Efforts to develop "explainable black box models," while appealing for their potential to let us continuing using the same tools we always have and to creation explanations after the fact, are inherently flawed. As Rudin notes in my single favorite passage from her paper:

Explainable ML methods provide explanations that are not faithful to what the original model computes. Explanations must be wrong. They cannot have perfect fidelity with respect to the original model. If the explanation was completely faithful to what the original model computes, the explanation would equal the original model, and one would not need the original model in the first place, only the explanation. (In other words, this is a case where the original model would be interpretable.) This leads to the danger that any explanation method for a black box model can be an inaccurate representation of the original model in parts of the feature space.

An inaccurate (low-fidelity) explanation model limits trust in the explanation, and by extension, trust in the black box that it is trying to explain. An explainable model that has a 90% agreement with the original model indeed explains the original model most of the time. However, an explanation model that is correct 90% of the time is wrong 10% of the time. If a tenth of the explanations are incorrect, one cannot trust the explanations, and thus one cannot trust the original black box. If we cannot know for certain whether our explanation is correct, we cannot know whether to trust either the explanation or the original model.

With this motivation in mind, in this exercise, we will use a cutting edge interpretable modeling framework to model credit risk using data from the 14th Pacific-Asia Knowledge Discovery and Data Mining conference (PAKDD 2010). This data covers the period of 2006 to 2009, and "comes from a private label credit card operation of a Brazilian credit company and its partner shops." (The competition was won by TIMi, who purely by coincidence helped me complete my PhD dissertation research!).

We will be working with Generalized Additive Models (GAMs) (not to be confused with Generalized *Linear* Models (GLMs) — GLMs are a special case of GAMs). In particular, we will be using the pyGAM, though this is far from the only GAM implementation out there. mvgam in R is probably considered the gold standard, as it was developed by a pioneering researcher of GAMs.

statsmodels also has an implementation, and GAM is also hiding in plain sight behind many other tools, like Meta's Prophet time series forecasting library (which is GAM-based).

```
In []: # import xlrd
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.model_selection import GridSearchCV
from pygam import LogisticGAM, s, f
from pygam.datasets import default
from sklearn.metrics import accuracy_score
from sklearn.metrics import confusion_matrix
import warnings
warnings.filterwarnings("ignore")
```

Data Prep

Exercise 1

The PADD 2010 data is in this repository. You can find column names in PAKDD2010_VariablesList.XLS and the actual data in PAKDD2010_Modeling_Data.txt.

Note: you may run into a string-encoding issue loading the PAKDD2010_Modeling_Data.txt data. All I'll say is that most latin-based languages used latin8 as a text encoding prior to broad adoption of UTF-8. (Don't know about UTF? Check out this video!)

columns_file_path = "https://raw.githubusercontent.com/nickeubank/MIDS_Data/master/PAKDD%202010/PAKDD2010_Varia

Load the data (including column names).

df_columns = pd.read_excel(columns_file_path, header=None)

In []: # column names

```
column_names = df_columns[1].tolist()
In [ ]: # find duplicate
        for i in range(len(column_names)):
             if column_names[i] in column_names[:i]:
                 print(column_names[i])
        # Two EDUCATION_LEVEL
        # Keep the first one to be EDUCATION_LEVEL
        # Change the seconde one to MATE_EDUCATION_LEVEL
        second_edu_level_idx = [
             i for i, value in enumerate(column_names) if value == "EDUCATION_LEVEL"
        1[-1]
        column names[second edu level idx] = "MATE EDUCATION LEVEL"
       EDUCATION LEVEL
In [ ]: # data
        data_file_path = "https://media.githubusercontent.com/media/nickeubank/MIDS_Data/master/PAKDD%202010/PAKDD2010_
        df_data = pd.read_csv(
             data_file_path, encoding="latin1", sep="\t", header=None, names=column_names[1:]
In [ ]: df_data.head()
Out[]:
            ID_CLIENT CLERK_TYPE PAYMENT_DAY APPLICATION_SUBMISSION_TYPE QUANT_ADDITIONAL_CARDS POSTAL_ADDRES
         0
                    1
                                 С
                                                5
                                                                              Web
                                                                                                            0
                    2
                                                15
                                                                             Carga
                                                                                                            0
         2
                    3
                                 С
                                                5
                                                                              Web
                                                                                                            0
                                 С
                                                                                                            0
         3
                    4
                                               20
                                                                              Web
         4
                    5
                                 С
                                               10
                                                                              Web
                                                                                                            0
        5 rows × 54 columns
In [ ]: df_columns.head()
Out[]:
                0
                                              1
                                                                                        2
                                                                                                                         3
         0 Var_ld
                                                                            Var_Description
                                                                                                               Field_Content
                                        Var_Title
                                       ID_CLIENT Sequential number for the applicant (to be use... 1-50000, 50001-70000, 70001-90000
         1
                1
         2
                2
                                    CLERK_TYPE
                                                                              Not informed
                                                                                                                         С
         3
                                   PAYMENT_DAY
                                                 Day of the month for bill payment, chosen by t...
                                                                                                              1,5,10,15,20,25
         4
                4 APPLICATION_SUBMISSION_TYPE
                                                  Indicates if the application was submitted via...
                                                                                                                 Web, Carga
```

Exercise 2

There are a few variables with a lot of missing values (more than half missing). Given the limited documentation for this data it's a little hard to be sure why, but given the effect on sample size and what variables are missing, let's go ahead and drop them. You you end up dropping 6 variables.

Hint: Some variables have missing values that aren't immediately obviously.

(This is not strictly necessary at this stage, given we'll be doing more feature selection down the line, but keeps things easier knowing we don't have to worry about missingness later.)

```
In [ ]: # df_data.describe()
In [ ]: df data["APPLICATION SUBMISSION TYPE"] = df data["APPLICATION SUBMISSION TYPE"].replace(
            "0", np.nan
        df_data["SEX"] = df_data["SEX"].replace(["N", " "], np.nan)
        df_data["MARITAL_STATUS"] = df_data["MARITAL_STATUS"].replace(0, np.nan)
        df_data["OCCUPATION_TYPE"] = df_data["OCCUPATION_TYPE"].replace(0.0, np.nan)
        df_data["RESIDENCE_TYPE"] = df_data["RESIDENCE_TYPE"].replace(0.0, np.nan)
        df_data["RESIDENCIAL_ZIP_3"] = df_data["RESIDENCIAL_ZIP_3"].replace("#DIV/0!", np.nan)
        df_data["PROFESSIONAL_STATE"] = df_data["PROFESSIONAL_STATE"].replace(" ", np.nan)
        df_data["PROFESSIONAL_PHONE_AREA_CODE"] = df_data[
            "PROFESSIONAL PHONE AREA CODE"
        ].replace(" ", np.nan)
In [ ]: # df data.isna().sum().sort values(ascending=False)
In []: missing = df data.isna().mean()
        missing_50_col = list(missing[missing > 0.5].index)
        missing_50_col
Out[]: ['PROFESSIONAL STATE',
          'PROFESSIONAL CITY'
          'PROFESSIONAL BOROUGH',
          'PROFESSIONAL_PHONE_AREA_CODE',
          'MATE_PROFESSION_CODE',
          'MATE_EDUCATION_LEVEL']
In []: # drop
        df1 = df_data.drop(columns=missing_50_col)
        # df1
```

Exercise 3

Let's start off by fitting a model that uses the following variables:

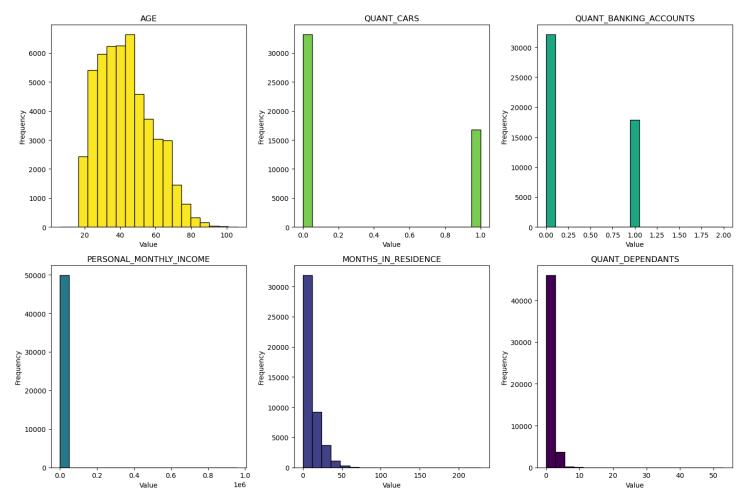
```
"QUANT_DEPENDANTS",
"QUANT_CARS",
"MONTHS_IN_RESIDENCE",
"PERSONAL_MONTHLY_INCOME",
"QUANT_BANKING_ACCOUNTS",
"AGE",
"SEX",
"MARITAL_STATUS",
"OCCUPATION_TYPE",
"RESIDENCE_TYPE",
"RESIDENCIAL_STATE",
"RESIDENCIAL_CITY",
"RESIDENCIAL_BOROUGH",
"RESIDENCIAL_ZIP 3"
```

(GAMs don't have any automatic feature selection methods, so these are based on my own sense of features that are likely to matter. A fully analysis would entail a few passes at feature refinement)

Plot and otherwise characterize the distributions of all the variables we may use. If you see anything bananas, adjust how terms enter your model. Yes, pyGAM has flexible functional forms, but giving the model features that are engineered to be more substantively meaningful (e.g., taking log of income) will aid model estimation.

You should probably do something about the functional form of *at least* PERSONAL_MONTHLY_INCOME , and QUANT_DEPENDANTS .

```
In [ ]: df1 = df1[
                "QUANT_DEPENDANTS",
                "QUANT_CARS",
                "MONTHS_IN_RESIDENCE",
                "PERSONAL_MONTHLY_INCOME",
                "QUANT_BANKING_ACCOUNTS",
                "AGE",
                "SEX",
                "MARITAL_STATUS",
                "OCCUPATION_TYPE",
                "RESIDENCE_TYPE",
                "RESIDENCIAL_STATE",
                "RESIDENCIAL_CITY",
                "RESIDENCIAL_BOROUGH",
                "RESIDENCIAL_ZIP_3",
                "TARGET_LABEL_BAD=1",
In [ ]: # df1.isna().sum()
In [ ]: # df1.info()
In [ ]: import matplotlib.pyplot as plt
        import numpy as np
        quant_vars = [
            "AGE",
            "QUANT CARS",
            "QUANT BANKING ACCOUNTS",
            "PERSONAL_MONTHLY_INCOME",
            "MONTHS_IN_RESIDENCE",
            "QUANT_DEPENDANTS",
        colors = ["#fde725", "#7ad151", "#22a884", "#2a788e", "#414487", "#440154"]
        fig, axes = plt.subplots(2, 3, figsize=(15, 10))
        axes = axes.flatten()
        for i, (column, color) in enumerate(zip(quant_vars, colors)):
            ax = axes[i]
            ax.hist(
                df1[column],
                bins=np.linspace(df1[column].min(), df1[column].max(), 20),
                color=color,
                edgecolor="black",
            ax.set_title(column)
            ax.set_xlabel("Value")
            ax.set_ylabel("Frequency")
        plt.tight_layout()
        plt.show()
```



```
In [ ]: import matplotlib.pyplot as plt
        import numpy as np
        qual_vars2 = [
            "SEX",
            "MARITAL_STATUS",
            "OCCUPATION_TYPE",
            "RESIDENCE_TYPE",
            "RESIDENCIAL_STATE",
            # "RESIDENCIAL_BOROUGH",
        fig, axes = plt.subplots(2, 3, figsize=(20, 10))
        axes = axes.flatten()
        cmap = plt.get_cmap("viridis")
        for i, var in enumerate(qual_vars2):
            ax = axes[i]
            var_counts = df1[var].value_counts()
            categories = var_counts.index
            colors = cmap(np.linspace(0, 1, len(categories)))
            var_counts.plot(kind="bar", ax=ax, color=colors, edgecolor="black")
            ax.set_title(var)
            ax.set_xlabel("Category")
            ax.set_ylabel("Count")
        plt.tight_layout()
        plt.show()
```

```
20000
                                              15000
15000
                                                                                         10000
       Š 15000
                                                                                         8000
                                                10000
                                                                                         6000
        10000
                                                                                         4000
                                                 5000
                                                                                         2000
                                                      2.0
                        RESIDENCE_TYPE
                                                                RESIDENCIAL_STATE
                                                                                          1.0
        35000
                                                                                          0.8
        30000
                                                                                          0.6
        25000
                                                                                          0.4
                                                                                          0.2
         5000
                                                              In [ ]: # Change to categorical variable
         for i in ["MARITAL_STATUS", "OCCUPATION_TYPE", "RESIDENCE_TYPE"]:
             df1[i] = df1[i].astype("Int64").astype("category")
         for i in ["SEX", "RESIDENCIAL_STATE", "RESIDENCIAL_CITY", "RESIDENCIAL_ZIP_3"]:
             df1[i] = df1[i].astype("category")
In []:
        # functional form of `PERSONAL_MONTHLY_INCOME` and `QUANT_DEPENDANTS`
         df1["LOG_PERSONAL_MONTHLY_INCOME"] = df1["PERSONAL_MONTHLY_INCOME"].apply(
             lambda x: np.log(x + 1)
         df1["LOG_MONTHS_IN_RESIDENCE"] = df1["MONTHS_IN_RESIDENCE"].apply(
             lambda x: np.log(x + 1)
         # QUANT DEPENDANTS
         df1["QUANT_DEPENDANTS_BINARY"] = df1["QUANT_DEPENDANTS"].apply(
             lambda x: 1 if x > 2 else 0
In [ ]: df1.drop(columns=["PERSONAL_MONTHLY_INCOME", "MONTHS_IN_RESIDENCE"], inplace=True)
In [ ]: # df1.info()
```

MARITAL_STATUS

20000

OCCUPATION TYPE

16000

14000

Exercise 4

SEX

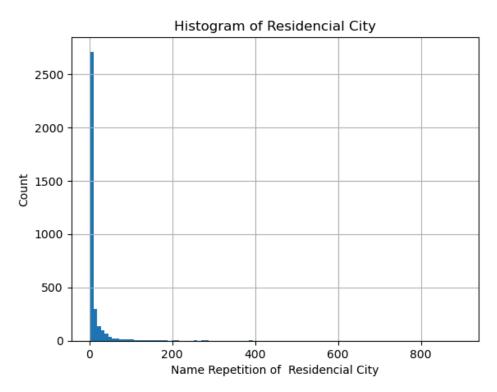
30000

25000

Geographic segregation means residency data often contains LOTS of information. But there's a problem with RESIDENCIAL_CITY and RESIDENCIAL_BOROUGH . What is the problem?

In any real project, this would be something absolutely worth resolving, but for this exercise, we'll just drop all three string RESIDENCIAL_ variables.

```
In []: df1["RESIDENCIAL_CITY"].value_counts().hist(bins=100)
    plt.title("Histogram of Residencial City")
    plt.xlabel("Name Repetition of Residencial City")
    plt.ylabel("Count")
    plt.show()
```



```
In [ ]: qual_vars = [
            "SEX",
            "MARITAL_STATUS",
            "OCCUPATION_TYPE",
            "RESIDENCE_TYPE",
            "RESIDENCIAL_STATE",
            "RESIDENCIAL_CITY",
            "RESIDENCIAL_BOROUGH",
            "RESIDENCIAL_ZIP_3",
        ]
        for column_name in qual_vars:
            total_unique = df1[column_name].nunique()
            print(
                f"Total number of unique observations in the '{column_name}' variable:",
                total_unique,
       Total number of unique observations in the 'SEX' variable: 2
       Total number of unique observations in the 'MARITAL_STATUS' variable: 7
       Total number of unique observations in the 'OCCUPATION_TYPE' variable: 5
       Total number of unique observations in the 'RESIDENCE_TYPE' variable: 5
       Total number of unique observations in the 'RESIDENCIAL_STATE' variable: 27
       Total number of unique observations in the 'RESIDENCIAL_CITY' variable: 3529
```

The total number of unique observations in the 'RESIDENCIAL_CITY' is 3529, and for 'RESIDENCIAL_BOROUGH' is 14511. By doing a review of these unique values, we can see that most of them point to the same borough but with different spelling, use of caps or use of complete name. These lack of consistency on spelling results in different observations in the dataset that refer to the same city or borough.

Total number of unique observations in the 'RESIDENCIAL_BOROUGH' variable: 14511 Total number of unique observations in the 'RESIDENCIAL_ZIP_3' variable: 1480

```
In []: # drop RESIDENCIAL_ variable
    new_df = df1.filter(regex=r"^(?!RESIDENCIAL_).*")
# new_df

In []: # drop missing values
    new_df = new_df.drop(columns="QUANT_DEPENDANTS").dropna()

In []: # solve categorical data
    for var in [
        "QUANT_CARS",
```

```
"QUANT_BANKING_ACCOUNTS",
   "QUANT_DEPENDANTS_BINARY",
]:
    new_df[var] = new_df[var].astype("category")

categorical_variables = [
    "QUANT_CARS",
    "QUANT_BANKING_ACCOUNTS",
    "SEX",
    "MARITAL_STATUS",
    "OCCUPATION_TYPE",
    "RESIDENCE_TYPE",
    "QUANT_DEPENDANTS_BINARY",
]

for var in categorical_variables:
    new_df[var] = new_df[var].cat.codes
```

```
In [ ]: # new_df.info()
```

Model Fitting

In []: |y_pred = gam.predict(X_test.to_numpy())

Exercise 5

First, use train_test_split to do an 80/20 split of your data. Then, using the TARGET_LABEL_BAD variable, fit a classification model on this data. Optimize with gridsearch. Use splines for continuous variables and factors for categoricals.

At this point we'd *ideally* be working with 11 variables. However pyGAM can get a little slow with factor features with lots of values + lots of unique values (e.g., 50,000 observations and the *many* values of RESIDENCIAL_ZIP takes about 15 minutes on my computer). In that configuration, you should get a model fit in 10-15 seconds.

So let's start by fitting a model that also excludes RESIDENCIAL_ZIP.

```
In [ ]: # Train test split\
        X = new_df.loc[:, new_df.columns != "TARGET_LABEL_BAD=1"]
        y = new_df["TARGET_LABEL_BAD=1"]
        X_train, X_test, y_train, y_test = train_test_split(
            X, y, test_size=0.2, random_state=42
In [ ]: X_train_arr = X_train.to_numpy()
        y_train_arr = y_train.to_numpy()
        # lambdas
        lams = np.logspace(-3, 3, 4)
        gam = LogisticGAM(f(0) + f(1) + s(2) + f(3) + f(4) + f(5) + f(6) + s(7) + s(8) + f(9))
        gam.gridsearch(X_train_arr, y_train_arr, lam=lams)
         0% (0 of 4) |
                                                | Elapsed Time: 0:00:00 ETA:
        25% (1 of 4) |#####
                                                | Elapsed Time: 0:00:01 ETA:
        50% (2 of 4) |############
                                                | Elapsed Time: 0:00:02 ETA:
                                                                               0:00:02
        75% (3 of 4) |################
                                                | Elapsed Time: 0:00:03 ETA:
                                                                               0:00:01
       100% (4 of 4) | #################### | Elapsed Time: 0:00:04 Time: 0:00:04
Out[]: LogisticGAM(callbacks=[Deviance(), Diffs(), Accuracy()],
           fit intercept=True, max iter=100,
           terms=f(0) + f(1) + s(2) + f(3) + f(4) + f(5) + f(6) + s(7) + s(8) + f(9) + intercept,
           tol=0.0001, verbose=False)
In [ ]: print("The best lambda (grid search) is:")
        print(gam.lam[0][0])
       The best lambda (grid search) is:
       10.0
```

```
print("Accuracy:", accuracy_score(y_test, y_pred))
```

Accuracy: 0.7404054054054054

Exercise 6

Create a (naive) confusion matrix using the predicted values you get with predict() on your test data. Our stakeholder cares about two things:

- · maximizing the number of people to whom they extend credit, and
- the false negative rate (the share of people identified as "safe bets" who aren't, and who thus default).

How many "good bets" does the model predict (true negatives), and what is the False Omission Rate (the share of predicted negatives that are false negatives)?

Looking at the confusion matrix, how did the model maximize accuracy?

```
In []: def cm_for(y_test, y_pred, process=True):
    ConfuMat = confusion_matrix(y_test, y_pred)
    TN, FP, FN, TP = ConfuMat.ravel()
    FOR = FN / (FN + TN) if (FN + TN) > 0 else 0
    if process:
        print(f"TN, FP, FN, TP: {TN, FP, FN, TP}")
        print(f"False Omission Rate: {FOR} ")

    return (TN, FP, FN, TP), FOR
```

```
In [ ]: col, FOR = cm_for(y_test, y_pred)
TN, FP, FN, TP: (5474, 1, 1920, 5)
False Omission Rate: 0.2596700027048959
```

There are 5474 "good bets" or True Negatives (people who would not default and thus we would extend credit)

The False Omission Rate, or predicted negatives that are false negatives, is around 0.2597. Thus the proportion that were 'safe' bets but that would default over all the predicted negatives.

The model maximizes the accuracy through predicting more people who would not default and minimizing the false positives. In the training set, there are more people whou would not default (TN+FN=5474+1920=7394)than would default, so the model predicts more majority class in the testing period and less minority class (7394 vs. 6).

Exercise 7

Suppose your stakeholder wants to minimize false negative rates. How low of a False Omission Rate (the share of predicted negatives that are false negatives) can you get (assuming more than, say, 10 true negatives), and how many "good bets" (true negatives) do they get at that risk level?

```
Hint: use predict_proba()
```

Note: One *can* use class weights to shift the emphasis of the original model fitting, but for the moment let's just play with predict_proba() and thresholds.

```
In []: # predict probability
y_prob = gam.predict_proba(X_test)

# Find the best threshold to minimize FOR and maximize Good Bets
# from 0% to 100%
thre_range = np.linspace(0, 1, 101)
GB = []
FORS = []

for thre in thre_range:
    y_pred_thre = (y_prob > thre).astype("int")

    (TN, FP, FN, TP), FOR = cm_for(y_test, y_pred_thre, process=False)
```

```
FORS.append(FOR)
In [ ]: # good bets
         plt.plot(thre_range, GB)
         plt.xlabel("Threshold")
         plt.ylabel("True Negative")
         plt.show()
           5000
           4000
        True Negative
           3000
          2000
           1000
              0
                   0.0
                               0.2
                                            0.4
                                                         0.6
                                                                     0.8
                                                                                  1.0
                                               Threshold
In [ ]: plt.plot(thre_range, FORS)
         plt.xlabel("Threshold")
         plt.ylabel("False Omission Rate")
         plt.show()
           0.40
           0.35
           0.30
       False Omission Rate
           0.25
           0.20
           0.15
           0.10
           0.05
           0.00
                               0.2
                                           0.4
                                                                     0.8
                  0.0
                                                        0.6
                                                                                  1.0
                                              Threshold
In [ ]: # Find min FOR and corresponding good bets
         min_FOR = min(x for x in FORS if x > 0.0)
```

min_indices = [i for i, x in enumerate(FORS) if x == min_FOR]

GB.append(TN)

```
corresponding_GB = [GB[i] for i in min_indices]
print(f"Best threshold: {thre_range[min_indices][0]}")
print(f"Minimum FOR: {min_FOR}")
print(f"Corresponding GB values: {corresponding_GB[0]}")
```

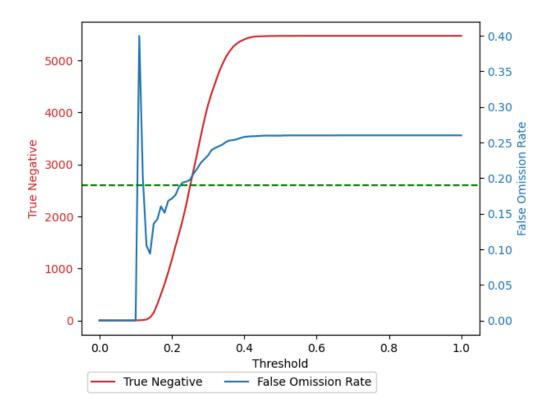
Best threshold: 0.14 Minimum FOR: 0.09375 Corresponding GB values: 58

We can minimize the False Omission Rate through changing the threshold value, the best threshold is 0.14. According to the result, the minimum False Omission Rate is 0.09375, where the corresponding "good bets" or True Negatives is 58.

Exercise 8

If the stakeholder wants to maximize true negatives and can tolerate a false omission rate of 19%, how many true negatives will they be able to enroll?

```
In [ ]: fig, ax1 = plt.subplots()
        color = "tab:red"
        ax1.set xlabel("Threshold")
        ax1.set_ylabel("True Negative", color=color)
        ax1.plot(thre_range, GB, color=color, label="True Negative")
        ax1.tick_params(axis="y", labelcolor=color)
        ax2 = ax1.twinx()
        color = "tab:blue"
        ax2.set_ylabel("False Omission Rate", color=color)
        ax2.plot(thre_range, FORS, color=color, label="False Omission Rate")
        ax2.tick_params(axis="y", labelcolor=color)
        ax2.axhline(y=0.19, color="green", linestyle="--")
        lines, labels = ax1.get_legend_handles_labels()
        lines2, labels2 = ax2.get_legend_handles_labels()
        ax2.legend(
            lines + lines2, labels + labels2, loc="upper left", bbox_to_anchor=(0, -0.1), ncol=3
        fig.tight_layout()
        plt.show()
```



```
In []: sele_FOR = [x for x in FORS if x > 0.0 and x < 0.19]
    sele_indices = [i for i, x in enumerate(FORS) if x in sele_FOR]
    sele_corresponding_GB = [GB[i] for i in sele_indices]
    max_sele_gb = max(sele_corresponding_GB)

In []: print(f"The max true negatives needed to be enrolled: {max_sele_gb}")</pre>
```

The max true negatives needed to be enrolled: 1670

Let's See This Interpretability!

We're using GAMs for their interpretability, so let's use it!

Exercise 9

Plot the partial dependence plots for all your continuous factors with 95% confidence intervals (I have three, at this stage).

If you get an error like this when generating partial_dependence errors:

```
----> pdep, confi = gam.partial_dependence(term=i, X=XX, width=0.95)
```

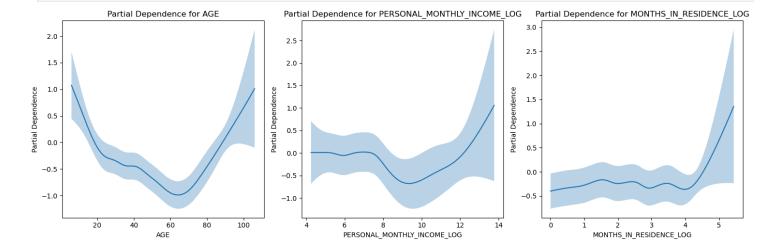
ValueError: X data is out of domain for categorical feature 4. Expected data on [1.0, 2.0], but found data on [0.0, 0.0]

it's because you have a variable set as a factor that doesn't have values of 0 . pyGAM is assuming 0 is the excluded category. Just recode the variable to ensure 0 is used to identify one of the categories.

```
cont_vars = {"AGE": 2, "PERSONAL_MONTHLY_INCOME_LOG": 7, "MONTHS_IN_RESIDENCE_LOG": 8}

In []: # Plot the partial dependence plots
fig, axs = plt.subplots(1, 3, figsize=(15, 5))

for ax, (feature, i) in zip(axs, cont_vars.items()):
    XX = gam.generate_X_grid(term=i)
    pdep, confi = gam.partial_dependence(term=i, X=XX, width=0.95)
    ax.plot(XX[:, i], pdep)
    ax.fill_between(XX[:, i], confi[:, 0], confi[:, 1], alpha=0.3)
    ax.set_xlabel(feature)
    ax.set_ylabel("Partial Dependence")
    ax.set_title(f"Partial Dependence for {feature}")
```



Exercise 10

continuous (float)

plt.tight_layout()

plt.show()

How does the partial correlation with respect to age look?

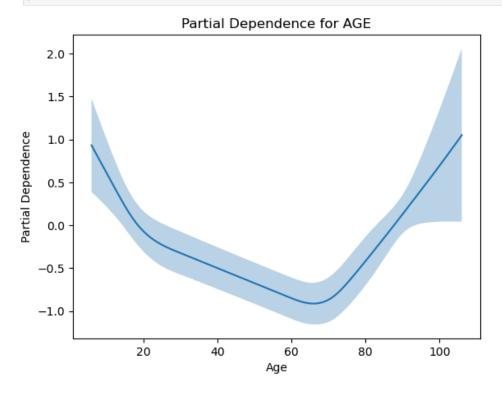
Based on the plot of the partial dependence of age, we can find that the partial correlation is "U-shape". It reveals that as the age increase, the likelihood of default decreases. At the age around 65, the result reaches the lowest likelihood to default. After reaching 65, the trend reverses, and the likelihood of default starts to increase with further increases in age.

Exercise 11

Refit your model, but this time impose monotonicity or concavity/convexity on the relationship between age and credit risk (which makes more sense to you?). Fit the model and plot the new partial dependence.

According to the "U-shape" curve, we should use convexity.

```
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Out[]: LogisticGAM(callbacks=[Deviance(), Diffs(), Accuracy()],
           fit_intercept=True, max_iter=100,
           terms = f(0) + f(1) + s(2) + f(3) + f(4) + f(5) + f(6) + s(7) + s(8) + f(9) + intercept
           tol=0.0001, verbose=False)
In []: XX = gam.generate_X_grid(term=2)
        pdep, confi = gam 1.partial dependence(term=2, X=XX, width=0.95)
        plt.plot(XX[:, 2], pdep)
        plt.fill_between(XX[:, 2], confi[:, 0], confi[:, 1], alpha=0.3)
        plt.xlabel("Age")
        plt.ylabel("Partial Dependence")
        plt.title(f"Partial Dependence for AGE")
        plt.show()
```



Exercise 12

Functional form constraints are often about fairness or meeting regulatory requirements, but they can also prevent overfitting.

Does this change the number of "true negatives" you can enroll below a false omission rate of 19%?

```
In []: # predict
y_prob1 = gam_1.predict_proba(X_test)

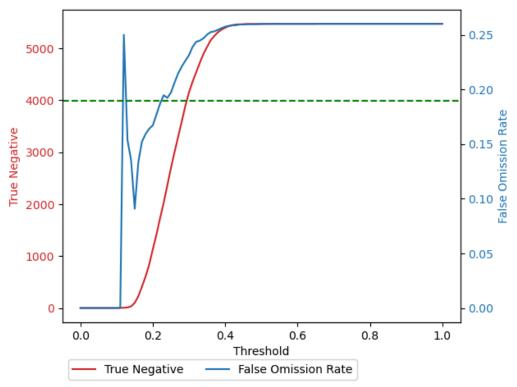
# Find the best threshold to minimize FOR and maximize Good Bets
# from 0% to 100%
thre_range = np.linspace(0, 1, 101)
GB1 = []
FORS1 = []

for thre in thre_range:
    y_pred_thre = (y_prob1 > thre).astype("int")

    (TN, FP, FN, TP), FOR = cm_for(y_test, y_pred_thre, process=False)

    GB1.append(TN)
    FORS1.append(FOR)
```

```
In [ ]: fig, ax1 = plt.subplots()
        color = "tab:red"
        ax1.set_xlabel("Threshold")
        ax1.set_ylabel("True Negative", color=color)
        ax1.plot(thre_range, GB1, color=color, label="True Negative")
        ax1.tick_params(axis="y", labelcolor=color)
        ax2 = ax1.twinx()
        color = "tab:blue"
        ax2.set_ylabel("False Omission Rate", color=color)
        ax2.plot(thre_range, FORS1, color=color, label="False Omission Rate")
        ax2.tick_params(axis="y", labelcolor=color)
        ax2.axhline(y=0.19, color="green", linestyle="--")
        lines, labels = ax1.get_legend_handles_labels()
        lines2, labels2 = ax2.get_legend_handles_labels()
        ax2.legend(
            lines + lines2, labels + labels2, loc="upper left", bbox_to_anchor=(0, -0.1), ncol=3
        fig.tight_layout()
        plt.show()
```



```
In []: sele_FOR1 = [x for x in FORS1 if x > 0.0 and x < 0.19]
    sele_indices1 = [i for i, x in enumerate(FORS1) if x in sele_FOR1]
    sele_corresponding_GB1 = [GB1[i] for i in sele_indices1]
    max_sele_gb1 = max(sele_corresponding_GB1)
    print(f"The max true negatives needed to be enrolled: {max_sele_gb1}")</pre>
```

The max true negatives needed to be enrolled: 1727

Yes. The number of "true negatives" we can enroll below FOR of 19% changes from 1670 to 1727.

In the preceding exercises, we allowed pyGAM to choose its own smoothing parameters / coefficient penalties. This makes life easy, but it isn't always optimal, especially because when it does so, it picks the same smoothing penalty (the lambda in summary()) for all terms.

(If you haven't seen them let, penalities are designed to limit overfitting by, basically, "penalizing" big coefficients on different terms. This tends to push models towards smoother fits.)

To get around this, we can do a grid or random search. This is definitely a little slow, but let's give it a try!

Then following the model given in the docs linked above, let's do a random search. Make sure your initial random points has a shape of $100 \times (\text{the number of terms in your model})$.

```
In []: # 10 terms
        num_terms = 10
        init_lams = 10 ** (np.random.rand(100, num_terms) * 6 - 3)
In [ ]: # modeling
        loggam = LogisticGAM(
            f(0)
            + f(1)
            + s(2, constraints="convex")
            + f(3)
            + f(4)
            + f(5)
            + f(6)
            + s(7)
            + s(8)
            + f(9)
        ).gridsearch(X_train_arr, y_train_arr, lam=init_lams)
```

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```

In []: # model summary
loggam.summary()

LogisticGAM

Distribution: BinomialDist Effective DoF: 36.2386 Link Function: LogitLink Log Likelihood: -16631.1563 Number of Samples: 29599 AIC: 33334.7899 AICc: 33334.8862 UBRE: 3.1272 Scale: 1.0 Pseudo R-Squared: 0.0185

Feature Function	Lambda	Rank	EDoF	P > x	Sig. Code
	=======================================				== =======
f(0)	[0.2025]	2	2.0	2.73e-01	
f(1)	[151.0413]	3	0.8	1.64e-01	
s(2)	[0.0015]	20	4.7	0.00e+00	***
f(3)	[0.0014]	2	1.0	1.17e-06	***
(4)	[0.0748]	7	6.0	6.75e-12	***
(5)	[0.0491]	5	3.9	2.02e-09	***
(6)	[0.4062]	5	4.0	1.22e-02	*
(7)	[55.0576]	20	4.3	4.65e-03	**
(8)	[9.9562]	20	8.6	3.25e-03	**
(9)	[0.0065]	2	1.0	3.48e-04	***
ntercept		1	0.0	4.88e-01	

Significance codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1

WARNING: Fitting splines and a linear function to a feature introduces a model identifiability problem which can cause p-values to appear significant when they are not.

WARNING: p-values calculated in this manner behave correctly for un-penalized models or models with known smoothing parameters, but when smoothing parameters have been estimated, the p-values are typically lower than they should be, meaning that the tests reject the null too readily.

Exercise 14

How many true negatives can you get now at a less than 19% False Omission Rate?

```
thre_range = np.linspace(0, 1, 101)
GB1 = []
FORS1 = []

for thre in thre_range:
    y_pred_thre = (y_prob1 > thre).astype("int")
    (TN, FP, FN, TP), FOR = cm_for(y_test, y_pred_thre, process=False)
    GB1.append(TN)
    FORS1.append(FOR)

sele_FOR1 = [x for x in FORS1 if x > 0.0 and x < threshold]
sele_indices1 = [i for i, x in enumerate(FORS1) if x in sele_FOR1]
sele_corresponding_GB1 = [GB1[i] for i in sele_indices1]
max_sele_gb1 = max(sele_corresponding_GB1)
print(f"The max true negatives needed to be enrolled: {max_sele_gb1}")</pre>
```

```
In [ ]: find_tn(loggam, X_test, 0.19)
```

The max true negatives needed to be enrolled: 1717

Exercise 15

Add an interaction term between age and personal income.

```
In [ ]: from pygam import LogisticGAM, s, te
        # 12 terms
        num\_terms = 12
        init_lams = 10 ** (np.random.rand(100, num_terms) * 6 - 3)
        # fit interaction
        gam_2 = LogisticGAM(
            f(0)
            + f(1)
            + s(2, constraints="convex")
            + f(3)
            + f(4)
            + f(5)
            + f(6)
            + s(7)
            + s(8)
            + f(9)
            + te(2, 7)
        gam_2.gridsearch(X_train_arr, y_train_arr, lam=init_lams)
```

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```

```
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```

In []: # model summary gam_2.summary()

LogisticGAM

Distribution: BinomialDist Effective DoF: 32.8102 Link Function: LogitLink Log Likelihood: -16622.4071Number of Samples: 29599 AIC: 33310.4347 AICc: 33310.5143 UBRE: 3.1263 Scale: 1.0 Pseudo R-Squared: 0.019

Feature Function	Lambda	Rank	EDoF	P > x	Sig. Code
======================================	[0.0015]	= ======= 2	2.0	4.35e-01	== =======
f(1)	[0.1617]	3	2.0	1.79e-01	
s(2)	[0.8744]	20	5.3	0.00e+00	***
f(3)	[5.0818]	2	1.0	9.24e-07	***
f(4)	[148.2355]	7	3.5	1.06e-12	***
f(5)	[0.008]	5	4.0	1.85e-09	***
f(6)	[0.0013]	5	4.0	3.03e-02	*
s(7)	[34.0676]	20	4.9	1.34e-03	**
5(8)	[523.0063]	20	4.0	2.71e-03	**
f(9)	[1.473]	2	1.0	6.64e-04	***
te(2, 7)	[44.91 38.8613]	100	1.2	1.26e-05	***
intercept		1	0.0	3.90e-01	

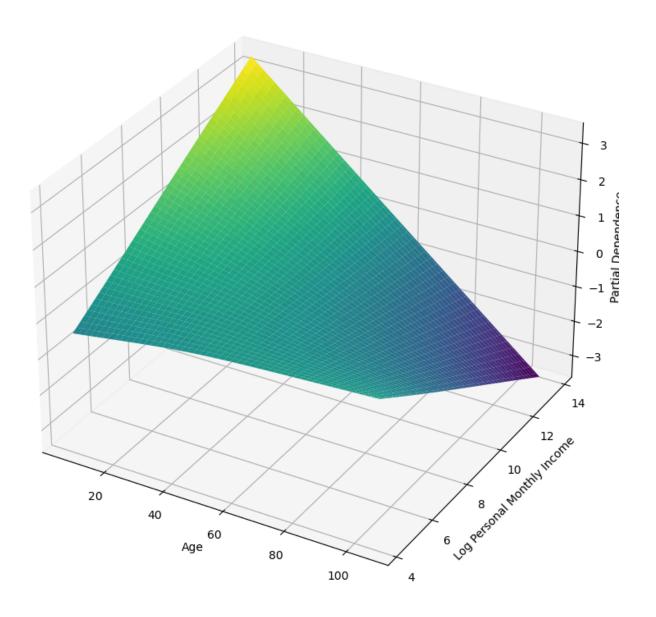
Significance codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1

WARNING: Fitting splines and a linear function to a feature introduces a model identifiability problem which can cause p-values to appear significant when they are not.

WARNING: p-values calculated in this manner behave correctly for un-penalized models or models with known smoothing parameters, but when smoothing parameters have been estimated, the p-values are typically lower than they should be, meaning that the tests reject the null too readily.

```
In []: plt.rcParams["figure.figsize"] = (12, 8)
    XX = gam_2.generate_X_grid(term=10, meshgrid=True)
    Z = gam_2.partial_dependence(term=10, X=XX, meshgrid=True)
    ax = plt.axes(projection="3d")
    ax.plot_surface(XX[0], XX[1], Z, cmap="viridis")
    ax.set_ylabel("Log Personal Monthly Income")
    ax.set_xlabel("Age")
    ax.set_zlabel("Partial Dependence", labelpad=1)
    ax.set_title("Partial Dependence of Interaction Term (Age and Income)")
    plt.tight_layout()
    plt.show()
```

Partial Dependence of Interaction Term (Age and Income)



Exercise 17

Finally, another popular interpretable model is the ExplainableBoostingClassifier. You can learn more about it here, though how much sense it will make to you may be limited if you aren't familiar with gradient boosting yet. Still, at least one of your classmates prefers it to pyGAM, so give it a try using this code:

```
from interpret.glassbox import ExplainableBoostingClassifier
from interpret import show
import warnings
```

```
ebm = ExplainableBoostingClassifier()
        ebm.fit(X_train, y_train)
        with warnings.catch_warnings():
           warnings.simplefilter("ignore")
           ebm_global = ebm.explain_global()
           show(ebm_global)
           ebm_local = ebm.explain_local(X_train, y_train)
           show(ebm_local)
In [ ]: from interpret.glassbox import ExplainableBoostingClassifier
        from interpret import show
        import warnings
        ebm = ExplainableBoostingClassifier()
        ebm.fit(X_train, y_train)
        with warnings.catch_warnings():
            warnings.simplefilter("ignore")
            ebm_global = ebm.explain_global()
            show(ebm_global)
            ebm_local = ebm.explain_local(X_train, y_train)
            show(ebm_local)
```





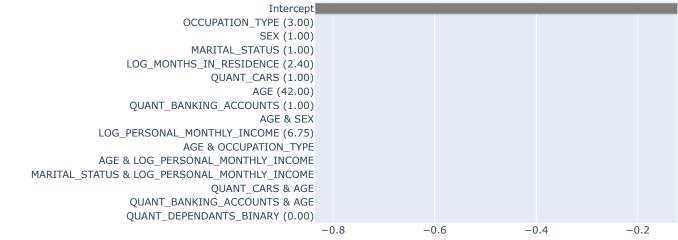
Select Component to Graph

0 : Actual (1) | Predicted (0) | PrScore (0.677)

× ▼

ExplainableBoostingClassifier_1 [0]

Local Explanation (Actual Class: 1 | Predicted Class: 0 Pr(y = 0): 0.677 | Pr(y = 1): 0.323)



Сс