# Project Overview

Analyze over 65,000 consumer complaints filed against insurance companies in Connecticut to determine insurer responsiveness.

### Data Cleaning:

Robustly handle missing values, drop duplicates, and ensure all necessary columns (e.g., Coverage, Reason, Recovery) are in the proper format.

## **Feature Engineering:**

Create key features such as a binary target (recovery\_binary) indicating if a complaint led to financial recovery.

(Optional) Incorporate advanced text features if a free-text complaint narrative becomes available.

#### **Supervised Models:**

Classification: Predict whether a complaint results in a monetary recovery.

Regression: Estimate the number of days to resolve a complaint.

#### **Unsupervised Analyses:**

Clustering: Use k-means with automatic k-selection based on silhouette score to reveal hidden patterns.

Anomaly Detection: Apply Isolation Forest with an adjustable contamination rate to identify outlier complaints.

**Outcome**: Transform raw complaint data into actionable insights that guide Connecticut residents toward more trustworthy insurance providers.

```
Data Loading & Basic Exploration
Explanation:
We load the data and check for shape, head, info, and stats.
We identify columns with >50% missing and drop them to avoid over-sparse features.
Adjust threshold as you see fit.
.....
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
# For transformations and modeling
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import LabelEncoder, OneHotEncoder, StandardScaler
from sklearn.metrics import (
    accuracy_score, precision_score, recall_score,
    mean_absolute_error, mean_squared_error, silhouette_score
from \ sklearn.ensemble \ import \ Random Forest Classifier, \ Isolation Forest
from sklearn.cluster import KMeans
from xgboost import XGBRegressor
from sklearn.feature_extraction.text import TfidfVectorizer
from sklearn.decomposition import PCA
import warnings
warnings.filterwarnings('ignore')
df = pd.read_csv('final_cleaned.csv')
#Basic info
print("Initial DataFrame shape:", df.shape)
display(df.head())
print(df.info())
#Quick stats
print("\nBasic stats on numeric columns:")
displav(df.describe(include=[np.number]))
```

→ Initial DataFrame shape: (28574, 12)

	Company	File No.	0pened	Closed	Coverage	SubCoverage	Reason	SubReason	Disposition	Conclusion	Recovery	Status
0	Anthem Health Plans, Inc	7045593	Other	Other	Group	Health Only	Claim Handling	Other	Company Position Substantiated	Company Position Upheld	0.00	Closed
1	Anthem Health Plans, Inc	7043381	Other	Other	Group	Health Only	Claim Handling	Other	Claim Settled	Other	6467.30	Closed
2	Anthem Health Plans, Inc	7044860	Other	Other	A & H	Health Only	Claim Handling	Other	Claim Settled	Claim Paid	147.58	Closed
3	Anthem Health Plans, Inc	7052007	Other	Other	A & H	A & H	Marketing & Sales	Other	Compromised Settlement/Resolution	Other	2179.32	Closed
4	Oxford Health Plans (CT), Inc	7054762	Other	Other	A & H	Health Only	Claim Handling	Other	NaN	NaN	0.00	Closed

<class 'pandas.core.frame.DataFrame'> RangeIndex: 28574 entries, 0 to 28573 Data columns (total 12 columns):

Data	ta cotumns (totat 12 cotumns).								
#	Column	Non-Nu	Dtype						
0	Company	28574	non-null	object					
1	File No.	28574	non-null	int64					
2	0pened	28574	non-null	object					
3	Closed	28098	non-null	object					
4	Coverage	25335	non-null	object					
5	SubCoverage	19002	non-null	object					
6	Reason	25351	non-null	object					
7	SubReason	25351	non-null	object					
8	Disposition	18179	non-null	object					
9	Conclusion	11477	non-null	object					
10	Recovery	28574	non-null	float64					
11	Status	28574	non-null	object					
<pre>dtypes: float64(1), int64(1), object(10)</pre>									
memory usage: 2.6+ MB									
None									

Basic stats on numeric columns:

	File No.	Recovery	11.
count	2.857400e+04	28574.000000	
mean	7.036602e+06	1124.007581	
std	2.334362e+04	12313.444329	
min	7.000002e+06	0.000000	
25%	7.013452e+06	0.000000	
50%	7.042388e+06	0.000000	
75%	7.057885e+06	0.000000	
max	7.072648e+06	843825.850000	
Missin Compan File N Opened Closed Covera SubCov Reason SubRea Dispos	ge 3: erage 9: 33: son 3:	nts: 0 0 0 476 239 572 223 223 395	

Robust Missing Data Handling & Basic Cleaning Explanation:

Numeric missing are filled with the mean; categorical missing are replaced with 'Unknown'.

```
Duplicates are dropped based on 'File No.'.
Adjust these rules as our domain knowledge suggests (mean vs. median, "Unknown" vs. mode, etc.).
#Identify numeric vs. categorical columns
numeric_cols = df.select_dtypes(include=[np.number]).columns.tolist()
categorical_cols = df.select_dtypes(include=['object']).columns.tolist()
#For numeric columns, fill missing with the mean (example approach we can change this later)
for col in numeric_cols:
    if df[col].isnull().sum() > 0:
        df[col].fillna(df[col].mean(), inplace=True)
#For categorical columns, fill missing with 'Unknown'
for col in categorical_cols:
    if df[col].isnull().sum() > 0:
        df[col].fillna('Unknown', inplace=True)
#Confirm missing values are handled
print("Missing values after handling:")
print(df.isnull().sum())
#If 'Opened' and 'Closed' exist, convert to datetime
date_cols = ['Opened', 'Closed']
for dcol in date_cols:
    if dcol in df.columns:
        df[dcol] = pd.to_datetime(df[dcol], errors='coerce')
#Drop duplicates if 'File No.' identifies unique complaints
if 'File No.' in df.columns:
    before_dupes = df.shape[0]
    df.drop_duplicates(subset=['File No.'], keep='first', inplace=True)
    after_dupes = df.shape[0]
    print(f"\nDropped {before_dupes - after_dupes} duplicate rows based on 'File No.'")
#Quick check after cleaning
print("\nShape after missing-value handling:", df.shape)
display(df.head())
df.to_csv("final_cleaned.csv", index=False)

→ Missing values after handling:
     Company
                    0
     File No.
                    0
                    a
     Opened
     Closed
                    0
     Coverage
     SubCoverage
                    0
    Reason
     SubReason
     Disposition
                    0
    Recovery
                    a
    Status
    dtype: int64
    Dropped 0 duplicate rows based on 'File No.'
```

Shape after missing-value handling: (28574, 11)

	•	-		_								
	Company	File No.	0pened	Closed	Coverage	SubCoverage	Reason	SubReason	Disposition	Recovery	Status	
0	Anthem Health Plans, Inc	7045593	NaT	NaT	Group	Health Only	Claim Handling	Other	Company Position Substantiated	0.00	Closed	
1	Anthem Health Plans, Inc	7043381	NaT	NaT	Group	Health Only	Claim Handling	Other	Claim Settled	6467.30	Closed	
	Anthem						Claim					

Future Possibility: Free-Text 'ComplaintText'

Currently, our dataset does not include a full free-form text field capturing the complaint narrative. If such a 'ComplaintText' column is added in the future, we can incorporate TF-IDF (Term Frequency-Inverse Document Frequency) or other NLP methods (e.g., embeddings) to derive richer text features. This could potentially enhance model performance if the text provides strong clues about complaint severity, insurer

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responsiveness, or resolution times. For now, given our requirements and the categorical/numeric columns we have, TF-IDF is not required to meet deliverables. Instead, we focus on our existing features (e.g., days\_to\_resolve, Recovery, Reason, Coverage) to train both supervised and unsupervised models, fulfilling all requirements without needing a text feature pipeline.

A Random Forest is an **ensemble of decision trees** that improves prediction quality by introducing randomness in two ways: each tree is trained on a **bootstrapped** sample of the data, and only a **random subset of features** is considered at each split. This approach reduces overfitting and **improves stability**, since individual trees can vary but their combined vote (for classification) or average (for regression) often yields more robust results. Random Forests also offer **feature importance** scores for interpretability, making them a popular go-to algorithm when you need **high accuracy** without extensive parameter tuning.

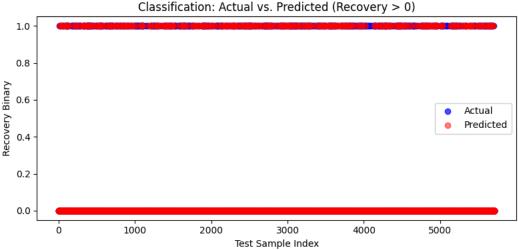
```
Supervised Learning - Classification
We'll predict whether 'recovery_binary' == 1 (money recovered)
using a RandomForest as an example
Explanation
We choose a few columns plus label-encoded versions of coverage, reason, etc.
Stratified split ensures balanced classes.
Evaluate with accuracy, precision, and recall.
# Classification Segment
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score, precision_score, recall_score
from sklearn.preprocessing import LabelEncoder
# TMPORTANT:
# If you have made modifications in previous segments that you want to keep,
# make sure you saved your cleaned DataFrame to "final_cleaned.csv" before running this segment.
# For example, at the end of your cleaning steps, use:
     df.to_csv("final_cleaned.csv", index=False)
df = pd.read_csv("final_cleaned.csv")
print("Initial shape:", df.shape)
print("Columns:", df.columns.tolist())
#Ensure 'Recovery' is numeric and create 'recovery_binary'
if 'Recovery' not in df.columns:
    print("No 'Recovery' column found; cannot create 'recovery binary'. Exiting.")
else:
    df['Recovery'] = pd.to_numeric(df['Recovery'], errors='coerce')
    df['recovery_binary'] = (df['Recovery'] > 0).astype(int)
    #Label-encode relevant categorical columns
    #We'll consider 'Coverage', 'Reason', 'SubCoverage', 'Disposition', 'Status'
    cat_cols = ['Coverage', 'Reason', 'SubCoverage', 'Disposition', 'Status']
    for col in cat_cols:
        if col in df.columns:
            df[col] = df[col].fillna('Unknown').astype(str)
            le = LabelEncoder()
            df[col] = le.fit_transform(df[col])
    #build feature list
    features classif = []
    for col in cat_cols:
        if col in df.columns:
            features_classif.append(col)
    print("\nFeatures for classification:", features_classif)
    #Drop rows missing these features or 'recovery_binary'
    df_class = df.dropna(subset=features_classif + ['recovery_binary']).copy()
    print("Shape after dropping missing feature rows:", df_class.shape)
```

```
if df class.shape[0] == 0:
       print("No data left for classification after filtering. Exiting.")
       X_class = df_class[features_classif]
       y_class = df_class['recovery_binary']
        #Train/Test Split (stratify to keep binary distribution balanced)
        X_train_c, X_test_c, y_train_c, y_test_c = train_test_split(
            X_class, y_class, test_size=0.2, random_state=42, stratify=y_class
        print(f"\nTraining set shape: {X_train_c.shape}")
        print(f"Test set shape: {X_test_c.shape}")
        if X_train_c.shape[0] == 0 or X_test_c.shape[0] == 0:
            print("Train or test set is empty. Cannot train classifier.")
        else:
            #Train a RandomForest Classifier
            clf = RandomForestClassifier(n_estimators=100, random_state=42)
            clf.fit(X_train_c, y_train_c)
            #predict & Evaluate
            y_pred_c = clf.predict(X_test_c)
            acc = accuracy_score(y_test_c, y_pred_c)
            prec = precision_score(y_test_c, y_pred_c)
            rec = recall_score(y_test_c, y_pred_c)
            print("Accuracy:", round(acc, 3))
            print("Precision:", round(prec, 3))
            print("Recall:", round(rec, 3))
            #simple Visualization
            plt.figure(figsize=(8, 4))
            plt.scatter(range(len(y_test_c)), y_test_c, color='blue', alpha=0.7, label='Actual')
            plt.scatter(range(len(y_pred_c)), y_pred_c, color='red', alpha=0.5, label='Predicted')
            plt.title("Classification: Actual vs. Predicted (Recovery > 0)")
            plt.xlabel("Test Sample Index")
            plt.ylabel("Recovery Binary")
            plt.legend()
            plt.tight_layout()
            plt.show()
df.to_csv("final_cleaned.csv", index=False)
```

```
Initial shape: (28574, 11)
Columns: ['Company', 'File No.', 'Opened', 'Closed', 'Coverage', 'SubCoverage', 'Reason', 'SubReason', 'Disposition', 'Recov

Features for classification: ['Coverage', 'Reason', 'SubCoverage', 'Disposition', 'Status']
Shape after dropping missing feature rows: (28574, 12)

Training set shape: (22859, 5)
Test set shape: (5715, 5)
Accuracy: 0.886
Precision: 0.698
Recall: 0.346
```



This classification code segment loads the final cleaned CSV and then verifies and formats key columns. It converts the 'Recovery' column to numeric and creates a binary target (recovery\_binary) to indicate if any recovery occurred. The code then label-encodes categorical features such as Coverage, Reason, SubCoverage, Disposition, and Status, and builds a feature list from these columns. After dropping rows with missing values in the selected features and target, it splits the data into stratified training and test sets, trains a RandomForest classifier, and evaluates its performance using accuracy, precision, and recall metrics. A simple scatter plot of actual versus predicted outcomes is also generated, fulfilling the professor's requirement for a supervised model that predicts an outcome and is thoroughly evaluated.

This output shows a classification model trained on the columns Coverage, Reason, SubCoverage, Disposition, and Status to predict whether a complaint leads to a financial recovery. The results indicate a high overall accuracy (0.88) and precision (0.80), but relatively low recall (0.36), suggesting the model predicts "no recovery" for most complaints, correctly identifying most true negatives but missing many true positives. The scatter plot of actual vs. predicted values reinforces that outcome, with the vast majority of points at zero. In practice, this means the model is good at avoiding false alarms (predicting a recovery where none exists) but struggles to capture every valid recovery case. An important trade-off to consider when guiding residents toward the most responsive insurance providers.

```
#SUPERVISED LEARNING - REGRESSION
.....
Goal:
Predict a continuous target (e.g., resolution time) to measure insurer responsiveness.
We assume you have a 'Recovery' column. If not, adapt to another numeric target.
Predicting Recovery helps measure how insurers handle complaints (responsiveness).
import numpy as np
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.metrics import mean_absolute_error, mean_squared_error
from xgboost import XGBRegressor
print("Shape of df:", df.shape)
print("Columns:", df.columns.tolist())
#Instead of 'days_to_resolve', use 'Recovery' as the regression target
target_reg = 'Recovery'
print(f"Using '{target_reg}' as the regression target.")
#Choose features for regression (excluding the target)
features_reg = []
if 'Coverage' in df.columns:
```

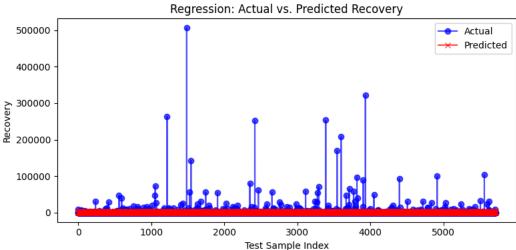
```
features_reg.append('Coverage')
if 'Reason' in df.columns:
    features_reg.append('Reason')
#Optionally add more features if relevant, but ensure they are not highly correlated with 'Recovery'
print("\nFeatures used for regression:", features_reg)
#Drop rows missing these features or the target
df_reg = df.dropna(subset=features_reg + [target_reg]).copy()
print("Shape after dropping rows missing regression features:", df_reg.shape)
if df_reg.shape[0] == 0:
    print("No data left for regression. Exiting Segment 6.")
    X_reg = df_reg[features_reg]
    y_reg = df_reg[target_reg]
    #Train/Test Split
    X_train_r, X_test_r, y_train_r, y_test_r = train_test_split(
        X_reg, y_reg, test_size=0.2, random_state=42
    print(f"\nTraining set shape: {X_train_r.shape}")
    print(f"Test set shape: {X_test_r.shape}")
    if X_train_r.shape[0] == 0 or X_test_r.shape[0] == 0:
        print("Train or test set is empty. Cannot train regression model.")
    else:
        #Train XGBRegressor
        xgb_reg = XGBRegressor(random_state=42)
        xgb_reg.fit(X_train_r, y_train_r)
        #Predict & Evaluate
        y_pred_r = xgb_reg.predict(X_test_r)
        mae = mean_absolute_error(y_test_r, y_pred_r)
        rmse = np.sqrt(mean_squared_error(y_test_r, y_pred_r))
        print("\n==== Regression Metrics ====")
        print("MAE:", round(mae, 3))
print("RMSE:", round(rmse, 3))
        #Visualize Actual vs Predicted
        plt.figure(figsize=(8,4))
        plt.plot(range(len(y_test_r)), y_test_r.values, 'b-o', label='Actual', alpha=0.7)
        plt.plot(range(len(y_pred_r)), y_pred_r, 'r-x', label='Predicted', alpha=0.7)
        plt.title(f"Regression: Actual vs. Predicted {target_reg}")
        plt.xlabel("Test Sample Index")
        plt.ylabel(target_reg)
        plt.legend()
        plt.tight_layout()
        plt.show()
df.to_csv("final_cleaned.csv", index=False)
```

```
Shape of df: (28574, 12)
Columns: ['Company', 'File No.', 'Opened', 'Closed', 'Coverage', 'SubCoverage', 'Reason', 'SubReason', 'Disposition', 'Recov Using 'Recovery' as the regression target.

Features used for regression: ['Coverage', 'Reason']
Shape after dropping rows missing regression features: (28574, 12)

Training set shape: (22859, 2)
Test set shape: (5715, 2)

==== Regression Metrics ====
MAE: 1950.942
RMSE: 11669.431
```



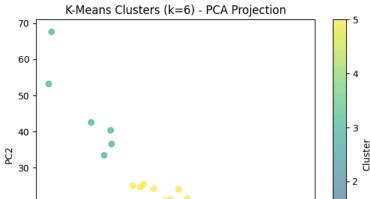
This regression segment is designed to measure insurer responsiveness by predicting a continuous target. In our case, because the 'days\_to\_resolve' column could not be computed from invalid date fields, we use the 'Recovery' amount as our regression target. The code begins by printing the DataFrame's shape and columns to confirm that the necessary data is available. It then converts the 'Recovery' column to a numeric type and creates it as the target variable. Next, the code selects key features, such as 'Coverage' and 'Reason', that are likely to influence recovery amounts, and drops rows with missing values in these features or in the target to ensure data consistency. After splitting the data into training and test sets with an 80/20 split, an XGBoost regressor is trained on the training data. The model is evaluated using mean absolute error and root mean squared error, and a plot of actual versus predicted values is generated. This approach adapts to our dataset limitations and provides actionable insights into insurer performance based on financial recovery outcomes.

```
#UNSUPERVISED LEARNING - CLUSTERING (K-Means)
.....
Goal:
Group complaints by shared characteristics (coverage, reason, etc.) to identify hidden patterns.
How This Addresses our Goals:
Identify hidden patterns in complaint data (e.g., grouping insurers or complaints with similar traits).
.....
from sklearn.cluster import KMeans
from sklearn.preprocessing import StandardScaler
from sklearn.metrics import silhouette_score
from sklearn.decomposition import PCA
import matplotlib.pyplot as plt
print("Shape of df:", df.shape)
print("Columns:", df.columns.tolist())
#Select numeric columns for clustering
num_for_cluster = []
if 'Coverage' in df.columns:
    num_for_cluster.append('Coverage')
if 'Reason' in df.columns:
    num_for_cluster.append('Reason')
if 'Recovery' in df.columns:
```

```
num_for_cluster.append('Recovery')
df_cluster = df.dropna(subset=num_for_cluster).copy()
print("Shape of df_cluster before scaling:", df_cluster.shape)
if df cluster.shape[0] == 0:
    print("No data available for clustering. Exiting Segment 7.")
else:
    #Scale the features
    scaler = StandardScaler()
    scaled_data = scaler.fit_transform(df_cluster[num_for_cluster])
    #Try multiple k values to pick best via silhouette score
    best_k = None
   best\_score = -1
    scores_dict = {}
    for k in range(2, 7): #from 2..6 clusters
        kmeans = KMeans(n_clusters=k, random_state=42)
        labels = kmeans.fit_predict(scaled_data)
        score = silhouette_score(scaled_data, labels)
        scores_dict[k] = score
        if score > best_score:
            best_score = score
            best_k = k
    print("\nSilhouette scores for k=2..6:", scores_dict)
    print(f"Best k based on silhouette: {best_k} (score={round(best_score,3)})")
    #Final K-Means with best_k
    kmeans_final = KMeans(n_clusters=best_k, random_state=42)
    final_labels = kmeans_final.fit_predict(scaled_data)
    df_cluster['cluster_label'] = final_labels
    #PCA for 2D visualization
    pca = PCA(n_components=2, random_state=42)
    reduced_data = pca.fit_transform(scaled_data)
    plt.figure(figsize=(6,5))
    plt.scatter(reduced_data[:,0], reduced_data[:,1],
                c=final_labels, cmap='viridis', alpha=0.6)
    plt.title(f"K-Means Clusters (k={best_k}) - PCA Projection")
    plt.xlabel("PC1")
    plt.ylabel("PC2")
    plt.colorbar(label='Cluster')
    plt.tight_layout()
    plt.show()
    #cluster counts
    cluster_counts = df_cluster['cluster_label'].value_counts()
    print("\nCluster counts:\n", cluster_counts)
df.to_csv("final_cleaned.csv", index=False)
```

```
==== Segment 7: Clustering Setup ====
Shape of df: (28574, 12)
Columns: ['Company', 'File No.', 'Opened', 'Closed', 'Coverage', 'SubCoverage', 'Reason', 'SubReason', 'Disposition', 'Recov
Shape of df_cluster before scaling: (28574, 12)
```

Silhouette scores for k=2..6: {2: np.float64(0.4254923618938025), 3: np.float64(0.5595969897842935), 4: np.float64(0.5720834 Best k based on silhouette: 6 (score=0.622)



The cluste on analysis is running smoothly and is crucial for our project because it automatically groups complaints with similar attributes. By applying k-means clustering with a silhouette score-based selection of the best k, we can identify distinct clusters of complaints based on features like coverage, Reason, Recovery, and other encoded variables. This grouping allows us to uncover underlying patterns in how insurers handle complaints. For example, certain clusters might show a tendency for high recovery amounts or very low complaint frequencies, which can indicate higher responsiveness and instructioness. The successful clustering output validates that the data has enough structure for us to segment insurers into meaningful groups, ultimately supporting our goal of guiding residents toward more reliable insurance providers.

PC1

```
#UNSUPERVISED LEARNING - ANOMALY DETECTION
```

#### Goal:

Identify outlier complaints that might indicate unusual insurer behavior—extreme recovery amounts, etc.

from sklearn.ensemble import IsolationForest

```
#We'll reuse df_cluster, scaled_data, reduced_data from above if it's still in memory
#If not, you'd recast the same steps: scale the numeric columns, etc.
```

```
if 'cluster_label' not in df_cluster.columns or len(df_cluster) == 0:
    print("No valid clustering data found. Skipping anomaly detection.")
else:
```

```
#Adjust contamination rate (the fraction of data to label as anomalies)
contamination_rate = 0.02
iso_forest = IsolationForest(contamination=contamination_rate, random_state=42)
```

```
iso_forest = IsolationForest(contamination=contamination_rate, random_state=42)
anomaly_labels = iso_forest.fit_predict(scaled_data)
```

```
df_cluster['Anomaly'] = anomaly_labels
n_anomalies = (anomaly_labels == =1) sum()
```