```
1 # Author: Sai Sri Surya Bandaru
     # CSCE 5222 - Project 2
  3 # Pancreatic Cancer Survival Analysis
  4 # April 2024
     # Imports
  2 import pandas as pd
  3
     import numpy as np
      import matplotlib.pyplot as plt
     import seaborn as sns
     from pathlib import Path
      from sklearn.model_selection import train_test_split
     from sklearn.metrics import mean_absolute_error, mean_squared_error
     from sklearn.preprocessing import StandardScaler, OneHotEncoder
 10
     from sklearn.compose import ColumnTransformer
 11
     from sklearn.pipeline import Pipeline
     from sklearn.linear model import ElasticNet
     from sklearn.svm import SVR
 14
     from sklearn.tree import DecisionTreeRegressor
 15 from lightgbm import LGBMRegressor
 16 from xgboost import XGBRegressor
     import joblib, os
  1 # Setup paths
     # Get the project root directory
     BASE DIR = Path(os.getcwd()).parent
     DATA_DIR = BASE_DIR / "data"
     MODEL_DIR = BASE_DIR / "models"
     # Print paths to verify
  8
     print(f"Base directory: {BASE_DIR}")
     print(f"Data directory: {DATA_DIR}")
 10
     print(f"Model directory: {MODEL_DIR}")
 12
 13
     # Verify directories exist
     DATA DIR.mkdir(parents=True, exist ok=True)
     MODEL_DIR.mkdir(parents=True, exist_ok=True)
 15
 16
 17
     # Load data function
 18
     def load_data(use_vae=False):
 19
         file = "vae_augmented_data.csv" if use_vae else "synthetic_data.csv"
          return pd.read_csv(DATA_DIR / file)
 20
 21
     # Build preprocessing pipeline
 22
 23
     def build preprocessor(df):
       num_cols = df.select_dtypes(include=["float64", "int64"]).columns.difference(["SurvivalMonths"])
 25
          cat_cols = df.select_dtypes(include=["object"]).columns
 26
         preproc = ColumnTransformer(
             transformers=[
                  ("num", StandardScaler(), num_cols),
 28
 29
                  ("cat", OneHotEncoder(handle_unknown="ignore"), cat_cols),
 30
              ]
 31
          return preproc
Base directory: c:\Users\vckkr\OneDrive - Chaitanya UNT\OneDrive - UNT System\Documents\3rd-Party-Assignments\2025\04-April\CSCE 522
Data directory: c:\Users\vckkr\OneDrive - Chaitanya UNT\OneDrive - UNT System\Documents\3rd-Party-Assignments\2025\04-April\CSCE 522
    Model directory: c:\Users\vckkr\OneDrive - Chaitanya UNT\OneDrive - UNT System\Documents\3rd-Party-Assignments\2025\04-April\CSCE 52
  1 # Plot feature distributions
 2 def plot feature distributions(df, title):
       num_cols = df.select_dtypes(include=["float64", "int64"]).columns.difference(["SurvivalMonths"])
       n_{cols} = 3
 5
      n_rows = (len(num_cols) + n_cols - 1) // n_cols
       plt.figure(figsize=(15, 5*n_rows))
 Ω
       for i, col in enumerate(num_cols, 1):
        plt.subplot(n_rows, n_cols, i)
 9
           sns.histplot(data=df, x=col, kde=True)
10
 11
          plt.title(f"{col} Distribution")
12
       plt.suptitle(title)
13
       plt.tight_layout()
       plt.show()
15
16 # Plot survival by stage
17 def plot survival by stage(df, title):
       plt.figure(figsize=(10, 6))
```

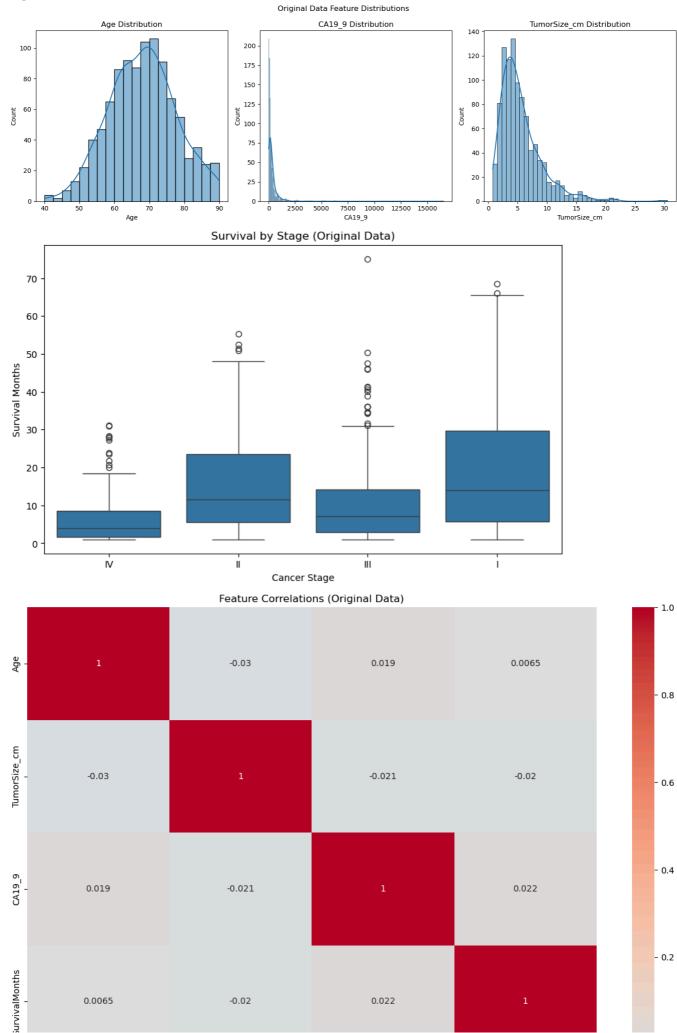
```
sns.boxplot(data=df, x="Stage", y="SurvivalMonths")
      plt.title(title)
20
      plt.xlabel("Cancer Stage")
21
      plt.ylabel("Survival Months")
22
23
      plt.show()
24
25 # Plot correlation heatmap
26 def plot_correlation(df, title):
      plt.figure(figsize=(12, 8))
      corr = df.select_dtypes(include=["float64", "int64"]).corr()
28
      sns.heatmap(corr, annot=True, cmap="coolwarm", center=0)
29
      plt.title(title)
31
      plt.tight_layout()
32
      plt.show()
1 # Train models and get scores
2 def train_models(df, use_vae=False):
      X = df.drop(columns="SurvivalMonths")
      y = df["SurvivalMonths"]
5
      Xtr, Xte, ytr, yte = train_test_split(
6
          X, y, test_size=0.2, random_state=42, stratify=df["Stage"]
8
9
      pre = build preprocessor(df)
10
11
12
      # Define models
13
      models = {
          "ElasticNet": ElasticNet(alpha=0.1, l1_ratio=0.5, random_state=42),
14
          "SVR":
                         SVR(C=10, gamma="scale"),
15
          "DecisionTree":DecisionTreeRegressor(max_depth=5, random_state=42),
16
          "LightGBM": LGBMRegressor(n_estimators=200, learning_rate=0.05),
17
                         XGBRegressor(n_estimators=300, learning_rate=0.05, max_depth=4),
          "XGBoost":
18
19
20
21
      scores = \{\}
22
      for name, est in models.items():
         pipe = Pipeline([("prep", pre), ("model", est)])
23
24
          pipe.fit(Xtr, ytr)
25
26
          y_pred = pipe.predict(Xte)
          scores[name] = {
27
              "MAE": mean absolute error(yte, y pred),
28
29
              "MSE": mean_squared_error(yte, y_pred),
31
32
          # Save model
          out_dir = MODEL_DIR / "vae" if use_vae else MODEL_DIR
33
34
          out_dir.mkdir(parents=True, exist_ok=True)
35
          joblib.dump(pipe, out_dir / f"{name}_model.pkl")
36
37
      return pd.DataFrame(scores).T
1 # First, check if we need to generate the data
2 from src.synthetic_data import generate_original_data, generate_synthetic_patients
4 # Generate data if it doesn't exist
5 if not (DATA DIR / "synthetic data.csv").exists():
      print("Generating original dataset...")
      original_df = generate_original_data(1000)
      original_df.to_csv(DATA_DIR / "synthetic_data.csv", index=False)
8
      print("Original dataset saved")
9
10
11 if not (DATA_DIR / "vae_augmented_data.csv").exists():
      print("Generating VAE-augmented dataset...")
      vae_df = generate_synthetic_patients(original_df, 5000)
13
      vae_df.to_csv(DATA_DIR / "vae_augmented_data.csv", index=False)
14
      print("VAE-augmented dataset saved")
15
16
17 # Now load the data
18 print("Loading datasets...")
19 original_df = load_data(use_vae=False)
20 vae df = load data(use vae=True)
21
22 # Run analysis
23 print("\nTraining on original data...")
24 original_scores = train_models(original_df, use_vae=False)
25 print("\nOriginal data scores:")
26 print(original_scores)
28 print("\nTraining on VAE data...")
```

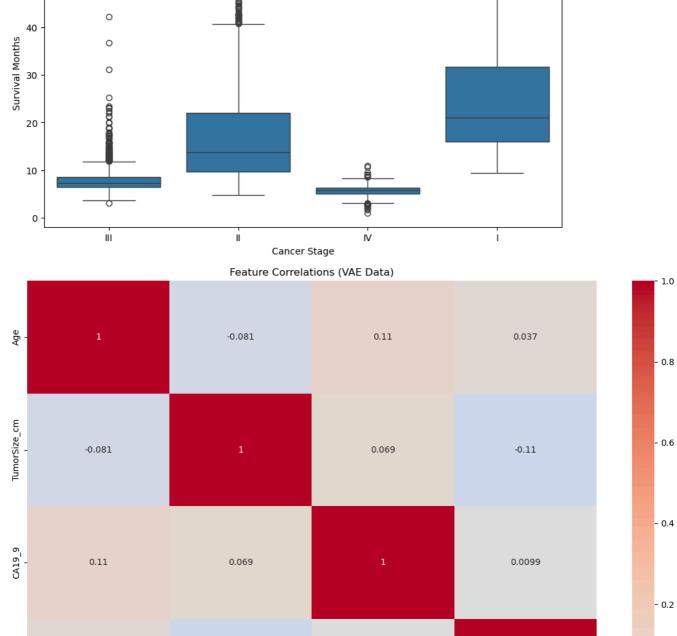
```
29 vae_scores = train_models(vae_df, use_vae=True)
30 print("\nVAE data scores:")
31 print(vae_scores)

→ Generating original dataset...
    Original dataset saved
    Generating VAE-augmented dataset...
    Epoch 20/150, Loss: 4.6211
    Epoch 40/150, Loss: 4.4997
    Epoch 60/150, Loss: 4.3596
    Epoch 80/150, Loss: 4.3186
    Epoch 100/150, Loss: 4.3031
    Epoch 120/150, Loss: 4.2156
    Epoch 140/150, Loss: 4.3525
    VAE-augmented dataset saved
    Loading datasets...
    Training on original data...
    [LightGBM] [Info] Auto-choosing row-wise multi-threading, the overhead of testing was 0.000481 seconds.
    You can set `force row wise=true` to remove the overhead.
    And if memory is not enough, you can set `force_col_wise=true`.
    [LightGBM] [Info] Total Bins 778
    [LightGBM] [Info] Number of data points in the train set: 800, number of used features: 11
    [LightGBM] [Info] Start training from score 10.872500
    [LightGBM] [Warning] No further splits with positive gain, best gain: -inf
    [LightGBM] [Warning] No further splits with positive gain, best gain: -inf
    [LightGBM] [Warning] No further splits with positive gain, best gain: -inf
    [LightGBM] [Warning] No further splits with positive gain, best gain: -inf
    [LightGBM] [Warning] No further splits with positive gain, best gain: -inf
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 1 # Run visualizations for original data
```

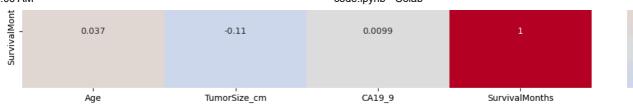
```
2 print("Original Data Visualizations:")
3 plot_feature_distributions(original_df, "Original Data Feature Distributions")
4 plot_survival_by_stage(original_df, "Survival by Stage (Original Data)")
5 plot_correlation(original_df, "Feature Correlations (Original Data)")
6
7 # Run visualizations for VAE data
8 print("\nVAE-Augmented Data Visualizations:")
9 plot_feature_distributions(vae_df, "VAE Data Feature Distributions")
10 plot_survival_by_stage(vae_df, "Survival by Stage (VAE Data)")
11 plot_correlation(vae_df, "Feature Correlations (VAE Data)")
```

→ Original Data Visualizations:



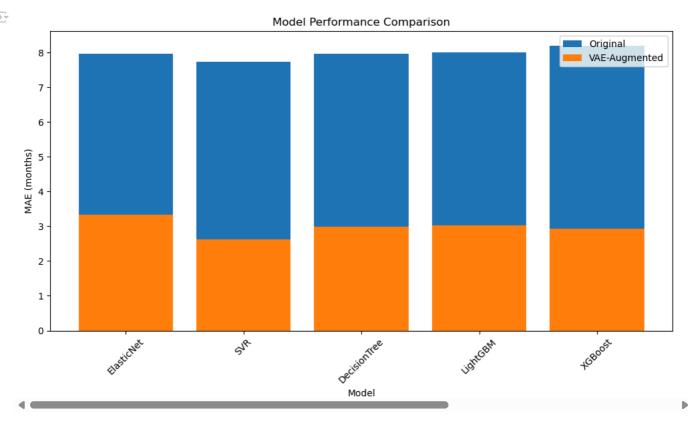


hs



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```
1 # Plot results
2 plt.figure(figsize=(10, 6))
3 plt.bar(original_scores.index, original_scores["MAE"], label="Original")
4 plt.bar(vae_scores.index, vae_scores["MAE"], label="VAE-Augmented")
5 plt.title("Model Performance Comparison")
6 plt.xlabel("Model")
7 plt.ylabel("MAE (months)")
8 plt.legend()
9 plt.xticks(rotation=45)
10 plt.tight_layout()
11 plt.show()
```



```
1 # Detailed evaluation metrics
 2 def print_detailed_metrics(scores_df, title):
      print(f"\n{title}")
      print("-" * 50)
      print("Mean Absolute Error (MAE):")
5
 6
      print(scores_df["MAE"].sort_values())
      print("\nMean Squared Error (MSE):")
      print(scores_df["MSE"].sort_values())
8
9
      print("\nBest Model by MAE:", scores_df["MAE"].idxmin())
      print("Best Model by MSE:", scores_df["MSE"].idxmin())
10
11
12 # Print metrics for both datasets
13 print_detailed_metrics(original_scores, "Original Data Model Performance")
14 print_detailed_metrics(vae_scores, "VAE-Augmented Data Model Performance")
15
16 # Calculate improvement percentages
17 mae_improvement = ((original_scores["MAE"] - vae_scores["MAE"]) / original_scores["MAE"] * 100)
18 \ \mathsf{mse\_improvement} = ((\mathsf{original\_scores}[\ \mathsf{"MSE"}] \ - \ \mathsf{vae\_scores}[\ \mathsf{"MSE"}]) \ / \ \mathsf{original\_scores}[\ \mathsf{"MSE"}] \ * \ 100)
19
20 print("\nImprovement with VAE-Augmented Data:")
21 print("-" * 50)
22 print("MAE Improvement (%):")
23 print(mae_improvement.sort_values(ascending=False))
24 print("\nMSE Improvement (%):")
25 print(mse_improvement.sort_values(ascending=False))
    Original Data Model Performance
    Mean Absolute Error (MAE):
    SVR
                    7.738537
    ElasticNet
                    7.964600
    DecisionTree
                   7.966413
    LightGBM
                    8.009767
    XGBoost
                    8.202568
    Name: MAE, dtype: float64
    Mean Squared Error (MSE):
```