



# Predictive Modeling of Diabetes Risk Using Health Behavior Indicators

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- Course: Clinical Decision Support Modeling
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# Introduction

- Diabetes affects over 34 million Americans and is a leading cause of morbidity, mortality, and healthcare costs. This project develops an artificial intelligence-driven Clinical Decision Support (CDS) system that predicts diabetes risk using health behavior indicators from the CDC Behavioral Risk Factor Surveillance System (BRFSS 2015).
- **Key Achievement:** XGBoost model achieved 82% ROC-AUC and 86% accuracy, outperforming baseline logistic regression and Random Forest classifiers.

# Problem Statement

- Diabetes affects millions of adults and remains a major cause of morbidity, mortality, and rising healthcare costs.
- Traditional screening methods often rely only on clinical or laboratory measures, missing key behavioral and lifestyle risk factors.
- As a result, many high-risk individuals remain undiagnosed or identified too late for effective prevention.
- There is a critical need for an accurate, interpretable, and scalable model that can predict diabetes risk using self-reported health behavior indicators.
- This project aims to address this gap by developing machine learning models evaluated on both balanced and real-world imbalanced datasets to support early detection and inform clinical decision-making



# Literature Review

Research consistently demonstrates that **behavioral and lifestyle indicators** such as obesity, hypertension, physical inactivity, and poor nutrition are key predictors of diabetes.

- **Rahman et al. (2023)** reported improved predictive performance and clinician trust when combining ML models with explainable AI, using SHAP to show feature contributions.
- **Ullah & Lee (2024)** demonstrated that ensemble models such as Random Forest and XGBoost outperform traditional linear models in diabetes risk stratification.
- **Xie et al. (2019)** used BRFSS data to successfully develop accurate diabetes prediction models, validating its suitability for population-level analytics.
- **Wang et al. (2024)** highlighted the importance of optimized behavioral features for diabetes questionnaires and risk scoring systems.

## Gaps This Project Addresses

Integration of *both* balanced and real-world imbalanced datasets.

Resampling (SMOTE) + model tuning.

Focus on interpretability for CDS use cases.

End-to-end framework suitable for healthcare deployment.

# Dataset Description

BRFSS 2015 dataset (CDC) with 400,000+ responses

- Includes health behaviors, chronic conditions, and lifestyle indicators

Datasets Used:

- Balanced Binary Dataset (equal diabetes / no diabetes)
- Imbalanced Binary Dataset (real-world distribution)

Feature Categories:

- Clinical: BMI, HighBP, HighChol
- Behavioral: Smoking, Physical Activity, Diet
- Socioeconomic: Education, Income
- Self-Reported Health: General, Physical, Mental Health
- Features strongly correlate with diabetes risk

# EDA

Datasets checked for missing values –  
none detected

- Balanced dataset: 50/50 diabetes distribution
- Imbalanced dataset reflects real-world majority class
- Class distribution

Key Variable Patterns:

- Higher BMI, poor General Health, HighBP, HighChol --higher diabetes risk
- Behavioral factors (diet, smoking, activity) show meaningful variation

Correlations:

- Strong positive: BMI, HighBP, HighChol
- Negative associations: Physical Activity, Fruit/Veg intake

```
# Split into train/test
X_train, X_test, y_train, y_test = train_test_split(
    X, y, test_size=0.2, stratify=y, random_state=42
)
print(f"Train shape: {X_train.shape}, Test shape: {X_test.shape}")

Train shape: (202944, 21), Test shape: (50736, 21)
```

```
#Handle Imbalance
from imblearn.over_sampling import SMOTE

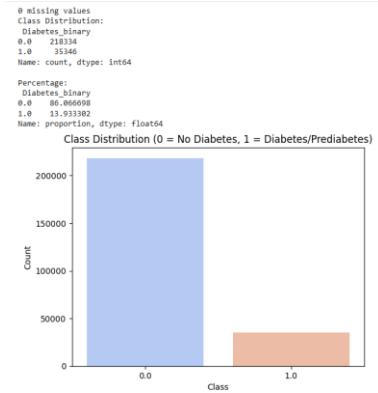
sm = SMOTE(random_state=42)
X_res, y_res = sm.fit_resample(X_train, y_train)

print("After SMOTE:", y_res.value_counts())

After SMOTE: Diabetes_binary
0    174667
1    174667
Name: count, dtype: int64
```

```
print(df.columns)

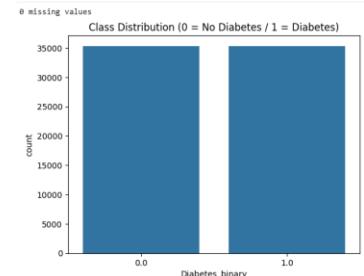
Index(['Diabetes_binary', 'HighBP', 'HighChol', 'CholCheck', 'BMI', 'Smoker',
       'Stroke', 'HeartDiseaseorAttack', 'PhysActivity', 'Fruits', 'Veggies',
       'HyalAlcoholConsump', 'AnyHealthcare', 'NoDocbCost', 'GenHlth',
       'MentHlth', 'PhysHlth', 'DiffWalk', 'Sex', 'Age', 'Education',
       'Income'],
      dtype='object')
```



```
#Explore the dataset
print(df.isna().sum(), "missing values")

# Summary stats
df.describe().head(10)

# Check class balance
sns.countplot(x='Diabetes_binary', data=df)
plt.title("Class Distribution (0 = No Diabetes / 1 = Diabetes)")
plt.show()
```



# Methodology

## Used machine learning to predict diabetes vs. no diabetes

- Loaded dataset and previewed to inspect structure and quality
- Cleaned and encoded data, then split into train/test sets
- Worked with balanced data and imbalanced data using SMOTE

### Models Used:

- Logistic Regression
- Random Forest
- XGBoost

### Evaluation:

- ROC-AUC, Accuracy, Precision, Recall, F1-score
- Hyperparameter tuning (GridSearchCV)
- SHAP explainability for feature impact

```
# Logistic Regression Model
logreg = LogisticRegression(max_iter=1000)
logreg.fit(X_train, y_train)
y_pred_logreg = logreg.predict(X_test)
y_prob_logreg = logreg.predict_proba(X_test)[:,1]

# Random Forest Classifier
rf = RandomForestClassifier(n_estimators=100, random_state=42)
rf.fit(X_train, y_train)
y_pred_rf = rf.predict(X_test)
y_prob_rf = rf.predict_proba(X_test)[:,1]

# XGBoost Classifier
xgb = XGBClassifier(eval_metric='logloss')
xgb.fit(X_train, y_train)
y_pred_xgb = xgb.predict(X_test)
y_prob_xgb = xgb.predict_proba(X_test)[:,1]
```

```
, y_test = train_test_split
shape, " Test:", X_test.shape
: (14139, 21)
```

```
#Handle Imbalance
from imblearn.over_sampling import SMOTE
sm = SMOTE(random_state=42)
X_res, y_res = sm.fit_resample(X_train, y_train)
print("After SMOTE:", y_res.value_counts())
```

```
After SMOTE: Diabetes_binary
0    174667
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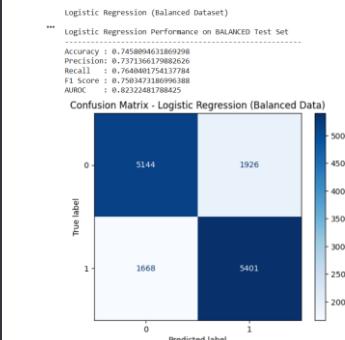
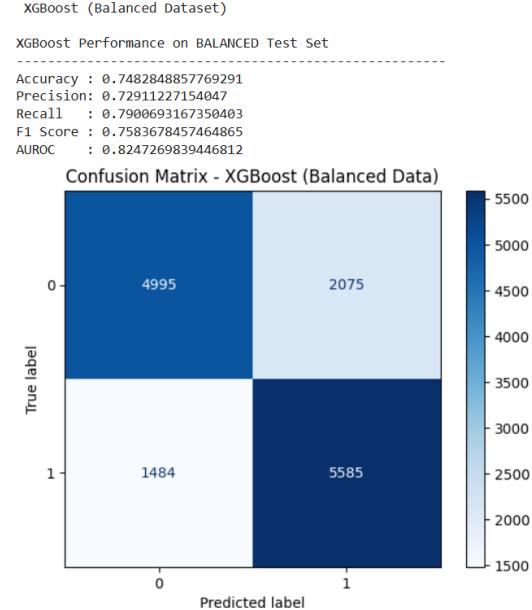
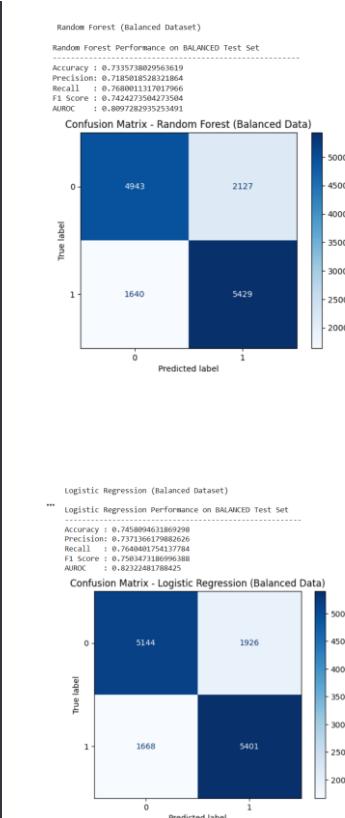
```
# balanced binary dataset
read_csv('/content/diabetes_binary_5050split_health_indicators_BRFSS2015.csv')
data
shape, df.shape
head(), df.head()
692, 22
diabetes_binary  HighBP  HighChol  CholCheck  BMI  Smoker  Stroke  HeartDiseaseorAttack
0.0      1.0      0.0      1.0   26.0      0.0      0.0      0.0
0.0      1.0      1.0      1.0   26.0      1.0      1.0      0.0
0.0      0.0      0.0      1.0   26.0      0.0      0.0      0.0
0.0      1.0      1.0      1.0   28.0      1.0      0.0      0.0
0.0      0.0      0.0      1.0   29.0      1.0      0.0      0.0
columns
```

# Balanced Dataset Results

- LR: Strong baseline performance
- RF: Better recall for diabetic class
- XGBoost: Best overall AUC and F1

## Feature Importance

- Both models agree that health-related features (BMI, General Health, Physical Health, Blood Pressure, Cholesterol) are important.
- Random Forest spreads importance across more features, while XGBoost puts more weight on a few top predictors



```
# Evaluation metrics
def print_metrics(y_true, y_pred, y_prob):
    print('Accuracy:', accuracy_score(y_true, y_pred))
    print('Precision:', precision_score(y_true, y_pred))
    print('Recall:', recall_score(y_true, y_pred))
    print('F1 Score:', f1_score(y_true, y_pred))
    print('AUROC:', roc_auc_score(y_true, y_prob))

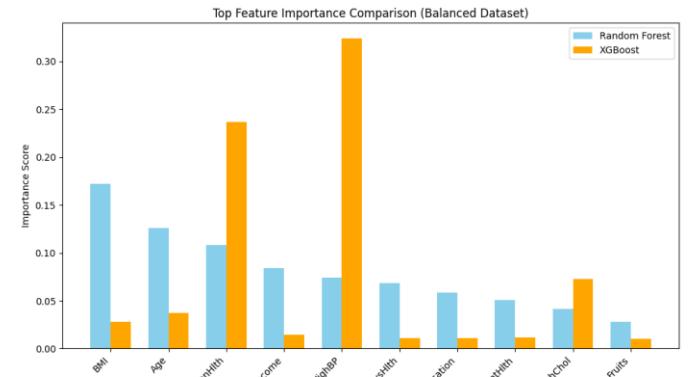
print("Logistic Regression Performance:")
print_metrics(y_test, y_pred_logreg, y_prob_logreg)

print("Random Forest Performance:")
print_metrics(y_test, y_pred_rf, y_prob_rf)

print("XGBoost Performance:")
print_metrics(y_test, y_pred_xgb, y_prob_xgb)

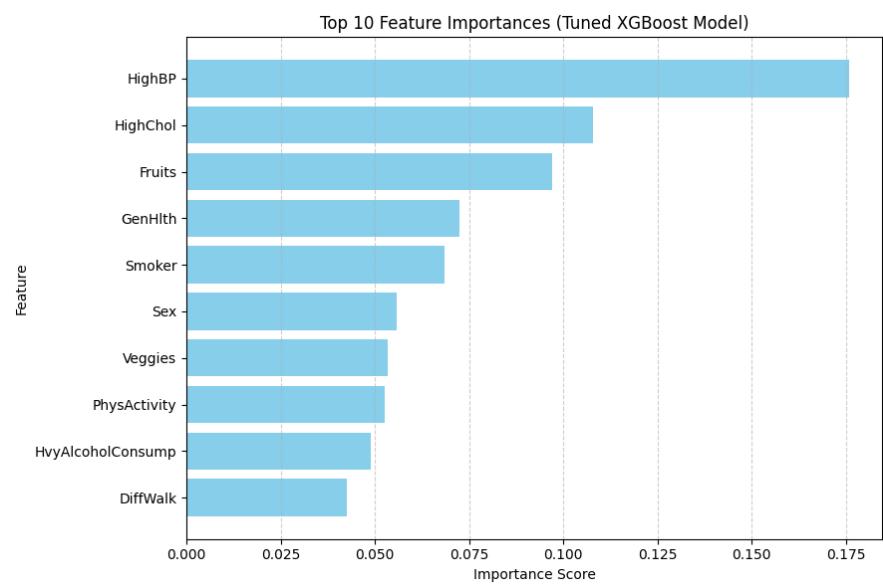
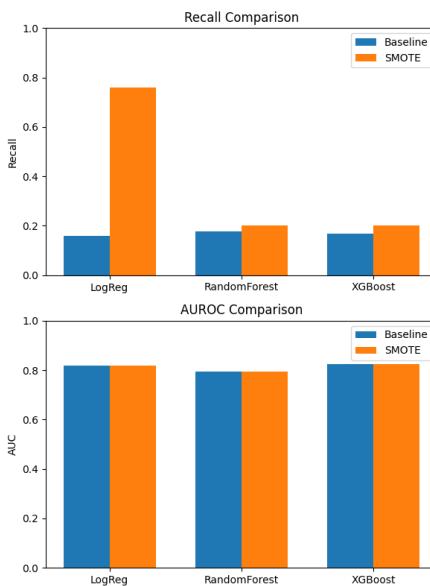
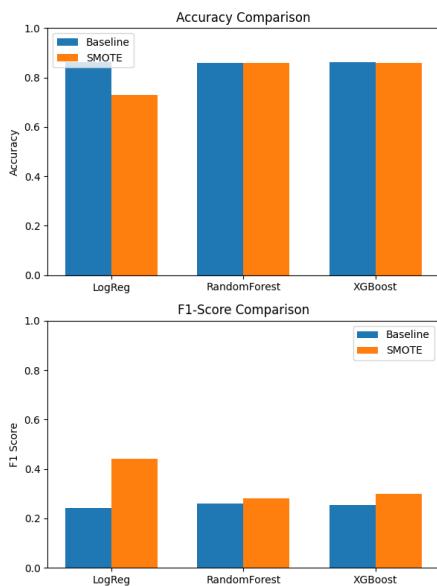
# Random Forest Performance:
Accuracy: 0.733578809563619
Precision: 0.715051852821806
Recall: 0.768001317037966
F1 Score: 0.7424273504273504
AUROC: 0.80972829325253491

# XGBoost Performance:
Accuracy: 0.7482848857769291
Precision: 0.72911227154047
Recall: 0.7900693167358403
F1 Score: 0.7583678457464865
AUROC: 0.8247269839446812
```



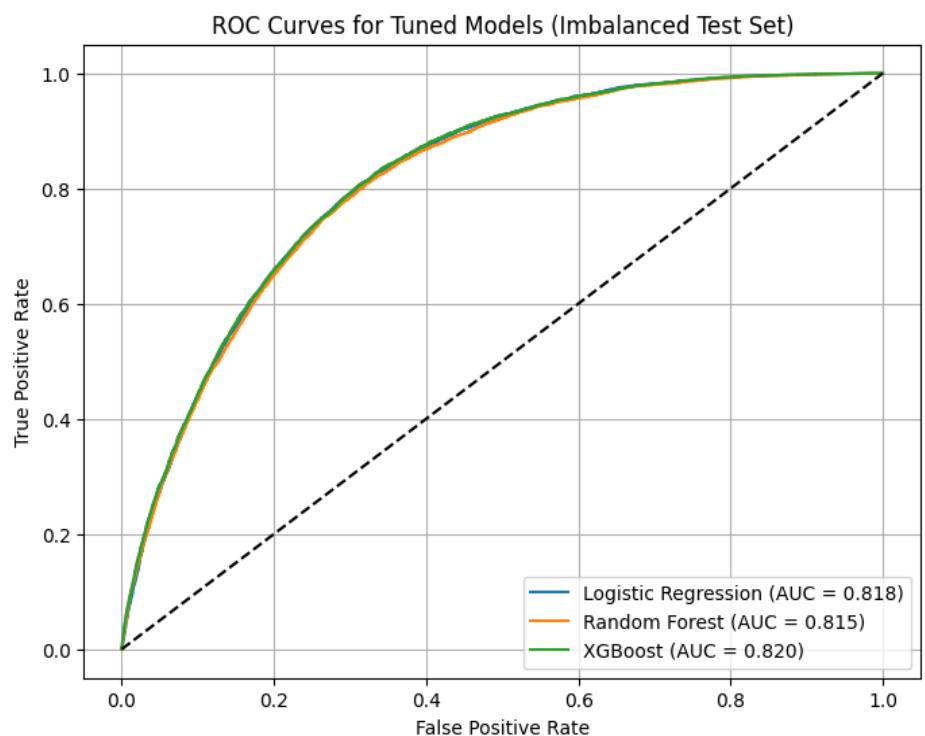
# Imbalanced Dataset Results

Baseline SMOTE



# ROC Curve Comparison

- **XGBoost (AUC = 0.820)** *Best generalization to real-world data.*
- **Logistic Regression (AUC = 0.818)** Very stable, consistent performance.
- **Random Forest (AUC = 0.815)** Slight decline despite strong CV results (overfitting on SMOTE data).



# Hyperparameter Tuning

- GridSearchCV used for LR, RF, XGBoost
- Improved calibration
- Enhanced model stability
- Better discrimination across classes
- Random Forest achieved 97%

```
# Logistic Regression Tuning
print("Tuning Logistic Regression")
lr_param_grid = {
    'C': [0.01, 0.1, 1, 10, 100],
    'solver': ['lbfgs', 'saga']
}
lr_grid_search = GridSearchCV(
    LogisticRegression(max_iter=2000, random_state=42),
    lr_param_grid,
    cv=cv,
    scoring='roc_auc',
    n_jobs=-1, verbose=1
)
lr_grid_search.fit(X_res, y_res)

best_lr_smote_tuned = lr_grid_search.best_estimator_
print("Best parameters for Logistic Regression:", lr_grid_search.best_params_)
print("Best cross-validation ROC-AUC for Logistic Regression:", lr_grid_search.best_score_)

# Random Forest Tuning
print("Tuning Random Forest")
rf_param_grid = {
    'n_estimators': [100, 200],
    'max_depth': [10, 20],
    'min_samples_split': [2, 5]
}
rf_grid_search = GridSearchCV(
    RandomForestClassifier(random_state=42),
    rf_param_grid,
    cv=cv,
    .....
)

# Tuning KNN
Fitting 3 folds for each of 10 candidates, totalling 30 fits
Best parameters for KNN: {'n_neighbors': 10, 'weights': 'uniform'}
Best cross-validation ROC-AUC for KNN: 0.8304770740839595

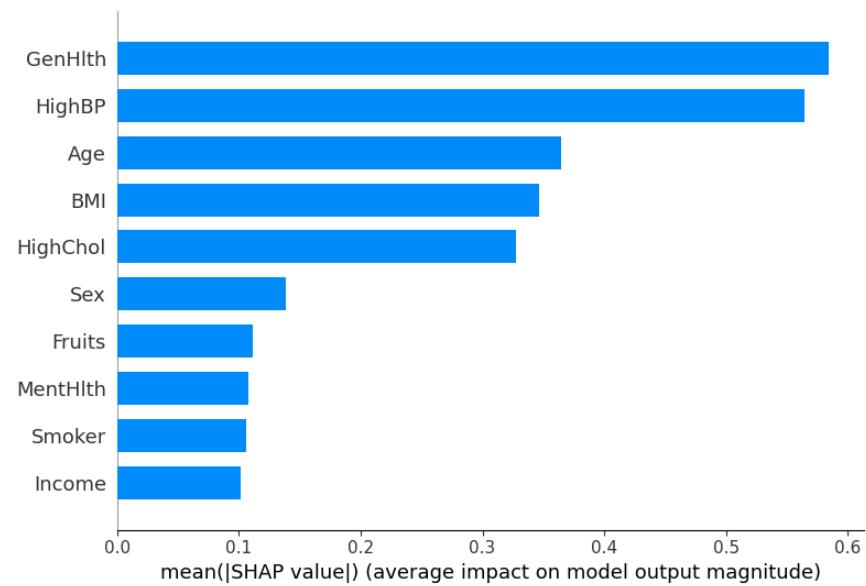
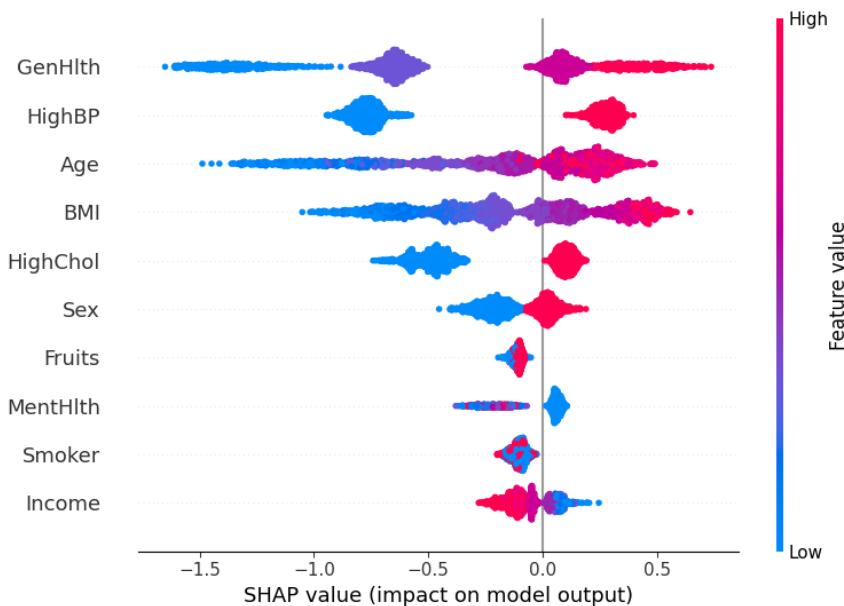
# Tuning KNN
Fitting 3 folds for each of 8 candidates, totalling 48 fits
Best parameters for Random Forest: {'max_depth': 20, 'min_samples_split': 2, 'n_estimators': 200}
Best cross-validation ROC-AUC for Random Forest: 0.9700221203639625

# Tuning KNN
Fitting 3 folds for each of 32 candidates, totalling 96 fits
Best parameters for KNN: {'colsample_bytree': 0.7, 'learning_rate': 0.05, 'max_depth': 5, 'n_estimators': 200, 'subsample': 0.7}
Best cross-validation ROC-AUC for KNN: 0.99051521204041
```

Model	Best Hyperparameters	Best CV ROC-AUC	Notes
<b>Logistic Regression</b>	C = 0.01, solver = 'saga'	<b>0.8305</b>	Performs well but limited by linear decision boundary
<b>Random Forest</b>	n_estimators = 200, max_depth = 20, min_samples_split = 2	<b>0.9708</b>	Best-performing tuned model
<b>XGBoost</b>	learning_rate = 0.05, max_depth = 5, n_estimators = 200, subsample = 0.7, colsample_bytree = 0.7	<b>0.9601</b>	Excellent performance, competitive with RF

# Model Interpretability (SHAP)

- SHAP reveals how each feature influences the model's prediction of diabetes risk.
- Positive SHAP values → increase predicted risk; negative values → decrease predicted risk.
- Colors indicate feature values: **red = high**, **blue = low**.



- Machine learning methods can reliably predict diabetes risk using self-reported behavioral and clinical indicators. SHAP explainability confirms that the model aligns with clinical knowledge, making it suitable for future development into an interpretable clinical decision support tool.

## Conclusion

# Future Work



Test advanced models (LightGBM, CatBoost) and calibrated probability outputs.



Incorporate additional clinical or behavioral data for improved accuracy.



Explore alternative imbalance strategies and fairness auditing.



Build a deployable CDS tool with interactive SHAP explanations.



Validate predictions with clinicians and real-world patient data.





THANK YOU

# References

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