Predicting Diabetes

Project 4

(UTOR-VIRT-DATA-PT-06-2023-U-LOLC-MWTH(B))

Contributors (Group 4):

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The Topic: Diabetes

- Diabetes affects the health of millions of people and puts an enormous financial burden on the US economy
- Early diagnosis of diabetes can lead to lifestyle changes and more effective treatment
- Predictive models for diabetes risk can be an important tool for public health officials.

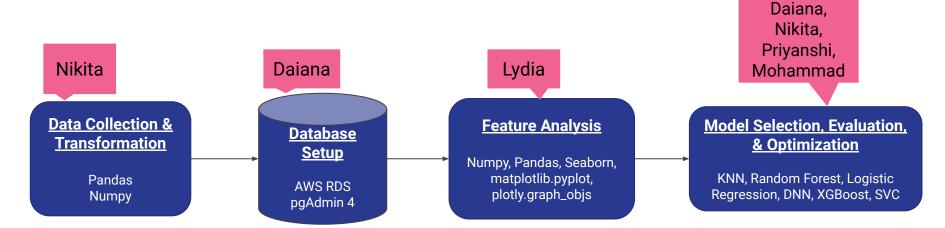


The Goal

Develop a machine learning model to identify individuals that either have diabetes or are high-risk for developing diabetes using screening information.



Division of Work





The Dataset



- Data for this project is taken from 2022 survey data off the CDCs website from their Behavioral Risk Factor Surveillance System (BRFSS) sector.
- The data contains information about U.S. residents health-related risk behaviors, chronic health conditions, and use of preventive services
- https://www.cdc.gov/brfss/annual_data/an
 nual_2022.html
- 326 features (columns) and 445,132 records for 2022

Category	Renamed-as	Label/Question	Value	Null/Refused
_STATE	STATE	-State FIPS Code	-Integer [1-78]	
DISPCODE	DISPCODE	- Final Disposition	1100 : Completed Interview 1200 : Partial Complete Interview	
SEXVAR	GENDER	-Sex of Respondent	1: MALE 2: FEMALE	
_INCOMG1	INCOME	-Income categories (Computed income categories)	Integer [1-7]	9: Don't Know/refused
HEIGHT3	HEIGHT	-About how tall are you without shoes? (Height in Feet and Inches)	200 - 711 : ft/inches 9061 - 9998 : m/cm	7777 & 9999 : Don't Know/refused BLANK
WTKG3	WEIGHT	-Computed Weight in Kilograms (Reported in kilograms)	FLOAT [2300 - 29500]	BLANK
_BMI5CAT	ВМІ	-Computed body mass index categories (Four-categories of BMI)	1: Underweight 2: Normal Weight 3: Over Weight 4: Obese	BLANK
_RACE1	RACE	-Computed Race-Ethnicity grouping (Race/ethnicity categories)	1: White 2: Black 3: Indian/ Alaskan Native 4: Asian 5: Hawaiian/Pacific Islander 7: Multiracial 8: Hispanic	9: Don't Know/refused BLANK

Data Collection & Transformation

	STATE	SEXVAR	_	INCOMG1	HEIGHT3	WTKG3 _B	MI5CAT _F	RACE1 _A	GEG5YR DIABE	TE4 PRE	DIAB2					
0	1.0	2.0)	9.0	9999.0	NaN	NaN	1.0	13.0	1.0	NaN		(115	122	r	ows x 33 columns)
1	1.0	2.0)	3.0	503.0	6804.0	3.0	1.0	13.0	3.0	NaN		(445	132		ows x 33 coluitilis)
2	1.0	2.0)	6.0	502.0	6350.0	3.0	1.0	8.0	3.0	NaN	***				
3	1.0	2.		STATE	SEYWAD	INCOMG1	HEIGHT3	WTKG3	RMISCAT	PACE1	AGE	FG5VD	DIABETE4	I PRENIA	R2	
4	1.0	2.			FEMALE	NaN	9999.0		NaN			13.0			laN	 Categorical values was
5	1.0	1.														
6	1.0	2.			FEMALE	3.0	503.0	6804.0	Over_Weight	White		13.0	0.0) 1	laN	categories
7	1.0	2.	2	1.0	FEMALE	6.0	502.0	6350.0	Over_Weight	White		8.0	0.0) 1	laN	
8	1.0	2.	3	1.0	FEMALE	NaN			Normal Walahi			MaN	0.0		IaM	
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			6	1.0	FEMALE	5.0	ŧ PE	RSDOC3	: [1. 2	. nan	0.]					actagorica ware convert
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			8	1.0	FEMALE	5.0			: [nan 2		0.	3.	6. 5.	4.]		
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			J	1.0	LEWINEL	3.0	Di	ABEYE1	: [nan 4	1. 1.	2.		0.]			All the "No/Never
									: [nan 0		3.	5.	4. 1.	2.]		
									: [nan 6	7						converted to 0
									: [0. 1	-						
									: [0. 1							
							CI	/DSTRK3	: [0. 1	. nan]						
							HA	VARTH4	: [0. 1	. nan]						
							D1	FFWALK	: [0. 1	. nan]						

Categorical values were converted to categories

'Don't know/Not sure' or 'Refused' categories were converted to NaN values

> All the "No/Never" categories were converted to 0

Data Collection & Transformation

NaN White 13.0 1.0	NaN	NaN	ID	STATE	GENDER	INCOME	WEIGHT	ВМІ	RACE	AGE	DIABETES			NAL_DOC C		HRT_ATTACK
Over Weight White 13.0 0.0	NaN	NaN	1		FEMALE		6804.0	Over_Weight		13.0	0.0	0.0		2.0	0.0	0.0
DIABETE4 799	NaN	NaN	2		FEMALE		6350.0 5398.0	Over_Weight Normal Weight	White	5.0	0.0	2.0		2.0	1.0	0.0
Noi PREDIAB2 234587	NaN	NaN	6		FEMALE		6260.0			13.0	0.0	0.0		1.0	1.0	0.0
No	NaN	NaN	7	1.0	FEMALE	5.0	7348.0	Over_Weight	White	13.0	0.0	0.0	***	1.0	1.0	0.0
PHYSHLTH 8139	***	2	445126	78.0	MALE	5.0	10433.0	Obese	White	3.0	0.0	0.0	2001	0.0	2.0	0.0
MENTHLTH 6742	NaN	WaN			FEMALE		6985.0	Over_Weight		1.0	0.0	0.0		0.0	2.0	0.0
DIABTYPE 343296	NaN	NaN 4	445128	78.0	FEMALE	5.0	8301.0	Over_Weight	Black	7.0	0.0	2.0		2.0	1.0	0.0
TOTINDA 802	NaN	Ivalv	445130 445131	78.0 78.0	MALE		10886.0 6350.0	Obese Normal Weight	Black	11.0	0.0	0.0		0.0	0.0	0.0
	NaN	NaN		1.0	U.U	0.0	U.U	Never	U.U	COILE	ge_N /2.					
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PRIMINSR 12334	NaN	Gen	eral	Infori	matio	n Dataf	rame	(246050	rows	× 2		ns): To	predic	t if som	eone h	as diabete
1101	NaN	Gen		Infori gender		n Dataf		(246050)			A- one- EYE					
PRIMINSR 12334	NaN	_	ID	GENDER	AGE	ВМ	II <mark>DIAB</mark> E	TES DIABTYPE	INSULI	N_Y/N	A- one- EYE C_test	EXAM1 DI	ABEYE1 DIA	AB_MNGMT I	FEETSORE	PERSONAL_DOC
PRIMINSR 12334 PERSDOC3 3116 CHECKUP1 4289	NaN		ID		AGE 13.0	BM Over_Weigh	II <mark>DIABE</mark>		INSULI		A- one- EYE				FEETSORE	PERSONAL_DOC
PRIMINSR 12334 PERSDOC3 3116 CHECKUP1 4289 PDIABTS1 241504	NaN		ID 55982	GENDER FEMALE	AGE 13.0 13.0	ВМ	II DIABE	TES DIABTYPE	INSULI	N_Y/N 0.0	A- one- C_test	EXAM1 DIJ	ABEYE1 DIA	AB_MNGMT 0.0	FEETSORE	PERSONAL_DOO
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PRIMINSR 12334 PERSDOC3 3116 CHECKUP1 4289 PDIABTS1 241504	NaN	_	ID 55982 55988 55992	GENDER FEMALE MALE	AGE 13.0 13.0 11.0 11.0	BM Over_Weigh Over_Weigh Over_Weigh	II DIABE	1.0 1.0 1.0 2.0 1.0 2.0	INSULI	0.0 0.0 0.0	A- one- C_test	1.0 1.0 2.0	1.0 1.0 2.0	0.0 6.0 0.0	0.0 0.0 0.0	PERSONAL_DOC 1.6 1.6 1.6
PRIMINSR 12334 PERSDOC3 3116 CHECKUP1 4289 PDIABTS1 241504 INSULIN1 342543	NaN	_	55982 55988 55992 55995	GENDER FEMALE MALE MALE MALE	AGE 13.0 13.0 11.0 11.0	BM Over_Weigh Over_Weigh Over_Weigh	II DIABE	TES DIABTYPE 1.0 1.0 1.0 2.0 1.0 2.0 1.0 2.0	INSULI	N_Y/N 0.0 0.0 0.0 0.0	A- one- C_test	1.0 1.0 2.0 3.0	1.0 1.0 2.0 3.0	0.0 6.0 0.0 3.0	0.0 0.0 0.0 0.0	PERSONAL_DOC 1.0 1.0 1.0 2.0
PRIMINSR 12334 PERSDOC3 3116 CHECKUP1 4289 PDIABTS1 241504 INSULIN1 342543 CHKHEMO3 343074 EYEEXAM1 342727	NaN	4	ID 55982 55988 55992 55995 56001	GENDER FEMALE MALE MALE MALE MALE MALE	13.0 13.0 11.0 11.0 12.0	Over_Weigh Over_Weigh Over_Weigh Over_Weigh Obese	II DIABE	1.0 1.0 1.0 1.0 1.0 2.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1	INSULI	N_Y/N 0.0 0.0 0.0 0.0 1.0 0.0	Anone EYE 0.0 2.0 2.0 1.0 4.0 4.0	1.0 1.0 2.0 3.0 2.0	1.0 1.0 2.0 3.0 2.0	0.0 6.0 0.0 3.0 0.0	0.0 0.0 0.0 0.0 0.0 0.0	PERSONAL_DOC 1.0 1.0 1.0 2.0
PRIMINSR 12334 PERSDOC3 3116 CHECKUP1 4289 PDIABTS1 241504 INSULIN1 342543 CHKHEMO3 343074 EYEEXAM1 342727 DIABEYE1 343884	NaN	4 4	1D 55982 55988 55992 55995 56001 45060	GENDER FEMALE MALE MALE MALE MALE MALE MALE MALE	13.0 13.0 11.0 11.0 12.0 11.0 N	Over_Weigh Over_Weigh Over_Weigh Obese Jormal_Weigh Over_Weigh	II DIABE	1.0 1.0 1.0 1.0 1.0 2.0 1.0 2.0 1.0 2.0 1.0 2.0 1.0 2.0 1.0 2.0 1.0 2.0 1.0 2.0 1.0 2.0 1.0 2.0 1.0 2.0 1.0 2.0 1.0 2.0 1.0 2.0	INSULI	0.0 0.0 0.0 0.0 1.0 0.0	Anones C_test	1.0 1.0 2.0 3.0 2.0 2.0	1.0 1.0 2.0 3.0 2.0 2.0	0.0 6.0 0.0 3.0 0.0 0.0 6.0	0.0 0.0 0.0 0.0 0.0 0.0 0.0	PERSONAL_DOC 1.0 1.0 1.0 2.0 2.0
PRIMINSR 12334 PERSDOC3 3116 CHECKUP1 4289 PDIABTS1 241504 INSULIN1 342543 CHKHEMO3 343074 EYEEXAM1 342727	NaN	4 4 4 4	ID 55982 55988 55992 55995 56001	GENDER FEMALE MALE MALE MALE MALE MALE	13.0 13.0 11.0 11.0 12.0 11.0 N 10.0	Over_Weigh Over_Weigh Over_Weigh Over_Weigh Obese	t DIABE	1.0 1.0 1.0 1.0 1.0 2.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1	INSULI	N_Y/N 0.0 0.0 0.0 0.0 1.0 0.0	Anone EYE 0.0 2.0 2.0 1.0 4.0 4.0	1.0 1.0 2.0 3.0 2.0	1.0 1.0 2.0 3.0 2.0	0.0 6.0 0.0 3.0 0.0	0.0 0.0 0.0 0.0 0.0 0.0	PERSONAL_DOC 1.0 1.1 1.1 2.1 2.0

Had a lot of NaN values: Questions only relevant to diabetic patients

Diabetic Dataframe (9975 rows x 17 columns): To predict the type of diabetes

Database Setup & Access



Creation of diabetes-database

hostname:

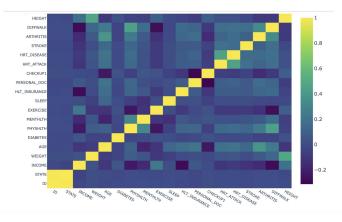
diabetes-dataset.cwpas6tssjkb.us-east-1.rds .amazonaws.com Link diabetes-database to pgAdmin & create user accounts

Psycopg2, Sqlalchemy

example query to grab all of the columns
sql_query = "SELECT * FROM general_info"
df = pd.read_sql_query(sql_query, conn)
df.head()

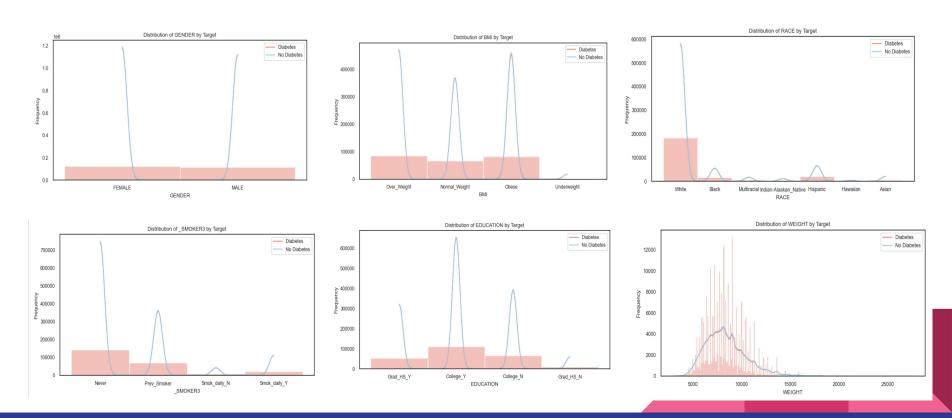
Exploratory Data Analysis

- Univariate Analysis
- Bivariate Analysis
- Correlation Matrix

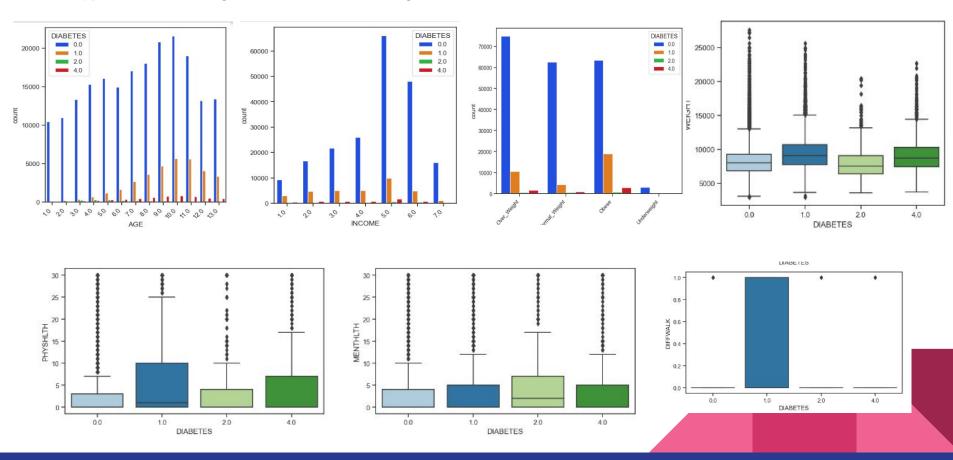




Exploratory Data Analysis



Exploratory Data Analysis



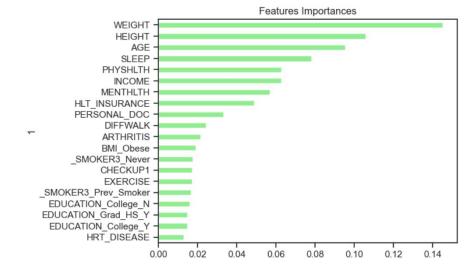
Feature Analysis

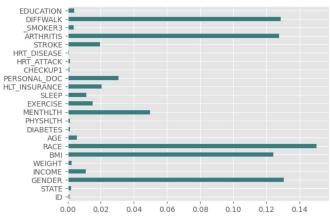
LIBRARIES

- # Statistics
- Pandas, numpy, scipy.stats, category_encoders, sklearn.feature_selection
- # Plots
- Seaborn, matplotlib.pyplot, plotly.

FEATURE SELECTION

- Mutual Information
- Chi-Square Test (categorical variable)
- Correlation Coefficient



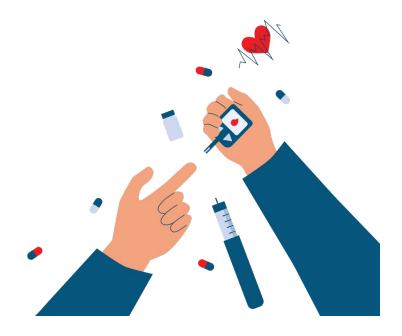


Model Selection in the selection in the





- Binary classification: has diabetes, no diabetes
 - K-Nearest Neighbours (k=2)
 - Random Forest
 - Deep Neural Network
 - Logistic Regression
- Binary classification: type 1, type 2
 - Support Vector Machine
 - XGBoost
 - Random Forest



Libraries used: Numpy, pandas, sklearn, tensorflow, train_test_split

Model Building & Evaluation weight in the second second





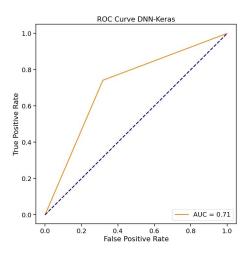
- **Evaluation metrics:**
 - Confusion matrix
 - Accuracy
 - F1 score
 - Precision
 - Recall 0
- Resampling techniques:
 - Random Under Sampling
 - RandomOversampler



Model Results - Dataset 1

GENDER	INCOME	WEIGHT	BMI	RACE	AGE	DIABETES
FEMALE	3.0	6804.0	Over_Weight	White	13.0	0.0
FEMALE	6.0	6350.0	Over_Weight	White	8.0	0.0
FEMALE	3.0	5398.0	Normal_Weight	White	5.0	0.0
FEMALE	5.0	6260.0	Normal_Weight	Black	13.0	0.0
FEMALE	5.0	7348.0	Over_Weight	White	13.0	0.0

	Accuracy	Precision	F1 Score	Recall	Model Type	Resample
0	0.611921	0.679167	0.522215	0.424187	KNN	50/50 Split
1	0.699871	0.687136	0.709722	0.733843	Random Forest	50/50 Split
2	0.701783	0.692934	0.708444	0.724665	DNN	50/50 Split
3	0.759709	0.288277	0.284663	0.281139	KNN	Oversampled Data
4	0.813698	0.422642	0.322615	0.260874	Random Forest	Oversampled Data
5	0.679645	0.310654	0.434937	0.724978	DNN	Oversampled Data
6	0.712012	0.700226	0.720223	0.741396	DNN	Keras
7	0.817811	0.360821	0.147173	0.092439	KNN	Reduced Features



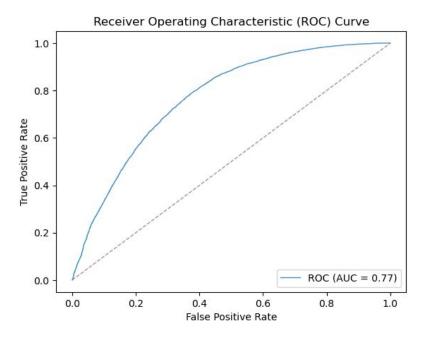
Classification	Report
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	===				
	1	precision	recall	f1-score	support
0.	0	0.73	0.68	0.70	10461
1.	0	0.70	0.74	0.72	10460
accurac	У			0.71	20921
macro av	g	0.71	0.71	0.71	20921
weighted av	g	0.71	0.71	0.71	20921

Model Results - Dataset 2

	DISPCODE	Diabetes	Smoker	CHD_1	CHD_2	Alcohol	GeneralHealth	MentalHealth	PhysicalHealth	Sex	Age	Education	Income
0	1100.0	1.0	4.0	2.0	2.0	1.0	2.0	88.0	88.0	2.0	13.0	4.0	99.0
1	1100.0	3.0	4.0	2.0	2.0	1.0	1.0	88.0	88.0	2.0	13.0	2.0	5.0
2	1100.0	3.0	4.0	2.0	2.0	1.0	2.0	3.0	2.0	2.0	8.0	4.0	10.0
3	1100.0	3.0	2.0	2.0	2.0	1.0	1.0	88.0	88.0	2.0	14.0	2.0	77.0
4	1100.0	3.0	4.0	2.0	2.0	1.0	4.0	88.0	2.0	2.0	5.0	3.0	5.0
				***			jun;						
445127	1100.0	3.0	4.0	2.0	2.0	9.0	3.0	3.0	88.0	2.0	1.0	2.0	1.0
445128	1100.0	3.0	4.0	2.0	2.0	1.0	1.0	2.0	2.0	2.0	7.0	4.0	7.0
445129	1100.0	3.0	1.0	2.0	2.0	9.0	5.0	30.0	30.0	2.0	10.0	2.0	77.0
445130	1100.0	3.0	4.0	2.0	1.0	1.0	2.0	88.0	88.0	1.0	11.0	3.0	8.0
445131	1100.0	3.0	3.0	2.0	2.0	9.0	2.0	1.0	88.0	1.0	5.0	1.0	4.0

	precision	recall	f1-score	support
0.0	0.69	0.70	0.70	9300
1.0	0.71	0.70	0.70	9566
accuracy			0.70	18866
macro avg	0.70	0.70	0.70	18866
weighted avg	0.70	0.70	0.70	18866



Model Results - Dataset 3- Random Forest

Predicted 0 Predicted 1

Actual 0	49902	1282
Actual 1	9026	1303

Accuracy Score: 0.8324256661193569

Classification Report

	precision	recall	f1-score	support
0	0.85	0.97	0.91	51184
1	0.50	0.13	0.20	10329
accuracy			0.83	61513
macro avg	0.68	0.55	0.55	61513
weighted avg	0.79	0.83	0.79	61513

Predicted 0 Predicted 1

Actual 0	46372	4774
Actual 1	714	50244

Accuracy Score : 0.9462508814542036

Classification Report

CIGODITICACIO	cpo. c			
	precision	recall	f1-score	support
Ø	0.98	0.91	0.94	51146
1	0.91	0.99	0.95	50958
accuracy			0.95	102104
macro avg	0.95	0.95	0.95	102104
weighted avg	0.95	0.95	0.95	102104

Y=df_dummies["DIABETIC"]
Y.value_counts()

0 204208 1 41842

Name: DIABETIC, dtype: int64

Model Results - Dataset 4

Two models provided the best results, XGB and random Forest For category

```
In [119]: 

| Sup. classifier_1 = spb_SEMCLosifier(objective 'multi-sefesa', mar_class2, random_tates42) |
| X_train_X_test_v_v_rain_x_test = train_test_split(v_v_t)_t test_sizend_f, random_tates42) |
| X_train_X_test_v_v_rain_x_test = train_test_split(v_v_t)_t test_sizend_f, random_tates42) |
| X_train_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_sizend_f_test_
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```
In [146]: xgb classifier = xgb.XGBClassifier(objective='multi:softmax', num class=2, random state=42)
         X_train, X_test, y_train, y_test = train_test_split(X, y_1, test_size=0.2, random_state=42)
         # Train the model on the training data
         xgb_classifier.fit(X_train, y_train)
        # Make predictions on the test data
        y_pred = xgb_classifier.predict(X_test)
        # Funluate the model
        accuracy = accuracy_score(y_test, y_pred)
        print(f"Accuracy: {accuracy}")
         print("Classification Report:")
        print(classification_report(y_test, y_pred))
         Accuracy: 0.9027568922305764
         Classification Report:
                     precision recall f1-score support
                         0.47
                                0 13 0 20
                        0.91
                                  0.98 0.95
                         0 60 0 56
           macro avg
                                            0.58
                         0.87
                                   0.90
```

```
In [14]: candom forest_l = Bandom forestClassifier(m|estinators=100, random_state=42)

# Train the model on the training data
random_forest_l=fift(fresh_scaled, y_rain)

# Nable predictions on the test data
y_rand = random_forest_rest(rest_scaled)

# Evaluate the model

# accuracy = accuracy_scare(y_test, y_pred)
print("Classification Report")

print("Classification Report")

print("Classification Report")

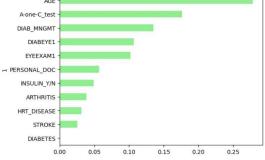
# Accuracy = accuracy_scare(y_test, y_pred)

# Accuracy = accuracy_scare(y_test, y_pred)
# Accuracy = accuracy_scare(y_test, y_pred)
# Accuracy = accuracy_scare(y_test, y_pred)
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# Accuracy = accuracy_scare(y_test, y_pred)
# Accuracy = accuracy_scare(y_test, y_pred)
# Accuracy = accuracy_scare(y_test, y_pred)
# Accuracy = accuracy_scare(y_test, y_pred)
```

```
In [77]: | from sklearn.ensemble import RandomForestClassifier
In [78]: random forest = RandomForestClassifier(n estimators=100, random state=42)
        # Train the model on the training data
        random forest.fit(X train, v train)
        # Make predictions on the test data
        y_pred = random_forest.predict(X_test)
        accuracy = accuracy_score(y_test, y_pred)
        print(f"Accuracy: {accuracy}")
        print("Classification Report:")
        print(classification report(y test, y pred))
        Accuracy: 0.8982456140350877
        Classification Report:
                    precision recall f1-score support
                1.0 0.41 0.14 0.20
                        0.91
                                0.98
                         0.66 0.56
                                           0.57
           macro avg
                                                     1995
        weighted avg
                         0.87 0.90
```

Model Results - Dataset 4

Two models provided the best results, XGB and random Forest for category



```
In [139]: random forest 2 = RandomForestClassifier(n estimators=100, random state=42)
          # Train the model on the training data
          random forest 2.fit(X train, y train)
          # Make predictions on the test data
          y pred = random forest 2.predict(X test)
          # Evaluate the model
          accuracy = accuracy_score(y_test, y_pred)
          print(f"Accuracy: {accuracy}")
          print("Classification Report:")
          print(classification report(y test, y pred))
          Accuracy: 0.8957393483709273
          Classification Report:
                        precision
                                      recall f1-score
                                                        support
                   1.0
                             0.39
                                        0.15
                                                 0.22
                                                             191
                   2.0
                             0.92
                                       0.97
                                                 0.94
                                                            1804
              accuracy
                                                 0.90
                                                            1995
             macro ave
                             0.65
                                        0.56
                                                 0.58
                                                            1995
          weighted avg
                             0.86
                                        0.90
                                                 0.87
                                                            1995
```

Summary of Results

		Diabetes Type Prediction		
	Dataset 1	Dataset 2	Dataset 3	Dataset 4
Accuracy	0.712012	0.70083	0.9462	0.8957
F1 Score	0.700226	0.70183	0.94	0.94
Precision	0.720223	0.71076	0.91	0.92
Recall	0.741396	0.70062	0.99	0.97
Model	-Hyperparameter tuning -Undersampled dataset -Neural Network -Subset of the data	- Equal Dataset - Over and Under Dataset	-Random Forest -Random Oversampling -Entire dataset (34 features)	-Scaling the dataset -Importance dataset formation and random forest model

Conclusion

- Preparing the data is very important for achieving good results
- More features lead to better model performance
- Having a balanced dataset is important to have higher recall and precision for the minority class
- Some features that may not look significant may play a big role in classification
- <u>Future improvements</u>: adding more data for the diabetic patients including sugar level, cholesterol, etc

Questions?





Thank you!
Hope to see you all soon!!