

DIABETES CLASSIFICATION

Group 6

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Id	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
1	6	148	72	35	0	33.6	0.627	50	1
2	1	85	66	29	0	26.6	0.351	31	0
3	8	183	64	0	0	23.3	0.672	32	1







SKIN THICKNESS

Triceps skinfold thickness measured in millimeters (mm).



Diastolic blood pressure measured in millimeters of mercury (mm Hg)



A GLUCOSE

Plasma glucose concentration measured over 2 hours in an oral glucose tolerance test.



Number of times the individual has been pregnant.

INSULIN

2-Hour serum insulin levels measured in micro international units per milliliter (mu U/ml).



Body mass index, calculated as weight in kilograms divided by height in meters squared (kg/m²)

DIABETES PEDIGREE FUNCTION 🚪

A genetic score representing the likelihood of diabetes.



2,768 SAMPLES

WITHIN **8 FEATURES**

Age of the individual in years.











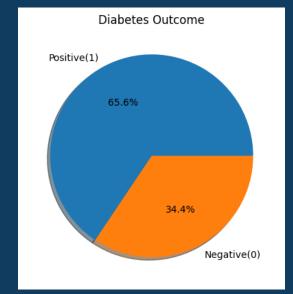
HOW MANY DUPLICATED SAMPLES?

0

HOW MANY MISSING VALUES?

Id	0
Pregnancies	0
Glucose	0
BloodPressure	0
SkinThickness	0
Insulin	0
BMI	0
DiabetesPedigreeFunction	0
Age	0
Outcome	0
dtype: int64	

DIABETES OUTCOME:





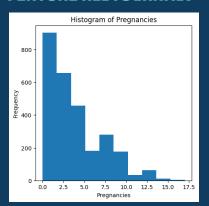


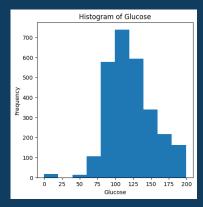


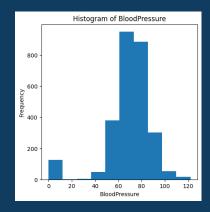
TOTAL DATASET OVERVIEW

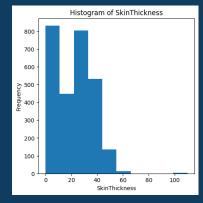
EDA

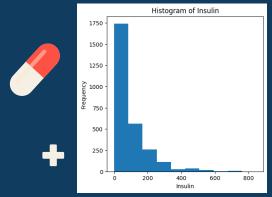
FEATURE HISTOGRAMS:

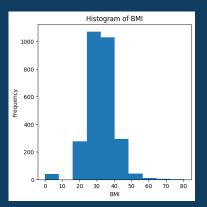


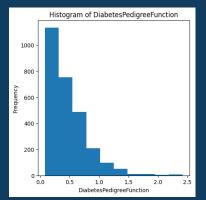


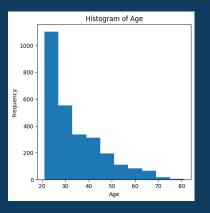


















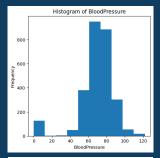
REMOVING DATA

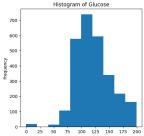
1.1) Blood Pressure (missing 125)
Diastolic BP is the latter number when you take a blood pressure test.
Recommend measure for healthy people is more than 80 [1].

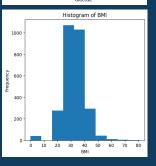
1.2) Glucose Tolerance Test (missing 18)
The recommend measure is the range between 70-140
Lower than the range considered as Hypoglycemia
More than the range considered impaired glucose tolerance, or diabetes [2].

1.3) BMI (missing 39)
Recommend measure is the range between 18.5 to 25 [3]
Lower than average means underweight
Greater than average means overweight, or obesity

The missing data in these features were very small compared to the total number (182 vs 2768). We considered removing it as the first approach.







^[1] NIH National Institute on Aging (NIA). High Blood Pressure and Older Adults. U.S. Department of Health and Human Services, National Institutes of Health. Retrieved September 22, 2023, from www.nia.nih.gov

^[2] Rao SS, Disraeli P, McGregor T. Impaired glucose tolerance and impaired fasting glucose. Am Fam Physician. 2004;69(8):1961-1968.

^[3] Centers for disease control and prevention (CDC). Healthy Weight, Nutrition, and Physical Activity. Division of Nutrition, Physical Activity, and Obesity, National Center for Chronic Disease Prevention and Health Promotion. Retrieved September 22, 2023, from www.cdc.gov



IMPUTATION

2.1) Insulin Test (missing 1330)

Insulin moves glucose from the blood into cells, particularly into muscles and fat tissue

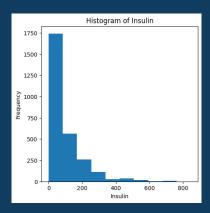
The average value of serum insulin metabolic parameter is 14.3±6.4 μU/mI [1].

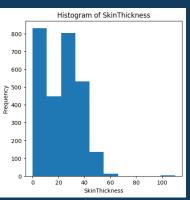
2.2) Tricept Skinfold Thickness (missing 800)

The average TSF thickness is 14.3 - 31.7 mm.

The world smallest TSF is 10 mm [2].

The missing values of these two features were approximately 50% of the dataset. Removing all of these values may affect the model performance. Thus, we suggested imputation method instead, and replaced 0 with median values.





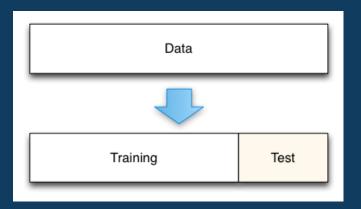


DATA SPLITTING AND FEATURE SCALING

Dataset is separated into Feature Variable (X) and Target Variable (y)

Data Splitting

- Train Data 80%
- Test Data 20%



Feature Scaling

- Features were scaled down with "StandardScaler" from scikit-learn.
- Standardize the features to 0 mean and 1 standard deviation



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LOGISTIC REGRESSION

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Parameter	max_iter =100
Accuracy	76.02%

APPLYING GRIDSEARCHCU HYPERPARAMETER

Best parameter	max_iter = 1000
Accuracy	79.21% (↑)



- The maximum tier has been increased to 1000, and the accuracy has been improved
- Using this model with GridSearchCV is recommended



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DEFAULT

Parameter	criterion = gini and splitter = best
Accuracy	98.90%

APPLYING GRIDSEARCHCU HYPERPARAMETER

Best parameter	criterion = entropy and splitter = random
Accuracy	95.67% (👃)



- The criteria and splitter were changed to entropy and random, resulting in lower accuracy
- Using this model without GridSearchCV is recommended



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RANDOM FOREST

DEFAULT

Parameter	criterion = gini and n_estimators = 100
Accuracy	99.53%

APPLYING GRIDSEARCHCU HYPERPARAMETER

Best parameter	criterion = gini and n_estimators = 200
Accuracy	96.06%(👃)



- Increasing n_estimators to 200 did not improve accuracy
- Using this model without GridSearchCV is recommended



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DEFAULT

Parameter	kernel = rbf
Accuracy	74.61%

APPLYING GRIDSEARCHCU HYPERPARAMETER

Best parameter	kernel = rbf
Accuracy	84.54% (↑)



- The parameter does not change, but the accuracy increases
- Using this model with GridSearchCV is recommended







STOCHASTIC GRADIENT DESCENT

DEFAULT

Parameter	alpha = 0.0001, loss = hinge, penalty = I2, max_tier = 1000
Accuracy	82.76%

APPLYING GRIDSEARCHCU HYPERPARAMETER

Best parameter	alpha = 0.01, loss = modified_huber, penalty = None, max_tier = 5000
Accuracy	80.21% (👃)



- The alpha increased to 0.01, loss changed to modified_huber, the penalty changed to none, and max_tier increased to 5000, but the accuracy decreased
- Using this model with GridSearchCV is recommended





RANDOMLY REMOVE FEATURE

	LOGISTIC REGRESSION	DECISION TREE	RANDOM FOREST	SGD	SVM
PREGNANCIES	76.97%	97.85%	98.37%	70.61%	77.31%
GLUCOSE	70.23%	97.32%	97.70%	67.60%	70.23%
BLOOD PRESSURE	77.32%	97.37%	98.23%	70.42%	77.12%
SKIN THICKNESS	77.70%	97.65%	98.13%	74.77%	77.02%
INSULIN	77.36%	97.61%	98.13%	74.77%	77.02%
вмі	77.02%	97.56%	97.89%	70.56%	77.26%
DIABETES PEDIGREE FUNCTION	77.12%	97.56%	98.42%	71.04%	77.17%
AGE	77.26%	97.08%	97.99%	69.70%	77.02%







CONCLUSIONS

- Diabetes Prediction Dataset is the dataset of our choice
- Pre-processing was done ensuring minimal drops
- Classification on the dataset was implemented using various approaches
 - Default parameter values were first obtained
 - Using GridSearchCV, the optimal results were obtained and compared
 - Features were then randomly dropped to see their effect on the model
 - Based on the obtained results, Random Forest is found to be the best approach







THANKS!

Do you have any questions? Feel free to ask us..







