Stroke Prediction



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What is a Stroke?

Stroke is a **disease that affects the arteries leading to and within the brain**. It is the number 5 cause of death and a leading cause of disability in the United States. A stroke occurs when a blood vessel that carries oxygen and nutrients to the brain is either blocked by a clot or bursts (or ruptures).

What are the effects of stroke?

The brain is an extremely complex organ that controls various body functions. If a stroke occurs and blood flow can't reach the region that controls a particular body function, that part of the body won't work as it should.



Project Target-

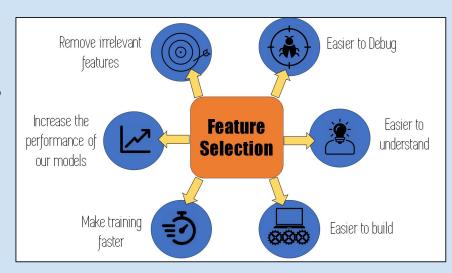
In order to try to reduce reduce stroke deaths, we will try to predict which people are more likely to have a stroke.

For this project we used Kaggle dataset, which includes 11 clinical features for predicting stroke events.

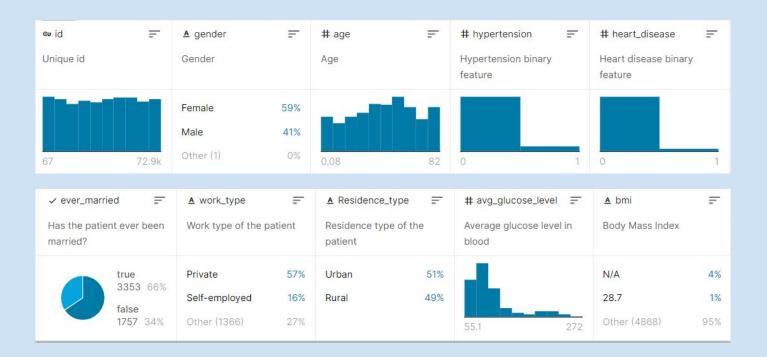


The Features-

- 1. Gender
- 2. Age
- 3. Hypertension binary feature
- 4. Heart disease binary feature
- 5. Has the patient ever been married?
- 6. Work type of the patient
- 7. Residence type of the patient
- 8. Average glucose level in blood
- 9. Body Mass Index
- 10. Smoking status of the patient
- 11. Stroke event



A Little Bit About Our Data...



```
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
data = pd.read_csv("/content/healthcare-dataset-stroke-data.csv")
data.head()
```

	id	gender	age	hypertension	heart_disease	ever_married	work_type	Residence_type	avg_glucose_level	bmi	smoking_status	stroke
0	9046	Male	67.0	0	1	Yes	Private	Urban	228.69	36.6	formerly smoked	1
1	51676	Female	61.0	0	0	Yes	Self-employed	Rural	202.21	NaN	never smoked	1
2	31112	Male	80.0	0	1	Yes	Private	Rural	105.92	32.5	never smoked	1
3	60182	Female	49.0	0	0	Yes	Private	Urban	171.23	34.4	smokes	1
4	1665	Female	79.0	1	0	Yes	Self-employed	Rural	174.12	24.0	never smoked	1



Lets Explore Our Data-

```
<bound method DataFrame.info of</pre>
                                            id gender
                                                        age ...
                                                                  bmi smoking status stroke
                  Male 67.0 ...
                                   36.6 formerly smoked
          51676 Female 61.0
                                            never smoked
                Female 49.0 ... 34.4
                                                  smokes
                Female 79.0 ... 24.0
                                            never smoked
                                            never smoked
                                            never smoked
                Female 35.0
                                            never smoked
                  Male 51.0 ... 25.6
                                        formerly smoked
          44679 Female 44.0 ... 26.2
                                                 Unknown
    [5110 rows x 12 columns]>
[34] data.isnull().sum()
    id
    gender
    hypertension
    heart disease
    ever married
    work type
    Residence type
    avg glucose level
                         201
    smoking status
    stroke
    dtype: int64
```

As we can see the 'bmi' column holds some missing values.

So we need to fill the null values:



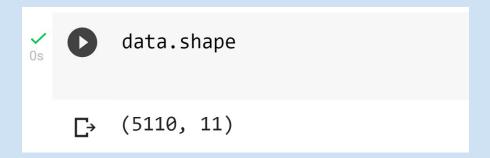
data['bmi'].fillna(data['bmi'].mean(), inplace = True)



Removing an Unnecessary Column-

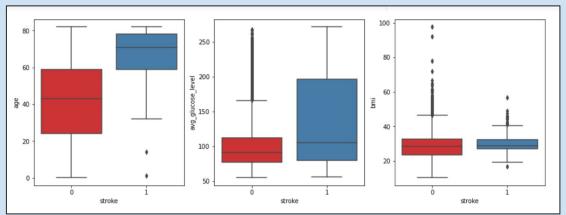
After we explored the data, we decided to remove one column- 'id', because the Id of the person is not a relevant information for our model.

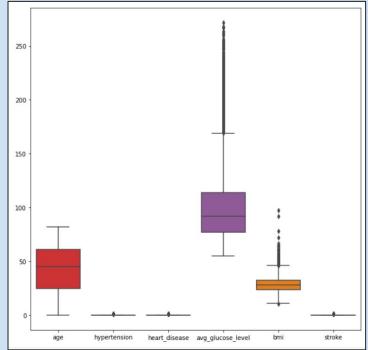
So now we have 11 attributes.



Outliers-

We used subplot function to find the outliers in our data.





Algorithms-

We want to know, based on the 11 features, what are the chances to of having a stroke.

We will use the following techniques:

- Logistic Regression
- SVM
- Decision Trees
- K- nearest neighbour

Train and Test-

We split our data into train and test,

```
x_{train}, x_{test}, y_{train}, y_{test} = train_{test}, y_{test}, y_{test}, y_{train}, y_{test} = 0.2, train_{test}, y_{train}
```

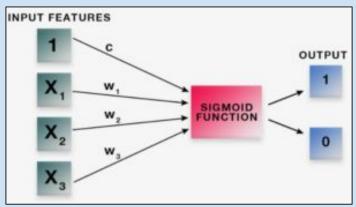
In order to scale the train and test features ,we used StandardScaler - StandardScaler removes the mean and scales each feature/variable to unit variance. This operation is performed feature-wise in an independent way. StandardScaler can be influenced by outliers (if they exist in the dataset) since it involves the estimation of the empirical mean and standard deviation of each feature.

After that, we used fit_transform() on the x_train variable and transform on the x_test variable.

Logistic Regression-

```
√ [382] #Logistic regression-

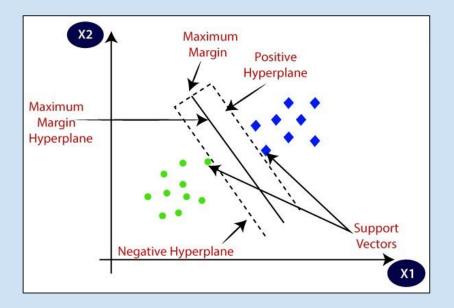
        from sklearn.linear model import LogisticRegression
       from sklearn.metrics import accuracy_score
       logistic_reg = LogisticRegression()
        logistic_reg.fit(x_train_std, y_train)
       y_pred_logistic_reg = logistic_reg.predict(x_test_std)
        accuracy logistic reg = accuracy score(y test, y pred logistic reg)
        print("Logistic Regression accuracy = " + str(accuracy_logistic_reg*100) + "%")
        Logistic Regression accuracy = 94.71624266144813%
```



SVM-

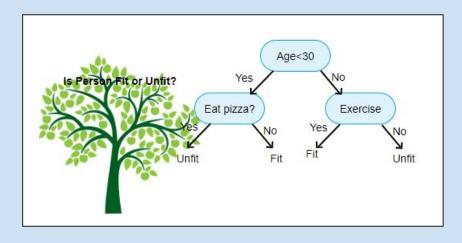
```
✓ [383] #SVM-

        from sklearn.svm import SVC
        svc = SVC()
        svc.fit(x_train_std, y_train)
        y_pred_svc = svc.predict(x_test_std)
        ac_svc = accuracy_score(y_test, y_pred_svc)
        print("SVM accuracy = " + str(ac_svc*100) + "%")
        SVM accuracy = 94.71624266144813%
```



Decision Tree-

```
[390] #Decision tree-
       from sklearn.tree import DecisionTreeClassifier
       decision tree = DecisionTreeClassifier()
       decision tree.fit(x train std. v train)
       decision_tree.feature_importances_
      array([0.04505728, 0.20077799, 0.03420359, 0.02100343, 0.00581771,
             0.03533076, 0.02881657, 0.29332933, 0.26733965, 0.06832369])
/ [386] x train.columns
      Index(['gender', 'age', 'hypertension', 'heart_disease', 'ever_married',
              'work type', 'Residence type', 'avg glucose level', 'bmi',
             'smoking status'l,
            dtype='object')
/ [387] from sklearn.metrics import accuracy score
      y pred decision tree = decision tree.predict(x test_std)
       accuracy_decision_tree = accuracy_score(y_test, y_pred_decision_tree)
       print("decision tree accuracy = " + str(accuracy decision tree*100) + "%")
      decision tree accuracy = 91.6829745596869%
```



K- Nearest Neighbour-

```
#K-Nearest-Neighbour

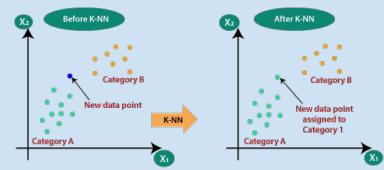
from sklearn.neighbors import KNeighborsClassifier
knn = KNeighborsClassifier()

knn.fit(x_train_std, y_train)

y_pred_knn = knn.predict(x_test_std)

accuracy_knn = accuracy_score(y_test, y_pred_knn)
print("K-nearest neighbour accuracy = " + str(accuracy_knn*100) + "%")

K-nearest neighbour accuracy = 94.71624266144813%
```



Results-

Algorithms-	Logistic regression	SVM	Decision Tree	K- Nearest Neighbour
Accuracy-	94.7162426	94.7162426	91.68297455	94.71624266
	6144813%	6144813%	96869%	144813%

Could We Use Other Algorithms?

Of course.

We could use other algorithms and make the necessary adjustments, and then choose the algorithm with the best accuracy to predict the chances of having a stroke.

מקורות וחומרי עזר-

- https://www.kaggle.com/fedesoriano/stroke-prediction-dataset
- https://towardsdatascience.com/scale-standardize-or-normalize-with-scikit-le-arn-6ccc7d176a02
- https://github.com/riddhi-jain
- https://machinelearningmastery.com/how-to-use-statistics-to-identify-outlier-s-in-data