Data preprocessing

These are the factors

A: Sex (0:male, 1:female)

B: age

C: hypertension (0:doesn't have,1:has)

D: heart_disease (0:doesn't have,1:has)

E: ever_married (0:no,1:yes)

F: work_type is ignored (0:"children", 1:"Govtjov", 2:"Never worked", 3:"Private", 4:"Self-employed)

G: Residence_type (0:Rural,1:Urban)

H: avg_glucose_level

I: BMI

J: smoking_status (0:never smoked, 1:formerly smoked, 2:smokes) "unknown" is ignored

K: stroke(0:no stroke,1:has a stroke)

```
In [127... import pandas as pd
    dataset = pd.read_csv("stroke_data2.csv")
    dataset = dataset.sample(frac=1)
    dataset.reset_index(drop=True, inplace=True)
    dataset
```

Out[127]:		sex	age	hypertension	heart_disease	ever_married	work_type	Residence_type	avg_glucose_level	bmi
	0	1	79	1	0	1	4	1	221.79	30.0
	1	1	47	0	0	0	4	0	154.08	20.2
	2	1	63	0	0	1	4	1	93.58	25.1
	3	1	31	0	0	0	3	0	77.52	21.9
	4	1	9	0	0	1	2	1	64.99	33.2
	•••									
	40844	0	56	0	1	1	4	1	91.92	35.9
	40845	0	49	0	0	1	4	0	65.47	24.1
	40846	1	25	0	0	1	4	0	100.98	28.2
	40847	0	28	0	0	1	4	1	86.94	41.1
	40848	1	42	0	0	1	4	0	95.36	25.1

40849 rows × 11 columns

There are 40849 rows of patients' data. After shuffling the rows, we choose the first 40000 rows as the training data, and 800 rows as validation data, and the last 10 rows as testing data.

```
In [128... from sklearn.preprocessing import StandardScaler

x = dataset.iloc[:,:-1]
y= dataset.iloc[:,-1]
x_train = x.iloc[:40000]
y_train = y.iloc[:40000]
x_valid = x.iloc[40039:40839]
```

```
y_valid = y.iloc[40039:40839]
x_test = x.iloc[40839:]
y_test = y.iloc[40839:]

scaler = StandardScaler()
x_train = scaler.fit_transform(x_train)
x_valid = scaler.transform(x_valid)
x test = scaler.transform(x test)
```

Logistic Regression

```
In [129... from sklearn.linear_model import LogisticRegression
    model_LR = LogisticRegression(penalty='12', C=0.0001, solver="lbfgs")
    model_LR.fit(x_train, y_train)

Out[129]: LogisticRegression
    LogisticRegression(C=0.0001)
```

Multilayer Perceptron

DecisionTreeClassifier

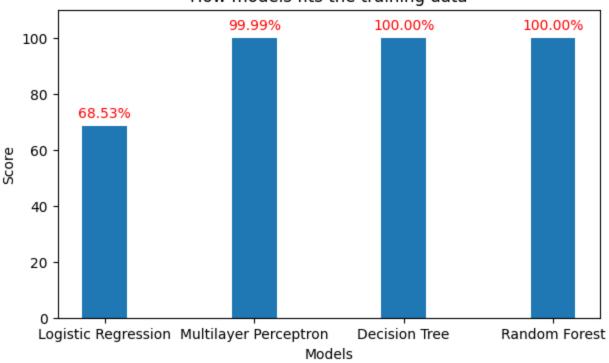
Random Forest

Evaluation

How models fits the training data

```
import matplotlib.pyplot as plt
In [182...
         import numpy as np
         labels = ["Logistic Regression", "Multilayer Perceptron", "Decision Tree", "Random Fores
         trainScore = np.array([model LR.score(x train, y train), model MLP.score(x train, y train),
         trainScore = trainScore*100
         size = len(labels)
         plt.figure(figsize=(7, 4))
         for i in range(size):
             plt.text(i,trainScore[i]+3,f'{trainScore[i]:.2f}%',ha = 'center',color='red')
         plt.xlabel('Models')
         plt.ylabel('Score')
         plt.title('How models fits the training data')
         plt.bar(range(size), trainScore, width=0.3)
         plt.xticks(range(size), labels)
         plt.ylim(0,110)
         plt.show()
```

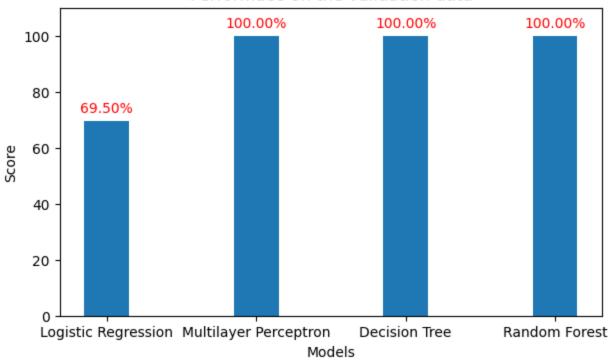
How models fits the training data



The performance of models on the validation data

```
In [183...
    trainScore = np.array([model_LR.score(x_valid,y_valid),model_MLP.score(x_valid,y_valid),
    trainScore = trainScore*100
    size = len(labels)
    plt.figure(figsize=(7, 4))
    for i in range(size):
        plt.text(i,trainScore[i]+3,f'{trainScore[i]:.2f}%',ha = 'center',color='red')
    plt.xlabel('Models')
    plt.ylabel('Score')
    plt.title('Performace on the validation data')
    plt.bar(range(size),trainScore,width=0.3)
    plt.xticks(range(size), labels)
    plt.ylim(0,110)
    plt.show()
```

Performace on the validation data



We conclude that random forest classifier model works the best. Therefore, we choose random forest model for our final testing.

Final testing

```
y final pred = pd.Series(model MLP.predict(x test))
In [260...
         y final prob = pd.DataFrame(model MLP.predict proba(x test)).iloc[:,1]
         leng = len(y final pred)
         cnt = 0
         for i in range(leng):
             print(f"Patient #{i+1}: Probability of having stroke is {y final prob.iloc[i]*100:.2
             if y final pred.iloc[i] == 0:
                 print("predict that do not have a stroke.", end=" ")
             elif y final pred.iloc[i] == 1:
                 print("predict that have a stroke.", end=" ")
             if y final pred.iloc[i] == y test.iloc[i]:
                 cnt+=10
                 print('\033[32m'+"We predicted correctly."+'\033[0m')
             else:
                 print('\033[31m'+"We didn't predict correctly."+'\033[0m')
         plt.figure(figsize=(3, 3))
         plt.pie([cnt, 100 - cnt], labels=[f'Correctly', f'Non-Correctly'],colors=['blue', 'red']
         plt.title('Accuracy of Prediction')
         plt.legend([f'correctly: {cnt}%', f'incorrectly{100 - cnt}%'],bbox to anchor=(0.7,0.3))
         plt.show()
        Patient #1: Probability of having stroke is 0.00%, predict that do not have a stroke. We
         predicted correctly.
        Patient #2: Probability of having stroke is 0.04%, predict that do not have a stroke. We
         predicted correctly.
        Patient #3: Probability of having stroke is 100.00%, predict that have a stroke. We pred
        icted correctly.
        Patient #4: Probability of having stroke is 0.00%, predict that do not have a stroke. We
         predicted correctly.
         Patient #5: Probability of having stroke is 0.00%, predict that do not have a stroke. We
```

predicted correctly.

Patient #6: Probability of having stroke is 100.00%, predict that have a stroke. We predicted correctly.

Patient #7: Probability of having stroke is 0.00%, predict that do not have a stroke. We predicted correctly.

Patient #8: Probability of having stroke is 100.00%, predict that have a stroke. We predicted correctly.

Patient #9: Probability of having stroke is 100.00%, predict that have a stroke. We predicted correctly.

Patient #10: Probability of having stroke is 0.00%, predict that do not have a stroke. We predicted correctly.

Accuracy of Prediction

