# **Ensemble Learning**

Agenda:

Ensemble learning

Bagging Boosting Stacking

- Random Forest
- Use Diabetes to perform RF
- Improve the model using GridSearch SV
- Build the best RF model
- Most important features
- Conclude

# Accessing the data

```
In [2]: import numpy as np
    import pandas as pd
    import seaborn as sns
    import matplotlib.pyplot as plt
```

```
In [3]: db=pd.read_csv('diabetes.csv')
db
```

Out[3]:		Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	ВМІ	DiabetesPedigreeFunction	Age	Outcome
	0	6	148	72	35	0	33.6	0.627	50	1
	1	1	85	66	29	0	26.6	0.351	31	0
	2	8	183	64	0	0	23.3	0.672	32	1
	3	1	89	66	23	94	28.1	0.167	21	0
	4	0	137	40	35	168	43.1	2.288	33	1
	763	10	101	76	48	180	32.9	0.171	63	0
	764	2	122	70	27	0	36.8	0.340	27	0
	765	5	121	72	23	112	26.2	0.245	30	0
	766	1	126	60	0	0	30.1	0.349	47	1
	767	1	93	70	31	0	30.4	0.315	23	0

768 rows × 9 columns

### Separating into target and features

```
In [4]: y=db['Outcome']
y.value_counts()
```

Out[4]: 0 500 1 268

Name: Outcome, dtype: int64

```
In [5]: X=db.drop(['Outcome'],axis=1)
X
```

Out[5]:	Pregnancies Gluc		Glucose	BloodPressure	SkinThickness	Insulin	вмі	DiabetesPedigreeFunction	Age
	0	6	148	72	35	0	33.6	0.627	50
	1	1	85	66	29	0	26.6	0.351	31
	2	8	183	64	0	0	23.3	0.672	32
	3	1	89	66	23	94	28.1	0.167	21
	4	0	137	40	35	168	43.1	2.288	33
	763	10	101	76	48	180	32.9	0.171	63
	764	2	122	70	27	0	36.8	0.340	27
	765	5	121	72	23	112	26.2	0.245	30
	766	1	126	60	0	0	30.1	0.349	47
	767	1	93	70	31	0	30.4	0.315	23

768 rows × 8 columns

### **Standardisation**

### Splitting the data to train and test

### **Building the model- Random Forest Classifier**

```
In [8]: from sklearn.ensemble import RandomForestClassifier
    rfc=RandomForestClassifier(random_state=100)
    rfc.fit(X_train,y_train)
```

Out[8]: RandomForestClassifier(random\_state=100)

# **Model performance**

```
In [9]: from sklearn.metrics import classification_report, confusion_matrix
        cm=confusion_matrix(y_test,rfc.predict(X_test))
        report=classification_report(y_test,rfc.predict(X_test))
        print('CM:\n',cm)
        print('Report:\n',report)
        CM:
         [[85 16]
         [26 27]]
        Report:
                                     recall f1-score
                       precision
                                                        support
                   0
                           0.77
                                      0.84
                                                0.80
                                                           101
                           0.63
                                      0.51
                                                0.56
                                                            53
                                                0.73
                                                           154
            accuracy
           macro avg
                           0.70
                                      0.68
                                                0.68
                                                           154
        weighted avg
                           0.72
                                      0.73
                                                           154
```

## Hyper parameter tuning using GridSearchCV

### **Best parameters**

```
In [12]: rfc_gs.best_params_
Out[12]: {'criterion': 'entropy', 'n_estimators': 122}
```

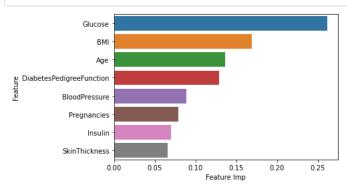
### Building a revised RF model

```
In [13]: rfc_new=RandomForestClassifier(n_estimators=122,
                                         criterion='entropy',random_state=10)
In [14]: rfc_new.fit(X_train,y_train)
Out[14]: RandomForestClassifier(criterion='entropy', n_estimators=122, random_state=10)
In [15]: # Perfromance
         cm=confusion_matrix(y_test,rfc_new.predict(X_test))
         report=classification_report(y_test,rfc_new.predict(X_test))
         print('CM:\n',cm)
         print('Report:\n',report)
         CM:
          [[84 17]
          [26 27]]
         Report:
                                      recall f1-score
                        precision
                                                         support
                     0
                             0.76
                                       0.83
                                                 0.80
                                                            101
                             0.61
                                       0.51
                                                 0.56
                                                             53
                                                 0.72
                                                            154
             accuracy
            macro avg
                             0.69
                                       0.67
                                                 0.68
                                                            154
         weighted avg
                             0.71
                                       0.72
                                                 0.71
                                                            154
```

### Important features

```
In [16]: rfc_new.feature_importances_
Out[16]: array([0.07865853, 0.26144447, 0.08867384, 0.06591887, 0.07007926,
                  0.16953733, 0.12920921, 0.13647848])
In [17]: # Creating a DF
          df=pd.DataFrame({'Feature':X.columns,'Feature Imp':rfc_new.feature_importances_})
Out[17]:
                             Feature Feature Imp
           0
                         Pregnancies
                                       0.078659
                                       0.261444
           1
                             Glucose
           2
                       BloodPressure
                                       0.088674
           3
                        SkinThickness
                                        0.065919
            4
                                        0.070079
           5
                                BMI
                                       0.169537
           6 DiabetesPedigreeFunction
                                       0.129209
                                       0.136478
In [20]: df=df.sort_values(['Feature Imp'],ascending=False)
          df
Out[20]:
                             Feature Feature Imp
           1
                             Glucose
                                       0.261444
           5
                                BMI
                                       0.169537
           7
                                Age
                                       0.136478
                                       0.129209
           6
              DiabetesPedigreeFunction
           2
                                       0.088674
                       BloodPressure
           0
                                       0.078659
                         Pregnancies
                                       0.070079
           4
                              Insulin
           3
                        SkinThickness
                                       0.065919
In [21]: # Adding one column
          df['Feature Imp Cum']=df['Feature Imp'].cumsum()
Out[21]:
                             Feature Feature Imp Feature Imp Cum
                             Glucose
                                        0.261444
                                                        0.261444
           5
                                BMI
                                       0.169537
                                                        0.430982
           7
                                       0.136478
                                                        0.567460
                                Age
                                       0.129209
                                                        0.696669
           6
              DiabetesPedigreeFunction
           2
                       BloodPressure
                                       0.088674
                                                        0.785343
                                       0.078659
                                                        0.864002
           0
                         Pregnancies
                                       0.070079
                                                        0.934081
                              Insulin
           3
                        SkinThickness
                                       0.065919
                                                        1.000000
```

In [22]: sns.barplot(x=df['Feature Imp'],y=df['Feature'],data=df);



In [ ]: