

## module 불러오기

In [ ]:

```
1 # data load
2 from sklearn.datasets import load_breast_cancer
3
4 # train test split
5 from sklearn.model_selection import train_test_split
6
7 # model
8 from sklearn.tree import DecisionTreeClassifier
9 from sklearn.ensemble import RandomForestClassifier
10 from sklearn.svm import SVC
11 from sklearn.linear_model import SGDClassifier
12 from sklearn.linear_model import LogisticRegression
13
14 # report
15 from sklearn.metrics import classification_report
```

## breast cancer 데이터 불러오기

In [ ]:

```
1 breast_cancer = load_breast_cancer()
2 print(breast_cancer)
```

In [ ]:

```
1 breast_cancer.feature_names
```

Out[16]:

```
array(['mean radius', 'mean texture', 'mean perimeter', 'mean area',
      'mean smoothness', 'mean compactness', 'mean concavity',
      'mean concave points', 'mean symmetry', 'mean fractal dimension',
      'radius error', 'texture error', 'perimeter error', 'area error',
      'smoothness error', 'compactness error', 'concavity error',
      'concave points error', 'symmetry error',
      'fractal dimension error', 'worst radius', 'worst texture',
      'worst perimeter', 'worst area', 'worst smoothness',
      'worst compactness', 'worst concavity', 'worst concave points',
      'worst symmetry', 'worst fractal dimension'], dtype='<U23')
```

In [ ]:

```
1 breast_cancer.target_names
```

Out[3]:

```
array(['malignant', 'benign'], dtype='<U9')
```



In [ ]:

```
1 # RandomForest
2 model_random_forest = RandomForestClassifier()
3 model_random_forest.fit(X_train, y_train)
```

Out[7]:

```
RandomForestClassifier(bootstrap=True, ccp_alpha=0.0, class_weight=None,
                        criterion='gini', max_depth=None, max_features='auto',
                        max_leaf_nodes=None, max_samples=None,
                        min_impurity_decrease=0.0, min_impurity_split=None,
                        min_samples_leaf=1, min_samples_split=2,
                        min_weight_fraction_leaf=0.0, n_estimators=100,
                        n_jobs=None, oob_score=False, random_state=None,
                        verbose=0, warm_start=False)
```

In [ ]:

```
1 # SVM
2 model_svc = SVC()
3 model_svc.fit(X_train, y_train)
```

Out[8]:

```
SVC(C=1.0, break_ties=False, cache_size=200, class_weight=None, coef0=0.0,
    decision_function_shape='ovr', degree=3, gamma='scale', kernel='rbf',
    max_iter=-1, probability=False, random_state=None, shrinking=True,
    tol=0.001, verbose=False)
```

In [ ]:

```
1 # SGD
2 model_sgd = SGDClassifier()
3 model_sgd.fit(X_train, y_train)
```

Out[9]:

```
SGDClassifier(alpha=0.0001, average=False, class_weight=None,
              early_stopping=False, epsilon=0.1, eta0=0.0, fit_intercept=True,
              l1_ratio=0.15, learning_rate='optimal', loss='hinge',
              max_iter=1000, n_iter_no_change=5, n_jobs=None, penalty='l2',
              power_t=0.5, random_state=None, shuffle=True, tol=0.001,
              validation_fraction=0.1, verbose=0, warm_start=False)
```

In [ ]:

```
1 # Logistic Regression
2 model_LR = LogisticRegression()
3 model_LR.fit(X_train, y_train)
```

/usr/local/lib/python3.7/dist-packages/sklearn/linear\_model/\_logistic.py:940: ConvergenceWarning: lbfgs failed to converge (status=1):  
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

Increase the number of iterations (max\_iter) or scale the data as shown in:

<https://scikit-learn.org/stable/modules/preprocessing.html> (<https://scikit-learn.org/stable/modules/preprocessing.html>)

Please also refer to the documentation for alternative solver options:

[https://scikit-learn.org/stable/modules/linear\\_model.html#logistic-regression](https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression) ([https://scikit-learn.org/stable/modules/linear\\_model.html#logistic-regression](https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression))  
extra\_warning\_msg=\_LOGISTIC\_SOLVER\_CONVERGENCE\_MSG)

Out[10]:

```
LogisticRegression(C=1.0, class_weight=None, dual=False, fit_intercept=True,
                    intercept_scaling=1, l1_ratio=None, max_iter=100,
                    multi_class='auto', n_jobs=None, penalty='l2',
                    random_state=None, solver='lbfgs', tol=0.0001, verbose=0,
                    warm_start=False)
```

## 모델 사용해보고 평가하기

In [ ]:

```
1 # Decision Tree
2 y_pred = model_tree.predict(X_test)
3 print(classification_report(y_pred, y_test))
```

	precision	recall	f1-score	support
0	0.92	0.88	0.90	41
1	0.93	0.96	0.95	73
accuracy			0.93	114
macro avg	0.93	0.92	0.92	114
weighted avg	0.93	0.93	0.93	114

In [ ]:

```
1 # Random Forest
2 y_pred = model_random_forest.predict(X_test)
3 print(classification_report(y_pred, y_test))
```

	precision	recall	f1-score	support
0	0.87	0.94	0.91	36
1	0.97	0.94	0.95	78
accuracy			0.94	114
macro avg	0.92	0.94	0.93	114
weighted avg	0.94	0.94	0.94	114

In [ ]:

```
1 # SVM
2 y_pred = model_svc.predict(X_test)
3 print(classification_report(y_pred, y_test))
```

	precision	recall	f1-score	support
0	0.77	0.97	0.86	31
1	0.99	0.89	0.94	83
accuracy			0.91	114
macro avg	0.88	0.93	0.90	114
weighted avg	0.93	0.91	0.92	114

In [ ]:

```
1 # SGD
2 y_pred = model_sgd.predict(X_test)
3 print(classification_report(y_pred, y_test))
```

	precision	recall	f1-score	support
0	0.92	0.77	0.84	47
1	0.85	0.96	0.90	67
accuracy			0.88	114
macro avg	0.89	0.86	0.87	114
weighted avg	0.88	0.88	0.87	114

In [ ]:

```
1 # Logistic Regression
2 y_pred = model_LR.predict(X_test)
3 print(classification_report(y_pred, y_test))
```

	precision	recall	f1-score	support
0	0.87	0.94	0.91	36
1	0.97	0.94	0.95	78
accuracy			0.94	114
macro avg	0.92	0.94	0.93	114
weighted avg	0.94	0.94	0.94	114

Random Forest, logistic Regression을 사용한다. 암은 반드시 발견되어야 하고, 한명의 환자도 놓치면 안 되며 recall이 안정적인 모델을 택하여야한다.

In [ ]:

```
1
```

In [ ]:

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1
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In [ ]:

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