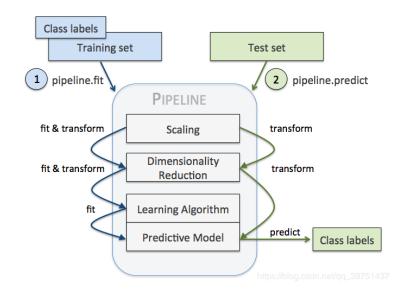
모형진단과 교차검증 (실습)

Evaluation & Cross Validation

파이프라인 실습 #1

- 파이프 라인(Pipeline)
 - 사이킷런의 Pipeline 클래스는 연속된 변환을 순차적으로 처리할 수 있는 기능을 제공하는 유용한 래퍼(Wrapper) 도구



파이프라인 실습 #1

- 데이터셋: 유방암 데이터
- 학습/시험 데이터: X, 학습/시험 데이터 라벨: Y

① 데이터 로드

```
1 # https://scikit-learn.org/stable/modules/generated/sklearn.datasets.load_breast_cancer.html
2 # Dimensionality: 30, Classes: 2
3 # 212(M-유방암)-label('0'), 357(B-정상인) - label('1')
4 from sklearn.datasets import load_breast_cancer
5 cancer = load_breast_cancer()
6 X = cancer.data
7 Y = cancer.target
```

② 데이터 분할 (학습/테스트)

```
1 # https://scikit-learn.org/stable/modules/generated/sklearn.model_selection.train_test_split.htm
2
3 # 학습 데이터 분할
4 from sklearn.model_selection import train_test_split
5 X_train, X_test, Y_train, Y_test = train_test_split(X,Y,test_size=0.2, stratify=Y, random_state=1)
```

파이프라인 실습 #1

- 데이터셋: 유방암 데이터
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③ 파이프라인 모듈 설계

```
# https://scikit-learn.org/stable/modules/generated/sklearn.pipeline.Pipeline.html
2 # 파이파라인 기능을 이용한 모듈 설계

4 from sklearn.preprocessing import StandardScaler
5 from sklearn.decomposition import PCA
6 from sklearn.linear_model import LogisticRegression
7 from sklearn.pipeline import make_pipeline

8 pipeline = make_pipeline(StandardScaler(), PCA(n_components=4), LogisticRegression())
```

④ 모델 학습 및 평가

```
1 # 모델 학습 및 테스트
2 pipeline.fit(X_train,Y_train)
3 Y_Pred = pipline.predict(X_test)
0.9736842105263158

1 # 모델 평가
2 from sklearn.metrics import accuracy_score
3 accuracy_score(Y_test, Y_Pred)
```

0.9736842105263158

- 데이터셋: 유방암 데이터
- 학습/시험 데이터: X, 학습/시험 데이터 라벨: Y

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```

- 데이터셋: 유방암 데이터
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4 from sklearn.preprocessing import StandardScaler
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7 from sklearn.pipeline import make_pipeline
8
9 pipeline = make_pipeline(StandardScaler(), PCA(n_components=4), LogisticRegression())
```

④ 교차검증

```
1 # https://scikit-learn.org/stable/modules/generated/sklearn.model_selection.cross_validate.html#
2
3 from sklearn.model_selection import cross_validate
4 scores = cross_validate(pipeline, X_train, Y_train, cv=10, return_train_score=True)
```

- 데이터셋: 유방암 데이터
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⑤ 과적합 여부 분석

```
1 import numpy as np
2
3 print('CV Validation Accuracy scores: ', scores['train_score'])
4 print('CV Validation Accuracy: %.3f +/- %.3f' %(np.mean(scores['train_score']), np.std(scores['train_score'])))
CV Validation Accuracy scores: [0.96577017 0.96577017 0.96577017 0.96577017 0.96821516 0.96585366 0.97073171 0.96829268 0.97560976 0.96585366]
CV Validation Accuracy: 0.968 +/- 0.003

1 import numpy as np
2
3 print('CV Validation Accuracy scores: ', scores['test_score'])
4 print('CV Validation Accuracy: %.3f +/- %.3f' %(np.mean(scores['test_score']), np.std(scores['test_score'])))
CV Validation Accuracy scores: [0.97826087 0.97826087 0.95652174 1. 0.95652174 0.97777778 0.93333333 0.95555556 0.91111111 1. ]
CV Validation Accuracy: 0.965 +/- 0.027
```

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⑥ 최적 모델 탐색

```
1 from sklearn.model_selection import GridSearchCV
2
3 parameters = {}
4 gs = GridSearchCV(pipeline, parameters, scoring='accuracy',cv=10)
5 gs.fit(X_train,Y_train)
```

```
1 best = gs.best_estimator_
1 gs.cv_results_
```

```
('mean fit time': array([0.00662093]),
'mean score time': array([0.00051041]),
'mean test score': array([0.9647343]),
'params': [{}],
'rank test score': array([1], dtype=int32),
'split0 test score': array([0.97826087]),
'split1 test score': array([0.97826087]),
'split2 test score': array([0.95652174]),
'split3 test score': array([1.]),
'split4 test score': array([0.95652174]),
'split5 test score': array([0.97777778]),
'split6 test score': array([0.93333333]),
'split7 test score': array([0.95555556]),
'split8 test score': array([0.91111111]),
'split9 test score': array([1.]),
'std fit time': array([0.00164769]),
'std score time': array([3.95498696e-05]),
'std test score': array([0.02665336])}
```

- 데이터셋: 유방암 데이터
- 학습/시험 데이터: X, 학습/시험 데이터 라벨: Y

⑦ 최적 모델 평가

```
1 from sklearn.metrics import accuracy_score
2
3 Y_train_Pred = best.predict(X_train)
4 accuracy_score(Y_train, Y_train_Pred)
5
```

0.967032967032967

```
1 from sklearn.metrics import accuracy_score
2
3 Y_test_Pred = best.predict(X_test)
4 accuracy_score(Y_test, Y_test_Pred)
5
```

0.9736842105263158

최적 모델 찾기 실습 #1

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5 cancer = load_breast_cancer()
6 X = cancer.data
7 Y = cancer.target
```

② 데이터 분할 (학습/테스트)

```
1 # https://scikit-learn.org/stable/modules/generated/sklearn.model_selection.train_test_split.htm
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3 # 학습 데이터 분할
4 from sklearn.model_selection import train_test_split
5 X_train, X_test, Y_train, Y_test = train_test_split(X,Y,test_size=0.2, stratify=Y, random_state=1)
```

최적 모델 찾기 실습 #1

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③ 파이프라인 모듈 설계

```
1 # https://scikit-learn.org/stable/modules/generated/sklearn.pipeline.Pipeline.html
2 # 파이파라인 기능을 이용한 모듈 설계
3
4 from sklearn.preprocessing import StandardScaler
5 from sklearn.svm import SVC
6 from sklearn.pipeline import Pipeline
7 from sklearn.model_selection import GridSearchCV
8
9 # 파라미터 Parsing
10 estimators = [('normalization', StandardScaler()), ('clf', SVC())]
11 pipe = Pipeline(estimators)
```

④ GridSearch 파라미터 세팅

```
1 # https://scikit-learn.org/stable/modules/generated/sklearn.model_selection.ParameterGrid.html#sklearn.model_selection.ParameterGrid
2
3 from sklearn.model_selection import ParameterGrid
4 grid = [{'clf_kernel': [['linear']], 'clf_C': [[0.001], [0.01], [0.1], [1], [10], [1000]]},
5 {'clf_kernel': [['rbf']], 'clf_gamma': [[0.001], [0.01], [0.1], [1], [10], [1000]], 'clf_C': [[0.001], [0.01], [0.1], [1], [10], [1000]]}}
6 grid_param = ParameterGrid(grid)
8 list(grid_param)
```

최적 모델 찾기 실습 #1

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⑤ GridSearch 정의 및 최적 모델 탐색

```
1 gs = GridSearchCV(pipe, grid_param, scoring='accuracy', cv=10, n_jobs=1)

1 gs.fit(X_train,Y_train)

1 print(gs.best_score_)

0.9758454106280192

1 print(gs.best_params_)
{'clf_C': 10, 'clf_gamma': 0.001, 'clf_kernel': 'rbf'}
```

⑥ 최적 모델 평가

```
1 best_model = gs.best_estimator_
2 Y_test_pred = best_model.predict(X_test)

1 from sklearn.metrics import accuracy_score
2
3 Y_test_Pred = best_model.predict(X_test)
4 accuracy_score(Y_test, Y_test_Pred)
```

0.9824561403508771