



# TA Session 2

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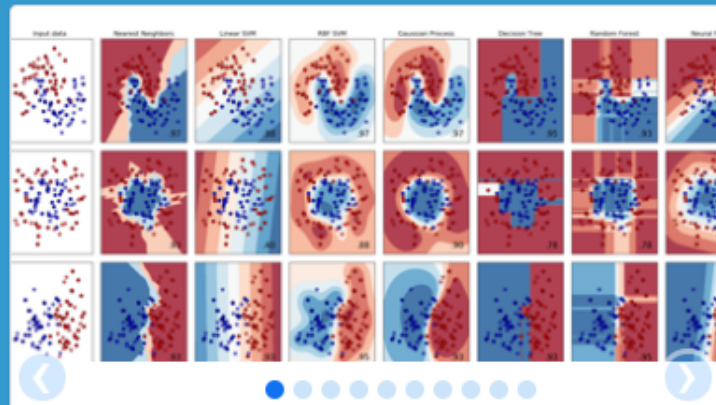
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Wonjong Rhee, Hyunghun Cho, Daeyoung Choi

Seoul National University

Graduate School of Convergence Science and Technology

Applied Data Science Lab.



# scikit-learn

*Machine Learning in Python*

- Simple and efficient tools for data mining and data analysis
- Accessible to everybody, and reusable in various contexts
- Built on NumPy, SciPy, and matplotlib
- Open source, commercially usable - BSD license

## Classification

Identifying to which category an object belongs to.

**Applications:** Spam detection, Image recognition.

**Algorithms:** SVM, nearest neighbors, random forest, ...

[— Examples](#)

## Regression

Predicting a continuous-valued attribute associated with an object.

**Applications:** Drug response, Stock prices.

**Algorithms:** SVR, ridge regression, Lasso, ...

[— Examples](#)

## Clustering

Automatic grouping of similar objects into sets.

**Applications:** Customer segmentation, Grouping experiment outcomes

**Algorithms:** k-Means, spectral clustering, mean-shift, ...

[— Examples](#)

## Dimensionality reduction

Reducing the number of random variables to consider.

**Applications:** Visualization, Increased efficiency

**Algorithms:** PCA, feature selection, non-negative matrix factorization.

[— Examples](#)

## Model selection

Comparing, validating and choosing parameters and models.

**Goal:** Improved accuracy via parameter tuning

**Modules:** grid search, cross validation, metrics.

[— Examples](#)

## Preprocessing

Feature extraction and normalization.

**Application:** Transforming input data such as text for use with machine learning algorithms.

**Modules:** preprocessing, feature extraction.

[— Examples](#)

## sklearn.model\_selection: Model Selection

**User guide:** See the [Cross-validation: evaluating estimator performance](#), [Tuning the hyper-parameters of an estimator](#) and [Learning curve](#) sections for further details.

### Splitter Classes

<code>model_selection.GroupKFold ([n_splits])</code>	K-fold iterator variant with non-overlapping groups.
<code>model_selection.GroupShuffleSplit ([...])</code>	Shuffle-Group(s)-Out cross-validation iterator
<code>model_selection.KFold ([n_splits, shuffle, ...])</code>	K-Folds cross-validator
<code>model_selection.LeaveOneGroupOut ()</code>	Leave One Group Out cross-validator
<code>model_selection.LeavePGroupsOut (n_groups)</code>	Leave P Group(s) Out cross-validator
<code>model_selection.LeaveOneOut ()</code>	Leave-One-Out cross-validator
<code>model_selection.LeavePOut (p)</code>	Leave-P-Out cross-validator
<code>model_selection.PredefinedSplit (test_fold)</code>	Predefined split cross-validator
<code>model_selection.RepeatedKFold ([n_splits, ...])</code>	Repeated K-Fold cross validator.
<code>model_selection.RepeatedStratifiedKFold ([...])</code>	Repeated Stratified K-Fold cross validator.
<code>model_selection.ShuffleSplit ([n_splits, ...])</code>	Random permutation cross-validator
<code>model_selection.StratifiedKFold ([n_splits, ...])</code>	Stratified K-Folds cross-validator
<code>model_selection.StratifiedShuffleSplit ([...])</code>	Stratified ShuffleSplit cross-validator
<code>model_selection.TimeSeriesSplit ([n_splits, ...])</code>	Time Series cross-validator

## sklearn.ensemble : Ensemble Methods

The `sklearn.ensemble` module includes ensemble-based methods for classification, regression and anomaly detection.

**User guide:** See the [Ensemble methods](#) section for further details.

<code>ensemble.AdaBoostClassifier</code> ([...])	An AdaBoost classifier.
<code>ensemble.AdaBoostRegressor</code> ([base_estimator, ...])	An AdaBoost regressor.
<code>ensemble.BaggingClassifier</code> ([base_estimator, ...])	A Bagging classifier.
<code>ensemble.BaggingRegressor</code> ([base_estimator, ...])	A Bagging regressor.
<code>ensemble.ExtraTreesClassifier</code> ([...])	An extra-trees classifier.
<code>ensemble.ExtraTreesRegressor</code> ([n_estimators, ...])	An extra-trees regressor.
<code>ensemble.GradientBoostingClassifier</code> ([loss, ...])	Gradient Boosting for classification.
<code>ensemble.GradientBoostingRegressor</code> ([loss, ...])	Gradient Boosting for regression.
<code>ensemble.IsolationForest</code> ([n_estimators, ...])	Isolation Forest Algorithm
<code>ensemble.RandomForestClassifier</code> ([...])	A random forest classifier.
<code>ensemble.RandomForestRegressor</code> ([...])	A random forest regressor.
<code>ensemble.RandomTreesEmbedding</code> ([...])	An ensemble of totally random trees.
<code>ensemble.VotingClassifier</code> (estimators[, ...])	Soft Voting/Majority Rule classifier for unfitted estimators.

## Model validation

<code>model_selection.cross_validate</code> (estimator, X)	Evaluate metric(s) by cross-validation and also record fit/score times.
<code>model_selection.cross_val_predict</code> (estimator, X)	Generate cross-validated estimates for each input data point
<code>model_selection.cross_val_score</code> (estimator, X)	Evaluate a score by cross-validation
<code>model_selection.learning_curve</code> (estimator, X, y)	Learning curve.
<code>model_selection.permutation_test_score</code> (...)	Evaluate the significance of a cross-validated score with permutations
<code>model_selection.validation_curve</code> (estimator, ...)	Validation curve.

# Task 1: LOOCV and K-fold cross validation



- ISLR/Chapter 5.ipynb 을 실행하고 결과를 확인해 보세요.

# Task 2: Tree based methods

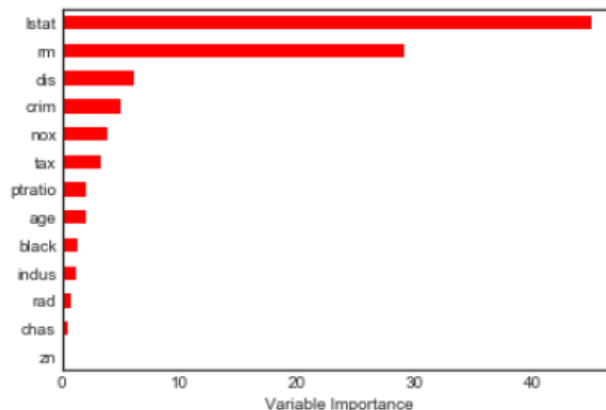


- ISLR/Chapter 8.ipynb 을 실행하고 결과를 확인해 보세요.

# Task 3: combining Task 1 and 2



- 1. Chapter 8.ipynb에 있는 Boston dataset을 이용하여 Chapter 5.ipynb의 Figure 5.2와 Figure 5.4 (오른쪽 10-fold CV만) 를 재현하세요. 알고리즘은 RandomForest를 사용하며 10-fold CV, hyperparameter는 feature의 개수(1~10개)로 합니다.
- 2. 1번에 해당하는 ISLR의 text를 읽고 1번의 결과를 설명하시오. 책의 관련 내용 (178p, 182p와 그 주변 page들) 을 읽고 이해하여 적으시면 됩니다.
- 3. Boston dataset으로 RandomForest 모델을 만든 후 다음과 같이 feature importance 그림을 그려보세요. Feature의 개수, tree의 개수, data split은 자유입니다.



- 4. (advanced) 1번을 GradientBoostingRegressor function을 이용해 재현해보세요. Hyperparameter는 n\_estimators로 합니다. (range는 자유)



- Task는 구현 자체보다는 **결과에 대한 분석과 이해를 중심으로 학습** 하시기 바랍니다.
- 제공된 2018\_fall\_session\_2\_task.ipynb 파일에 코드를 작성하고 설명을 단 뒤 ipynb 파일을 제출해주세요.
- 제출시간은 진행상황을 보가며 정하도록 하겠습니다.
- 다음의 이메일로 [choid@snu.ac.kr](mailto:choid@snu.ac.kr) (최대영 조교) 로 제출하시고 보내실 때 반드시 **조별 대표자 성명을 기록**해주시기 바랍니다.