

# Artificial Intelligence HW2 Report

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## 實作簡介

使用環境為 Python 3.8.7，使用的第三方套件請參照 Appendix. B.

### ➤ Decision Tree

訓練時，從根節點出發，將 data 和對應的 label 傳入建出一棵樹。

每個節點在一開始的時候會先檢查此節點是否應為葉節點。檢查的依據為：傳入的 label 中只剩下一個 unique 的值、節點深度抵達限制或傳入的 sample 數量低於限制。

若此節點應為葉節點，則從 label 中選出出現次數最多的最為此節點的回答；若非葉節點則挑出幾個 feature，在納入考慮的 feature 中挑出 Gini's impurity 最小的 threshold 值作為此節點在之後預測時的分割依據，並將 data 和 label 以此 threshold 分為兩組作為兩個 child node 的 data 和 label 進行遞迴建樹。

#### ✓ Feature bagging (attribute bagging)

在分割節點時只考慮部分的 feature，而不計算所有 feature，選擇方式為隨機挑選  $\min(F(m), m_{splitable})$  個 feature。其中  $m$  為 feature 數量， $F(m)$  為  $\sqrt{m}$  (在 extremely random 的情況為 1)。

預測時則根據傳入的 data 尋訪建立好的樹，抵達葉節點時給出葉節點的答案。

### ➤ Random Forest

訓練時，使用傳入的 data 和 label 分別訓練  $n$  棵 Decision Tree。

預測時，根據每棵樹對同一筆 data 的回答，選出最多棵樹給出的答案 (majority vote)。

#### ✓ Tree bagging

在訓練時將 training set 分為跟樹數量同樣的  $n$  組，訓練第  $i$  棵樹的時後不使用第  $i$  組 data，來做到每棵樹使用不一樣的 subset。

## 結果統計

以下為使用 5 棵樹、深度限制 8、sample 數無限制的 Random Forest 執行 10 個 episode 的 K-fold 驗證(k=8)的平均結果：

| Dataset           | Iris    | Bezdek Iris | Breast Cancer | Glass  | Ionosphere | Wine    |
|-------------------|---------|-------------|---------------|--------|------------|---------|
| Training Accuracy | 100.00% | 100.00%     | 91.91%        | 97.38% | 98.61%     | 100.00% |
| Testing Accuracy  | 94.35%  | 94.08%      | 76.43%        | 72.08% | 92.28%     | 96.30%  |
| Time Elapsed      | < 3s    | < 3s        | 9s            | 22s    | 82s        | 13s     |

## 實驗探討

### ➤ 執行時間

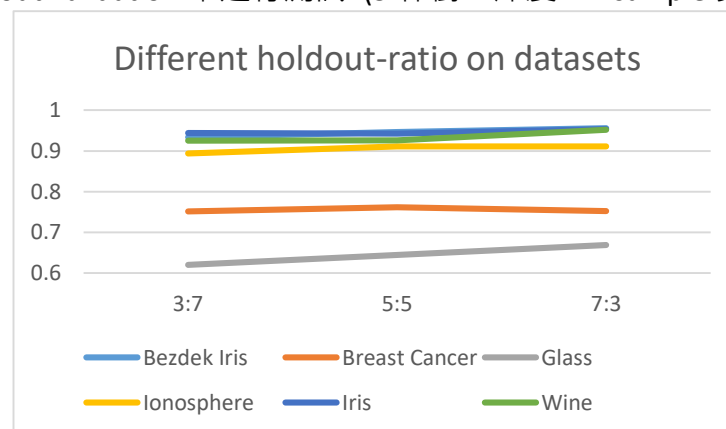
執行時間我認為應該跟 dataset 的 feature 數量有直接的關係。因為 feature 的多寡會影響到深度時有沒有剩餘的 feature 能夠再繼續分下去。由上結果統計部分可以發現，Iris (4 個 feature)的執行時間較 Ionosphere (34 個 feature)來的快非常多，應該是因為使用 Iris 執行時數並沒有長的過深的必要。

### ➤ Categorical Feature 的處理

雖然 spec 只要求我們做 Real attributes 的 dataset，但我還是有處理 categorical 的部分。處理方式是將 categorical 的 feature 進行 one-hot，將該 feature 中的每個類別分離出來變為 0 或 1，如此一來在選取 threshold 時只會選到 0.5，這個節點的意義就能轉為「feature X 的值是否為 Y？」。

### ➤ Relative sizes of the training and validation subsets

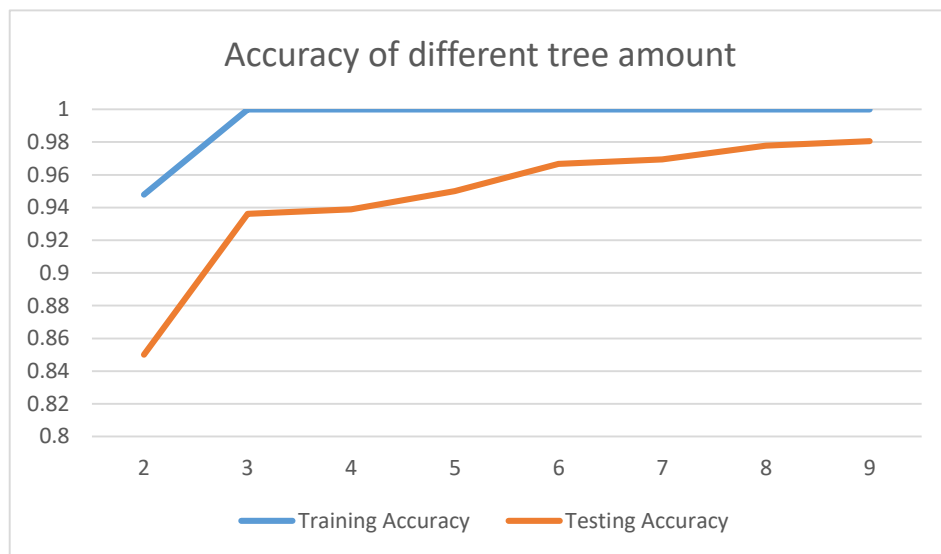
使用 Holdout-validation 來進行測試 (5 棵樹、深度 4、sample 數無限制)：



可以看到，隨著 training data 的增加，大部分的 dataset 都有著些微的表現提升，沒有顯著差距的原因我認為應該是 dataset 本身不是很複雜的關係。

➤ **Number of trees in the forest**

使用 Wine 資料集來進行測試 (Kfold(k=8)、深度 8、sample 數無限制)：



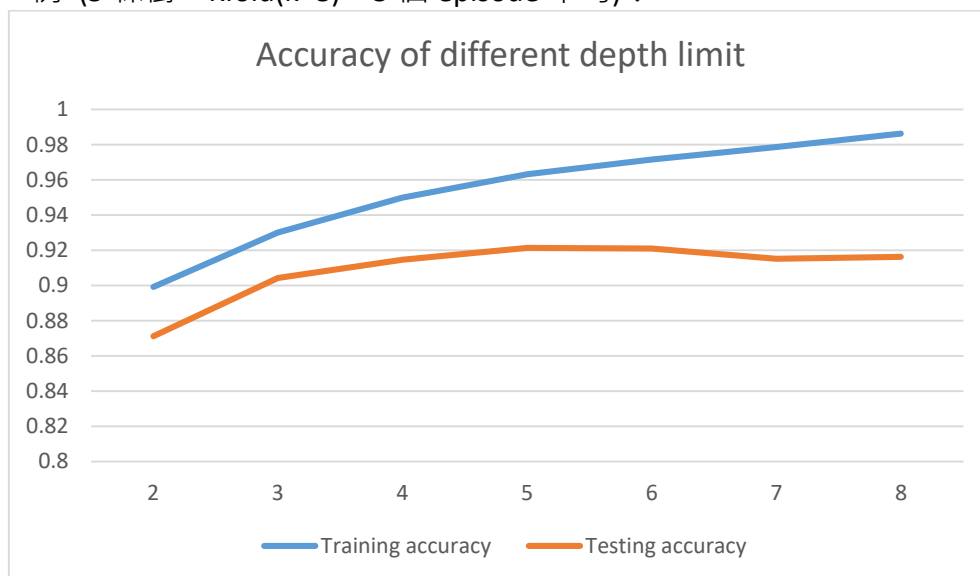
可以看到，提升樹的數量對於提升 testing accuracy 有著明顯的幫助。

➤ **Parameters used during tree induction**

在選擇 feature 的部分，就如實作簡介部分提到的，我預設是從可切割的  $m$  個 feature 中隨機挑選  $\sqrt{m}$  個出來選擇最佳 threshold。至於 extremely random forest 的部分我認為和 depth limit 有些許關係，所以我將其放在後面討論。

➤ **Depth Limit**

深度限制會直接影響較為複雜的資料集的表現。舉 Ionosphere 資料集為例 (5 棵樹、Kfold(k=8)、5 個 episode 平均)：

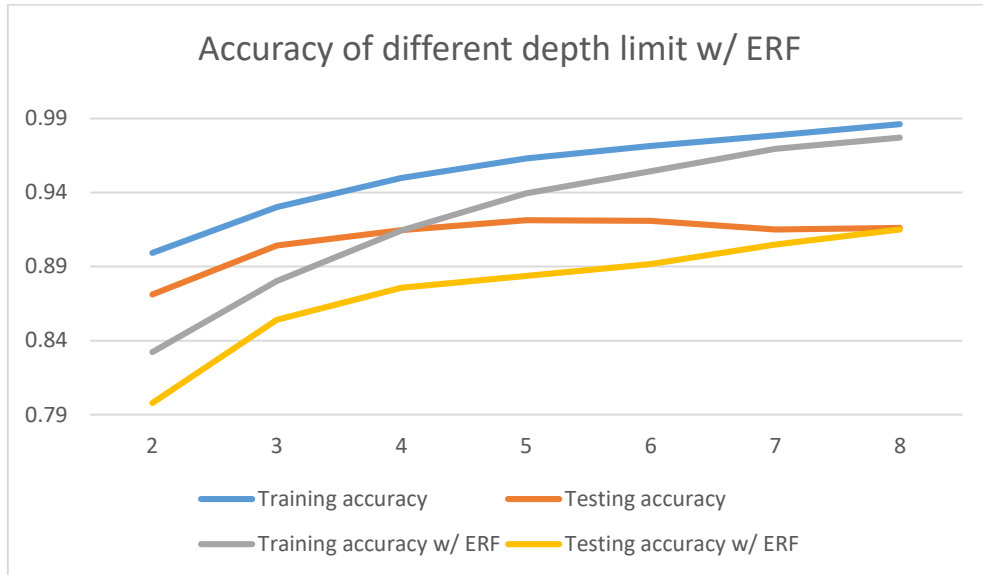


可以看到，增加限制深度可以很好的提升準確度，尤其是在 training accuracy 方面，深度的增加讓模型可以更好的切割資料，使 out-of-bag error 下降。

✓ **Extremely Random Forest**

Extremely Random Forest 因為在選擇 feature 時完全隨機，所以深度會直接影響其能夠選到有意義的 feature 的機率 (單個 node 的機率不變，但選擇的次數變多了)

以下是上面 Ionosphere 資料集的實驗加上 Extremely RF 的結果：



可以看出，depth limit 越大，Extremely Random Forest 和普通 Random Forest 的表現差距就越小。

## Appendix 附錄

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- A. GitHub 連結
- B. pip-requirements
- C. main.py
- D. models.py
- E. utils.py

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### Appendix.A Github 連結

<https://github.com/hm-ysjiang/ArtificialIntelligence-HW2>

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### Appendix.B pip-requirements

```
numpy>=1.20.1  
pandas>=1.2.3  
requests  
tqdm
```

## Appendix.C main.py

```
import numpy as np
import tqdm

from models import DecisionTree, RandomForest
from utils import Dataset, compute_accuracy

def model_provider(use_forest=True):
    n_trees = 5
    tree_bagging = True
    feature_bagging = True
    depth_lim = 8
    min_samples = 0

    if use_forest:
        return RandomForest(n_trees, tree_bagging, feature_bagging, d
epth_lim, min_samples)
    return DecisionTree(feature_bagging, depth_lim, min_samples)

def run(dataset, use_forest=True, use_holdout=False, holdout_ratio=0.
8, kfold=8, episodes=5):
    assert use_holdout or kfold >= 1

    accu_scores_train = []
    accu_scores = []
    with tqdm.tqdm(total=episodes * (1 if use_holdout else kfold)) as
pg:
        for _ in range(episodes):
            if use_holdout:
                train_data, train_label, test_data, test_label = data
set.holdout(
                    holdout_ratio)
                model = model_provider(use_forest)
                model.train(train_data, train_label)
                accu_scores_train.append(
```

```

        compute_accuracy(train_label, model.predict(train
_data)))

    accu_scores.append(
        compute_accuracy(test_label, model.predict(test_d
ata)))

    pg.update(1)
else:
    for train_data, train_label, test_data, test_label in
dataset.kfold(kfold):
        model = model_provider(use_forest)
        model.train(train_data, train_label)
        accu_scores_train.append(
            compute_accuracy(train_label, model.predict(t
rain_data)))

        accu_scores.append(
            compute_accuracy(test_label, model.predict(te
st_data)))

        pg.update(1)
print('Training data accuracy: %.2f%%' %
      (100 * np.average(accu_scores_train)))
print('Testing data accuracy: %.2f%%' %
      (100 * np.average(accu_scores)))

if __name__ == '__main__':
    # Uncomment this line to enable extreme DecisionTree
    # DecisionTree._F_bagging_policy = lambda x: 1

    run(Dataset.Wine, use_forest=True, use_holdout=False, holdout_rat
io=0.8, kfold=8, episodes=10)

```

## Appendix.D models.py

```
import numpy as np

from utils import compute_gini, kfold_indices

class DecisionTree:
    class Node:
        def __init__(self, depth=0):
            self.depth = depth
            self.feature = None
            self.threshold = None
            self.label = None
            self.child1 = None
            self.child2 = None

        def split(self, data, label, feature_bagging, depth_lim, min_
samples):
            # Set node label if only one label presents
            if np.unique(label).shape[0] == 1:
                self.label = label[0]
                return

            # Set node label if limit reached
            if self.depth >= depth_lim or label.shape[0] <= min_sampl
es:
                self.label = np.argmax(np.bincount(label))
                return

            n_samples, n_features = data.shape
            # Consider only sqrt(n_features) features
            n_consider = DecisionTree._F_bagging_policy(n_features)
            # Find splitable features
            splitable = list(filter(lambda feature: np.unique(data[:,
feature]).shape[0] > 1,
                                list(range(n_features))))
            n_splitable = len(splitable)
            if n_splitable > 0:
                min_gini = None
```



```

        # Iterate through the features chosen
        for feature in (np.random.choice(splitable, min(n_consider, n_splitable), replace=False) if feature_bagging else range(n_features)):

            # Sorted unique elements
            values = np.unique(data[:, feature])
            for idx in range(len(values) - 1):
                # Try midpoints between each two unique values

                threshold = (values[idx] + values[idx+1]) / 2

                # Compute total Gini impurity
                g1 = data[:, feature] <= threshold
                g2 = np.invert(g1)
                n1 = g1.sum()
                n2 = n_samples - n1
                gini = n1 * compute_gini(label[g1]) \
                    + n2 * compute_gini(label[g2])

                # Updates
                if self.feature is None or min_gini is None or gini < min_gini:
                    min_gini = gini
                    self.feature = feature
                    self.threshold = threshold
            else: # If no splitable feature
                self.label = np.argmax(np.bincount(label))
                return

        # Split the data into two groups and continue the split of children
        g1 = data[:, self.feature] <= self.threshold
        g2 = np.invert(g1)
        self.child1 = DecisionTree.Node(self.depth+1)
        self.child1.split(data[g1], label[g1],
                        feature_bagging, depth_lim, min_samples
        )

        self.child2 = DecisionTree.Node(self.depth+1)

```

```

        self.child2.split(data[g2], label[g2],
                           feature_bagging, depth_lim, min_samples
)

def __call__(self, data):
    if self.label is not None:
        return self
    else:
        return self.child1 if data[self.feature] <= self.threshold else self.child2

def _F_bagging_policy(x):
    return max(1, round(np.sqrt(x)))

def __init__(self, feature_bagging=True, depth_lim=8, min_samples=0):
    """Initialize a Decision Tree Classifier

    Args:
        feature_bagging (bool, optional): Enable Feature-
        bagging or not. Defaults to True.
        depth_lim (int, optional): The depth limit of the tree. Defaults to 8.
        min_samples (int, optional): The minimum amount of samples in each node. Defaults to 0.
    """
    assert depth_lim > 0
    assert min_samples >= 0

    self.feature_bagging = feature_bagging
    self.depth_lim = depth_lim
    self.min_samples = min_samples
    self.root = DecisionTree.Node()
    self.trained = False

def train(self, data, label):
    """
    Args:

```

```

        data (Iterable): An iterable contains training data, the
dimension should be (samples, features)
        label (Iterable): An iterable contains training data, the
dimension should be (samples, )
    """
    assert not self.trained, 'This tree has already been trained!'
    ,

    self.root.split(data, label, self.feature_bagging,
                    self.depth_lim, self.min_samples)
    self.trained = True

def predict(self, data):
    """
    Args:
        data (Iterable): An iterable contains testing data, the d
imension should be (samples, features)

    Returns:
        numpy.ndarray: An array of dim (samples, ), contains the
predictions of each input
    """
    assert self.trained, 'This tree has not been trained yet!'

    res = [self.root] * data.shape[0]
    for _ in range(self.depth_lim):
        res = [node(data[idx]) for idx, node in enumerate(res)]
    return np.array([node.label for node in res])

class RandomForest:
    def __init__(self, n_tree=5, tree_bagging=True, feature_bagging=T
rue, depth_lim=8, min_samples=0):
        """Initialize a Random Forest Classifier

    Args:
        n_tree (int, optional): The number of trees in this fores
t. Defaults to 5.
        tree_bagging (bool, optional): Enable Tree-
bagging or not. Defaults to True.

```

```

        feature_bagging (bool, optional): Enable Feature-
bagging or not. Defaults to True.
        depth_lim (int, optional): The depth limit of each tree.
Defaults to 8.
        min_samples (int, optional): The minimum amount of sample
s in each tree's node. Defaults to 0.
    """
    assert n_tree >= 1
    assert tree_bagging or feature_bagging

    self.n_tree = n_tree
    self.tree_bagging = tree_bagging
    self.trees = [DecisionTree(feature_bagging, depth_lim, min_sa
mples)
                    for _ in range(n_tree)]
    self.trained = False

    def train(self, data, label):
        """
        Args:
            data (Iterable): An iterable contains training data, the
dimension should be (samples, features)
            label (Iterable): An iterable contains training data, the
dimension should be (samples, )
        """
        assert not self.trained, 'This forest has already been traine
d!'

        if not type(data) is np.ndarray:
            data = np.array(data)
        if not type(label) is np.ndarray:
            label = np.array(label)

        if self.tree_bagging:
            for idx, fold_idx in enumerate(kfold_indices(self.n_tree,
label.shape[0])):
                self.trees[idx].train(data[fold_idx[0]], (label[fold_
idx[0]]))
        else:

```

```

        [tree.train(data, label) for tree in self.trees]
    self.trained = True

def predict(self, data):
    """
    Args:
        data (Iterable): An iterable contains testing data, the dimension should be (samples, features)

    Returns:
        numpy.ndarray: An array of dim (samples, ), contains the predictions of each input
    """
    assert self.trained, 'This forest has not been trained yet!'
    if not type(data) is np.ndarray:
        data = np.array(data)
    return np.array([np.argmax(np.bincount(votes)) for votes in np.array([tree.predict(data) for tree in self.trees]).T])

```

## Appendix.E utils.py

```
from pathlib import Path
import numpy as np
import pandas as pd
import requests as req

def compute_gini(clazz):
    return 1 - sum([(c/clazz.shape[0]) ** 2 for c in np.bincount(clazz)])

def compute_accuracy(a, b):
    try:
        iter(a)
        a = np.array(a)
    except TypeError:
        a = np.array([a])
    try:
        iter(b)
        b = np.array(b)
    except TypeError:
        b = np.array([b])
    assert len(a.shape) == 1 and len(b.shape) == 1 and a.shape == b.shape, \
        'a and b should have the same dimension of (n, )'
    return (a == b).sum() / a.shape[0]

def kfold_indices(k, dimension):
    fold_size = dimension // k
    remainder = dimension % k
    fold_sizes = [fold_size + 1 if _ < remainder else fold_size for _
in range(k)]
    counter = 0
    for fold in fold_sizes:
        test_fold = np.array([True if counter <= x < counter + fold else False
```

```

        for x in range(dimension)])

    counter += fold
    yield np.invert(test_fold), test_fold

class Dataset:
    FTYPE_REAL = 0
    FTYPE_CATEGORICAL = 1
    FTYPE_UNUSED = 2
    FTYPE_CLASS = 3

    BezdekIris: 'Dataset' = None
    BreastCancer: 'Dataset' = None
    Glass: 'Dataset' = None
    Ionosphere: 'Dataset' = None
    Iris: 'Dataset' = None
    Wine: 'Dataset' = None

    def __init__(self, filepath, feature_types, header=None, delim=',',
        dl_url=None):
        """Initialize a Dataset

        Args:
            filepath (str): path to the csv file
            feature_types (Iterable / Callable): An iterable contains
            feature type of each column, or a callable that gives corresponding
            feature type from column index
            header (Iterable, optional): The header of the csv file.
            Defaults to None.
            delim (str, optional): The string used to separate column
            s in the csv file. Defaults to ','.
            dl_url (str, optional): The url of the data file to download
            if the file does not present. Defaults to None.
        """
        # Read csv in
        fp = Path(filepath)
        if not fp.exists():
            if dl_url is not None:
                print('Downloading dataset %s' % filepath)

```

```

        res = req.get(dl_url)
        if not fp.parent.exists():
            fp.parent.mkdir(parents=True)
            with open(filepath, 'wb') as file:      # Write content with LF instead of CRLF
                file.write(res.text.encode())
        else:
            raise FileNotFoundError(
                'Cannot find file %s, and no dl_url provided.' %
filepath)

df = pd.read_csv(filepath, sep=delim, header=header)
# Sanity check the feature types
feature_types = Dataset._sanity_check_ftypes(
    df.shape[1], feature_types)
# Drop unused columns
unused_features = [x[0] for x in filter(
    lambda x: x[1] == Dataset.FTYPE_UNUSED, enumerate(feature
_types))]
df.drop(columns=unused_features, inplace=True)
# Split feature and class label
class_label = [x[0] for x in filter(
    lambda x: x[1] == Dataset.FTYPE_CLASS, enumerate(feature_
types))]
target_classes = df[class_label].to_numpy().reshape(-1)
df.drop(columns=class_label, inplace=True)
# One-hot categorical features
cate_features = [x[0] for x in filter(
    lambda x: x[1] == Dataset.FTYPE_CATEGORICAL, enumerate(fe
ature_types))]
df = pd.get_dummies(df, columns=cate_features).astype('float3
2')

# Encode the class labels
self._class_dict = {}
self._r_class_dict = []
_class = []
for clazz in target_classes:
    if clazz not in self._class_dict:
        self._class_dict[clazz] = len(self._r_class_dict)

```



```

        self._r_class_dict.append(clazz)
        _class.append(self._class_dict[clazz])
# Set final results
self._class = np.array(_class)
self._data = df.to_numpy()

def holdout(self, train_test_ratio=0.7, shuffle=True):
    """Holdout validation

    Args:
        train_test_ratio (float, optional): The ratio of train data to split the dataset. Defaults to 0.7.
        shuffle (bool, optional): Should the data be shuffled. Defaults to True.

    Returns:
        tuple: (train_data, train_labels, test_data, test_labels)
    """
    assert 0 <= train_test_ratio <= 1, 'Train-Test ratio should be in [0, 1]'
    data, class_ = self._shuffle() \
        if shuffle else (self._data.copy(), self._class.copy())
    sep = int(self._class.shape[0] * train_test_ratio)
    return data[:sep, :], class_[:sep], data[sep:, :], class_[sep:]

def kfold(self, k=3, shuffle=True):
    """K-Fold validation

    Args:
        k (int, optional): The 'K' in kfold. Defaults to 3.
        shuffle (bool, optional): Should the data be shuffled. Defaults to True.

    Yields:
        tuple: (train_data, train_labels, test_data, test_labels)
    """
    data, class_ = self._shuffle() \

```

```

        if shuffle else (self._data.copy(), self._class.copy())
    res = []
    for train, test in kfold_indices(k, class_.shape[0]):
        res.append((data[train, :], class_[train],
                    data[test, :], class_[test]))
    return res

def convert_label(self, x):
    try:
        iter(x)
        # Handle x as a list
        return np.array([self._r_class_dict[_] for _ in x])
    except TypeError:
        # Handle x as a single value
        return self._r_class_dict[x]

@property
def data(self):
    return self._data.copy()

@property
def clazz(self):
    return self._class.copy()

def _shuffle(self):
    tmp = np.concatenate((self._class.reshape(-
1, 1), self._data), axis=1)
    np.random.shuffle(tmp)
    return tmp[:, 1:], tmp[:, 0].astype(np.int32)

def _sanity_check_ftypes(n_features, ftypes):
    if hasattr(ftypes, '__call__'):
        ftypes = [ftypes(x) for x in range(n_features)]
    else:
        assert len(ftypes) == n_features, \
            'The feature_types length does not match with the input data. Expecting %d, got %d' \
            % (n_features, len(ftypes))

```

```

        ftypes = list(ftypes)
        for idx, x in enumerate(ftypes):
            if not 0 <= x <= 3:
                ftypes[idx] = Dataset.FTYPE_REAL
                print('Detect an undefined feature type: %d, this value will be changed to FTYPE_REAL. Check Dataset.FTYPE_* for proper usage.' % x)
        return tuple(ftypes)

    def __len__(self):
        return self._class.shape[0]

    def __getitem__(self, idx):
        return self._data[idx].reshape(1, -1), self._class[idx]

Dataset.BezdekIris = Dataset('./data/iris/bezdekiris.data',
                             lambda x: Dataset.FTYPE_REAL if x != 4 else Dataset.FTYPE_CLASS,
                             dl_url='https://archive.ics.uci.edu/ml/machine-learning-databases/iris/bezdekIris.data')

Dataset.BreastCancer = Dataset('./data/breast-cancer/breast-cancer.data',
                                [Dataset.FTYPE_CATEGORICAL] * 6 +
                                [Dataset.FTYPE_REAL] +
                                [Dataset.FTYPE_CATEGORICAL] * 2 +
                                [Dataset.FTYPE_CLASS],
                                dl_url='https://archive.ics.uci.edu/ml/machine-learning-databases/breast-cancer/breast-cancer.data')

Dataset.Glass = Dataset('./data/glass/glass.data',
                        [Dataset.FTYPE_UNUSED] +
                        [Dataset.FTYPE_REAL] * 9 +
                        [Dataset.FTYPE_CLASS],
                        dl_url='https://archive.ics.uci.edu/ml/machine-learning-databases/glass/glass.data')

```

```
Dataset.Ionosphere = Dataset('./data/ionosphere/ionosphere.data',
                              lambda x: Dataset.FTYPE_CLASS if x == 34
                              else Dataset.FTYPE_REAL,
                              dl_url='https://archive.ics.uci.edu/ml/m
achine-learning-databases/ionosphere/ionosphere.data')

Dataset.Iris = Dataset('./data/iris/iris.data',
                       [Dataset.FTYPE_REAL,
                        Dataset.FTYPE_REAL,
                        Dataset.FTYPE_REAL,
                        Dataset.FTYPE_REAL,
                        Dataset.FTYPE_CLASS],
                       dl_url='https://archive.ics.uci.edu/ml/machine
-learning-databases/iris/iris.data')

Dataset.Wine = Dataset('./data/wine/wine.data',
                       lambda x: Dataset.FTYPE_CLASS if x == 0 else D
ataset.FTYPE_REAL,
                       dl_url='https://archive.ics.uci.edu/ml/machine
-learning-databases/wine/wine.data')
```