```
In [136... import warnings
         warnings.filterwarnings('ignore')
         # Complex pytorch
         import torch
         import torch.nn as nn
         import torch.nn.functional as F
         from torch.utils.data import DataLoader
         from torchvision import datasets, transforms
         from complexPyTorch.complexLayers import *
         from complexPyTorch.complexFunctions import *
         from torch geometric.nn import GCNConv
         from torch_geometric.data import Data
         # Plot
         import matplotlib.pyplot as plt
         import seaborn as sns
         import time
         # Load Data
         import numpy as np
         import json
         import os
         import math
         import librosa
         import pathlib
         from scipy.spatial.distance import cdist
         from torch.utils.data import Dataset
         from sklearn.model_selection import train_test_split
         import random
         import pandas as pd
         # MFCCS
         from scipy.io import wavfile
         import scipy.fftpack as fft
         from scipy.signal import get_window
         import librosa
         import librosa.display
         import IPython.display as ipd
         import scipy as spp
         # CV
         from sklearn.model_selection import cross_val_score, KFold
In [139... def custom_cross_val(model, X, y, k=5):
             np.random.seed(42)
             indices = np.arange(len(X))
```

```
In [139...

def custom_cross_val(model, X, y, k=5):
    np.random.seed(42)
    indices = np.arange(len(X))
    np.random.shuffle(indices)
    splits = np.array_split(indices, k)
    accuracies = []
    for i in range(k):
        test_indices = splits[i]
        train_indices = np.concatenate([splits[j] for j in range(k) if j
        X_train, y_train = X.iloc[train_indices], y.iloc[train_indices]
        X_test, y_test = X.iloc[test_indices], y.iloc[test_indices]
        model.fit(X_train.to_numpy(), y_train.to_numpy())
        y_pred = model.predict(X_test.to_numpy())
        accuracy = np.mean(y_pred == y_test.to_numpy())
```

```
accuracies.append(accuracy)
return accuracies
```

#### **Create Data**

```
In [ ]: DATASET_PATH = "Data/train"
        SAMPLE RATE = 22050
        TRACK DURATION = 30 # measured in seconds
        SAMPLES PER TRACK = SAMPLE RATE * TRACK DURATION
        BATCH SIZE = 32
        NUM_EPOCHS = 50
        genre_list = os.listdir(DATASET_PATH)
        if '.DS_Store' in genre_list: genre_list.remove('.DS_Store')
        genre_mappings = dict(zip(genre_list, range(len(genre_list))))
        print(genre mappings)
In [ ]: class TimeDomainFeatures:
            @staticmethod
            def amplitude_envelope(signal, frame_size, hop_length):
                 res = []
                for i in range(0, len(signal), hop_length):
                     cur_portion = signal[i:i + frame_size]
                     ae_val = max(cur_portion)
                     res.append(ae val)
                 return np.array([np.nanmean(res), np.nanvar(res)])
            @staticmethod
            def RMS_energy(signal, frame_size, hop_length):
                 res = []
                for i in range(0, len(signal), hop_length):
                     cur portion = signal[i:i + frame size]
                     rmse_val = np.sqrt(1 / len(cur_portion) * sum(i**2 for i in c
                     res.append(rmse_val)
                 return np.array([np.nanmean(res), np.nanvar(res)])
            @staticmethod
            def crest_factor(signal, frame_size, hop_length):
                 res = []
                for i in range(0, len(signal), hop_length):
                     cur_portion = signal[i:i + frame_size]
                     rmse_val = np.sqrt(1 / len(cur_portion) * sum(i ** 2 for i in
                     crest_val = max(np.abs(cur_portion)) / rmse_val
                     res.append(crest_val)
                return np.array([np.nanmean(res), np.nanvar(res)])
            @staticmethod
            def ZCR(signal, frame_size, hop_length):
                def num_sign_changes(signal):
                     for i in range(0, len(signal) - 1):
                         if (signal[i] * signal[i + 1] < 0): res += 1</pre>
                     return res
                 res = []
                for i in range(0, len(signal), hop_length):
                     cur_portion = signal[i:i + frame_size]
                     zcr_val = num_sign_changes(cur_portion)
```

```
res.append(zcr_val)
return np.array([np.nanmean(res), np.nanvar(res)])
```

### 0.1 Real Frequncy Domain

```
In [ ]: class FreqDomainFeatures:
            @staticmethod
            def normalize audio(audio):
                audio = audio / np.max(np.abs(audio))
                return audio
            @staticmethod
            def compute_spectrogram(signal, frame_size, hop_length):
                signal = FreqDomainFeatures.normalize_audio(signal)
                spec = librosa.stft(signal, n_fft=frame_size, hop_length=hop_leng
                return np.abs(spec).T
            @staticmethod
            def band_energy_ratio(spec, split_freq = 2048):
                def find_split_freq_bin(spec, split_freq):
                     range_of_freq = SAMPLE_RATE / 2
                    change_per_bin = range_of_freq / spec.shape[0]
                    split_freq_bin = split_freq / change_per_bin
                    return int(np.floor(split_freq_bin))
                split_freq_bin = find_split_freq_bin(spec.T, split_freq)
                res = []
                for sub_arr in spec:
                    low_freq_density = sum(i ** 2 for i in sub_arr[:split_freq_bi
                    high_freq_density = sum(i ** 2 for i in sub_arr[split_freq_bi
                    ber_val = low_freq_density / high_freq_density
                    res.append(ber_val)
                return np.array([np.nanmean(res), np.nanvar(res)])
            @staticmethod
            def spectral_centroid(spec):
                def sc(arr):
                    res = 0
                    for i in range(0, len(arr)):
                         res += i*arr[i]
                    return res/sum(arr)
                res = []
                for sub_arr in spec:
                    sc_val = sc(sub_arr)
                     res.append(sc_val)
                return np.array([np.nanmean(res), np.nanvar(res)])
            @staticmethod
            def spectral_bandwidth(spec):
                def sc(arr):
                    res = 0
                    for i in range(0, len(arr)):
                         res += i*arr[i]
                    return res/sum(arr)
                def sb(arr):
                    res = 0
                    sc_val = sc(arr)
                    for i in range(0, len(arr)):
```

```
res += (abs(i - sc val))*arr[i]
        return res/sum(arr)
    res = []
    for sub_arr in spec:
        sb_val = sb(sub_arr)
        res.append(sb val)
    return np.array([np.nanmean(res), np.nanvar(res)])
@staticmethod
def spectral_flatness(spec):
    res = []
    for sub_arr in spec:
        geom_mean = np.exp(np.log(sub_arr).mean())
        ar_mean = np.mean(sub_arr)
        sl_val = geom_mean/ar_mean
        res.append(sl_val)
    return np.array([np.nanmean(res), np.nanvar(res)])
```

```
In [ ]: class GenreTimeFreqDomain(Dataset):
            def __init__(self, train_path, frame_size=1024, hop_length=512, num_s
                cur path = pathlib.Path(train path)
                self.files = []
                for i in list(cur path.rglob("*.wav")):
                     for j in range(num_segments):
                         self.files.append([j, i])
                self.frame_size = frame_size
                self.hop length = hop length
                self.training = training
                self.samples_per_segment = int(SAMPLES_PER_TRACK / num_segments)
                self.num_segments = num_segments
            def apply_augmentations(self, signal):
                # Apply augmentations to the audio signal
                if random.random() < 0.5:</pre>
                     signal = librosa.effects.pitch_shift(signal, sr=SAMPLE_RATE,
                if random.random() < 0.5:</pre>
                     signal = librosa.effects.time_stretch(signal, rate=random.uni
                return signal
            def __len__(self):
                 return len(self.files)
            def adj_shape(self, features):
                if features.shape[0] < 130:</pre>
                     features = np.pad(features, (0, 130 - features.shape[0]), mod
                else:
                     features = features[:130]
                return features
            def get_time_domain(self, cur_signal):
                ae = TimeDomainFeatures.amplitude_envelope(cur_signal, self.frame
                 rmse = TimeDomainFeatures.RMS_energy(cur_signal, self.frame_size,
                cf = TimeDomainFeatures.crest_factor(cur_signal, self.frame_size,
                 zcr = TimeDomainFeatures.ZCR(cur_signal, self.frame_size, self.ho
                return np.concatenate([ae, rmse, cf, zcr])
            def get_freq_domain(self, cur_signal):
                spec = FreqDomainFeatures.compute_spectrogram(cur_signal, self.fr
                ber = FreqDomainFeatures.band_energy_ratio(spec)
```

```
sc = FreqDomainFeatures.spectral centroid(spec)
    sb = FreqDomainFeatures.spectral_bandwidth(spec)
    sf = FreqDomainFeatures.spectral_flatness(spec)
    return np.concatenate([ber, sc, sb, sf])
def __getitem__(self, idx):
    cur file = self.files[idx]
    d = cur_file[0]
    file_path = cur_file[1]
    target = genre_mappings[str(file_path).split("/")[2]]
    signal, sample_rate = librosa.load(file_path, sr=SAMPLE_RATE)
    start = self.samples_per_segment * d
    finish = start + self.samples_per_segment
    cur_signal = signal[start:finish]
    # if self.training: cur_signal = self.apply_augmentations(cur_sig
    td_features = self.get_time_domain(cur_signal)
    fd_features = self.get_freq_domain(cur_signal)
    return torch.tensor(np.array([td features, fd features]).flatten(
```

```
In [ ]: train_dataset = GenreTimeFreqDomain("Data/train/", training = True)
        test_dataset = GenreTimeFreqDomain("Data/test/", training = False)
        column names = []
        for j in ["ae", "rmse", "cf", "zcr", "ber", "sc", "sb", "sf"]:
            for i in ["mean", "var"]:
                column_names.extend([f"{j}_{i}"])
        # Add label and set columns
        column names.extend(['label'])
        print(column names)
        def create_dataframe(dataset):
            features_list = []
            labels list = []
            for i in range(len(dataset)):
                features, label = dataset[i]
                features_list.append(features)
                labels_list.append(label)
                if i%10 == 0: print(i)
            df = pd.DataFrame(features_list) # Flatten the features
            df['label'] = labels_list
            return df
        train_df = create_dataframe(train_dataset)
        train df.columns = column names
        print("-"*75)
        test_df = create_dataframe(test_dataset)
        test_df.columns = column_names
        tr_df = train_df.applymap(lambda x: x.numpy() if hasattr(x, 'numpy') else
        te_df = test_df.applymap(lambda x: x.numpy() if hasattr(x, 'numpy') else
        tr_df.to_csv("train_tff.csv", index = False)
        te_df.to_csv("test_tff.csv", index = False)
```

### 0.2 Complex Frequncy Domain

```
In [ ]: class FreqDomainFeatures:
            @staticmethod
            def normalize_audio(audio):
                audio = audio / np.max(np.abs(audio))
                return audio
            @staticmethod
            def compute_spectrogram(signal, frame_size, hop_length):
                 signal = FreqDomainFeatures.normalize audio(signal)
                spec = librosa.stft(signal, n_fft=frame_size, hop_length=hop_leng
                 return (spec).T
            @staticmethod
            def band_energy_ratio(spec, split_freq = 2048):
                def find_split_freq_bin(spec, split_freq):
                     range_of_freq = SAMPLE_RATE / 2
                     change_per_bin = range_of_freq / spec.shape[0]
                     split freq bin = split freq / change per bin
                     return int(np.floor(split_freq_bin))
                split_freq_bin = find_split_freq_bin(spec.T, split_freq)
                res = []
                for sub arr in spec:
                     low_freq_density = sum(i ** 2 for i in sub_arr[:split_freq_bi
                     high freq density = sum(i ** 2 for i in sub arr[split freq bi
                     ber_val = low_freq_density / high_freq_density
                     res.append(ber val)
                return np.array([np.nanmean(res), np.nanvar(res)])
            @staticmethod
            def spectral centroid(spec):
                def sc(arr):
                     res = 0
                     for i in range(0, len(arr)):
                         res += i*arr[i]
                     return res/sum(arr)
                 res = []
                for sub_arr in spec:
                     sc_val = sc(sub_arr)
                     res_append(sc_val)
                 return np.array([np.nanmean(res), np.nanvar(res)])
            @staticmethod
            def spectral_bandwidth(spec):
                def sc(arr):
                     res = 0
                     for i in range(0, len(arr)):
                         res += i*arr[i]
                     return res/sum(arr)
                def sb(arr):
                     res = 0
                     sc_val = sc(arr)
                     for i in range(0, len(arr)):
                         res += (np.abs(i - sc_val))*arr[i]
                    return res/sum(arr)
                 res = []
                for sub_arr in spec:
                     sb_val = sb(sub_arr)
                     res.append(sb_val)
```

```
return np.array([np.nanmean(res), np.nanvar(res)])

@staticmethod
def spectral_flatness(spec):
    res = []
    for sub_arr in spec:
        geom_mean = np.exp(np.log(sub_arr).mean())
        ar_mean = np.mean(sub_arr)
        sl_val = geom_mean/ar_mean
        res.append(sl_val)
    return np.array([np.nanmean(res), np.nanvar(res)])
```

```
In [ ]: class GenreTimeFreqDomain(Dataset):
            def __init__(self, train_path, frame_size=1024, hop_length=512, num_s
                cur path = pathlib.Path(train path)
                self.files = []
                for i in list(cur_path.rglob("*.wav")):
                    for j in range(num_segments):
                         self.files.append([j, i])
                self.frame_size = frame_size
                self.hop length = hop length
                self.training = training
                self.samples_per_segment = int(SAMPLES_PER_TRACK / num_segments)
                self.num_segments = num_segments
            def apply_augmentations(self, signal):
                # Apply augmentations to the audio signal
                if random.random() < 0.5:</pre>
                    signal = librosa.effects.pitch shift(signal, sr=SAMPLE RATE,
                if random.random() < 0.5:</pre>
                    signal = librosa.effects.time_stretch(signal, rate=random.uni
                return signal
            def len (self):
                return len(self.files)
            def adj_shape(self, features):
                if features.shape[0] < 130:</pre>
                    features = np.pad(features, (0, 130 - features.shape[0]), mod
                    features = features[:130]
                return features
            def get_time_domain(self, cur_signal):
                ae = TimeDomainFeatures.amplitude envelope(cur signal, self.frame
                rmse = TimeDomainFeatures.RMS_energy(cur_signal, self.frame_size,
                cf = TimeDomainFeatures.crest_factor(cur_signal, self.frame_size,
                zcr = TimeDomainFeatures.ZCR(cur_signal, self.frame_size, self.hd
                return np.concatenate([ae, rmse, cf, zcr])
            def get_freq_domain(self, cur_signal):
                spec = FreqDomainFeatures.compute_spectrogram(cur_signal, self.fr
                ber = FreqDomainFeatures.band_energy_ratio(spec)
                sc = FreqDomainFeatures.spectral_centroid(spec)
                sb = FreqDomainFeatures.spectral_bandwidth(spec)
                sf = FreqDomainFeatures.spectral_flatness(spec)
                return np.concatenate([ber, sc, sb, sf])
```

```
def __getitem__(self, idx):
    cur_file = self.files[idx]
    d = cur_file[0]
    file_path = cur_file[1]
    target = genre_mappings[str(file_path).split("/")[2]]
    signal, sample_rate = librosa.load(file_path, sr=SAMPLE_RATE)
    start = self.samples_per_segment * d
    finish = start + self.samples_per_segment
    cur_signal = signal[start:finish]
    # if self.training: cur_signal = self.apply_augmentations(cur_signal)
    td_features = self.get_time_domain(cur_signal)
    fd_features = self.get_freq_domain(cur_signal)
    return torch.tensor(np.array([td_features, fd_features]).flatten(
```

```
In [ ]: train_dataset = GenreTimeFreqDomain("Data/train/", training = True)
        test_dataset = GenreTimeFreqDomain("Data/test/", training = False)
        column_names = []
        for j in ["ae", "rmse", "cf", "zcr", "ber", "sc", "sb", "sf"]:
            for i in ["mean", "var"]:
                column_names.extend([f"{j}_{i}"])
        # Add label and set columns
        column names.extend(['label'])
        print(column names)
        def create_dataframe(dataset):
            features list = []
            labels_list = []
            for i in range(len(dataset)):
                features, label = dataset[i]
                features_list.append(features)
                labels_list.append(label)
                if i%10 == 0: print(i)
            df = pd.DataFrame(features_list) # Flatten the features
            df['label'] = labels_list
            return df
        train_df = create_dataframe(train_dataset)
        train_df.columns = column_names
        print("-"*75)
        test_df = create_dataframe(test_dataset)
        test_df.columns = column_names
        tr_df = train_df.applymap(lambda x: x.numpy() if hasattr(x, 'numpy') else
        te_df = test_df.applymap(lambda x: x.numpy() if hasattr(x, 'numpy') else
        tr_df.to_csv("train_tff_comp.csv", index = False)
        te_df.to_csv("test_tff_comp.csv", index = False)
```

# 1. Simple Decsion Tree with Real Valued Frequency domain Features

```
In [132... tr_df = pd.read_csv("train_tff.csv")
    te_df = pd.read_csv("test_tff.csv")
```

```
In [133... # Separate features and labels
         X_train = tr_df.drop('label', axis=1)
         y_train = tr_df['label']
         X_test = te_df.drop('label', axis=1)
         y_test = te_df['label']
 In [9]: from sklearn.model_selection import train_test_split
         from sklearn.linear_model import LogisticRegression, SGDClassifier, Ridge
         from sklearn.neighbors import KNeighborsClassifier, RadiusNeighborsClassi
         from sklearn.naive_bayes import GaussianNB, MultinomialNB, ComplementNB,
         from sklearn.tree import DecisionTreeClassifier
         from sklearn.ensemble import RandomForestClassifier, AdaBoostClassifier,
         from sklearn.svm import SVC, NuSVC, LinearSVC
         from sklearn.discriminant_analysis import LinearDiscriminantAnalysis, Qua
         from sklearn.ensemble import VotingClassifier
         from sklearn.metrics import accuracy_score
         # List of models to try
         models = [
              ('Logistic Regression', LogisticRegression()),
             ('SGD Classifier', SGDClassifier()),
             ('Ridge Classifier', RidgeClassifier()),
             ('Passive Aggressive Classifier', PassiveAggressiveClassifier()),
             ('K-Nearest Neighbors', KNeighborsClassifier()),
             ('Gaussian Naive Bayes', GaussianNB()),
             ('Multinomial Naive Bayes', MultinomialNB()),
             ('Complement Naive Bayes', ComplementNB()),
             ('Bernoulli Naive Bayes', BernoulliNB()),
             ('Decision Tree', DecisionTreeClassifier()),
             ('Random Forest', RandomForestClassifier()),
             ('AdaBoost', AdaBoostClassifier()),
             ('Gradient Boosting', GradientBoostingClassifier()),
             ('Support Vector Machine', SVC()),
             ('Nu-Support Vector Machine', NuSVC()),
             ('Linear Support Vector Machine', LinearSVC()),
             ('Linear Discriminant Analysis', LinearDiscriminantAnalysis()),
             ('Quadratic Discriminant Analysis', QuadraticDiscriminantAnalysis()),
             ('Voting Classifier', VotingClassifier(estimators=[
                  ('lr', LogisticRegression()),
                  ('rf', RandomForestClassifier()),
                 ('svc', SVC())
             1))
             # Add more models as needed
         # Loop through models
         for model_name, model in models:
             # Train the model
             model.fit(X_train, y_train)
             # Predict on test data
             y_pred = model.predict(X_test)
             # Calculate accuracy
             accuracy = accuracy_score(y_test, y_pred)
             # Print results
             print(f'Model: {model_name}')
             print(f'Accuracy: {accuracy}\n')
```

Model: Logistic Regression

Accuracy: 0.11625

Model: SGD Classifier Accuracy: 0.05875

Model: Ridge Classifier Accuracy: 0.364375

Model: Passive Aggressive Classifier

Accuracy: 0.21625

Model: K-Nearest Neighbors

Accuracy: 0.2625

Model: Gaussian Naive Bayes

Accuracy: 0.113125

Model: Multinomial Naive Bayes

Accuracy: 0.145

Model: Complement Naive Bayes

Accuracy: 0.110625

Model: Bernoulli Naive Bayes

Accuracy: 0.1

Model: Decision Tree Accuracy: 0.4125

Model: Random Forest Accuracy: 0.5425

Model: AdaBoost Accuracy: 0.420625

Model: Gradient Boosting

Accuracy: 0.539375

Model: Support Vector Machine

Accuracy: 0.099375

Model: Nu-Support Vector Machine

Accuracy: 0.110625

Model: Linear Support Vector Machine

Accuracy: 0.11

Model: Linear Discriminant Analysis

Accuracy: 0.421875

Model: Quadratic Discriminant Analysis

Accuracy: 0.34375

Model: Voting Classifier

Accuracy: 0.18375

In [134... class Node:
 def \_\_init\_\_(self, feature=None, threshold=None, left=None, right=None)

```
self.feature = feature
        self.threshold = threshold
        self.left = left
        self.right = right
        self.value = value
    def is leaf(self):
        return self.value is not None
class DecisionTree:
    def __init__(self, max_depth=100, min_samples_split=2):
        self.max depth = max depth
        self.min_samples_split = min_samples_split
        self.root = None
    def _is_finished(self, depth):
        if (depth >= self.max_depth
            or self.n class labels == 1
            or self.n samples < self.min samples split):</pre>
            return True
        return False
    def _entropy(self, y):
        proportions = np.bincount(y) / len(y)
        entropy = -np.sum([p * np.log2(p) for p in proportions if p > 0])
        return entropy
    def _create_split(self, X, thresh):
        left_idx = np.argwhere(X <= thresh).flatten()</pre>
        right_idx = np.argwhere(X > thresh).flatten()
        return left_idx, right_idx
    def _information_gain(self, X, y, thresh):
        parent_loss = self._entropy(y)
        left_idx, right_idx = self._create_split(X, thresh)
        n, n_left, n_right = len(y), len(left_idx), len(right_idx)
        if n_left == 0 or n_right == 0:
            return 0
        child_loss = (n_left / n) * self._entropy(y[left_idx]) + (n_right
        return parent_loss - child_loss
    def _best_split(self, X, y, features):
        split = {'score':- 1, 'feat': None, 'thresh': None}
        for feat in features:
            X_{feat} = X[:, feat]
            thresholds = np.unique(X_feat)
            for thresh in thresholds:
                score = self._information_gain(X_feat, y, thresh)
                if score > split['score']:
                    split['score'] = score
                    split['feat'] = feat
                    split['thresh'] = thresh
        return split['feat'], split['thresh']
    def _build_tree(self, X, y, depth=0):
```

```
self.n samples, self.n features = X.shape
                 self.n_class_labels = len(np.unique(y))
                 # stopping criteria
                 if self._is_finished(depth):
                     most common Label = np.argmax(np.bincount(y))
                     return Node(value=most_common_Label)
                 # get best split
                 rnd_feats = np.random.choice(self.n_features, self.n_features, re
                 best_feat, best_thresh = self._best_split(X, y, rnd_feats)
                 # grow children recursively
                 left_idx, right_idx = self._create_split(X[:, best_feat], best_th
                 left_child = self._build_tree(X[left_idx, :], y[left_idx], depth
                 right_child = self._build_tree(X[right_idx, :], y[right_idx], dep
                 return Node(best_feat, best_thresh, left_child, right_child)
             def traverse tree(self, x, node):
                 if node.is leaf():
                     return node.value
                 if x[node.feature] <= node.threshold:</pre>
                     return self._traverse_tree(x, node.left)
                 return self._traverse_tree(x, node.right)
             def fit(self, X, y):
                 self.root = self._build_tree(X, y)
             def predict(self, X):
                 predictions = [self._traverse_tree(x, self.root) for x in X]
                 return np.array(predictions)
In [35]: np.random.seed(42)
         model = DecisionTree(max_depth=10)
         model.fit(X_train.to_numpy(), y_train.to_numpy())
         y_pred = model.predict(X_test.to_numpy())
         accuracy = accuracy_score(y_test.to_numpy(), y_pred)
         print(f'Accuracy: {accuracy}\n')
        Accuracy: 0.451875
In [138... # CV:
         merged_df = pd.concat([tr_df, te_df], axis=0)
         X = merged_df.drop('label', axis=1)
         y = merged_df['label']
         np.random.seed(42)
         model = DecisionTree(max_depth=10)
         cv_results = custom_cross_val(model, X, y, k=5)
         for i, acc in enumerate(cv_results):
             print(f'Fold {i+1} Accuracy: {acc}')
         print(f'Mean Accuracy: {np.mean(cv_results)}')
        Fold 1 Accuracy: 0.551051051051051
        Fold 2 Accuracy: 0.5725725725725
        Fold 3 Accuracy: 0.5720720720720721
        Fold 4 Accuracy: 0.545045045045045
        Fold 5 Accuracy: 0.5565565565565
       Mean Accuracy: 0.5594594594595
```

## 2. Simple Decision Tree with Complex Valued Frequency Domain Features

```
In [172... | tr_df = pd.read_csv("train_tff_comp.csv")
         te_df = pd.read_csv("test_tff_comp.csv")
         import pandas as pd
         def df_csv_complex(df):
             result_df = df.copy() # Make a copy to avoid modifying the original
             result_df.iloc[:, :-1] = df.iloc[:, :-1].apply(lambda col: col.apply(
                 lambda val: torch.tensor((complex(val.strip('()'))), dtype=torch.
             return result df
         tr_df = df_csv_complex(tr_df)
         te_df = df_csv_complex(te_df)
         tr_df = train_df.applymap(lambda x: x.numpy() if hasattr(x, 'numpy') else
         te_df = test_df.applymap(lambda x: x.numpy() if hasattr(x, 'numpy') else
In [141... | # Separate features and labels
         X_train = tr_df.drop('label', axis=1)
         y train = tr df['label']
         X_test = te_df.drop('label', axis=1)
         y_test = te_df['label']
```

#### 2.1 Compare only real

```
In [145... class Node:
             def __init__(self, feature=None, threshold=None, left=None, right=None
                  self.feature = feature
                  self.threshold = threshold
                  self.left = left
                  self.right = right
                  self.value = value
             def is leaf(self):
                  return self.value is not None
         class DecisionTree:
             def __init__(self, max_depth=100, min_samples_split=2):
                  self.max_depth = max_depth
                  self.min_samples_split = min_samples_split
                  self.root = None
             def _is_finished(self, depth):
                  if (depth >= self.max_depth
                      or self.n_class_labels == 1
                      or self.n_samples < self.min_samples_split):</pre>
                      return True
                  return False
             def _entropy(self, y):
                  proportions = np.bincount(y) / len(y)
                  entropy = -np.sum([p * np.log2(p) for p in proportions if p > 0])
```

```
return entropy
def _create_split(self, X, thresh):
    left_idx = np.argwhere(X <= thresh).flatten()</pre>
    right_idx = np.argwhere(X > thresh).flatten()
    return left idx, right idx
def _information_gain(self, X, y, thresh):
    parent_loss = self._entropy(y)
    left_idx, right_idx = self._create_split(X, thresh)
    n, n_left, n_right = len(y), len(left_idx), len(right_idx)
    if n_left == 0 or n_right == 0:
        return 0
    child_loss = (n_left / n) * self._entropy(y[left_idx]) + (n_right
    return parent_loss - child_loss
def _best_split(self, X, y, features):
    split = {'score':- 1, 'feat': None, 'thresh': None}
    for feat in features:
       X_{feat} = X[:, feat]
        thresholds = np.unique(X feat)
        for thresh in thresholds:
            score = self._information_gain(X_feat, y, thresh)
            if score > split['score']:
                split['score'] = score
                split['feat'] = feat
                split['thresh'] = thresh
    return split['feat'], split['thresh']
def _build_tree(self, X, y, depth=0):
    self.n_samples, self.n_features = X.shape
    self.n_class_labels = len(np.unique(y))
    # stopping criteria
    if self._is_finished(depth):
        most_common_Label = np.argmax(np.bincount(y))
        return Node(value=most_common_Label)
    # get best split
    rnd_feats = np.random.choice(self.n_features, self.n_features, re
    best_feat, best_thresh = self._best_split(X, y, rnd_feats)
    # grow children recursively
    left_idx, right_idx = self._create_split(X[:, best_feat], best_th
    left_child = self._build_tree(X[left_idx, :], y[left_idx], depth
    right_child = self._build_tree(X[right_idx, :], y[right_idx], dep
    return Node(best_feat, best_thresh, left_child, right_child)
def _traverse_tree(self, x, node):
    if node.is_leaf():
        return node.value
    if x[node.feature] <= node.threshold:</pre>
        return self._traverse_tree(x, node.left)
    return self._traverse_tree(x, node.right)
```

```
def fit(self, X, y):
                 self.root = self._build_tree(X, y)
             def predict(self, X):
                 predictions = [self._traverse_tree(x, self.root) for x in X]
                 return np.array(predictions)
In [68]: np.random.seed(42)
         model = DecisionTree(max_depth=10)
         model.fit(X_train.to_numpy(), y_train.to_numpy())
         y_pred = model.predict(X_test.to_numpy())
         accuracy = accuracy_score(y_test.to_numpy(), y_pred)
         print(f'Accuracy: {accuracy}\n')
        Accuracy: 0.4575
In [175... # CV:
         merged_df = pd.concat([tr_df, te_df], axis=0)
         X = merged_df.drop('label', axis=1)
         y = merged df['label']
         np.random.seed(42)
         model = DecisionTree(max_depth=10)
         cv_results = custom_cross_val(model, X, y, k=5)
         for i, acc in enumerate(cv_results):
             print(f'Fold {i+1} Accuracy: {acc}')
         print(f'Mean Accuracy: {np.mean(cv_results)}')
        Fold 1 Accuracy: 0.5680680680680681
        Fold 2 Accuracy: 0.5640640640640641
        Fold 3 Accuracy: 0.5570570570570571
        Fold 4 Accuracy: 0.561061061061061
        Fold 5 Accuracy: 0.561061061061061
```

### 2.2 Compare only magnitude

Mean Accuracy: 0.5622622622624

```
In [176... class Node:
             def __init__(self, feature=None, threshold=None, left=None, right=Non
                  self.feature = feature
                  self.threshold = threshold
                  self.left = left
                  self.right = right
                  self.value = value
             def is_leaf(self):
                  return self.value is not None
         class DecisionTree:
             def __init__(self, max_depth=100, min_samples_split=2):
                  self.max_depth = max_depth
                  self.min_samples_split = min_samples_split
                  self.root = None
             def _is_finished(self, depth):
                  if (depth >= self.max_depth
                      or self.n_class_labels == 1
                      or self.n_samples < self.min_samples_split):</pre>
```

```
return True
    return False
def _entropy(self, y):
    proportions = np.bincount(y) / len(y)
    entropy = -np.sum([p * np.loq2(p) for p in proportions if p > 0])
    return entropy
def _create_split(self, X, thresh):
    left_idx = np.argwhere(np.abs(X) <= np.abs(thresh)).flatten()</pre>
    right_idx = np.argwhere(np.abs(X) > np.abs(thresh)).flatten()
    return left idx, right idx
def _information_gain(self, X, y, thresh):
    parent_loss = self._entropy(y)
    left_idx, right_idx = self._create_split(X, thresh)
    n, n_left, n_right = len(y), len(left_idx), len(right_idx)
    if n left == 0 or n right == 0:
        return 0
    child_loss = (n_left / n) * self._entropy(y[left_idx]) + (n_right
    return parent_loss - child_loss
def _best_split(self, X, y, features):
    split = {'score':- 1, 'feat': None, 'thresh': None}
    for feat in features:
        X_{\text{feat}} = X[:, \text{ feat}]
        thresholds = np.unique(X feat)
        for thresh in thresholds:
            score = self._information_gain(X_feat, y, thresh)
            if np.abs(score) > np.abs(split['score']) if split['score
                split['score'] = score
                split['feat'] = feat
                split['thresh'] = thresh
    return split['feat'], split['thresh']
def _build_tree(self, X, y, depth=0):
    self.n_samples, self.n_features = X.shape
    self.n_class_labels = len(np.unique(y))
    # stopping criteria
    if self._is_finished(depth):
        most_common_Label = np.argmax(np.bincount(y))
        return Node(value=most_common_Label)
    # get best split
    rnd_feats = np.random.choice(self.n_features, self.n_features, re
    best_feat, best_thresh = self._best_split(X, y, rnd_feats)
    # grow children recursively
    left_idx, right_idx = self._create_split(X[:, best_feat], best_th
    left_child = self._build_tree(X[left_idx, :], y[left_idx], depth
    right_child = self._build_tree(X[right_idx, :], y[right_idx], dep
    return Node(best_feat, best_thresh, left_child, right_child)
def _traverse_tree(self, x, node):
```

```
if node.is leaf():
                     return node.value
                 if x[node.feature] <= node.threshold:</pre>
                     return self._traverse_tree(x, node.left)
                 return self. traverse tree(x, node.right)
             def fit(self, X, y):
                 self.root = self._build_tree(X, y)
             def predict(self, X):
                 predictions = [self._traverse_tree(x, self.root) for x in X]
                 return np.array(predictions)
In [94]:
         np.random.seed(42)
         model = DecisionTree(max depth=10)
         model.fit(X_train.to_numpy(), y_train.to_numpy())
         y_pred = model.predict(X_test.to_numpy())
         accuracy = accuracy_score(y_test.to_numpy(), y_pred)
         print(f'Accuracy: {accuracy}\n')
        Accuracy: 0.3975
In [177... # CV:
         merged_df = pd.concat([tr_df, te_df], axis=0)
         X = merged_df.drop('label', axis=1)
         y = merged_df['label']
         np.random.seed(42)
         model = DecisionTree(max_depth=10)
         cv_results = custom_cross_val(model, X, y, k=5)
         for i, acc in enumerate(cv_results):
             print(f'Fold {i+1} Accuracy: {acc}')
         print(f'Mean Accuracy: {np.mean(cv_results)}')
        Fold 1 Accuracy: 0.47147147147145
        Fold 2 Accuracy: 0.45245245245245247
        Fold 3 Accuracy: 0.47097097097096
        Fold 4 Accuracy: 0.476976976977
        Fold 5 Accuracy: 0.4954954954955
       Mean Accuracy: 0.47347347347346
```