

Revision of Dataframe

Dataframe from the Previous model

```
In [1]: import pandas as pd
import numpy as np
from sklearn.preprocessing import LabelEncoder
df = pd.read_csv('df.csv')
print("df.shape",df.shape)
item_count = df["target"].value_counts()
print("Number of each species in dataframe:\n",item_count)
```

```
df.shape (3607, 1505)
Number of each species in dataframe:
A_luchuensis      318
A_trinidadensis   235
A_indologenus     232
A_welwitschiae    223
A_sclerotiicarbonarius  220
A_homomorphus     192
A_ibericus        160
A_japonicus       141
A_saccharolyticus 140
A_niger           140
A_vadensis        137
A_costaricaensis  130
A_heteromorphus   130
A_carbonarius     120
A_brasiliensis    120
A_aculeatinus     120
A_sclerotioniger  119
A_ellipticus      110
A_eucalypticola   110
A_aculeatus       100
A_floridensis     90
A_tubingensis     90
A_neoniger        80
A_brunneoviolaceus 80
A_uvarum          70
Name: target, dtype: int64
```

Create New Dataframe

```
In [2]: """
Creating individual dataframe of 'A_costaricaensis' , 'A_neoniger' , 'A_tubingensis',
"""
df_class6 = df[df['target'] == 'A_costaricaensis']
df_class16 = df[df['target'] == 'A_neoniger']
df_class22 = df[df['target'] == 'A_tubingensis']
df_class17 = df[df['target'] == 'A_niger']
df_class25 = df[df['target'] == 'A_welwitschiae']
```

```

"""
Select 80 data randomly from each dataframe of 'A_costaricaensis' , 'A_neoniger' and '
Select 140 data randomly from each dataframe of 'A_niger' and 'A_welwitschiae'
(the number 80,140 based on the minimum of data in each group)
"""

df_class6_rd = df_class6.sample(n = 80)
df_class16_rd = df_class16.sample(n = 80)
df_class22_rd = df_class22.sample(n = 80)
df_class17_rd = df_class17.sample(n = 140)
df_class25_rd = df_class25.sample(n = 140)


"""
Selecting all data from the dataframe in which 'target' is not
'A_costaricaensis', 'A_neoniger', 'A_tubingensis', 'A_niger' and 'A_welwitschiae'
"""

target_cut = ['A_costaricaensis', 'A_neoniger', 'A_tubingensis', 'A_niger', 'A_welwitschiae']
df_cut = df.loc[~df['target'].isin(target_cut)]
print("df after cut 5 species out.shape", df_cut.shape)


"""
Concatenate dataframe (samples randomly) of
'A_costaricaensis', 'A_neoniger', 'A_tubingensis', 'A_niger', 'A_welwitschiae'
"""

list_5 = [df_class6_rd, df_class16_rd, df_class22_rd, df_class17_rd, df_class25_rd]
df_com5 = pd.concat(list_5, axis=0, ignore_index=True)
print("\nShape of combined 5 species dataframe :", df_com5.shape)
item_counts_5 = df_com5["target"].value_counts()
print("Number of each species :\n", item_counts_5)


"""
Renamed A_costaricaensis and A_neoniger in combined dataframe as A_tubingensis
and renamed A_welwitschiae as A_niger

since A_costaricaensis and A_neoniger are synonyms of A_tubingensis
and A_welwitschiae is a synonyms of A_niger
(Bian et al. 2022)
"""

df_com5_rename = df_com5.replace({'A_costaricaensis': 'A_tubingensis',
                                   'A_neoniger': 'A_tubingensis',
                                   'A_welwitschiae': 'A_niger'})
print("\nShape of renamed dataframe :", df_com5_rename.shape)
print

item_counts = df_com5_rename["target"].value_counts()
print("Number of each species :\n", item_counts)


"""
Combine dataframe of cut dataframe and renamed dataframe
"""

list_2 = [df_cut, df_com5_rename]
df_all = pd.concat(list_2, axis=0, ignore_index=True)
df_all.to_csv('dataframe.csv', index=False) # Save to csv file
print("\nSave file as: dataframe.csv")
print("Shape of dataframe :", df_all.shape)

print("Number of each species in dataframe:\n", df_all["target"].value_counts())

```

```
df after cut 5 species out.shape (2944, 1505)
```

```
Shape of combined 5 species dataframe : (520, 1505)
```

```
Number of each species :
```

```

A_niger          140
A_welwitschiae   140
A_costaricaensis 80
A_neoniger       80
A_tubingensis    80
Name: target, dtype: int64
```

```
Shape of renamed dataframe : (520, 1505)
```

```
Number of each species :
```

```

A_niger          280
A_tubingensis    240
Name: target, dtype: int64
```

```
Save file as: dataframe.csv
```

```
Shape of dataframe : (3464, 1505)
```

```
Number of each species in dataframe:
```

```

A_luchuensis     318
A_niger           280
A_tubingensis     240
A_trinidadensis  235
A_indologenus    232
A_sclerotiicarbonarius 220
A_homomorphus    192
A_ibericus       160
A_japonicus      141
A_saccharolyticus 140
A_vadensis       137
A_heteromorphus  130
A_aculeatinus    120
A_brasiliensis   120
A_carbonarius    120
A_sclerotioniger 119
A_ellipticus     110
A_eucalypticola  110
A_aculeatus      100
A_floridensis    90
A_brunneoviolaceus 80
A_uvarum         70
Name: target, dtype: int64
```

Count no. of member in each set & Print class mapping encoder

```

In [3]: cols =[x for x in df.columns if x not in ['target']]
rowused = []
for i in range (len(df)):
    if i % 10 == 0:
        rowused.append('test')

    elif i % 10 == 1:
        rowused.append('validate')

    else:
```

```
rowused.append('train')

df['rowused'] = rowused
dd=df['rowused'].sample(len(df))
test_set=df[df['rowused']=='test']
validate_set=df[df['rowused']=='validate']
train_set=df[df['rowused']=='train']
print('Count test_set:\n',test_set['target'].value_counts())
print('\nCount validate_set:\n',validate_set['target'].value_counts())
print('\nCount train_set:\n',train_set['target'].value_counts())
print('-----')
label_encoder = LabelEncoder()
data_y = df.loc[:, 'target']
encoded_y = label_encoder.fit_transform(data_y.values.ravel())
label_encoder_name_mapping = dict(zip(label_encoder.classes_,label_encoder.transform([
print('Mapping of Label Encoded Classes:', label_encoder_name_mapping, sep="\n")
```

```

Count test_set:
  A_luchuensis          32
  A_trinidadensis       24
  A_indologenus         23
  A_sclerotiicarbonarius 22
  A_welwitschiae        22
  A_homomorphus         20
  A_ibericus            16
  A_saccharolyticus     14
  A_japonicus           14
  A_niger               14
  A_costaricaensis      13
  A_vadensis            13
  A_heteromorphus       13
  A_carbonarius         12
  A_brasiliensis        12
  A_sclerotioniger      12
  A_aculeatinus         12
  A_ellipticus          11
  A_eucalypticola       11
  A_aculeatus           10
  A_floridensis         9
  A_tubingensis         9
  A_neoniger            8
  A_brunneoviolaceus    8
  A_uvarum              7
Name: target, dtype: int64

```

```

Count validate_set:
  A_luchuensis          32
  A_trinidadensis       24
  A_indologenus         23
  A_sclerotiicarbonarius 22
  A_welwitschiae        22
  A_homomorphus         19
  A_ibericus            16
  A_saccharolyticus     14
  A_vadensis            14
  A_japonicus           14
  A_niger               14
  A_costaricaensis      13
  A_heteromorphus       13
  A_carbonarius         12
  A_brasiliensis        12
  A_sclerotioniger      12
  A_aculeatinus         12
  A_ellipticus          11
  A_eucalypticola       11
  A_aculeatus           10
  A_floridensis         9
  A_tubingensis         9
  A_neoniger            8
  A_brunneoviolaceus    8
  A_uvarum              7
Name: target, dtype: int64

```

```

Count train_set:
  A_luchuensis          254
  A_trinidadensis       187
  A_indologenus         186

```

A_welwitschiae	179
A_sclerotiicarbonarius	176
A_homomorphus	153
A_ibericus	128
A_japonicus	113
A_saccharolyticus	112
A_niger	112
A_vadensis	110
A_costaricaensis	104
A_heteromorphus	104
A_carbonarius	96
A_brasiliensis	96
A_aculeatinus	96
A_sclerotioniger	95
A_ellipticus	88
A_eucalypticola	88
A_aculeatus	80
A_floridensis	72
A_tubingensis	72
A_neoniger	64
A_brunneoviolaceus	64
A_uvarum	56

Name: target, dtype: int64

Mapping of Label Encoded Classes:

{'A_aculeatinus': 0, 'A_aculeatus': 1, 'A_brasiliensis': 2, 'A_brunneoviolaceus': 3, 'A_carbonarius': 4, 'A_costaricaensis': 5, 'A_ellipticus': 6, 'A_eucalypticola': 7, 'A_floridensis': 8, 'A_heteromorphus': 9, 'A_homomorphus': 10, 'A_ibericus': 11, 'A_indologenus': 12, 'A_japonicus': 13, 'A_luchuensis': 14, 'A_neoniger': 15, 'A_niger': 16, 'A_saccharolyticus': 17, 'A_sclerotiicarbonarius': 18, 'A_sclerotioniger': 19, 'A_trinidadensis': 20, 'A_tubingensis': 21, 'A_uvarum': 22, 'A_vadensis': 23, 'A_welwitschiae': 24}