

Protein classification

June 17, 2020

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In [1]: import pandas as pd
import numpy as np
from sklearn.preprocessing import OneHotEncoder

In [2]: data=pd.read_csv("/home/bscuser/data/train.csv")

In [3]: data

Out[3]:
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	Sequence	Active
0	DKWL	0
1	FCHN	0
2	KDQP	0
3	FNWI	0
4	NKRM	0
...
111995	GSME	0
111996	DLPT	0
111997	SGHC	0
111998	KIGT	0
111999	PGPT	0

```

[112000 rows x 2 columns]

In [5]: def split(word):
return [char for char in word]

In [14]: first=[]
second=[]
third=[]
fourth=[]
for index,row in data.iterrows():
    first.append(split(row[0])[0])
    second.append(split(row[0])[1])
    third.append(split(row[0])[2])
    fourth.append(split(row[0])[3])

In [21]: data_split=pd.DataFrame()
data_split["First"]=first
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data_split["Second"]=second
data_split["Third"]=third
data_split["Fourth"]=fourth
enc = OneHotEncoder(handle_unknown='ignore')
enc.fit(data_split)

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Out[21]: OneHotEncoder(categories='auto', drop=None, dtype=<class 'numpy.float64'>,
        handle_unknown='ignore', sparse=True)

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In [22]: enc.categories_

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Out[22]: [array(['A', 'C', 'D', 'E', 'F', 'G', 'H', 'I', 'K', 'L', 'M', 'N', 'P',
                'Q', 'R', 'S', 'T', 'V', 'W', 'Y'], dtype=object),
         array(['A', 'C', 'D', 'E', 'F', 'G', 'H', 'I', 'K', 'L', 'M', 'N', 'P',
                'Q', 'R', 'S', 'T', 'V', 'W', 'Y'], dtype=object),
         array(['A', 'C', 'D', 'E', 'F', 'G', 'H', 'I', 'K', 'L', 'M', 'N', 'P',
                'Q', 'R', 'S', 'T', 'V', 'W', 'Y'], dtype=object),
         array(['A', 'C', 'D', 'E', 'F', 'G', 'H', 'I', 'K', 'L', 'M', 'N', 'P',
                'Q', 'R', 'S', 'T', 'V', 'W', 'Y'], dtype=object)]

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In [23]: enc_data=enc.transform(data_split).toarray()

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In [25]: import numpy as np
         from sklearn.model_selection import train_test_split
         X=enc_data
         y=np.array(data["Active"])
         X_train, X_val, y_train, y_val = train_test_split(X, y, test_size=0.1, random_state=42)

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In [26]: from sklearn.pipeline import make_pipeline
         from sklearn.preprocessing import StandardScaler
         from sklearn.svm import SVC

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In [27]: clf = make_pipeline(StandardScaler(), SVC(gamma='auto'))
         clf.fit(X_train, y_train)

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Out[27]: Pipeline(memory=None,
                  steps=[('standardscaler',
                          StandardScaler(copy=True, with_mean=True, with_std=True)),
                          ('svc',
                           SVC(C=1.0, break_ties=False, cache_size=200, class_weight=None,
                                coef0=0.0, decision_function_shape='ovr', degree=3,
                                gamma='auto', kernel='rbf', max_iter=-1, probability=False,
                                random_state=None, shrinking=True, tol=0.001,
                                verbose=False))],
                  verbose=False)

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In [28]: from sklearn.metrics import f1_score
         y_pred=clf.predict(X_val)
         f1_score(y_val, y_pred, average='macro')

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Out [28]: 0.917703465156569

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In [30]: test_data=pd.read_csv("/home/bscuser/data/test.csv")
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In [31]: first=[]
        second=[]
        third=[]
        fourth=[]
        for index,row in test_data.iterrows():
            first.append(split(row[0])[0])
            second.append(split(row[0])[1])
            third.append(split(row[0])[2])
            fourth.append(split(row[0])[3])
```

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In [32]: test_data_split=pd.DataFrame()
        test_data_split["First"]=first
        test_data_split["Second"]=second
        test_data_split["Third"]=third
        test_data_split["Fourth"]=fourth
```

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In [33]: enc_test_data=enc.transform(test_data_split).toarray()
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In [34]: y_test=clf.predict(enc_test_data)
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In [36]: np.savetxt("test_answer.csv", y_test, delimiter=",")
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