

```

import pandas as pd
from sklearn import preprocessing
from sklearn.datasets import load_iris

iris = load_iris()
df = pd.DataFrame(iris.data, columns=iris.feature_names)
df['Species'] = pd.Categorical.from_codes(iris.target, iris.target_names)

print("Iris Dataset:")
print(df.head())

```

Iris Dataset:

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2

	Species
0	setosa
1	setosa
2	setosa
3	setosa
4	setosa

```
min_max_scaler = preprocessing.MinMaxScaler()
```

```
x = df.iloc[:, :-1]
```

```
x_scaled = min_max_scaler.fit_transform(x)
```

```
df_normalized = pd.DataFrame(x_scaled, columns=iris.feature_names)
```

```
print("\nNormalized Iris Dataset:")
print(df_normalized)
```

Normalized Iris Dataset:

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
0	0.222222	0.625000	0.067797	0.041667
1	0.166667	0.416667	0.067797	0.041667
2	0.111111	0.500000	0.050847	0.041667
3	0.083333	0.458333	0.084746	0.041667
4	0.194444	0.666667	0.067797	0.041667
...
145	0.666667	0.416667	0.711864	0.916667
146	0.555556	0.208333	0.677966	0.750000
147	0.611111	0.416667	0.711864	0.791667
148	0.527778	0.583333	0.745763	0.916667
149	0.444444	0.416667	0.694915	0.708333

[150 rows x 4 columns]

```
print("\nUnique values in Species column before Label Encoding:")
print(df['Species'].unique())
```

Unique values in Species column before Label Encoding:

['setosa', 'versicolor', 'virginica']

Categories (3, object): ['setosa', 'versicolor', 'virginica']

```
label_encoder = preprocessing.LabelEncoder()
```

```
df['Species'] = label_encoder.fit_transform(df['Species'])
print("\nUnique values in Species column after Label Encoding:")
print(df['Species'].unique())
```

Unique values in Species column after Label Encoding:

[0 1 2]

```
print("\nUnique values in Species column for One-Hot Encoding:")
print(df['Species'].unique())
features_df = df.drop(columns=['Species'])
enc = preprocessing.OneHotEncoder(sparse_output=False)
enc_df = pd.DataFrame(enc.fit_transform(df[['Species']]))
df_encoded = features_df.join(enc_df)
df_encoded.rename(columns={0: 'Iris-Setosa', 1: 'Iris-Versicolor', 2: 'Iris-
Virginica'}, inplace=True)
print("\nOne-Hot Encoded Iris Dataset:")
print(df_encoded)
```

Unique values in Species column for One-Hot Encoding:

[0 1 2]

One-Hot Encoded Iris Dataset:

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
\				
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
..
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

	Iris-Setosa	Iris-Versicolor	Iris-Virginica
0	1.0	0.0	0.0
1	1.0	0.0	0.0
2	1.0	0.0	0.0
3	1.0	0.0	0.0
4	1.0	0.0	0.0
..
145	0.0	0.0	1.0
146	0.0	0.0	1.0
147	0.0	0.0	1.0
148	0.0	0.0	1.0
149	0.0	0.0	1.0

[150 rows x 7 columns]

```
df_dummy_encoded = pd.get_dummies(df, drop_first=True)
```

```
print("\nDummy Encoded Iris Dataset:")
print(df_dummy_encoded)
```

Dummy Encoded Iris Dataset:

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
\				
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
..
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

	Species
0	0
1	0
2	0
3	0
4	0
..	...
145	2
146	2
147	2
148	2
149	2

[150 rows x 5 columns]

