

```
import pandas as pd
import seaborn as sns

dt = pd.read_csv('/content/iris.csv')

dt
```

Release notes

NameErrc ...



Please explain the error:

NameError: name 'dt' is not d



Hypothesis

The NameError: name 'dt' is not defined indicates that the variable dt was not found in the current scope when the code dt was executed. This is likely because the cell where dt = pd.read_csv('/content/iris.csv') was defined was not run, or its output was cleared from the environment.

Based on the provided source files and traceback, it appears that the cells were executed in the following order:

1. import pandas as pd\nimport seaborn as sns (Cell with source file ipython-input-3-36673dcb6ee0) - This imports the necessary libraries.
2. dt (Cell with source file ipython-input-5-36673dcb6ee0) - This cell attempts to display the dt variable. However, dt has not been defined yet.
3. dt = pd.read_csv('/content/irj (Cell with source file ipython-input-0-36673dcb6ee0) - This cell,



which *defines* `dt` , was executed *after* the cell that tried to display it.

Therefore, when the cell with `dt` was executed, the variable `dt` did not exist in the global namespace, leading to the `NameError` .

Suggested Changes

To fix this issue, ensure that the cell defining `dt` is executed before the cell that attempts to use it. The corrected order of execution should be:

1. Import libraries.
2. Load the CSV file into `dt` .
3. Display `dt` .

Since you are working in a Jupyter notebook, simply re-running the cells in the correct order should resolve the problem. Below is the code in the correct execution order, assuming each block is a separate cell in your notebook.



```
# Import necessary libraries
import pandas as pd
import seaborn as sns
```

[Use code with caution](#)



```
# Load the dataset into the variable
dt = pd.read_csv('/content/iris.csv')
```

[Use code with caution](#)

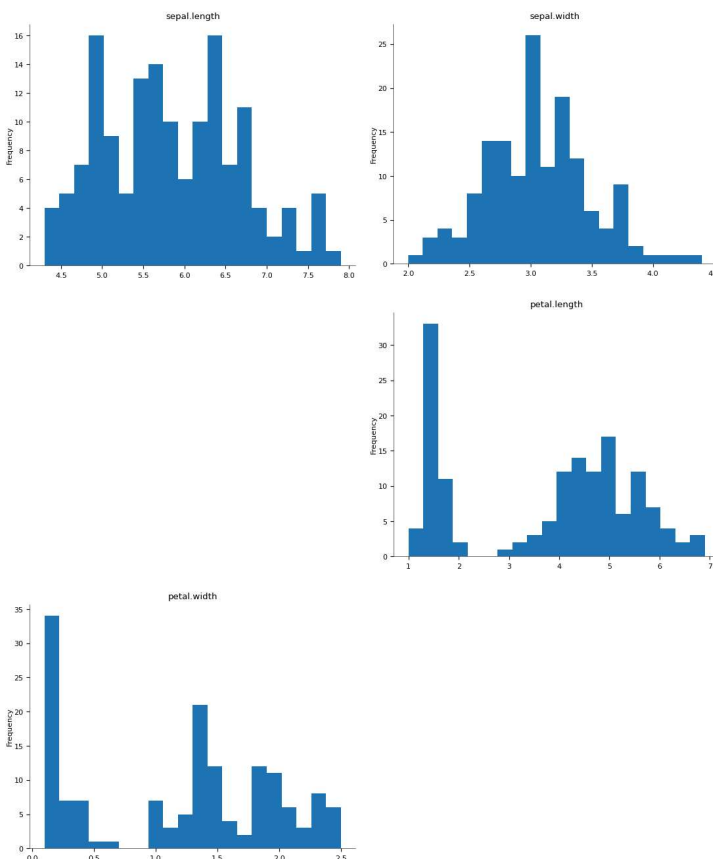


```
# Display the content of dt
dt
```

[Use code with caution](#)

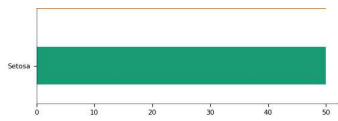
150 rows × 5 columns

Distributions

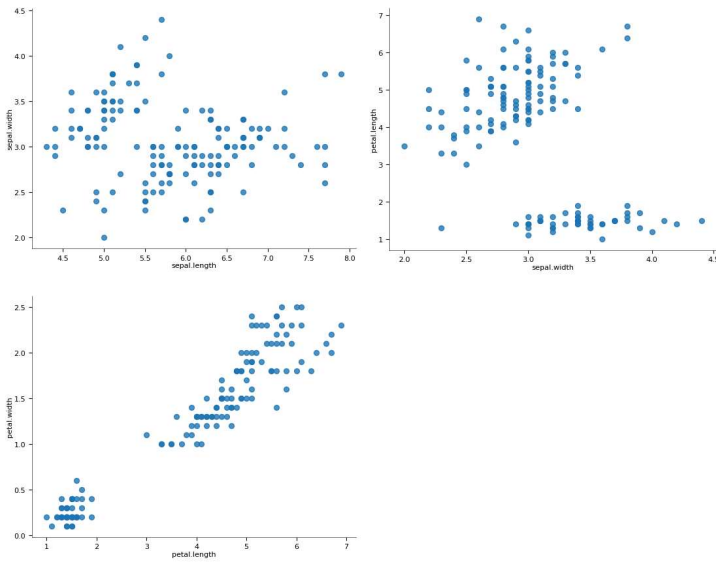


Categorical distributions

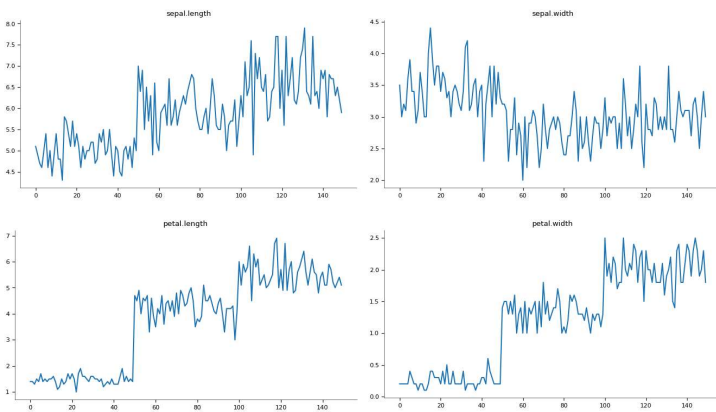




2-d distributions



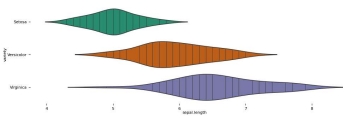
Values



Faceted distributions

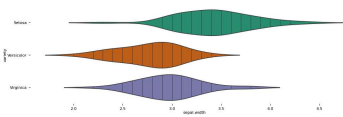
<string>:5: FutureWarning:

Passing `palette` without assigning `hue` is depre



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<string>:5: FutureWarning:

Passing `palette` without assigning `hue` is depre



Rate this answer  

Next
steps:

[Generate code with dt](#)
[View recommended plots](#)

```
dt.describe()
```



	sepal.length	sepal.width	petal.length	petal.width
count	150.000000	150.000000	150.000000	150.000000
mean	5.843333	3.057333	3.758000	1.593333
std	0.828066	0.435866	1.765298	0.469768
min	4.300000	2.000000	1.000000	0.100000
25%	5.100000	2.800000	1.600000	0.400000
50%	5.800000	3.000000	4.350000	1.300000
75%	6.400000	3.300000	5.100000	1.800000
max	7.900000	4.400000	6.900000	4.700000

```
dt.info()
```



```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
#   Column          Non-Null Count  Dtype
---  -
0   sepal.length    150 non-null    float64
1   sepal.width     150 non-null    float64
2   petal.length    150 non-null    float64
3   petal.width     150 non-null    float64
4   variety         150 non-null    object
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
```

```
dt['variety'].unique()
```



```
array(['Setosa', 'Versicolor', 'Virginica'],
      dtype=object)
```

```
from sklearn.preprocessing import LabelEncoder
```

```
l = LabelEncoder()
```

```
dt['variety'] = l.fit_transform(dt['variety'])
```

```
dt['variety'].unique()
```

```
array([0, 1, 2])
```

```
from sklearn.model_selection import train_test_split
```

```
x = dt.drop(['variety'],axis =1)
y = dt['variety']
```

x

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

150 rows × 4 columns

Next steps:

Generate code with x

 View recommended plots

```
dt.corr()
```

	sepal.length	sepal.width	petal.length
sepal.length	1.000000	-0.117570	0.871754
sepal.width	-0.117570	1.000000	-0.428440
petal.length	0.871754	-0.428440	1.000000
petal.width	0.817941	-0.366126	0.962865
variety	0.782561	-0.426658	0.949035

```
xtrain, xtest, ytrain, ytest = train_test_split(x,y, tes
```

```
xtest
```



	sepal.length	sepal.width	petal.length	petal
15	5.7	4.4	1.5	
98	5.1	2.5	3.0	
93	5.0	2.3	3.3	
19	5.1	3.8	1.5	
89	5.5	2.5	4.0	
37	4.9	3.6	1.4	
0	5.1	3.5	1.4	
94	5.6	2.7	4.2	
66	5.6	3.0	4.5	
97	6.2	2.9	4.3	
49	5.0	3.3	1.4	
132	6.4	2.8	5.6	
63	6.1	2.9	4.7	
58	6.6	2.9	4.6	
90	5.5	2.6	4.4	
142	5.8	2.7	5.1	
70	5.9	3.2	4.8	
53	5.5	2.3	4.0	
84	5.4	3.0	4.5	
62	6.0	2.2	4.0	
33	5.5	4.2	1.4	
122	7.7	2.8	6.7	
76	6.8	2.8	4.8	
77	6.7	3.0	5.0	
32	5.2	4.1	1.5	
123	6.3	2.7	4.9	
82	5.8	2.7	3.9	
136	6.3	3.4	5.6	
114	5.8	2.8	5.1	
3	4.6	3.1	1.5	