In [44]:

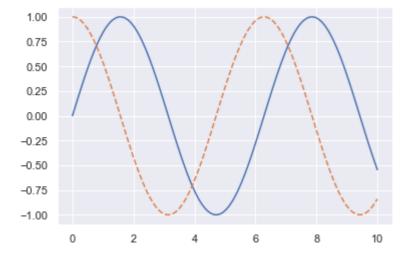
```
# maha ebrahim mohammed
# 4051350
# IA8G
# lab 5: Data Exploratory Analysis
import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
%matplotlib inline
sns.set(color_codes=True)
```

In [45]:

```
x = np.linspace(0, 10, 100)
fig = plt.figure()
plt.plot(x, np.sin(x), '-')
plt.plot(x, np.cos(x), '--')
```

Out[45]:

[<matplotlib.lines.Line2D at 0x238a462c400>]



In [47]:

```
from sklearn import datasets
iris = datasets.load_iris()
df = pd.read_csv(iris.filename)
df.head(5)
```

Out[47]:

	150	4	setosa	versicolor	virginica
0	5.1	3.5	1.4	0.2	0
1	4.9	3.0	1.4	0.2	0
2	4.7	3.2	1.3	0.2	0
3	4.6	3.1	1.5	0.2	0
4	5.0	3.6	1.4	0.2	0

In [48]:

df_with_filepath = pd.read_csv('C:\\anaconda3\\lib\\site-packages\\sklearn\\datasets\\data\ df_with_filepath.head()

Out[48]:

	5.1	3.5	1.4	0.2	0
0	4.9	3.0	1.4	0.2	0
1	4.7	3.2	1.3	0.2	0
2	4.6	3.1	1.5	0.2	0
3	5.0	3.6	1.4	0.2	0
4	5.4	3.9	1.7	0.4	0

In [14]:

df_with_filepath.head()

Out[14]:

	4.7	3.2	1.3	0.2	0
0	4.6	3.1	1.5	0.2	0
1	5.0	3.6	1.4	0.2	0
2	5.4	3.9	1.7	0.4	0
3	4.6	3.4	1.4	0.3	0
4	5.0	3.4	1.5	0.2	0

In [18]:

```
iris.feature_names
```

Out[18]:

```
['sepal length (cm)',
 'sepal width (cm)',
 'petal length (cm)',
 'petal width (cm)']
```

In [37]:

```
df_with_columns = pd.read_csv('C:\\anaconda3\\lib\\site-packages\\sklearn\\datasets\\data\\
df_with_columns.head()
```

Out[37]:

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

In [31]:

```
df = pd.read_csv(iris.filename, names = iris.feature_names)
df
```

Out[31]:

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
150.0	4.0	setosa	versicolor	virginica
5.1	3.5	1.4	0.2	0
4.9	3.0	1.4	0.2	0
4.7	3.2	1.3	0.2	0
4.6	3.1	1.5	0.2	0
6.7	3.0	5.2	2.3	2
6.3	2.5	5.0	1.9	2
6.5	3.0	5.2	2.0	2
6.2	3.4	5.4	2.3	2
5.9	3.0	5.1	1.8	2

151 rows × 4 columns

```
In [32]:
```

```
df.info()
<class 'pandas.core.frame.DataFrame'>
```

```
Float64Index: 151 entries, 150.0 to 5.9
Data columns (total 4 columns):
    Column
#
                       Non-Null Count Dtype
                       -----
    sepal length (cm) 151 non-null
 0
                                      float64
 1
    sepal width (cm)
                       151 non-null
                                      object
    petal length (cm) 151 non-null
                                      object
    petal width (cm)
                       151 non-null
                                      object
dtypes: float64(1), object(3)
memory usage: 5.9+ KB
```

In [38]:

```
print(df_with_columns.shape)
print(df_with_columns.columns)
```

```
(150, 5)
Index(['sepal length (cm)', 'sepal width (cm)', 'petal length (cm)',
       'petal width (cm)', 'species'],
      dtype='object')
```

In [39]:

```
df_with_columns.isnull()
```

Out[39]:

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	species
0	False	False	False	False	False
1	False	False	False	False	False
2	False	False	False	False	False
3	False	False	False	False	False
4	False	False	False	False	False
145	False	False	False	False	False
146	False	False	False	False	False
147	False	False	False	False	False
148	False	False	False	False	False
149	False	False	False	False	False

150 rows × 5 columns

In [40]:

```
print(df_with_columns["species"].value_counts())
     50
```

50 1 2 50

Name: species, dtype: int64

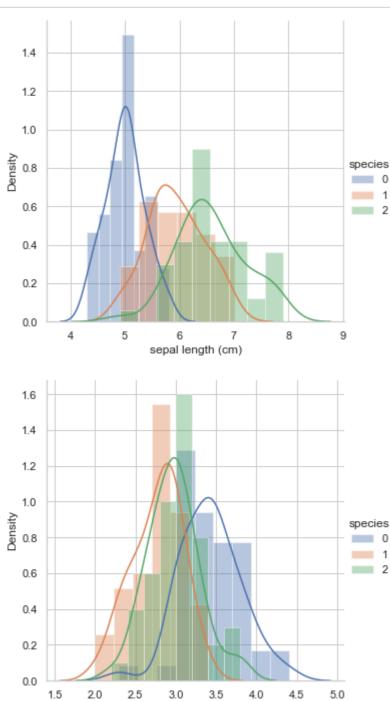
In [41]:

```
print(df_with_columns.describe())
```

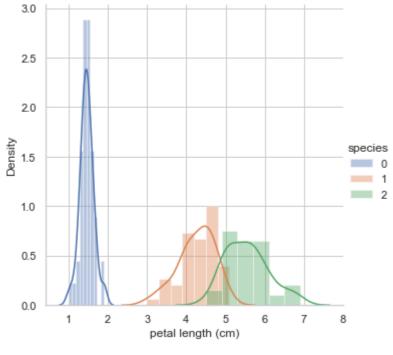
count mean std min 25% 50% 75% max	sepal length (cm) 150.000000 5.843333 0.828066 4.300000 5.100000 5.800000 6.400000 7.900000	sepal width (cm) 150.000000 3.057333 0.435866 2.000000 2.800000 3.000000 3.300000 4.400000	petal length (cm) 150.000000 3.758000 1.765298 1.000000 1.600000 4.350000 5.100000 6.900000	\
count mean std min 25% 50% 75% max	petal width (cm) 150.000000 1.199333 0.762238 0.100000 0.300000 1.300000 1.800000 2.500000	species 150.000000 1.000000 0.819232 0.000000 0.000000 2.000000 2.000000		

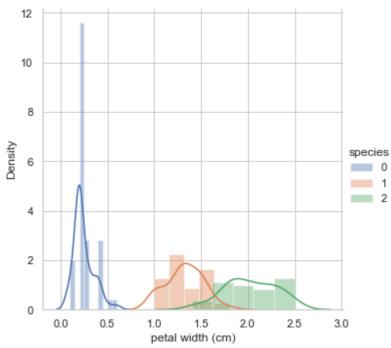
In [57]:

```
import warnings
warnings.simplefilter(action='ignore', category=FutureWarning)
for ojha, feature in enumerate(list(df_with_columns.columns)[:-1]):
    fg = sns.FacetGrid(df_with_columns, hue ='species', height=5)
    fg.map(sns.distplot, feature).add_legend()
plt.show()
```



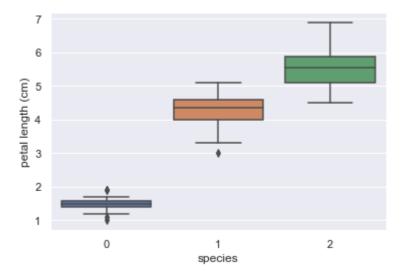
sepal width (cm)





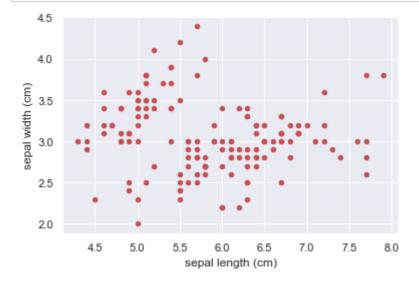
In [52]:

```
sns.boxplot(x='species',y='petal length (cm)', data=df_with_columns)
plt.show()
```



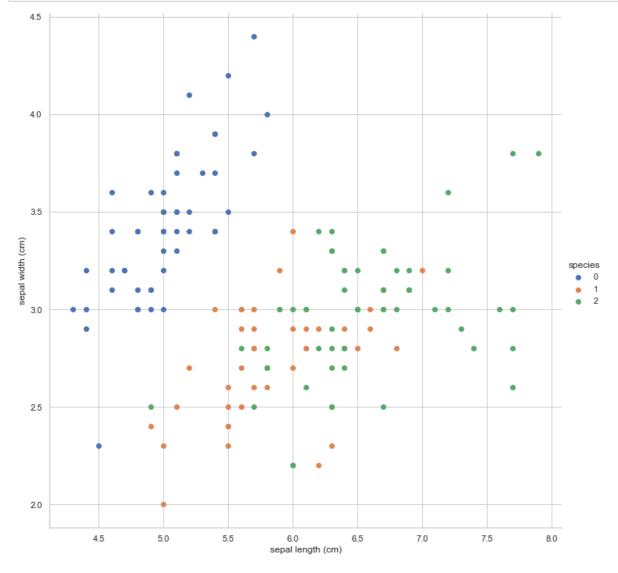
In [54]:

df_with_columns.plot(kind= 'scatter' ,x= 'sepal length (cm)', y= 'sepal width (cm)', c = 'r plt.show()



In [55]:

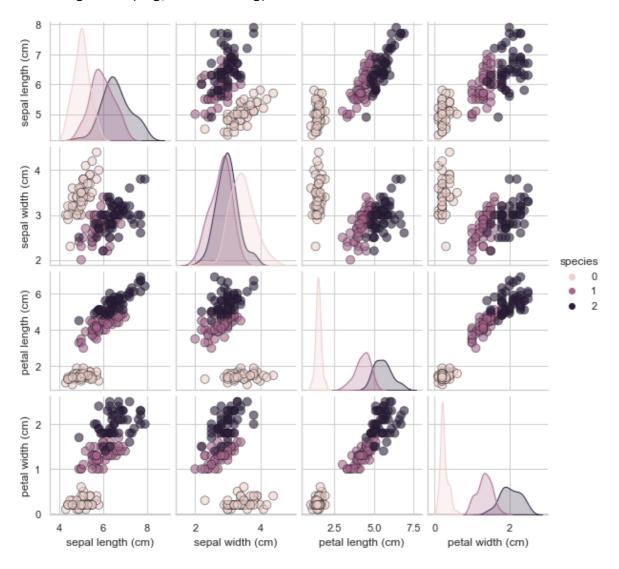
```
sns.set_style("whitegrid");
sns.FacetGrid(df_with_columns, hue="species", height=10) \
    .map(plt.scatter, "sepal length (cm)", "sepal width (cm)") \
    .add_legend();
plt.show();
```



In [56]:

```
sns.set_style("whitegrid");
sns.pairplot(df_with_columns, vars = iris.feature_names, hue = 'species', diag_kind='kde',
            plot_kws = {'alpha': 0.6, 's': 80, 'edgecolor': 'k'}, size = 2)
plt.show()
```

C:\anaconda3\lib\site-packages\seaborn\axisgrid.py:2076: UserWarning: The `s ize` parameter has been renamed to `height`; please update your code. warnings.warn(msg, UserWarning)



In []: