### **Table of Contents**

- 1 sns pairplot
- 2 LinearRegression
- 3 Gaussian Naive Bayes (GaussianNB)
- 4 iris dimensionality
- 5 Gaussian mixture model (GMM)

#### In [1]:

```
import seaborn as sns
import matplotlib.pyplot as plt
%matplotlib inline
```

In [2]:

import numpy as np

### sns - pairplot

In [3]:

```
iris = sns. load_dataset('iris')
iris. head()
```

Out[3]:

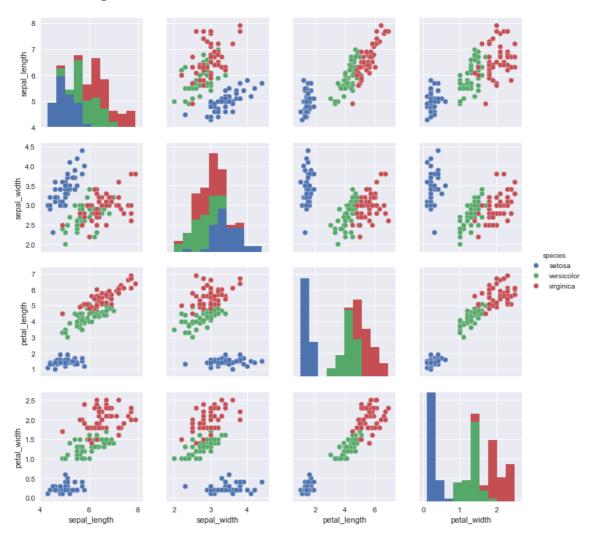
	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	setosa
1	4.9	3.0	1.4	0.2	setosa
2	4.7	3.2	1.3	0.2	setosa
3	4.6	3.1	1.5	0.2	setosa
4	5.0	3.6	1.4	0.2	setosa

### In [4]:

sns.set()
sns.pairplot(iris, hue='species', size=2.5)

#### Out[4]:

<seaborn.axisgrid.PairGrid at 0xae1dc50>



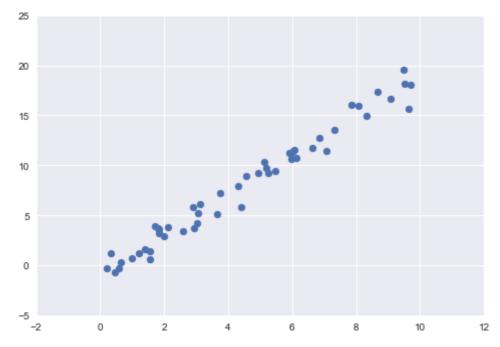
# LinearRegression

#### In [5]:

```
rng = np. random. RandomState(42)

x = 10* rng. rand(50)
y = 2*x -1 + rng. randn(50)

plt. scatter(x, y)
plt. xlim(-2, 12)
plt. ylim(-5, 25)
# plt. xlim(xmax=12)
# plt. ylim(ymax=25)
plt. show()
```



#### In [6]:

from sklearn.linear\_model import LinearRegression

#### In [7]:

```
model = LinearRegression(fit_intercept=True)
model
```

#### Out[7]:

LinearRegression(copy\_X=True, fit\_intercept=True, n\_jobs=1, normalize=False)

#### In [8]:

```
X = x[: , np.newaxis]
print(X.shape)
# X
```

(50, 1)

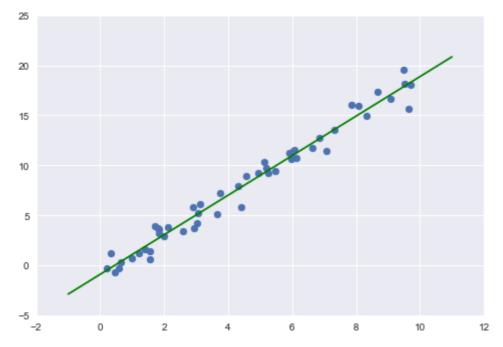
```
In [9]:
model.fit(X,y)
Out[9]:
LinearRegression(copy_X=True, fit_intercept=True, n_jobs=1, normalize=False)
In [10]:
model.coef_
Out[10]:
array([ 1.9776566])
In [11]:
model.intercept_
Out[11]:
-0.90331072553111635
In [12]:
xfit = np. linspace(-1, 11)
Xfit = xfit[:, np.newaxis]
yfit = model.predict(Xfit)
In [13]:
xfit.shape
Out[13]:
(50,)
```

#### In [14]:

```
plt.scatter(x, y)
plt.plot(xfit, yfit, color='g')

plt.xlim(-2, 12)
plt.ylim(-5, 25)

plt.show()
```



### Gaussian Naive Bayes (GaussianNB)

In [15]:

from sklearn.cross\_validation import train\_test\_split

C:\Program Files\Anaconda3\lib\site-packages\sklearn\cross\_validation.py:44: Depre cationWarning: This module was deprecated in version 0.18 in favor of the model\_se lection module into which all the refactored classes and functions are moved. Also note that the interface of the new CV iterators are different from that of this module. This module will be removed in 0.20.

"This module will be removed in 0.20.", DeprecationWarning)

In [16]:

from sklearn.model\_selection import train test split

```
In [17]:
```

```
X_iris = iris.drop('species', axis=1)
y_iris = iris['species']

# Xtrain, Xtest, ytrain, ytest = train_test_split(X_iris, y_iris, random_state=1, test_size=0.5)
Xtrain, Xtest, ytrain, ytest = train_test_split(X_iris, y_iris, random_state=1)
```

```
In [18]:
```

from sklearn.naive\_bayes import GaussianNB

```
In [19]:
```

```
model = GaussianNB()
model.fit(Xtrain, ytrain)
y_predict = model.predict(Xtest)
```

In [20]:

from sklearn.metrics import accuracy\_score

In [21]:

```
accuracy_score(ytest, y_predict)
```

Out[21]:

0.97368421052631582

### iris dimensionality

```
In [22]:
```

```
from sklearn.decomposition import PCA

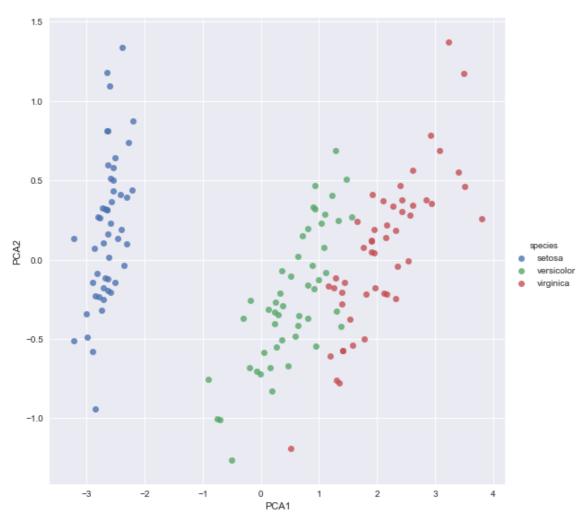
model = PCA(n_components=2)
model.fit(X_iris)
X_2D = model.transform(X_iris)
```

#### In [23]:

```
iris['PCA1'] = X_2D[:, 0]
iris['PCA2'] = X_2D[:, 1]
sns.lmplot('PCA1', 'PCA2', hue='species', data=iris, fit_reg=False, size=8)
```

#### Out[23]:

<seaborn.axisgrid.FacetGrid at 0xdc839b0>



## Gaussian mixture model (GMM)

```
In [24]:
```

```
# from sklearn.mixture import GMM
```

#### In [28]:

```
# model = GMM(n_components=3, covariance_type='full')
# model.fit(X_iris)
# y_gmm = model.predict(X_iris)
```