

```
!pip install pyspark
```

```
Looking in indexes: https://pypi.org/simple, https://us-python.pkg.dev/colab-w
Collecting pyspark
  Downloading pyspark-3.2.1.tar.gz (281.4 MB)
    |████████████████████████████████████████| 281.4 MB 34 kB/s
Collecting py4j==0.10.9.3
  Downloading py4j-0.10.9.3-py2.py3-none-any.whl (198 kB)
    |████████████████████████████████████████| 198 kB 53.7 MB/s
Building wheels for collected packages: pyspark
  Building wheel for pyspark (setup.py) ... done
  Created wheel for pyspark: filename=pyspark-3.2.1-py2.py3-none-any.whl size=
  Stored in directory: /root/.cache/pip/wheels/9f/f5/07/7cd8017084dce4e93e84e9
Successfully built pyspark
Installing collected packages: py4j, pyspark
Successfully installed py4j-0.10.9.3 pyspark-3.2.1
```

```
from pyspark.sql import SparkSession
spark=SparkSession.builder.appName('SparkML').getOrCreate()
```

```
spark
```

↳ **SparkSession - in-memory**

SparkContext

[Spark UI](#)

Version

v3.2.1

Master

local[*]

AppName

SparkML

```
df = spark.read.csv('IRIS.csv',header=True,inferSchema=True)
```

```
df.printSchema()
```

```
root
 |-- sepal_length: double (nullable = true)
 |-- sepal_width: double (nullable = true)
 |-- petal_length: double (nullable = true)
 |-- petal_width: double (nullable = true)
 |-- species: string (nullable = true)
```

```
df.show(5)
```

```
+-----+-----+-----+-----+-----+
|sepal_length|sepal_width|petal_length|petal_width|    species|
+-----+-----+-----+-----+-----+
|         5.1|         3.5|         1.4|         0.2| Iris-setosa|
|         4.9|         3.0|         1.4|         0.2| Iris-setosa|
```

4.7	3.2	1.3	0.2	Iris-setosa
4.6	3.1	1.5	0.2	Iris-setosa
5.0	3.6	1.4	0.2	Iris-setosa

only showing top 5 rows

```
#Preprocessing steps
```

```
from pyspark.ml.feature import StringIndexer, OneHotEncoder
```

```
# create object of StringIndexer class and specify input and output column
```

```
SI_species = StringIndexer(inputCol='species',outputCol='species1')
```

```
# transform the data
```

```
df = SI_species.fit(df).transform(df)
```

```
df.tail(5)
```

```
Row(sepal_length=6.7, sepal_width=3.0, petal_length=5.2, petal_width=2.3, spe
Row(sepal_length=6.3, sepal_width=2.5, petal_length=5.0, petal_width=1.9, spe
Row(sepal_length=6.5, sepal_width=3.0, petal_length=5.2, petal_width=2.0, spe
Row(sepal_length=6.2, sepal_width=3.4, petal_length=5.4, petal_width=2.3, spe
Row(sepal_length=5.9, sepal_width=3.0, petal_length=5.1, petal_width=1.8, spe
```

```
df.printSchema()
```

```
root
```

```
-- sepal_length: double (nullable = true)
-- sepal_width: double (nullable = true)
-- petal_length: double (nullable = true)
-- petal_width: double (nullable = true)
-- species: string (nullable = true)
-- species_Index: double (nullable = false)
-- species1: double (nullable = false)
```

```
from pyspark.ml.feature import VectorAssembler
```

```
fa1=VectorAssembler(inputCols=['sepal_length','petal_length','petal_width'],outputC
```

```
df1=fa1.transform(df)
```

```
df1.show()
```

sepal_length	sepal_width	petal_length	petal_width	species	newic1
5.1	3.5	1.4	0.2	Iris-setosa	[5.1,1.4,0.2]
4.9	3.0	1.4	0.2	Iris-setosa	[4.9,1.4,0.2]
4.7	3.2	1.3	0.2	Iris-setosa	[4.7,1.3,0.2]
4.6	3.1	1.5	0.2	Iris-setosa	[4.6,1.5,0.2]
5.0	3.6	1.4	0.2	Iris-setosa	[5.0,1.4,0.2]
5.4	3.9	1.7	0.4	Iris-setosa	[5.4,1.7,0.4]
4.6	3.4	1.4	0.3	Iris-setosa	[4.6,1.4,0.3]

	5.0		3.4		1.5		0.2		Iris-setosa		[5.0,1.5,0.2]	
	4.4		2.9		1.4		0.2		Iris-setosa		[4.4,1.4,0.2]	
	4.9		3.1		1.5		0.1		Iris-setosa		[4.9,1.5,0.1]	
	5.4		3.7		1.5		0.2		Iris-setosa		[5.4,1.5,0.2]	
	4.8		3.4		1.6		0.2		Iris-setosa		[4.8,1.6,0.2]	
	4.8		3.0		1.4		0.1		Iris-setosa		[4.8,1.4,0.1]	
	4.3		3.0		1.1		0.1		Iris-setosa		[4.3,1.1,0.1]	
	5.8		4.0		1.2		0.2		Iris-setosa		[5.8,1.2,0.2]	
	5.7		4.4		1.5		0.4		Iris-setosa		[5.7,1.5,0.4]	
	5.4		3.9		1.3		0.4		Iris-setosa		[5.4,1.3,0.4]	
	5.1		3.5		1.4		0.3		Iris-setosa		[5.1,1.4,0.3]	
	5.7		3.8		1.7		0.3		Iris-setosa		[5.7,1.7,0.3]	
	5.1		3.8		1.5		0.3		Iris-setosa		[5.1,1.5,0.3]	

```
+-----+-----+-----+-----+-----+-----+-----+-----+
```

only showing top 20 rows

```
df2=df1.select("newic1","species")
```

```
df2.show()
```

	newic1		species	
	[5.1,1.4,0.2]		Iris-setosa	
	[4.9,1.4,0.2]		Iris-setosa	
	[4.7,1.3,0.2]		Iris-setosa	
	[4.6,1.5,0.2]		Iris-setosa	
	[5.0,1.4,0.2]		Iris-setosa	
	[5.4,1.7,0.4]		Iris-setosa	
	[4.6,1.4,0.3]		Iris-setosa	
	[5.0,1.5,0.2]		Iris-setosa	
	[4.4,1.4,0.2]		Iris-setosa	
	[4.9,1.5,0.1]		Iris-setosa	
	[5.4,1.5,0.2]		Iris-setosa	
	[4.8,1.6,0.2]		Iris-setosa	
	[4.8,1.4,0.1]		Iris-setosa	
	[4.3,1.1,0.1]		Iris-setosa	
	[5.8,1.2,0.2]		Iris-setosa	
	[5.7,1.5,0.4]		Iris-setosa	
	[5.4,1.3,0.4]		Iris-setosa	
	[5.1,1.4,0.3]		Iris-setosa	
	[5.7,1.7,0.3]		Iris-setosa	
	[5.1,1.5,0.3]		Iris-setosa	

```
+-----+-----+-----+-----+-----+
```

only showing top 20 rows

```
df.show(5)
```

	sepal_length		sepal_width		petal_length		petal_width		species	
	5.1		3.5		1.4		0.2		Iris-setosa	
	4.9		3.0		1.4		0.2		Iris-setosa	
	4.7		3.2		1.3		0.2		Iris-setosa	
	4.6		3.1		1.5		0.2		Iris-setosa	

	5.0	3.6	1.4	0.2	Iris-setosa
only showing top 5 rows					

```

from pyspark.ml.clustering import KMeans
from pyspark.ml.evaluation import ClusteringEvaluator
train_data,test_data=df2.randomSplit([0.99,0.01])
applyml=KMeans(featuresCol='newic1', k=3)
applyml=applyml.fit(train_data)

```

```
applyml
```

```
KMeansModel: uid=KMeans_36cc25d0ed52, k=3, distanceMeasure=euclidean, numFeatu
```

```
applyml.transform(train_data).groupBy("prediction").count().show()
```

prediction	count
1	37
2	61
0	50

```
predict=applyml.transform(train_data)
```

```
predict.show()
```

newic1	species	prediction
[4.3,1.1,0.1]	Iris-setosa	0
[4.4,1.3,0.2]	Iris-setosa	0
[4.4,1.3,0.2]	Iris-setosa	0
[4.4,1.4,0.2]	Iris-setosa	0
[4.5,1.3,0.3]	Iris-setosa	0
[4.6,1.0,0.2]	Iris-setosa	0
[4.6,1.4,0.2]	Iris-setosa	0
[4.6,1.4,0.3]	Iris-setosa	0
[4.6,1.5,0.2]	Iris-setosa	0
[4.7,1.3,0.2]	Iris-setosa	0
[4.7,1.6,0.2]	Iris-setosa	0
[4.8,1.4,0.1]	Iris-setosa	0
[4.8,1.4,0.3]	Iris-setosa	0
[4.8,1.6,0.2]	Iris-setosa	0
[4.8,1.6,0.2]	Iris-setosa	0
[4.8,1.9,0.2]	Iris-setosa	0
[4.9,1.4,0.2]	Iris-setosa	0
[4.9,1.5,0.1]	Iris-setosa	0
[4.9,1.5,0.1]	Iris-setosa	0
[4.9,1.5,0.1]	Iris-setosa	0

only showing top 20 rows

```
predict.groupBy("species","prediction").count().show()
```

species	prediction	count
Iris-versicolor	2	47
Iris-setosa	0	50
Iris-virginica	1	35
Iris-virginica	2	14
Iris-versicolor	1	2

```
predict.
```

```
-----
IllegalArgumentExce... Traceback (most recent call last)
<ipython-input-316-dbd573341709> in <module>()
----> 1 predict.corr('species','prediction')

----- 2 frames -----
/usr/local/lib/python3.7/dist-packages/pyspark/sql/utils.py in deco(*a, **kw)
    115         # Hide where the exception came from that shows a
non-Pythonic
    116         # JVM exception message.
--> 117         raise converted from None
    118     else:
    119         raise

IllegalArgumentExce... requirement failed: Currently correlation
calculation for columns with dataType string not supported.
```

 0s completed at 6:14 PM

 