Machine Learning Assignment (Clustering)

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Load Data

```
In [1]:
         import pandas as pd
         import numpy as np
         # // Read Data and Evaluation files
         Artificial Eva = pd.read csv('Evaluations1.csv')
         Spiral_Eva =pd.read_csv('Evaluations2.csv')
         Path_Eva= pd.read_csv('Evaluations3.csv')
         Path Eva.head()
         Artificial_data = pd.read_csv('Artificial.csv')
         Artificial data = np.array(Artificial data.iloc[:, [0, 1]]);
         Spiral_data = pd.read_csv('Spiral.csv')
         Spiral_data = np.array(Spiral_data.iloc[:, [0, 1]]);
         Path data = pd.read csv('Path.csv')
         Path data = np.array(Path data.iloc[:, [0, 1]]);
         data= Artificial_data;
         def Load Data(Is slicing):
             Artificial data = pd.read csv('Artificial.csv')
             Spiral data = pd.read csv('Spiral.csv')
             Path_data = pd.read_csv('Path.csv')
             if Is slicing==1:
                 Artificial_data = np.array(Artificial_data.iloc[:, [0, 1]]);
                 Spiral data = np.array(Spiral data.iloc[:, [0, 1]]);
                 Path_data = np.array(Path_data.iloc[:, [0, 1]]);
             return (Artificial_data, Spiral_data, Path_data )
         Load Data(1)
         Artificial_Eva.head()
```

Out[1]:		Clustring Algorithm	Root MSE	DBI	Silhouette Score	Purity	Square Error
	0	K-Means	0.0	0.0	0.0	0.0	0.0
	1	Mini-Batch K-Means	0.0	0.0	0.0	0.0	0.0
	2	DBSCAN	0.0	0.0	0.0	0.0	0.0
	3	Spectral Clustering	0.0	0.0	0.0	0.0	0.0
	4	K means self implemenation	0.0	0.0	0.0	0.0	0.0

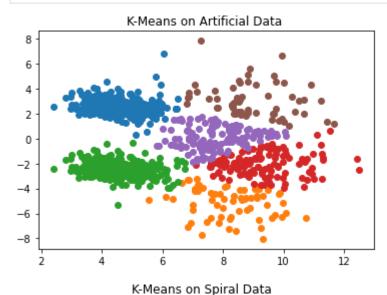
Evaluations Metrics

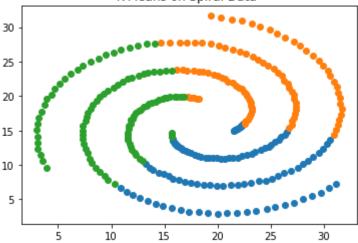
```
In [2]:
         import numpy as np
         import math
         from sklearn import metrics
         from sklearn.metrics import davies bouldin score
         from sklearn.metrics import silhouette score
         from sklearn.metrics import mean squared error
         def purity_score(y_true, y_pred):
             # compute contingency matrix (also called confusion matrix)
             contingency matrix = metrics.cluster.contingency matrix(y true, y pred)
             # return purity
             return np.sum(np.amax(contingency matrix, axis=0)) / np.sum(contingency matrix)
         def Measure_Evaluation(index, k, data= [], Eva_df= [], inertia=[], y_pred=[]):
             X = np.array(data.iloc[:, [0, 1]]);
             db_index = davies_bouldin_score(X, y_per[k])
             sh_score = silhouette_score(X, y_per[k], metric='euclidean')
             X = np.array(data.iloc[:, [2]])
             purity= purity_score(X,y_per[k])
             RMSE=math. sqrt(mean_squared_error(X, y_per[k]))
             Eva_df.at[index,'Root MSE']= RMSE
             Eva df.at[index,'DBI']=db index
             Eva_df.at[index,'Silhouette Score']=sh_score
             Eva_df.at[index, 'Purity']=purity
             Eva df.at[index, 'Square Error'] = inertia[k]
```

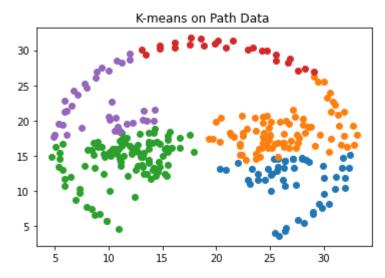
K - Means

```
In [3]:
         # k-means clustering
         from numpy import unique
         from numpy import where
         import numpy as np
         from sklearn.cluster import KMeans
         from matplotlib import pyplot as plt
         inertia= []
         y per= []
         def Kmeans ( msg, cluster, data= []):
             model = KMeans(n clusters=cluster, max iter=1000)
              # fit the model
             kmeans=model.fit(data)
             inertia.append(kmeans.inertia )
             yhat = model.predict(data)
             y_per.append(yhat)
             # retrieve unique clusters
             clusters = unique(yhat)
             # create scatter plot for samples from each cluster
             for cluster in clusters:
```

```
# get row indexes for samples with this cluster
        row ix = where(yhat == cluster)
        # create scatter of these samples
        plt.scatter(data[row_ix, 0], data[row_ix, 1])
    # show the plot
    plt.title(msg);
    plt.show()
Artificial_data,Spiral_data,Path_data= Load_Data(1)
Kmeans( "K-Means on Artificial Data ", 6, Artificial data)
Kmeans( "K-Means on Spiral Data ", 3,Spiral_data)
Kmeans( "K-means on Path Data ",5, Path_data)
# Evaluations
inertia=[0,0,0]
Artificial_data,Spiral_data,Path_data= Load_Data(0)
Measure_Evaluation(0, 0, Artificial_data, Artificial_Eva, inertia ,y_per)
Measure_Evaluation(0, 1,Spiral_data, Spiral_Eva, inertia, y_per)
Measure Evaluation(0, 2, Path data, Path Eva, inertia, y per)
Artificial_Eva.head()
# Spiral_Eva.head()
# Path Eva.head()
```







Out[3]:	Clustring Algorithm	Root MSE	DBI	Silhouette Score	Purity	Square Error
0	K-Means	2.719711	0.774641	0.563799	0.957341	0.0
1	Mini-Batch K-Means	0.000000	0.000000	0.000000	0.000000	0.0
2	DBSCAN	0.000000	0.000000	0.000000	0.000000	0.0
3	Spectral Clustering	0.000000	0.000000	0.000000	0.000000	0.0
4	K means self implemenation	0.000000	0.000000	0.000000	0.000000	0.0

```
In [4]: inertia
```

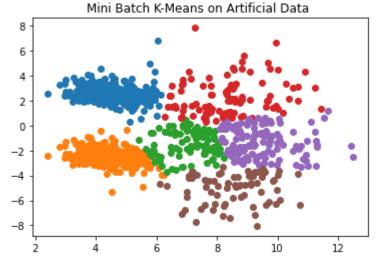
Out[4]: [0, 0, 0]

Mini-Batch K-Means

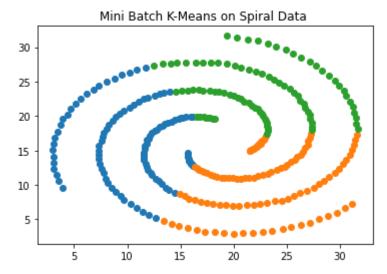
```
In [5]:
         # Mini batch k-means clustering
         from numpy import unique
         from numpy import where
         import numpy as np
         from sklearn.cluster import MiniBatchKMeans
         from matplotlib import pyplot as plt
         inertia= []
         y_per= []
         def Mini_Batch_KMeans ( msg, cluster, data= []):
             model = MiniBatchKMeans(n_clusters=cluster, max_iter=1000)
              # fit the model
             mini_batch_kmeans=model.fit(data)
             inertia.append(mini_batch_kmeans.inertia_)
             yhat = model.predict(data)
             y per.append(yhat)
             # retrieve unique clusters
             clusters = unique(yhat)
             # create scatter plot for samples from each cluster
             for cluster in clusters:
```

```
# get row indexes for samples with this cluster
        row ix = where(yhat == cluster)
        # create scatter of these samples
        plt.scatter(data[row_ix, 0], data[row_ix, 1])
    # show the plot
    plt.title(msg);
    plt.show()
Artificial_data,Spiral_data,Path_data= Load_Data(1)
Mini Batch KMeans ("Mini Batch K-Means on Artificial Data", 6, Artificial data)
Mini Batch KMeans ( "Mini Batch K-Means on Spiral Data ", 3, Spiral data)
Mini_Batch_KMeans( "Mini Batch K-means on Path Data ",5, Path_data)
# Evaluations
inertia=[0,0,0]
Artificial data, Spiral data, Path data = Load Data(0)
Measure_Evaluation(1, 0,Artificial_data, Artificial_Eva, inertia ,y_per)
Measure Evaluation(1, 1, Spiral data, Spiral Eva, inertia, y per)
Measure Evaluation(1, 2, Path data, Path Eva, inertia, y per)
Artificial Eva.head()
# Spiral Eva.head()
# Path Eva.head()
```

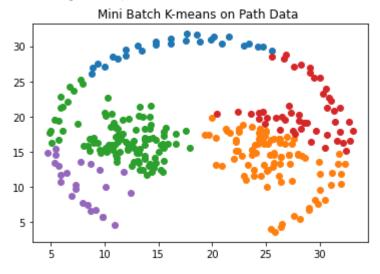
C:\ProgramData\Anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:887: UserWarning:
MiniBatchKMeans is known to have a memory leak on Windows with MKL, when there are less
chunks than available threads. You can prevent it by setting batch_size >= 1024 or by se
tting the environment variable OMP_NUM_THREADS=1
 warnings.warn(



C:\ProgramData\Anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:887: UserWarning:
MiniBatchKMeans is known to have a memory leak on Windows with MKL, when there are less
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C:\ProgramData\Anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:887: UserWarning:
MiniBatchKMeans is known to have a memory leak on Windows with MKL, when there are less
chunks than available threads. You can prevent it by setting batch_size >= 1024 or by se
tting the environment variable OMP_NUM_THREADS=1
 warnings.warn(



Out[5]:	Clustring Algorithm	Root MSE	DBI	Silhouette Score	Purity	Square Error
0	K-Means	2.719711	0.774641	0.563799	0.957341	0.0
1	Mini-Batch K-Means	2.838231	0.820054	0.519345	0.963294	0.0
2	DBSCAN	0.000000	0.000000	0.000000	0.000000	0.0
3	Spectral Clustering	0.000000	0.000000	0.000000	0.000000	0.0
4	K means self implemenation	0.000000	0.000000	0.000000	0.000000	0.0

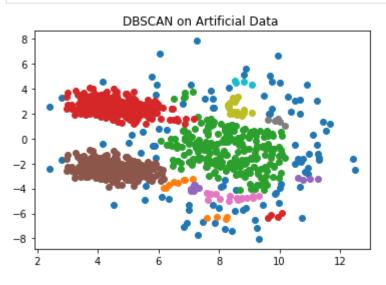
DBSCAN

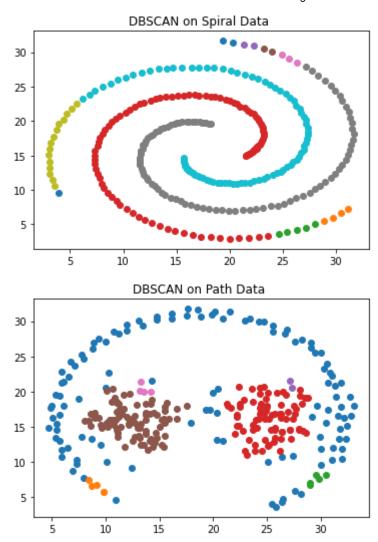
In [6]:

Mini batch k-means clustering
from numpy import unique
from numpy import where
import numpy as np
from sklearn.cluster import DBSCAN

localhost:8888/nbconvert/html/AssignmentMachineLearning.ipynb?download=false

```
from matplotlib import pyplot as plt
test=0
y_per= []
def DbScan ( msg,window size ,sample, data= []):
    model = DBSCAN(eps=window size, min samples=(sample))
     # fit the model
    test=model.fit(data)
    yhat = model.fit predict(data)
    y per.append(yhat)
    # retrieve unique clusters
    clusters = unique(yhat)
    # create scatter plot for samples from each cluster
    for cluster in clusters:
        # get row indexes for samples with this cluster
        row ix = where(yhat == cluster)
        # create scatter of these samples
        plt.scatter(data[row_ix, 0], data[row_ix, 1])
    # show the plot
    plt.title(msg);
    plt.show()
Artificial_data,Spiral_data,Path_data= Load_Data(1)
DbScan( "DBSCAN on Artificial Data ", 0.4,4, Artificial_data)
DbScan( "DBSCAN on Spiral Data ", 0.99999999,1.5, Spiral_data)
DbScan( "DBSCAN on Path Data ",1.333,4.1, Path_data)
# Evaluations
inertia= [0,0,0]
Artificial_data,Spiral_data,Path_data= Load_Data(0)
Measure Evaluation(2, 0, Artificial data, Artificial Eva, inertia, y per)
Measure_Evaluation(2, 1,Spiral_data, Spiral_Eva, inertia, y_per)
Measure Evaluation(2, 2, Path data, Path Eva, inertia, y per)
Artificial_Eva.head()
# Spiral Eva.head()
# Path Eva.head()
```





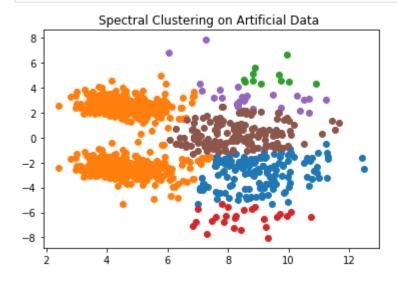
Out[6]:		Clustring Algorithm	Root MSE	DBI	Silhouette Score	Purity	Square Error
	0	K-Means	2.719711	0.774641	0.563799	0.957341	0.0
	1	Mini-Batch K-Means	2.838231	0.820054	0.519345	0.963294	0.0
	2	DBSCAN	2.559025	5.331623	0.255895	0.962302	0.0
	3	Spectral Clustering	0.000000	0.000000	0.000000	0.000000	0.0
	4	K means self implemenation	0.000000	0.000000	0.000000	0.000000	0.0

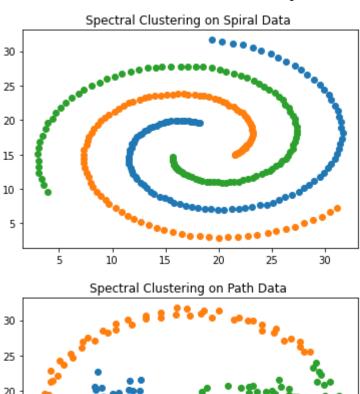
Spectral Clustering

```
In [7]: # Mini batch k-means clustering
    from numpy import unique
    from numpy import where
    import numpy as np
    from sklearn.cluster import SpectralClustering
    from matplotlib import pyplot as plt

y_per= []
    def Spectral_Clustering ( msg,cluster, data= []):
        model = SpectralClustering(n_clusters=cluster)
```

```
# fit model and predict clusters
    yhat = model.fit predict(data)
    y_per.append(yhat)
    # retrieve unique clusters
    clusters = unique(yhat)
    # create scatter plot for samples from each cluster
    for cluster in clusters:
        # get row indexes for samples with this cluster
        row ix = where(yhat == cluster)
        # create scatter of these samples
        plt.scatter(data[row_ix, 0], data[row_ix, 1])
    # show the plot
    plt.title(msg);
    plt.show()
# Load Data
Artificial data, Spiral data, Path data = Load Data(1)
Spectral_Clustering( "Spectral Clustering on Artificial Data ", 6, Artificial_data)
Spectral_Clustering( "Spectral Clustering on Spiral Data ",3,Spiral_data)
Spectral Clustering ("Spectral Clustering on Path Data", 3, Path data)
# Evaluations
inertia= [0,0,0]
Artificial data, Spiral data, Path data = Load Data(0)
Measure_Evaluation(3, 0,Artificial_data, Artificial_Eva, inertia ,y_per)
Measure Evaluation(3, 1, Spiral data, Spiral Eva, inertia, y per)
Measure_Evaluation(3, 2,Path_data, Path_Eva, inertia, y_per)
Artificial Eva.head()
# Spiral_Eva.head()
# Path Eva.head()
```





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	5 -		**			•	•••	
		5	10	15	20	25	30	_
Out[7]:		(Clustring A	lgorithm	Root MSE		DBI Silhoue	ett

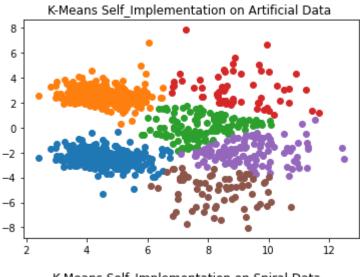
Out[7]:	Clustring Algorithm	Root MSE	DBI	Silhouette Score	Purity	Square Error
0	K-Means	2.719711	0.774641	0.563799	0.957341	0.0
1	Mini-Batch K-Means	2.838231	0.820054	0.519345	0.963294	0.0
2	DBSCAN	2.559025	5.331623	0.255895	0.962302	0.0
3	Spectral Clustering	2.432624	1.089437	0.273916	0.648810	0.0
4	K means self implemenation	0.000000	0.000000	0.000000	0.000000	0.0

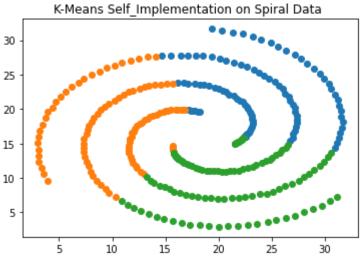
In []:

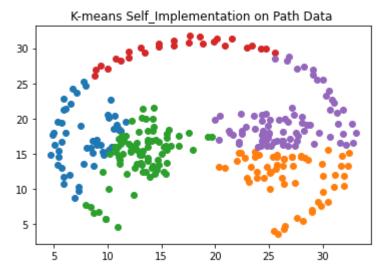
K means self implemenation

```
In [8]: #Loading the required modules
import pandas as pd
import numpy as np
from scipy.spatial.distance import cdist
from sklearn.datasets import load_digits
from sklearn.decomposition import PCA
from sklearn.cluster import KMeans
```

```
import matplotlib.pyplot as plt
# define dataset
inertia= []
y_per= []
def Kmeans Self Implementation ( msg, k, no of iterations, data= []):
    idx = np.random.choice(len(data), k, replace=False)
    #Randomly choosing Centroids
    centroids = data[idx, :] #Step 1
    distance2=0
    #finding the distance between centroids and all the data points
    distances = cdist(data, centroids ,'euclidean') #Step 2
    #Centroid with the minimum Distance
    yhat = np.array([np.argmin(i) for i in distances]) #Step 3
    #Repeating the above steps for a defined number of iterations
    for _ in range(no_of_iterations):
        centroids = []
        for idx in range(k):
            #Updating Centroids by taking mean of Cluster it belongs to
            temp cent = data[yhat==idx].mean(axis=0)
            centroids.append(temp cent)
        centroids = np.vstack(centroids) #Updated Centroids
        distances = cdist(data, centroids ,'euclidean')
        yhat = np.array([np.argmin(i) for i in distances])
    y per.append(yhat)
    inertia.append(distances)
    clusters = unique(yhat)
    # create scatter plot for samples from each cluster
    for cluster in clusters:
        # get row indexes for samples with this cluster
        row ix = where(yhat == cluster)
        # create scatter of these samples
        plt.scatter(data[row ix, 0], data[row ix, 1])
    # show the plot
    plt.title(msg);
    plt.show()
Artificial data, Spiral data, Path data = Load Data(1)
Kmeans Self Implementation("K-Means Self Implementation on Artificial Data ", 6,1000, A
Kmeans_Self_Implementation("K-Means Self_Implementation on Spiral Data ", 3,1000, Spira
Kmeans Self Implementation("K-means Self Implementation on Path Data ",5, 1000, Path da
# Evaluations
inertia[0]= 0
inertia[1]= 0
inertia[2]= 0
Artificial data, Spiral data, Path data = Load Data(0)
Measure Evaluation(4, 0, Artificial data, Artificial Eva, inertia, y per)
Measure Evaluation(4, 1, Spiral data, Spiral Eva, inertia, y per)
Measure_Evaluation(4, 2,Path_data, Path_Eva, inertia, y_per)
Artificial Eva.head()
# Spiral_Eva.head()
# Path Eva.head()
```



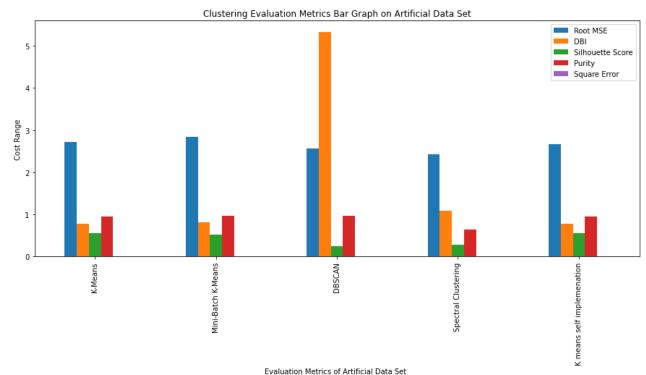




Out[8]:		Clustring Algorithm	Root MSE	DBI	Silhouette Score	Purity	Square Error
	0	K-Means	2.719711	0.774641	0.563799	0.957341	0.0
	1	Mini-Batch K-Means	2.838231	0.820054	0.519345	0.963294	0.0
	2	DBSCAN	2.559025	5.331623	0.255895	0.962302	0.0
	3	Spectral Clustering	2.432624	1.089437	0.273916	0.648810	0.0
	4	K means self implemenation	2.659216	0.779103	0.562926	0.956349	0.0

Result for Artificial data set

```
In [12]:
    from matplotlib import pyplot as plt
    Artificial_Eva.set_index('Clustring Algorithm').plot(kind = 'bar')
    plt.title("Clustering Evaluation Metrics Bar Graph on Artificial Data Set")
    plt.xlabel("Evaluation Metrics of Artificial Data Set ")
    plt.ylabel("Cost Range")
    plt.rcParams["figure.figsize"] = (15,6)
    plt.show()
    Artificial_Eva.head()
```

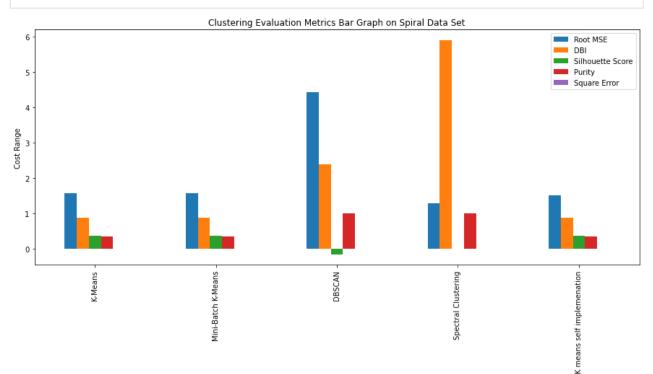


Out[12]:		Clustring Algorithm	Root MSE	DBI	Silhouette Score	Purity	Square Error
	0	K-Means	2.719711	0.774641	0.563799	0.957341	0.0
	1	Mini-Batch K-Means	2.838231	0.820054	0.519345	0.963294	0.0
	2	DBSCAN	2.559025	5.331623	0.255895	0.962302	0.0
	3	Spectral Clustering	2.432624	1.089437	0.273916	0.648810	0.0
	4	K means self implemenation	2.659216	0.779103	0.562926	0.956349	0.0

Result for Spiral data set

```
In [10]:
    from matplotlib import pyplot as plt
    Spiral_Eva.set_index('Clustring Algorithm').plot(kind = 'bar')
    plt.title("Clustering Evaluation Metrics Bar Graph on Spiral Data Set")
    plt.xlabel("Evaluation Metrics of Spiral Data Set ")
    plt.ylabel("Cost Range")
    plt.rcParams["figure.figsize"] = (15,6)
```

plt.show()
Spiral_Eva.head()

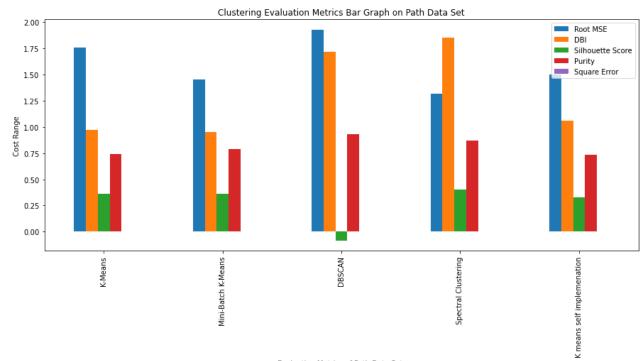


Evaluation	Metrics	of Spira	al Data Set

Out[10]:		Clustring Algorithm	Root MSE	DBI	Silhouette Score	Purity	Square Error
	0	K-Means	1.563243	0.886698	0.361981	0.350482	0.0
	1	Mini-Batch K-Means	1.565299	0.879688	0.359801	0.344051	0.0
	2	DBSCAN	4.436765	2.378795	-0.156053	0.996785	0.0
	3	Spectral Clustering	1.294311	5.899294	0.000785	1.000000	0.0
	4	K means self implemenation	1.515196	0.885007	0.361372	0.347267	0.0

Result for Path data set

```
In [11]:
    from matplotlib import pyplot as plt
    Path_Eva.set_index('Clustring Algorithm').plot(kind = 'bar')
    plt.title("Clustering Evaluation Metrics Bar Graph on Path Data Set")
    plt.xlabel("Evaluation Metrics of Path Data Set ")
    plt.ylabel("Cost Range")
    plt.rcParams["figure.figsize"] = (15,6)
    plt.show()
    Path_Eva.head()
```



Evaluation Metrics of Path Data Set

Out[11]:		Clustring Algorithm	Root MSE	DBI	Silhouette Score	Purity	Square Error
	0	K-Means	1.759827	0.971655	0.363569	0.742475	0.0
	1	Mini-Batch K-Means	1.450406	0.951018	0.359900	0.789298	0.0
	2	DBSCAN	1.926752	1.718659	-0.083031	0.933110	0.0
	3	Spectral Clustering	1.318761	1.849967	0.401102	0.869565	0.0
	4	K means self implemenation	1.502506	1.061061	0.329675	0.732441	0.0

In []: