# ScalableKMeansPlusPlus

April 30, 2015

### 1 Scalable K-means++

### 1.1 Outline of Background

This project is an implementation based on Bahmani, Moseley, Vattani, Kumar and Vassilvitskii's paper  $Scalable\ K\ Means++$  in 2012.

- 1. **K-means** remains one of the most popular data processing algorithms. This algorithm has been used in many fields such us machine learning, pattern recognition and bioinformatics. However, the original k-means algorithm with random initialization has lots of weakness. For example.
  - A proper initialization is crucial for reciveing successful results.
  - For large dataset, it may take long time to achive convergent.
- 2. **K-means++** algorithm achived the goal of finding proper initialization, k-means++ has a deterministic initialization process, however, with downside of its inherent sequentical nature, which limits its efficiency (O(n)) and applicability to big massive datasets.
- 3. k-means||, aka Scalable k-means++, which proposed by this paper, oversamples by sampling each points independently with a larger probability, which is intuitively equivalent to updating the distribution much less frequently, with efficiency ( $O(\log n)^*$ ), which forms k-means++ in both sequential and parallel settings.

In this project, I'll implement the  $Scalable\ K\ Means++$  algorithm, and compare it to the general  $K\ Means$  and  $K\ Means++$ .

### 1.2 Algorithm / Pseudocode

#### 1.2.1 Notations

Let  $X = \{x_1, ... x_n\}$  be the set of points in d-dimensional Euclidean space, and let k be a positive integer specifying the number of clusters. Let  $||x_i - x_j||$  denote the Euclidean distance between  $x_i$  and  $x_j$ . For a point x and a subset  $Y \subseteq X$  of points, the distance is defined as  $d(x, Y) = min_{y \in Y} ||x - y||$ . For a subset  $Y \subseteq X$  of points, let its centroid be given by

$$\operatorname{centroid}(Y) = \frac{1}{|Y|} \sum_{y \in Y} y$$

Let  $C = \{c_1, ..., c_k\}$  be the set of points and let  $Y \subseteq X$ . We define the cost of Y with respect to C as

$$\phi_Y(C) = \sum_{y \in Y} d^2(y, C) = \sum_{y \in Y} \min_{i=1,\dots,k} ||y - c_i||^2$$

### 1.2.2 k-means++(k) initialization

- 1.  $C \leftarrow$  sample a point uniformlt at random from X
- 2. while |C| < k do
- Sample  $x \in X$  with probability  $\frac{d^2(x,C)}{\phi_X(C)}$
- $C \leftarrow C \cup \{x\}$
- 3. end while

### 1.2.3 k-means||(k, l)| initialization

- 1.  $C \leftarrow$  sample a point uniformly at random from X
- 2.  $\psi \leftarrow \phi_X(C)$
- 3. for  $O(\log \psi)$  times do
- $C' \leftarrow$  sample each point  $x \in X$  independently with probability  $p_x = \frac{l \cdot d^2(x,C)}{\phi_X(C)}$
- $C \leftarrow C \cup C'$
- 4. end for
- 5. For  $x \in C$ , set  $w_x$  to be the number of points in X closer to x than any point in C
- 6. Recluster the weighted points in C into k clusters

### 1.3 Draft of unit test

### Will verify the code correctness using following tests:

- 1. Test the Cost function using examples in terms of:
- non-negative
- data = c
- the size of c
- 2. Test Sampling Probability function using examples in terms of:
- $0 \le Pi \le 1$
- sum = 1
- 3. Test K Means Plus Plus Parallel Probability function using examples in terms of:
- feasiblility
- labels attributes

### 1.4 Data simulation

Simulating sample data from three bivariate normal distribution.

$$\begin{pmatrix} x_1 \\ x_2 \end{pmatrix} \sim N_2 \begin{pmatrix} \begin{pmatrix} 0 \\ 1 \end{pmatrix}, \begin{pmatrix} 1 & 0.5 \\ 0 & 2 \end{pmatrix} \end{pmatrix}$$

$$\begin{pmatrix} y_1 \\ y_2 \end{pmatrix} \sim N_2 \begin{pmatrix} \begin{pmatrix} 3 \\ 7 \end{pmatrix}, \begin{pmatrix} 1 & 0.33 \\ 0.33 & 1 \end{pmatrix} \end{pmatrix}$$

$$\begin{pmatrix} z_1 \\ z_2 \end{pmatrix} \sim N_2 \begin{pmatrix} \begin{pmatrix} -8 \\ 2 \end{pmatrix}, \begin{pmatrix} 2 & 0.66 \\ 0.66 & 2 \end{pmatrix} \end{pmatrix}$$

```
In [48]: # Prepare
         #!/usr/bin/python
         from __future__ import division
         import os
         import sys
         import glob
         import random
         import sklearn
         import sklearn.cluster
         import numpy as np
         import pandas as pd
         import matplotlib.pyplot as plt
         from matplotlib.figure import Figure
         from matplotlib.axes import Subplot
         %matplotlib inline
         plt.style.use('ggplot')
In [49]: # Simulated "Real" Data Set
         class SimulatedData:
             def __init__(self, n, data):
                 self.data = data
                 self.n = n
             def DataSimulation(n):
                 mean1 = np.array([0, 1])
                 cov1 = np.array([[1, 0.5], [0.5, 2]])
                 mean2 = np.array([3, 7])
                 cov2 = np.array([[1, 0.33], [0.33, 1]])
                 mean3 = np.array([-8, 2])
                 cov3 = np.array([[2, 0.66], [0.66, 2]])
                 tmp = np.vstack((np.random.multivariate_normal(mean1, cov1, n),
                                   np.random.multivariate_normal(mean2, cov2, n),
                                   np.random.multivariate_normal(mean3, cov3, n)))
                 data = tmp[np.random.choice(range(3*n),size = 3*n, replace=False),]
                 return data
             def AsDataFrame(data):
                 df = pd.DataFrame(data,columns=["X","Y"])
                 return df
In [50]: data = SimulatedData.DataSimulation(5)
         data
Out[50]: array([[ 3.27244498,
                                 6.19079547],
                [ 2.83259403,
                                 5.31488359],
                [ -6.29887528,
                                 1.15665735],
                [ 2.26433884,
                                 4.68468001],
                [ 0.28221065,
                                 1.48472561],
                [ 4.05630379,
                                 6.34592495],
                [ 4.16797629,
                                 7.93527882],
                [-10.70344788, 1.63873811],
                [-1.34911855, -2.49537396],
```

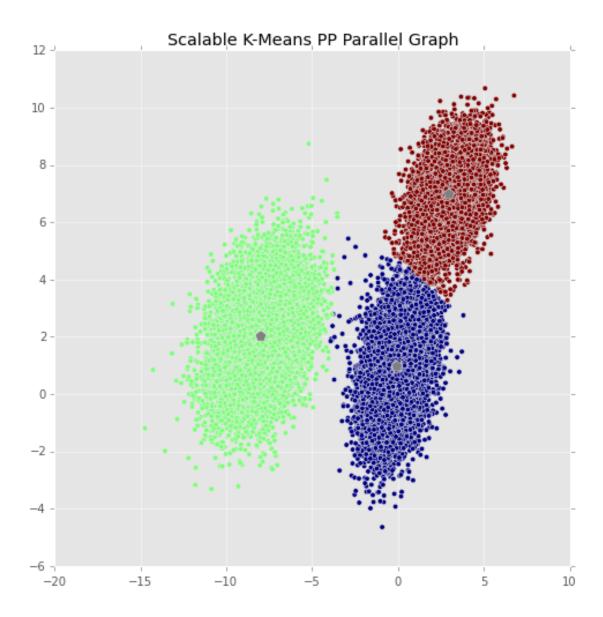
```
[ 1.52048191, 1.62214511],
[ -9.279794 , 1.20963785],
[ 1.07200651, 2.15463174],
[ -9.58562466, 0.61057195],
[ -5.51352065, 3.68254202],
[ 0.84596127, 2.51050088]])
```

### 1.5 K-Means || Code

#### 1.5.1 Navïe Version

```
In [51]: class ScalableKMeansPP:
             def __init__(self, data, k, 1):
                 self.data = data
                 self.k = k
                 self.1 = 1
             def KMeansParallel(data, k, 1):
                 N = data.__len__()
                 if k <= 0 or not(isinstance(k,int)) or 1 <= 0:</pre>
                     sys.exit()
                 # Then we start to Implement the algorithm
                 # 1. Sample one point uniformly at random from X
                 c = np.array(data[np.random.choice(range(N),1),])
                 # 2. To Cost function
                 phi = ScalableKMeansPP.CostFunction(c, data)
                 # 3. Looping
                 for i in range(np.ceil(np.log(phi)).astype(int)):
                     cPrime = data[ScalableKMeansPP.SamplingProbability(c,data,1) > np.random.uniform(s
                     c = np.concatenate((c, cPrime))
                 # End looping
                 # 7. For x in C, set w_x to be the number of pts closest to X
                 cMini = [np.argmin(np.sum((c-pts)**2,axis=1)) for pts in data];
                 closerPts = [cMini.count(i) for i in range(len(c))]
                 weight = closerPts/np.sum(closerPts)
                 # 8. Recluster the weighted points in C into k clusters
                 allC = data[np.random.choice(range(len(c)), size=1, p=weight),]
                 data_final = c
                 for i in range(k-1):
                     Probability = ScalableKMeansPP.SamplingProbability(allC,data_final,1) * weight
                     # choose next centroid
                     cPrimeFin = data[np.random.choice(range(len(c)), size=1, p=Probability/np.sum(Prob
                     allC = np.concatenate((allC,cPrimeFin))
                 KMeansPP = sklearn.cluster.KMeans(n_clusters=k, n_init=1, init=allC, max_iter=500, tol-
                 KMeansPP.fit(data);
                 return KMeansPP
             def SamplingProbability(c,data,l):
                 cost = ScalableKMeansPP.CostFunction(c,data)
                 return np.array([(min(np.sum((c-pts)**2,axis=1))) * 1 / cost for pts in data])
             def CostFunction(c,data):
                 return np.sum([min(np.sum((c-pts)**2,axis=1)) for pts in data])
```

```
In [52]: data = SimulatedData.DataSimulation(10000)
         ScalableKMeansPP.KMeansParallel(data,3,6)
Out[52]: KMeans(copy_x=True,
             init=array([[ 3.43704, 5.82661],
                [ 1.19989, 6.19787],
                [ 2.45681, 6.21065]]),
             max_iter=500, n_clusters=3, n_init=1, n_jobs=1,
             precompute_distances=True, random_state=None, tol=0.0001, verbose=0)
In [69]: k = 3
        start = timeit.default_timer()
        KMeansPP = ScalableKMeansPP.KMeansParallel(data=data, k=k, l=2*k); # paper suggesting using l=
        elapsed = timeit.default_timer() - start
        print(elapsed)
        df = SimulatedData.AsDataFrame(data)
        plt.figure(figsize=(8, 8));
        plt.scatter(df.X,df.Y,c=KMeansPP.labels_);
        plt.scatter(KMeansPP.cluster_centers_[:,0],KMeansPP.cluster_centers_[:,1], c='grey', s=100, ma
        plt.title("Scalable K-Means PP Parallel Graph");
        plt.show()
```



### 1.6 Algorithm Comparison

For algorithm comparison, I will using different sizes of same sample data to clustering using original k-means, k-means++, k-means||. Test their result and compare the efficiency.

We use "K-Means" and "K-Means++" methods from sklearn.cluster.KMeans package.

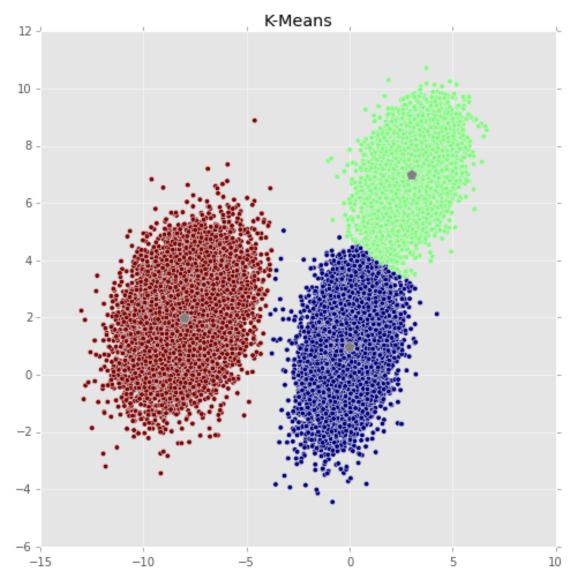
### 1.6.1 K-Means

```
In [75]: import timeit
   data1 = SimulatedData.DataSimulation(10000)
   start = timeit.default_timer()
   k = 3
   KMeans = sklearn.cluster.KMeans(n_clusters=3, init='random', n_init=10, max_iter=500, tol=0.00
   KMeans.fit(data1);
```

```
elapsed = timeit.default_timer() - start
print(elapsed)
```

### 0.12828452099347487

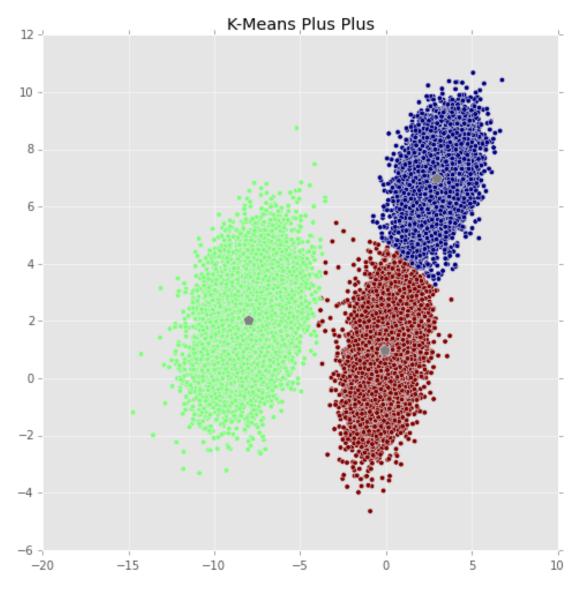
```
In [87]: KMeans.fit(data);
    df = SimulatedData.AsDataFrame(data)
    plt.figure(figsize=(8, 8));
    plt.scatter(df.X,df.Y,c=KMeans.labels_);
    plt.scatter(KMeans.cluster_centers_[:,0],KMeans.cluster_centers_[:,1], c='grey', s=100, marker
    plt.title("K-Means");
    plt.show()
```



### 1.6.2 K-Means++

```
k = 3
KMeansPlusPlus = sklearn.cluster.KMeans(n_clusters=3, init='k-means++', n_init=10, max_iter=50
KMeansPlusPlus.fit(data1);
elapsed = timeit.default_timer() - start
print(elapsed)
```

#### 0.08849192800698802



From the result of timeit, we can concluded that K-Means++ is much efficienct than the base K-Means. However, it seems that the navïe version of Scalable K-Means++ are slower than K-Means++, or even the base K-Means.

The reason would be, the package algorithms using parallel. I will try to optimization my code.

### 1.7 Optimization Strategies

- 1. Using alternative method to replace inefficent code in Python
- 2. Will try on large datasets, if still take too long. Could try to use other languages to write the looping part

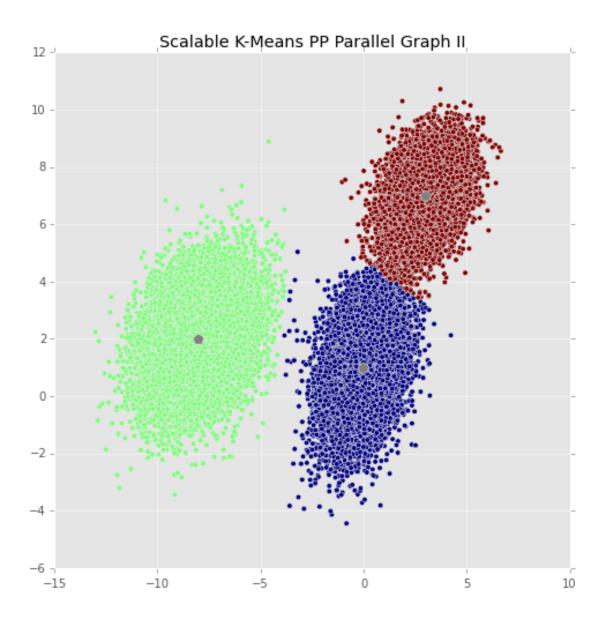
### 1.8 Optimization: Vectorized Version

```
In [79]: class VectorizedScalableKMeansPP(ScalableKMeansPP):
             def __init__(self, data, k, 1):
                 ScalableKMeansPP.__init__(self, data, k, 1)
             def KMeansParallel(data, k, 1):
                 N = data.__len__()
                 # 1. Sample one point uniformly at random from X
                 c = data[np.random.choice(range(data.shape[0]),1), :]
                 data__ = data[:,np.newaxis,:]
                 # 2. To Cost function
                 phi = ScalableKMeansPP.CostFunction(c, data)
                 # 3. Looping
                 for i in range(np.ceil(np.log(phi)).astype(int)):
                     d2 = (data_{-} - c) ** 2
                     distance = np.sum(d2, axis=2)
                     cMini = np.zeros(distance.shape)
                     cMini[range(distance.shape[0]), np.argmin(distance, axis=1)] = 1
                     min_dist = distance[cMini == 1]
                     phi = np.sum(min_dist)
                     for i, cPrime in enumerate(data):
                         Probability = l*min_dist[i]/phi
                         u = np.random.uniform(0,1)
                         if Probability >= u:
                             c = np.vstack([c, cPrime])
                 # End looping
                 # 7. For x in C, set w_x to be the number of pts closest to X
                 d2 = (data_{-} - c) ** 2
                 distance = np.sum(d2, axis=2)
                 cMini = np.zeros(distance.shape)
                 cMini[range(distance.shape[0]), np.argmin(distance, axis=1)] = 1
                 weight = np.array([np.count_nonzero(cMini[:, i]) for i in range(c.shape[0])]).reshape(
                 # 8. Recluster the weighted points in C into k clusters
                 allC = c[np.random.choice(range(c.shape[0]),1), :]
                 data_final = c
                 index = np.where(data_final==allC)[0]
                 data_final = np.delete(data_final,index[0],axis=0)
                 weight = np.delete(weight,index[0])
                 for i in range(k-1):
                     Probability = ScalableKMeansPP.SamplingProbability(allC,data_final,1) * weight
                     # choose next centroid
```

c = data\_final[np.random.choice(range(data\_final.shape[0]),size=1, p=Probability/n

```
index = np.where(data_final==c)[0]
                     allC = np.vstack([allC, c])
                     #Remove the selected center and its corresponding weight
                     data_final = np.delete(data_final,index[0],axis=0)
                     weight = np.delete(weight,index[0])
                 vKMeansPP = sklearn.cluster.KMeans(n_clusters=k, n_init=1, init=allC, max_iter=500, to
                 vKMeansPP.fit(data)
                 return vKMeansPP
In [80]: data = SimulatedData.DataSimulation(10000)
         VectorizedScalableKMeansPP.KMeansParallel(data,3,6)
Out[80]: KMeans(copy_x=True,
             init=array([[-2.33086, 0.71013],
                [ 3.51431, 7.72725],
                [-8.37233, 3.60263]]),
             max_iter=500, n_clusters=3, n_init=1, n_jobs=1,
             precompute_distances=True, random_state=None, tol=0.0001, verbose=0)
In [85]: start = timeit.default_timer()
         KMeansPP2 = VectorizedScalableKMeansPP.KMeansParallel(data=data, k=k, 1=2*k);
         elapsed = timeit.default_timer() - start
         print(elapsed)
         df = SimulatedData.AsDataFrame(data)
         plt.figure(figsize=(8, 8));
         plt.scatter(df.X,df.Y,c=KMeansPP2.labels_);
         plt.scatter(KMeansPP2.cluster_centers_[:,0], KMeansPP2.cluster_centers_[:,1], c='grey', s=100, s
         plt.title("Scalable K-Means PP Parallel Graph II");
         plt.show()
```

### 1.8935244909953326



## 1.9 Efficiency Comparsion: Profiling

```
In [ ]: ! pip install --pre line-profiler &> /dev/null
     ! pip install psutil &> /dev/null
     ! pip install memory_profiler &> /dev/null
```

In [18]: %load\_ext line\_profiler

In [ ]: %lprun -f ScalableKMeansPP.KMeansParallel ScalableKMeansPP.KMeansParallel(data = data, k=3, 1 =

In []: Timer unit: 1e-06 s

Total time: 13.3943 s

File: <ipython-input-125-d01bb3e3c9ef> Function: KMeansParallel at line 7

Line #	$\mathit{Hits}$	Time	Per Hit	% Time	Line Contents
7					def KMeansParallel(data, k, 1):
8	1	4	4.0	0.0	<pre>N = datalen()</pre>
9	1	2	2.0	0.0	<pre>if k &lt;= 0 or not(isinstance(k,int)) or</pre>
10					sys.exit()
11					# Then we start to Implement the algor
12					# 1. Sample one point uniformly at ran
13	1	3501	3501.0	0.0	c = np.array(data[np.random.choice(rang
14					# 2. To Cost function
15	1	270143	270143.0	2.0	<pre>phi = ScalableKMeansPP.CostFunction(c,</pre>
16					# 3. Looping
17	16	56	3.5	0.0	for i in range(np.ceil(np.log(phi)).as
18	15	12559007	837267.1	93.8	cPrime = data[ScalableKMeansPP.Sam
19	15	127	8.5	0.0	<pre>c = np.concatenate((c, cPrime))</pre>
20					# End looping
21					# 7. For $x$ in $C$ , set $w_x$ to be the num
22	1	369043	369043.0	2.8	<pre>cMini = [np.argmin(np.sum((c-pts)**2,a</pre>
23	1	174695	174695.0	1.3	<pre>closerPts = [cMini.count(i) for i in r</pre>
24	1	114	114.0	0.0	<pre>weight = closerPts/np.sum(closerPts)</pre>
25					# 8. Recluster the weighted points in
26	1	112	112.0	0.0	<pre>allC = data[np.random.choice(range(len</pre>
27	1	1	1.0	0.0	<pre>data_final = c</pre>
28	3	5	1.7	0.0	for i in range(k-1):
29	2	3734	1867.0	0.0	Probability = ScalableKMeansPP.Sam
30					# choose next centroid
31	2	278	139.0	0.0	cPrimeFin = data[np.random.choice(
32	2	13	6.5	0.0	allC = np.concatenate((allC,cPrime)
33	1	23	23.0	0.0	<pre>KMeansPP = sklearn.cluster.KMeans(n_cl</pre>
34	1	13424	13424.0	0.1	<pre>KMeansPP.fit(data);</pre>
35	1	2	2.0	0.0	return KMeansPP

In [ ]: Timer unit: 1e-06 s

Total time: 3.66441 s

File: <ipython-input-114-00df5cba66a8>
Function: KMeansParallel at line 5

Line #	Hits	Time	Per Hit	% Time	Line Contents
5					def KMeansParallel(data, k, 1):
6	1	7	7.0	0.0	<pre>N = datalen()</pre>
7					# 1. Sample one point uniformly at ran
8	1	3579	3579.0	0.1	<pre>c = data[np.random.choice(range(data.sl</pre>
9	1	4	4.0	0.0	<pre>data2 = data[:,np.newaxis,:]</pre>
10					# 2. To Cost function
11	1	264291	264291.0	7.2	<pre>phi = ScalableKMeansPP.CostFunction(c,</pre>
12					# 3. Looping
13	16	49	3.1	0.0	for i in range(np.ceil(np.log(phi)).as
14	15	505753	33716.9	13.8	d2 = (data2 - c) ** 2
15	15	254088	16939.2	6.9	<pre>distance = np.sum(d2, axis=2)</pre>
16	15	36050	2403.3	1.0	cMini = np.zeros(distance.shape)

17	15	86425	5761.7	2.4	<pre>cMini[range(distance.shape[0]), np</pre>
18	15	33310	2220.7	0.9	<pre>min_dist = distance[cMini == 1]</pre>
19	15	796	53.1	0.0	<pre>phi = np.sum(min_dist)</pre>
20	450015	577435	1.3	15.8	<pre>for i, cPrime in enumerate(data):</pre>
21	450000	646465	1.4	17.6	Probability = l*min_dist[i]/ph
22	450000	597961	1.3	16.3	<pre>u = np.random.uniform(0,1)</pre>
23	450000	508532	1.1	13.9	<pre>if Probability &gt;= u:</pre>
24	75	2341	31.2	0.1	<pre>c = np.vstack([c, cPrime])</pre>
25					# End looping
26					# 7. For $x$ in $C$ , set $w_{-}x$ to be the num
27	1	67089	67089.0	1.8	d2 = (data2 - c) ** 2
28	1	32498	32498.0	0.9	<pre>distance = np.sum(d2, axis=2)</pre>
29	1	5074	5074.0	0.1	cMini = np.zeros(distance.shape)
30	1	6751	6751.0	0.2	cMini[range(distance.shape[0]), np.arg
31	1	15825	15825.0	0.4	<pre>weight = np.array([np.count_nonzero(cM:</pre>
32					# 8. Recluster the weighted points in
33	1	87	87.0	0.0	allC = c[np.random.choice(range(c.shape
34	1	2	2.0	0.0	<pre>data_final = c</pre>
35	1	16	16.0	0.0	<pre>index = np.where(data_final==allC)[0]</pre>
36	1	48	48.0	0.0	<pre>data_final = np.delete(data_final,inde:</pre>
37	1	32	32.0	0.0	<pre>weight = np.delete(weight,index[0])</pre>
38	3	4	1.3	0.0	for i in range(k-1):
39	2	2686	1343.0	0.1	Probability = ScalableKMeansPP.Sam
40					# choose next centroid
41	2	140	70.0	0.0	<pre>c = data_final[np.random.choice(ra</pre>
42	2	17	8.5	0.0	<pre>index = np.where(data_final==c)[0]</pre>
43	2	43	21.5	0.0	allC = np.vstack([allC, c])
44					#Remove the selected center and it.
45	2	62	31.0	0.0	data_final = np.delete(data_final,:
46	2	55	27.5	0.0	<pre>weight = np.delete(weight,index[0]]</pre>
47	1	20	20.0	0.0	vKMeansPP = sklearn.cluster.KMeans(n_c)
48	1	16874	16874.0	0.5	vKMeansPP.fit(data)
49	1	3	3.0	0.0	return vKMeansPP

By profiling, it is clearlt that Vectored KMeansParallel function is significantly faster than the navie KMeansParallel function. (move than three times faster) The total time was improved from 13.4s to 3.66s.