session4_pyspark_solution

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1 [MIRACUM 2019][Session 4] Solution

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- This is prepared for a turorial Data analysis tools (Datenanalysewerkzeuge)

1.0.1 Protocol:

1. Create a Spark Context 2. Import required libraries 3. Load the cholera death data 4. Create a KMeans model 5. Assign centroids to the data 6. Visualize the result

```
[1]: from pyspark import SparkContext
sc = SparkContext()
sc
```

[1]: <SparkContext master=local[*] appName=pyspark-shell>

1.0.2 Protocol:

1. Create a Spark Context 2. Import required libraries 3. Load the cholera death data 4. Create a KMeans model 5. Assign centroids to the data 6. Visualize the result

```
[2]: import pyspark.mllib.clustering
from pyspark.mllib.clustering import KMeans, KMeansModel
import pandas as pd
import numpy as np
import folium
from folium import plugins
import matplotlib.pyplot as plt
%matplotlib inline
```

1.0.3 Protocol:

1. Create a Spark Context 2. Import required libraries 3. Load the cholera death data 4. Create a KMeans model 5. Assign centroids to the data 6. Visualize the result

```
[3]: raw = sc.textFile("data/cholera_deaths.csv")
header = sc.parallelize([raw.first()])
raw = raw.subtract(header)
raw.take(2)
```

[3]: ['1,2,-0.137883,51.513361', '13,1,-0.138378,51.513169']

```
[4]: line = raw.map(lambda x: x.split(','))
line.take(2)
```

[4]: [['1', '2', '-0.137883', '51.513361'], ['13', '1', '-0.138378', '51.513169']]

```
[5]: geo = line.map(lambda x: (float(x[2]), float(x[3])))
geo.take(2)
```

[5]: [(-0.137883, 51.513361), (-0.138378, 51.513169)]

1.0.4 Protocol:

1. Create a Spark Context 2. Import required libraries 3. Load the cholera death data 4. Create a KMeans model with k=3 5. Assign centroids to the data 6. Visualize the result

```
[6]: k = 3
model = KMeans.train(geo, k)
```

1.0.5 Protocol:

1. Create a Spark Context 2. Import required libraries 3. Load the cholera death data 4. Create a KMeans model 5. Assign centroids to the data 6. Visualize the result

```
[7]: clusters = model.predict(geo) clusters.take(20)
```

[7]: [2, 2, 2, 2, 2, 2, 2, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 0]

1.0.6 Protocol:

1. Create a Spark Context 2. Import required libraries 3. Load the cholera death data 4. Create a KMeans model 5. Assign centroids to the data 6. Visualize the result

```
[8]: # data preparatoin for VIZ
lat = geo.map(lambda x: float(x[0])).collect()
lon = geo.map(lambda x: float(x[1])).collect()
clusters = clusters.collect()
```

```
df_list = list(map(lambda x,y,z:[x,y,z], lon,lat,clusters))
      df_list[:5]
 [8]: [[51.513361, -0.137883, 2],
       [51.513169, -0.138378, 2],
       [51.513402, -0.139045, 2],
       [51.513676, -0.138887, 2],
       [51.512885, -0.138624, 2]]
 [9]: # the latitude and Longitude coordinates of the Soho District center
      SOHO_COORDINATES = (51.513578, -0.136722)
      map_soho = folium.Map(location = SOHO_COORDINATES, width = "100%", zoom_start = "
      →15) # max zoom: 18
      folium.TileLayer('cartodbpositron').add_to(map_soho) # stamentoner
      map soho
 [9]: <folium.folium.Map at 0x7f6cc21590f0>
[10]: def my_palettes(cluster):
          if cluster == 0:
              return 'green'
          elif cluster == 1:
              return 'yellow'
          elif cluster == 2:
              return 'red'
          elif cluster == 3:
              return 'blue'
          elif cluster == 4:
              return 'orange'
          elif cluster == 5:
              return 'skyblue'
          elif cluster == 6:
              return 'gray'
          elif cluster == 7:
              return 'purple'
          else:
              return 'black'
[11]: for i in range(0, len(df_list)):
          folium.RegularPolygonMarker(location = df_list[i][:2], \
                                      stroke = False, \
                                      fill_color = my_palettes(df_list[i][2]), \
                                      fill_opacity = 0.5, number_of_sides = 12,
       →radius = 3
                                     ).add_to(map_soho)
      map_soho
```

[11]: <folium.folium.Map at 0x7f6cc21590f0>

[]: