山东大学 计算机科学与技术 学院

信息检索与数据挖掘 课程实验报告

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实验题目: 聚类方法与评估

实验内容:

调用 sklearn 的方法,实现

KMeans,AffinityPropagation,MeanShift,SpectralClustering,AgglomerativeClustering,DBSCAN,高丝混合等方法,并用 nmi 进行评估

结果如下

load the data...

method meanshift
compute the result...
NMI:0.65015

method DBSCAN
compute the result...
NMI:0.707871

method k_means compute the result... NMI:0.798497

method Spectral compute the result... NMI:0.681795

method ward compute the result... NMI:0.67917

method Agglomerative compute the result... NMI:0.908023

method AP compute the result... NMI:0.782623

method GaussianMixture

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compute the result...
NMI:0.813259
from sklearn.feature_extraction.text import
TfidfTransformer,CountVectorizer
from sklearn.cluster import
KMeans, AffinityPropagation, MeanShift, SpectralClustering, AgglomerativeClust
ering, DBSCAN
from sklearn.mixture import GMM
from sklearn.metrics.cluster import normalized_mutual_info_score as NMI
import sklearn
import json
import nltk
import re
# 去除停用词
def cutstopwords(str):
   stopwords = {}.fromkeys([line.rstrip() for line in
open('estopwords.txt')])
   segs = str.replace('\n','').lower().split(' ')
   new str = ''
   for seg in segs:
      if seg not in stopwords:
          new_str = new_str + " " +seg
   return new_str
# 去除标点
def cutsyms(str):
   new_str = re.sub('[,.\'\"\t\n*_+=?/|!@#$%^&*()`~<>:;\-\[\]]'," ",str)
   return new_str
# 词干提取
def stemming(str):
   s = nltk.stem.SnowballStemmer('english')
   segs = str.replace('\n', '').lower().split(' ')
   new_str = ''
   for seg in segs:
      new_str = new_str + " " + s.stem(seg)
   return new_str
path = '/Users/apple/Desktop/ir/hw5/Homework5Tweets.txt'
file = open(path,'r',encoding='UTF-8',errors='ignore')
tweets = []
cluster = []
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text = []
count = 0
print("load the data...\n")
for line in file:
   tweets.append(json.loads(line))
   cluster.append(tweets[count]['cluster'])
   tweets[count]['text'] = tweets[count]['text'].lower()
   tweets[count]['text'] = cutsyms(tweets[count]['text'])
   tweets[count]['text'] = cutstopwords(tweets[count]['text'])
   tweets[count]['text'] = stemming(tweets[count]['text'])
   text.append(tweets[count]['text'])
   count = count + 1
file.close()
tv = TfidfTransformer()
vec = CountVectorizer()
mv = tv.fit transform(vec.fit transform(text))
tm = mv.toarray()
def showresult(result,cluster):
   print("show the result:")
   for i in range(len(result)):
      print("cluster of text %d:%d. and its truecluster:%d" % (i + 1,
result[i], cluster[i]))
print("method meanshift")
ms = MeanShift(bandwidth=0.4,bin_seeding=True,min_bin_freq=2)
print("compute the result...")
result = ms.fit_predict(tm)
#showresult(result,cluster)
nmi=NMI(cluster, result)
print("NMI:%g\n"%(nmi))
print("method DBSCAN")
DB = DBSCAN(eps=0.7,min_samples=1)
print("compute the result...")
result = DB.fit predict(tm)
#showresult(result,cluster)
nmi=NMI(cluster,result)
print("NMI:%g\n"%(nmi))
print("method k means")
num cluster = 150
km = KMeans(n_clusters=num_cluster, max_iter=300, n_init=40, init='k-
```

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means++')
print("compute the result...")
result = km.fit_predict(mv)
#showresult(result,cluster)
nmi=NMI(cluster, result)
print("NMI:%g\n"%(nmi))
print("method Spectral")
sc = SpectralClustering(n_clusters=max(cluster))
print("compute the result...")
result = sc.fit predict(tm)
#showresult(result, cluster)
nmi=NMI(cluster,result)
print("NMI:%g\n"%(nmi))
print("method ward")
wh = SpectralClustering(n_clusters=max(cluster))
print("compute the result...")
result = wh.fit_predict(tm)
#showresult(result,cluster)
nmi=NMI(cluster,result)
print("NMI:%g\n"%(nmi))
print("method Agglomerative")
ac = AgglomerativeClustering(n_clusters=max(cluster),linkage='average')
print("compute the result...")
result = ac.fit_predict(tm)
#showresult(result,cluster)
nmi=NMI(cluster, result)
print("NMI:%g\n"%(nmi))
print("method AP")
ap = AffinityPropagation()
print("compute the result...")
result = ap.fit_predict(mv)
#showresult(result,cluster)
nmi=NMI(cluster,result)
print("NMI:%g\n"%(nmi))
print("method GaussianMixture")
GM = GMM(n_components=150)
print("compute the result...")
result = GM.fit_predict(tm)
#showresult(result,cluster)
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nmi=NMI(cluster,result)
<pre>print("NMI:%g\n"%(nmi))</pre>
结论分析与体会: