# University of Missouri-Kansas City School of Computing and Engineering



ICP: ICP-06

**Course Name:** Big Data Analytics and Applications

Course ID: COMP-SCI 5542

Semester /Session: Spring 2022

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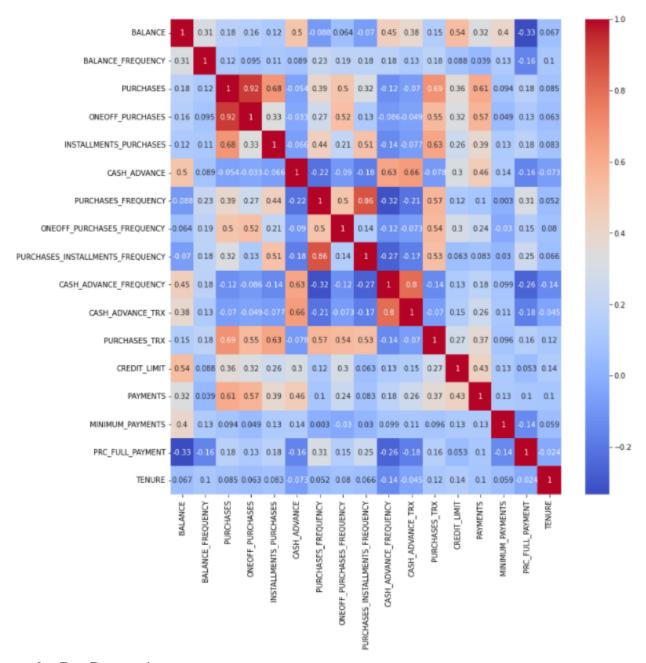
## **Description of the Problem:**

Unsupervised Learning help us solve problems with unlabeled data. A majority of data is unstructured and lacks labels. Unsupervised Learning can be used to group this kind of data into clusters. One most well known algorithm in unsupervised learning is KMeans clustering. This algorithm partitions data into groups based on similarity between data points. Similarity is found by calculating the Euclidean distance between these data points.

In this ICP, we have used a dataset with Credit Card usage among various users. Our goal is to segment these users into buckets so that we can target specific groups. For example, if we wanted to target a group of high spenders with an ad campaign, we would first need to realize who that high spending group consists of. We can't manually assign labels to all of the high spenders as that would take too much time. Instead we used unsupervised learning and clustering.

## **Description of Solution:**

- 1. Get the dataset
  - a. The credit card dataset is a commonly used dataset. We downloaded it from <a href="Kaggle">Kaggle</a> but you can also download it from the UCI repository.
- 2. Data Exploration
  - a. The traditional KMeans algorithm can only be used with numeric data so let's explore our dataset.
  - b. We can use df.info() to get the datatypes of each column. Every column is numeric except for the CustID column (type Object). Customer ID doesn't tell us anything useful about the customers spending so we drop that column.
  - c. The dataset also has about 300 rows that contain null values. Unlike Neural Networks, KMeans does not need a lot of data. We can drop these null rows.
  - d. We can use df.describe() to get statistics about the data. KMeans is sensitive to outliers so we pay close attention to the standard deviation. The standard deviation ranges from 0.2 to 3700. Before we use KMeans we will want to scale our data.
  - e. Let's get an understanding on the correlation in our data. We can do this by creating a heatmap using Seaborn. From this heatmap we can see that correlation exists in the data.



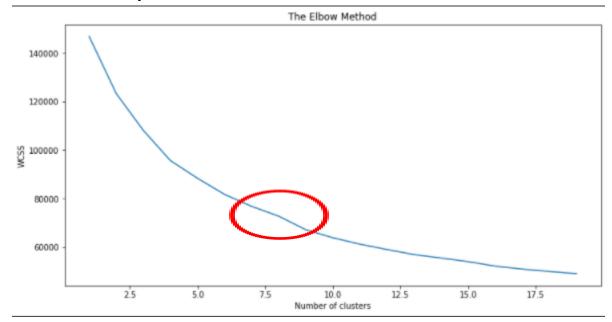
## 3. Pre-Processing

- a. We will pre-process the data by scaling it. The StandardScalar() method from Sci-Kit Learn scales the data so that it has a mean of 0 and a standard deviation of 1.
- b. We can see this by using df.describe() again.
- c. Our data has the shape (8636, 17), we will keep this in mind for the visualization step.

#### 4. KMeans

a. To determine the optimal number of clusters for our dataset, we will use the elbow method. The elbow method plots # clusters against the inertia.

- b. Ideally, we want minimal inertia but we also want to deal with a small number of clusters. We choose the number of clusters by visualizing the inertia decline and deciding where the slope starts to flatten out.
- c. To create this plot, we looped through 20 times and created/fit 20 different KMeans objects and measured their inertia. Here were our results.

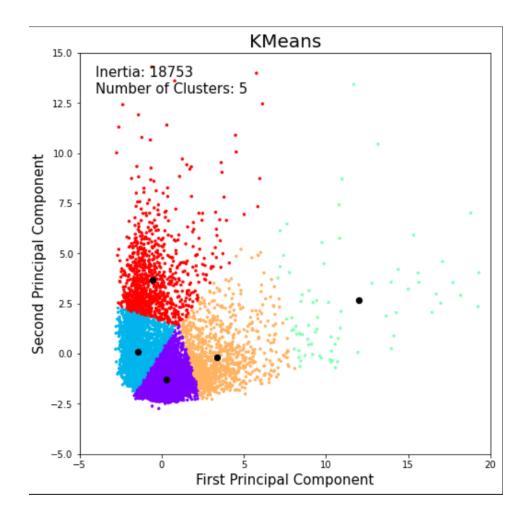


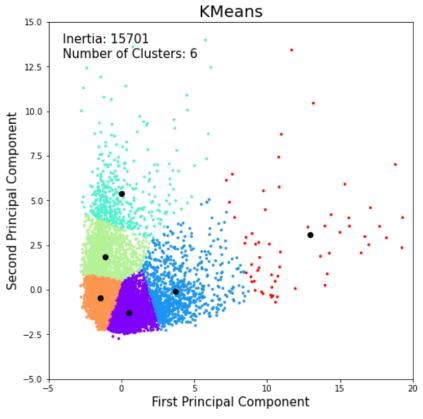
d. We can now recreate KMeans objects with 5 to 9 clusters. These objects can be used to predict the labels of our dataset as well as any new data that we have. We use the KMeans.fit() and KMeans.predict() methods.

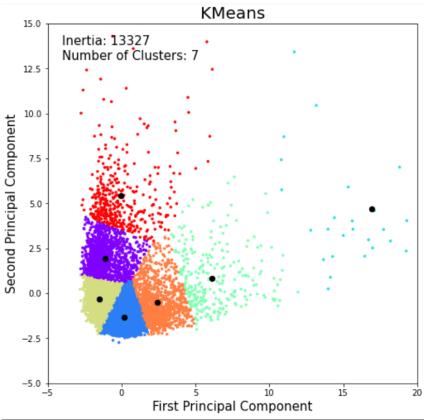
#### 5. Visualization

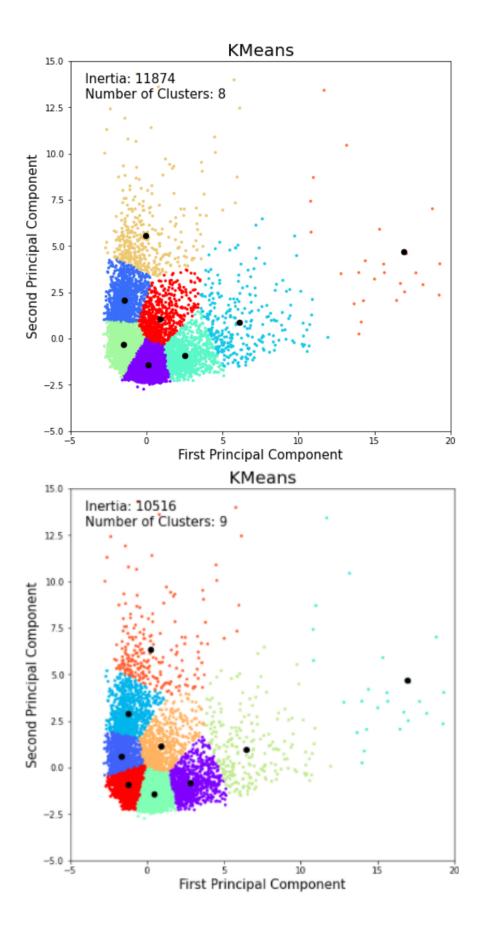
- a. Our dataset contains 17 dimensions as stated earlier. This is too many dimensions for humans to imagine. We need to get our data into a lower dimensional space.
- b. We can do this by using principal component analysis. This process creates a new dataset with columns that are calculated by mixing all of the old columns together. The PCA process does this while maximizing the variance of the dataset so that it can try to capture the characteristics of the original dataset. In this way, we can decompose our 17 dimensions into 2 dimensions without losing too much of the data.
- c. With the new PCA data, we ran KMeans algorithm again with 5 to 9 clusters. We then plotted the first 2 principal components along with the respective cluster label for each data point.
- d. The PCA KMeans will not perfectly reproduce the clusters from the original KMeans, but it is good enough for visualization purposes. Here is the resultant graph. Each cluster is grouped around a black dot which is the centroid of that group.

e. The optimal number of clusters is a subjective measure. Having too many clusters can make subsequent learning algorithms less efficient. For this exercise, we have decided that 9 clusters are the optimal amount of clusters because this has the lowest inertia.









## **Challenges:**

The biggest challenge of this ICP was finding a good dataset. Many of the datasets that we found online through Kaggle and UCI had already been pre-processed and labeled for supervised learning. This dataset was chosen because of the idea that we could group users together by their spending habits.

Another challenge we had was graphing the clusters. Our dataset had 17 columns or independent variables. It is impossible to graph this many dimensions. This is where PCA helped. The drawback of PCA is that we lose some of the data when we decompose our dataset. It will be important to keep in mind that a PCA KMeans model allows us to visualize these clusters. However if we have new data that we want to run through the model, we will not want to use the PCA model because some data is lost.

# **Learning Outcomes:**

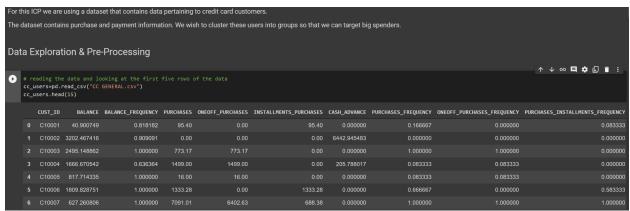
- 1. We learned when to apply unsupervised vs supervised learning.
- 2. We learned how to apply the Elbow method on a dataset.
- 3. We learned that you can use the KMeans++ initializer to get a better beginning iteration.
- 4. We learned that you can use PCA to graph a high-dimensional dataset

#### Resources:

https://www.kaggle.com/arjunbhasin2013/ccdata?select=CC+GENERAL.csv Kaggle data set

Video Link: <a href="https://youtu.be/PYujyuT0HDA">https://youtu.be/PYujyuT0HDA</a>

#### Screenshots:



We can see that we have fully numberical data besides the Customer ID. This ID does not tell us anything useful about spending habits so we will drops this column.

+ Code + Text

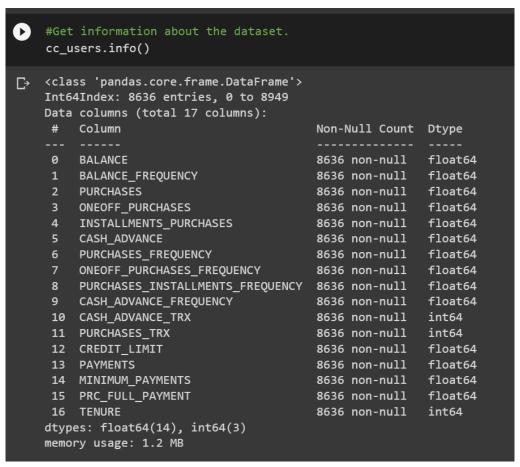
[49] cc\_users.drop(columns=['CUST\_ID'], inplace=True)

We have null values in the columns 'Credit Limit' and 'Minumum payments'.

Lets drop any rows with null data

[50] cc\_users.dropna(how='any', inplace=True)

We now have a uniform dataset with no null values



<pre># statistics of the data cc_users.describe()</pre>									
<b>□</b> •	BALANCE	BALANCE_FREQUENCY	PURCHASES	ONEOFF_PURCHASES	INSTALLMENTS_PURCHASES	CASH_ADVANCE	PURCHASES_FREQUENCY	ONEOFF_PURCHASES_FREQUENCY	PURCHASES_INSTALLMENTS_FREQUENCY C
cour	t 8636.000000	8636.000000	8636.000000	8636.000000	8636.000000	8636.000000	8636.000000	8636.000000	8636.000000
mea	1601.224893	0.895035	1025.433874	604.901438	420.843533	994.175523	0.496000	0.205909	0.368820
std	2095.571300		2167.107984	1684.307803	917.245182	2121.458303	0.401273	0.300054	0.398093
min	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000
25%	148.095189	0.909091	43.367500	0.000000	0.000000	0.000000		0.000000	0.000000
50%	916.855459	1.000000	375.405000	44.995000	94.785000	0.000000	0.500000	0.083333	0.166667
75%	2105.195853	1.000000	1145.980000	599.100000	484.147500	1132.385490	0.916667	0.333333	0.750000
max	19043.138560	1.000000	49039.570000	40761.250000	22500.000000	47137.211760	1.000000	1.000000	1.000000
%									

```
[53] import seaborn as sns
    plt.figure(figsize=(12,12))
    sns.heatmap(cc_users.corr(), annot=True, cmap='coolwarm')
```

```
KMeans uses Euclidian distance to calculate the cluster centroid.

We Scale our data because KMeans is sensitive to outliers. By scaling it, we can get better centroids and thus a better inertia score.

+ Code + Text

[54] from sklearn.preprocessing import StandardScaler
#Scale the data set and hold it in variable X by using fit_transform
scaler = StandardScaler()
X = scaler.fit_transform(cc_users)
X.shape

[8636, 17)

Let's see the statistics of our scaled data. This new data has a standard deviation of 1 for every column. This is from the scaling process where the Scalar transforms our data. This transformed data is called Standard data.
```

#### **KMeans**

Let's do KMeans clustering on our scaled data. We can iterate through different numbers of clusters and get an inertia score.

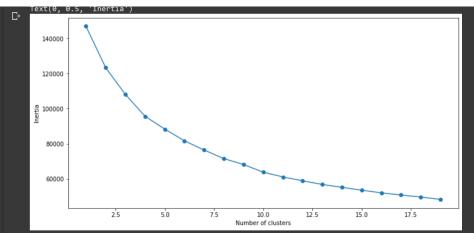
Our goal is to have meaningful clusters that characterize all of the subgroups in the data.

We want to avoid choosing too many clusters because this can lead to bloat. It is best to take the smallest amount of clusters that also has a good inertia score.

We can do this by using the elbow method.

```
[56] # fitting multiple k-means algorithms and storing the values in an empty list
    SSE = []
    for cluster in range(1,20):
        kmeans = KMeans(n_clusters = cluster, init='k-means++')
        kmeans.fit(X)
        SSE.append(kmeans.inertia_)

# converting the results into a dataframe and plotting them
    frame = pd.DataFrame({'Cluster':range(1,20), 'SSE':SSE})
    plt.figure(figsize=(12,6))
    plt.plot(frame['cluster'], frame['SSE'], marker='o')
    plt.xlabel('Number of clusters')
    plt.ylabel('Inertia')
```



From the elbow plot, we choose k=5 to k=9 clusters for the data. We then create this new KMeans model and predict labels with a new dataset.

From the elbow plot, we choose 9 clusters for the data. We then create this new KMeans model and predict labels with a new dataset.

```
# k means using 5 clusters and k-means++ initialization
n_clusters = 5
k_means_orig = KMeans( n_clusters = n_clusters, init='k-means++')
k_means_orig.fit(X)
pred = k_means_orig.predict(X)

[58] frame_orig = pd.DataFrame(X)
frame_orig['cluster'] = pred
frame_orig['cluster'].value_counts()

0 3139
3 3011
1 1248
2 980
4 258
Name: cluster, dtype: int64
```

#### Visualization

Our dataset contains 17 dimensions, but humans can only comprehend 2 or 3 dimensions.

We need to get our dataset into a lower dimension for visualization so we will use principal component analysis.

```
[59] from sklearn.decomposition import PCA

# Instantiate the pca object using 2 components
pca = PCA(n_components=2)

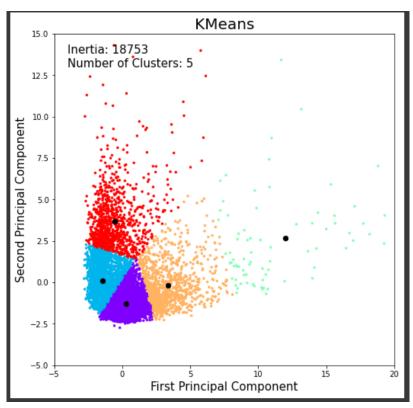
principalComponents = pca.fit_transform(X) #fit transform the data
```

Here we create another KMeans model and run the PCA data. This creates clusters for the lower dimensional data.

We chose to use 5 clusters according to the elbow method so we will again use 5 clusters for this reduced data.

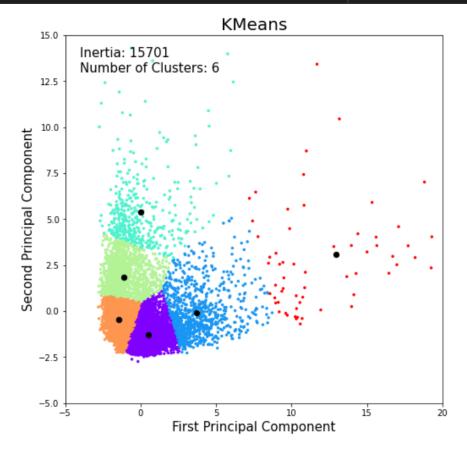
We can now visualize our clusters. The first 2 principal components are able to capture most of the variance in the original dataset, but these clusters may differ slightly from the clusters gathered in the initial KMeans calculation.

```
#plot the clusters
plt.figure(figsize=(8,8))
plt.scatter(principalComponents[:,0],principalComponents[:,1],c=k_means_5.labels_,cmap=plt.cm.get_cmap('rainbow'), s=6)
plt.scatter(k_means_5.cluster_centers_[:,0] , k_means_5.cluster_centers_[:,1] , s = 40, color = 'k')
plt.xlabel('First Principal Component', size=15)
plt.ylabel('Second Principal Component', size=15)
plt.axis([-5, 20, -5, 15])
plt.title('KMeans', size=20)
text = 'Inertia: {:.0f}\nNumber of Clusters: {}'.format(k_means_5.inertia_, k_means_5.n_clusters)
plt.text(-4, 13, text, size=15)
plt.show()
```

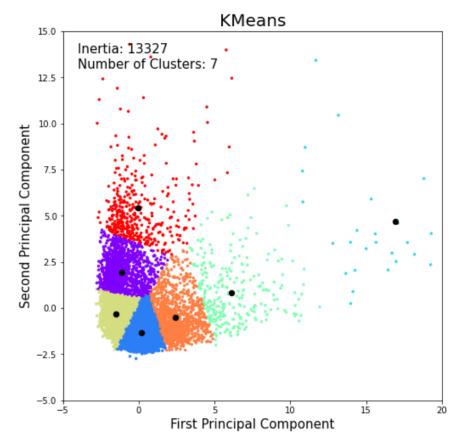


```
6 Clusters
Fit a kmeans model with 6 clusters
[63] # k means using 6 clusters and k-means++ initialization
     n_clusters = 6
     k_means_6 = KMeans( n_clusters = n_clusters, init='k-means++')
     k_means_6.fit(principalComponents)
     pred = k_means_6.predict(principalComponents)
Get value counts for 6 clusters
[64] frame_6 = pd.DataFrame(principalComponents)
     frame_6['cluster'] = pred
frame_6['cluster'].value_counts()
           2903
           2875
           1492
           906
            395
     Name: cluster, dtype: int64
```

```
#plot the clusters
plt.figure(figsize=(8,8))
plt.scatter(principalComponents[:,0],principalComponents[:,1],c=k_means_6.labels_,cmap=plt.cm.get_cmap('rainbow'), s=6)
plt.scatter(k_means_6.cluster_centers_[:,0] , k_means_6.cluster_centers_[:,1] , s = 40, color = 'k')
plt.xlabel('First Principal Component', size=15)
plt.ylabel('Second Principal Component', size=15)
plt.axis([-5, 20, -5, 15])
plt.title('KMeans', size=20)
text = 'Inertia: {:.0f}\nNumber of Clusters: {}'.format(k_means_6.inertia_, k_means_6.n_clusters)
plt.text(-4, 13, text, size=15)
plt.show()
```

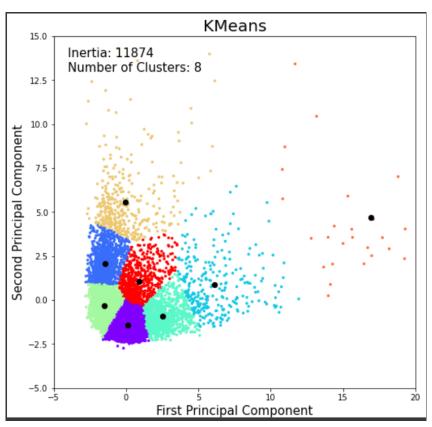


```
7 Clusters
Fit a model with 7 clusters
[66] # k means using 7 clusters and k-means++ initialization
     n_clusters = 7
     k_means_7 = KMeans( n_clusters = n_clusters, init='k-means++')
     k_means_7.fit(principalComponents)
     pred = k_means_7.predict(principalComponents)
Get value counts for 7 clusters
[67] frame_7 = pd.DataFrame(principalComponents)
     frame_7['cluster'] = pred
     frame_7['cluster'].value_counts()
          2705
          2689
          1375
          1172
           387
           278
           30
     Name: cluster, dtype: int64
```



```
8 Clusters
Fit a model with 8 clusters
[69] # k means using 8 clusters and k-means++ initialization
    n_clusters = 8
    k_means_8 = KMeans( n_clusters = n_clusters, init='k-means++')
    k_means_8.fit(principalComponents)
    pred = k_means_8.predict(principalComponents)
Get value counts for 8 clusters
[70] frame_8 = pd.DataFrame(principalComponents)
     frame_8['cluster'] = pred
     frame_8['cluster'].value_counts()
         2660
    0
        1020
          629
          30
    Name: cluster, dtype: int64
```

```
#plot the clusters
plt.figure(figsize=(8,8))
plt.scatter(principalComponents[:,0],principalComponents[:,1],c=k_means_8.labels__,cmap=plt.cm.get_cmap('rainbow'), s=6)
plt.scatter(k_means_8.cluster_centers_[:,0] , k_means_8.cluster_centers_[:,1] , s = 40, color = 'k')
plt.xlabel('First Principal Component', size=15)
plt.ylabel('Second Principal Component', size=15)
plt.axis([-5, 20, -5, 15])
plt.title('KMeans', size=20)
text = 'Inertia: {:.0f}\nNumber of Clusters: {}'.format(k_means_8.inertia_, k_means_8.n_clusters)
plt.text(-4, 13, text, size=15)
plt.show()
```



```
9 Clusters
Fit a KMeans model with 9 clusters
[72] # k means using 9 clusters and k-means++ initialization
     n_clusters = 9
     k_means_9 = KMeans( n_clusters = n_clusters, init='k-means++')
     k_means_9.fit(principalComponents)
     pred = k_means_9.predict(principalComponents)
Get value counts for 9 clusters
[73] frame_9 = pd.DataFrame(principalComponents)
     frame_9['cluster'] = pred
     frame_9['cluster'].value_counts()
          2038
          1770
           814
           805
           598
           225
           30
     Name: cluster, dtype: int64
```

```
#Plot the clusters
plt.figure(figsize=(8,8))
plt.scatter(principalComponents[:,0],principalComponents[:,1],c=k_means_9.labels_,cmap=plt.cm.get_cmap('rainbow'), s=6)
plt.scatter(k_means_9.cluster_centers_[:,0] , k_means_9.cluster_centers_[:,1] , s = 40, color = 'k')
plt.xlabel('First Principal Component', size=15)
plt.ylabel('Second Principal Component', size=15)
plt.axis([-5, 20, -5, 15])
plt.title('KMeans', size=20)
text = 'Inertia: {:.0f}\nNumber of Clusters: {}'.format(k_means_9.inertia_, k_means_9.n_clusters)
plt.text(-4, 13, text, size=15)
plt.show()
```

