```
In [1]:
         # k-means clustering
         from numpy import unique
         from numpy import where
         from sklearn.datasets import make_classification
         from sklearn.cluster import KMeans
         from matplotlib import pyplot
          import numpy as np
          import pandas as pd
          import seaborn as sns
          import matplotlib.pyplot as plt
         %matplotlib inline
          import pandas as pd
          import numpy as np
          import numpy as np
         from sklearn.cluster import KMeans
         from sklearn import datasets
         from sklearn.preprocessing import StandardScaler
In [2]:
         import warnings
         warnings.filterwarnings('ignore')
In [3]:
         # the combined data
         data_folder = './nhanes_input_data/'
         # import the CSV as a pandas dataframe
         df = pd.read csv( data folder + '0 dietaryIntakeDataForClassificationAndAnalysisData.cs
         df.shape
         (193805, 87)
Out[3]:
In [4]:
         df.head(5)
Out[4]:
                                                                                             SEQN -
                                                                                         Respondent
           RIDAGEYR_Age_in_years_at_screening URDACT_Albumin_creatinine_ratio_mg_g DataYear
                                                                                            sequence
                                                                                             number
                                                                                   2017-
         0
                                       53.0
                                                                            3.0
                                                                                             95405.0
                                                                                    2018
                                                                                   2017-
         1
                                       53.0
                                                                                             95405.0
                                                                            3.0
                                                                                    2018
                                                                                   2017-
         2
                                       53.0
                                                                            3.0
                                                                                             95405.0
                                                                                    2018
                                                                                   2017-
         3
                                       53.0
                                                                                             95405.0
                                                                            3.0
                                                                                    2018
```

53.0

4

2017-

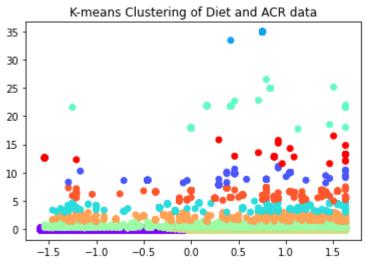
2018

95405.0

3.0

```
In [5]:
         # parameters to be used for KMeans clustring: centres
         # X and/or kdf will have only features we want to create cluster around
         kdf = df[
             Ε
                 'RIDAGEYR_Age_in_years_at_screening'
                 , 'URDACT Albumin creatinine ratio mg g'
             1
         ]
         X = kdf
         X[:5]
Out[5]:
          0
                                    53.0
                                                                      3.0
        1
                                    53.0
                                                                      3.0
        2
                                    53.0
                                                                      3.0
        3
                                    53.0
                                                                      3.0
                                    53.0
                                                                      3.0
In [6]:
         # ref: internet (not my code, using as a library)
         def clean_dataset(df):
             assert isinstance(df, pd.DataFrame), "df needs to be a pd.DataFrame"
             df.dropna(inplace=True)
             indices_to_keep = ~df.isin([np.nan, np.inf, -np.inf]).any(1)
             return df[indices_to_keep].astype(np.float64)
In [7]:
         # X has the features to cluster around (centres: Age, ACR). df has the complete data
         # after clustering is done using features in X, we find positions (index) for each data
         # in a cluster then we use those index positions to cluster the data from df
         X.shape, df.shape
        ((193805, 2), (193805, 87))
Out[7]:
In [8]:
         X = clean_dataset(X)
In [9]:
         # define the model
         model = KMeans(n clusters = 10) #,random state=0, n init="auto"
         # fit the model
         model.fit(X)
         #model.labels
        KMeans(n_clusters=10)
Out[9]:
```

```
In [10]:
          # Create csv files with the cluster daya
          # One csv for one Cluster
In [11]:
          howManyClusters = 10
          for clusterId in range (howManyClusters):
              ind_list = np.where(model.labels_ == clusterId )[0]
              cluster = df.iloc[ind list]
              cluster.to_csv('./nhanes_output_data/classifiedGroups/kmeanscluster/cluster-'
                             + str(clusterId) + '.csv');
In [12]:
          model.cluster_centers_
                   18.87596942,
         array([[
                                  12.04466503],
Out[12]:
                   51.67164179, 2496.77649254],
                   58.87096774, 9398.2316129 ],
                   50.85535466, 1061.83630042],
                   49.14736842, 5597.96115789],
                  49.44935831, 203.29598508],
                  60.84809966, 15.0886079],
                   57.20701513, 620.4038033 ],
                   59.28975741, 1682.01998652],
                   52.31707317, 3538.97792683]])
In [13]:
          # Scatter plot to see each cluster points visually
          std_data = StandardScaler().fit_transform(X)
          plt.scatter(std_data[:,0], std_data[:,1], c = model.labels_, cmap = "rainbow")
          plt.title("K-means Clustering of Diet and ACR data")
          plt.show()
```



## References:

print("Shape of cluster:", model.clustercenters.shape)

https://stackoverflow.com/questions/50297142/ccluster-points-after-kmeans-in-a-list-format

https://machinelearningmastery.com/clustering-algorithms-with-python/

https://stackoverflow.com/questions/50297142/ccluster-points-after-kmeans-in-a-list-format

https://scikitlearn.org/stable/modules/generated/sklearn.clus

https://datascience.stackexchange.com/questionk-means-clustering-over-multiple-columns

https://scikitlearn.org/stable/modules/generated/sklearn.clus

from sklearn.cluster import KMeans import numpy as np X = np.array([[1, 2], [1, 4], [1, 0], ... [10, 2], [10, 4], [10, 0]]) kmeans = KMeans(n\_clusters=2, random\_state=0, ninit="auto").fit(X) kmeans.labels array([1, 1, 1, 0, 0, 0], dtype=int32) kmeans.predict([[0, 0], [12, 3]]) array([1, 0], dtype=int32) kmeans.clustercenters array([[10., 2.], [1., 2.]])

```
In [ ]:
```