РК №2 по ТМО

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```
In [0]: from google.colab import drive
         import pandas as pd
         import numpy as np
         from sklearn.preprocessing import LabelEncoder
         from sklearn.cluster import KMeans
         from sklearn.metrics import adjusted_rand_score
         from sklearn.metrics import adjusted_mutual_info_score
         from sklearn.metrics import homogeneity_completeness_v_measure
         from sklearn.metrics import silhouette_score
         from sklearn.cluster import DBSCAN
         from sklearn.cluster import MeanShift
         import seaborn as sns
         import matplotlib.pyplot as plt
         from sklearn.cluster import Birch
         from sklearn.model_selection import GridSearchCV, KFold
In [0]: #Загружаю данные с гугл диска
         data = pd.read csv('/content/gdrive/My Drive/mushrooms.csv', sep=",")
In [4]: data.head()
Out[4]:
                                                                                          stalk-
                                                                                                  stalk-
                                                      gill-
                                                              gill- gill-
                                                                        gill-
                                                                            stalk- stalk- surface-
                                                                                                surface-
                                                                                                        colo
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                          сар-
                               сар-
            class
                                    bruises odor
                 shape
                       surface
                               color
                                                 attachment spacing
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                                                                       color
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                                  g
In [0]: data.size
Out[0]: 186852
In [0]: data.isnull().sum()
Out[0]: class
                                       0
                                       0
         cap-shape
                                       0
         cap-surface
         cap-color
                                       0
         bruises
                                       0
                                       0
         odor
         gill-attachment
                                       0
         gill-spacing
                                       0
         gill-size
                                       0
         gill-color
         stalk-shape
                                       0
         stalk-root
                                       0
         stalk-surface-above-ring
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         stalk-surface-below-ring
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         stalk-color-above-ring
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         stalk-color-below-ring
                                       0
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         veil-type
         veil-color
                                       0
         ring-number
                                       0
         ring-type
                                       0
         spore-print-color
                                       0
         population
                                       0
         habitat
         dtype: int64
```

```
In [0]: data.dtypes
 Out[0]: class
                                      object
         cap-shape
                                      object
         cap-surface
                                      object
         cap-color
                                      object
         bruises
                                      object
                                      object
         odor
         gill-attachment
                                      object
         gill-spacing
                                      object
         gill-size
                                      object
         gill-color
                                      object
         stalk-shape
                                      object
         stalk-root
                                      object
         stalk-surface-above-ring
                                      object
         stalk-surface-below-ring
                                      object
         stalk-color-above-ring
                                      object
         stalk-color-below-ring
                                      object
         veil-type
                                      object
         veil-color
                                      object
         ring-number
                                      object
         ring-type
                                      object
         spore-print-color
                                      object
         population
                                      object
         habitat
                                      object
         dtype: object
 In [0]: le = LabelEncoder()
'ring-type', 'spore-print-color', 'population', 'habitat']
         col y = 'class'
 In [0]: | for i in cols_x:
           data[i] = le.fit transform(data[[i]])
         data['class'] = le.fit_transform(data[['class']])
 In [0]: | data.head()
 Out[0]:
                                                                                      stalk-
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                              сар-
                                                    gill-
                                                           gill- gill-
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                                                                                    surface-
                                                                                           surface-
                                                                                                   colo
                                   bruises odor
            class
                                              attachment spacing
                 shape
                       surface color
                                                               size color
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                                                                               root
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                                            6
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          1
               0
                     5
                            2
                                 9
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                                            0
                                                      1
                                                             0
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                                        1
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          3
               1
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                            3
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                                                      1
                                                             0
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               O
                     5
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                                       0
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                                                             1
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In [10]: for i in cols_x2:
           print(data[i].unique())
         [1 0]
         [0 1]
         [1 0]
         [45271039108116]
         [3 2 1 4 0]
         [2 0 1 3]
         [2 0 3 1]
         [4 0 2 1 3]
         [3 2 0 4 5 1]
```

```
In [0]: plt.figure(figsize = (18,5))
             sns.heatmap(data.corr(method='pearson'), annot=True, fmt='.2f', square=False)
Out[0]: <matplotlib.axes._subplots.AxesSubplot at 0x7f0463568d68>
                                                                                                                                                - 0.9
                      cap-surface
                                                                                                                                                - 0.6
             gill-attachment
gill-spacing
gill-size
gill-color
stalk-shape
stalk-root
stalk-surface-above-ring
stalk-surface
                              -0.35
                                            -0.46
                                                                                                                                                 0.3
              stalk-surface-below-ring
               calk-surface-below-ring
stalk-color-below-ring
stalk-color-below-ring
veil-color
ring-number
ring-type
spore-print-color
                                                                                                                                                 0.0
                                                                                                                                                 -0.3
                                                                                       stalk-surface-above
                                                                                                stalk
                                                                                                    stalk
                                                                                           stalk
'population', ]
In [0]: X = data[cols x2]
             Y = data[col_y]
```

DBSCAN

```
In [0]: | temp_cluster_db = DBSCAN(eps=0.99).fit_predict(X)
In [0]: | ari = adjusted_rand_score(Y, temp_cluster_db)
        ami = adjusted_mutual_info_score(Y, temp_cluster_db)
        h, c, v = homogeneity_completeness_v_measure(Y, temp_cluster_db)
        sl = silhouette_score(X, temp_cluster_db)
        print('''ARI: \{\overline{0}\},
        AMI:{1},
        Homogeneity: {2},
        Completeness: {3},
        V-measure: {4},
        Silhouette:{5}'''.format(ari, ami, h, c, v, sl))
        /usr/local/lib/python3.6/dist-packages/sklearn/metrics/cluster/supervised.py:746: FutureW
        arning: The behavior of AMI will change in version 0.22. To match the behavior of 'v_meas
        ure_score', AMI will use average_method='arithmetic' by default.
          FutureWarning)
        ARI: 0.045137012766120144,
        AMI:0.15358923643298394,
        Homogeneity: 0.9728493007154457,
        Completeness: 0.15573378177437797,
        V-measure: 0.26848798825290127,
        Silhouette:0.9998172438350283
```

Mean Shift

```
In [0]: temp_cluster_ms = MeanShift().fit_predict(X)
```

```
In [0]: ari = adjusted rand score(Y, temp cluster ms)
        ami = adjusted_mutual_info_score(Y, temp_cluster_ms)
        h, c, v = homogeneity_completeness_v_measure(Y, temp_cluster_ms)
        sl = silhouette_score(X, temp_cluster_ms)
        print('''ARI: {0},
        AMI:{1},
        Homogeneity: {2}
        Completeness: {3},
        V-measure: {4},
        Silhouette: {5}'''.format(ari, ami, h, c, v, sl))
        /usr/local/lib/python3.6/dist-packages/sklearn/metrics/cluster/supervised.py:746: FutureW
        arning: The behavior of AMI will change in version 0.22. To match the behavior of 'v meas
        ure_score', AMI will use average_method='arithmetic' by default.
          FutureWarning)
        ARI: 0.3069603675011273,
        AMI:0.28856523801640643,
        Homogeneity: 0.39129684154183636,
        Completeness: 0.2886585628469996,
        V-measure: 0.33223115279908827,
        Silhouette: 0.4531091565478523
```

Birch

```
In [0]: temp_cluster_br = Birch().fit_predict(X)
In [20]: | ari = adjusted rand score(Y, temp cluster br)
         ami = adjusted mutual info score(Y, temp cluster br)
         h, c, v = homogeneity_completeness_v_measure(Y, temp_cluster_br)
         sl = silhouette_score(X, temp_cluster_br)
         print('''ARI: {\overline{0}},
         AMI:{1},
         Homogeneity: {2},
         Completeness: {3},
         V-measure: {4},
         Silhouette: {5}'''.format(ari, ami, h, c, v, sl))
         /usr/local/lib/python3.6/dist-packages/sklearn/metrics/cluster/supervised.py:746: FutureW
         arning: The behavior of AMI will change in version 0.22. To match the behavior of 'v meas
         ure_score', AMI will use average_method='arithmetic' by default.
           FutureWarning)
         ARI: 0.30438499877111097,
         AMI:0.269229255154706,
         Homogeneity: 0.361660376699665,
         Completeness: 0.26932602881101514,
         V-measure: 0.30873740601743555,
         Silhouette: 0.462697384729021
In [33]: n \text{ range } br = np.array(np.arange(0.01,1,0.1))
         tuned_parameters_br = [{'threshold': n_range_br}]
         tuned_parameters_br
Out[33]: [{'threshold': array([0.01, 0.11, 0.21, 0.31, 0.41, 0.51, 0.61, 0.71, 0.81, 0.91])}]
         br qs = GridSearchCV(Birch(), tuned parameters br, cv=KFold(n splits=5), scoring='adjust
         ed mutual info score')
         br_gs.fit(data[cols_x2], data[col_y])
         br_gs.best_params_
In [0]: | temp_cluster_br_gs = br_gs.best_estimator_.fit_predict(X)
```

```
In [44]: ari = adjusted_rand_score(Y, temp_cluster_br_gs)
ami = adjusted_mutual_info_score(Y, temp_cluster_br_gs)
h, c, v = homogeneity_completeness_v_measure(Y, temp_cluster_br_gs)
sl = silhouette_score(X, temp_cluster_br_gs)
print('''ARI: {0},
AMI:{1},
Homogeneity:{2},
Completeness: {3},
V-measure: {4},
Silhouette: {5}'''.format(ari, ami, h, c, v, sl))
```

/usr/local/lib/python3.6/dist-packages/sklearn/metrics/cluster/supervised.py:746: FutureW arning: The behavior of AMI will change in version 0.22. To match the behavior of 'v_meas ure_score', AMI will use average_method='arithmetic' by default. FutureWarning)

ARI: 0.13627703099355284, AMI:0.1765824873001168,

Homogeneity:0.20769636588400608, Completeness: 0.1767070717384534, V-measure: 0.1909525932082184, Silhouette: 0.2787015966914067