

# lab\_0

March 8, 2021

## 1 Adam Jochna

```
[3]: import pandas as pd
import matplotlib.pyplot as plt
from sklearn.preprocessing import StandardScaler
import seaborn as sns
from sklearn.decomposition import PCA
import time
```

```
[4]: PROJECT_PATH = '/home/adam/Desktop/ml_labs'
```

```
[5]: datasets_data = {
    'iris': {
        'columns': [
            'sepal_length',
            'sepal_width',
            'petal_length',
            'petal_width',
            'class'
        ],
        'y_column': 'class',
        'class_mapping': {
            0: 'Iris-setosa',
            1: 'Iris-versicolor',
            2: 'Iris-virginica'
        },
        'plot_0_cols': ['sepal_width', 'sepal_length']
    },
    'glass': {
        'columns': [
            'Id',
            'RI',
            'Na',
            'Mg',
            'Al',
            'Si',
            'K',
            'Ca',
            'Ba',
            'Fe'
        ]
    }
},
```

```

        'Ca',
        'Ba',
        'Fe',
        'type_of_glass'
    ],
    'y_column': 'type_of_glass',
    'class_mapping': None,
    'class_mapping': {
        1: 'building_windows_float_processed',
        2: 'building_windows_non_float_processed',
        3: 'vehicle_windows_float_processed',
        4: 'vehicle_windows_non_float_processed (none in this database)',
        5: 'containers',
        6: 'tableware',
        7: 'headlamps',
    },
    'plot_0_cols': ['RI', 'Si']
},
'wine': {
    'columns': [
        'class',
        'alcohol',
        'malic_acid',
        'ash',
        'alcalinity_of_ash',
        'magnesium',
        'total_phenols',
        'flavanoids',
        'nonflavanoid_phenols',
        'proanthocyanins',
        'color_intensity',
        'hue',
        'OD280/OD315_of_diluted_wines',
        'proline'
    ],
    'y_column': 'class',
    'class_mapping': {
        1: 'class_1',
        2: 'class_2',
        3: 'class_3',
    },
    'plot_0_cols': ['alcohol', 'malic_acid']
},
}

```

```
[6]: def perform_analysis(dataset_name):
    assert dataset_name in ['iris', 'glass', 'wine']
```

```

df = pd.read_csv(
    '{}/lab_0/datasets/{}/{}.data'.format(
        PROJECT_PATH,
        dataset_name,
        dataset_name
    ),
    header=None
)

df.columns = datasets_data[dataset_name]['columns']
y_col = datasets_data[dataset_name]['y_column']

x_cols = datasets_data[dataset_name]['columns'].copy()
x_cols.remove(datasets_data[dataset_name]['y_column'])

if 'Id' in x_cols:
    x_cols.remove('Id')

if dataset_name == 'iris':
    class_inv_mapping = {v: k for k, v in
    ↪datasets_data[dataset_name]['class_mapping'].items()}
    df[y_col] = df[y_col].apply(lambda x: class_inv_mapping[x])

df = df[x_cols + [y_col]]
df.columns = x_cols + ['class_idx']

class_counts = df['class_idx'].value_counts().to_frame()
class_counts = class_counts.reset_index()
class_counts.columns = ['class_idx', 'class_count']
class_counts['class_name'] = class_counts['class_idx'].apply(lambda x: ↪
    datasets_data[dataset_name]['class_mapping'][x])
class_counts = class_counts[['class_name', 'class_idx', 'class_count']]
class_counts['class_perc'] = class_counts['class_count']/
    ↪class_counts['class_count'].sum()*100

print('#'*30)
print('DATASET NAME: {}'.format(dataset_name))

print()
print('CLASS DISTRIBUTION ANALYSIS:')

print(class_counts)

print()
print('DATASET ATTRIBUTES:')

print(x_cols)

```

```

print()
print('DATASET ANALYSIS:')

print(df[x_cols].describe())

# SCATTER PLOT

plt.figure(figsize=(9, 9))

for class_idx in datasets_data[dataset_name]['class_mapping'].keys():
    df_class = df.loc[df['class_idx'] == class_idx]
    class_name = datasets_data[dataset_name]['class_mapping'][class_idx]
    col_0, col1 = datasets_data[dataset_name]['plot_0_cols']

    plt.scatter(df_class[col_0], df_class[col1], label=class_name)

plt.legend(loc='lower right')
plt.xlabel(col_0)
plt.ylabel(col1)

# PAIRGRID PLOT

df['class_name'] = df['class_idx'].apply(lambda x:_
    datasets_data[dataset_name]['class_mapping'][x])
g = sns.PairGrid(df[x_cols + ['class_name']], hue="class_name")
g.map_diag(plt.hist)
g.map_offdiag(plt.scatter)
g.add_legend()

# PCA SCATTER PLOT

x = df.loc[:, x_cols].values
y = df.loc[:, ['class_idx']].values
x = StandardScaler().fit_transform(x)

pca = PCA(n_components=2)
pca_components = pca.fit_transform(x)
df_pca = pd.DataFrame(
    data=pca_components,
    columns=['comp_0', 'comp_1'])
df_pca = pd.concat([df_pca, df[['class_idx']]], axis=1)

plt.figure(figsize=(9, 9))

for class_idx in datasets_data[dataset_name]['class_mapping'].keys():

```

```

df_class = df_pca.loc[df_pca['class_idx'] == class_idx]
class_name = datasets_data[dataset_name]['class_mapping'][class_idx]
col_0, col1 = 'comp_0', 'comp_1'

plt.scatter(df_class[col_0], df_class[col1], label=class_name)

plt.legend(loc='lower right')
plt.xlabel(col_0)
plt.ylabel(col1)

```

[7]: `perform_analysis(dataset_name='iris')`

```

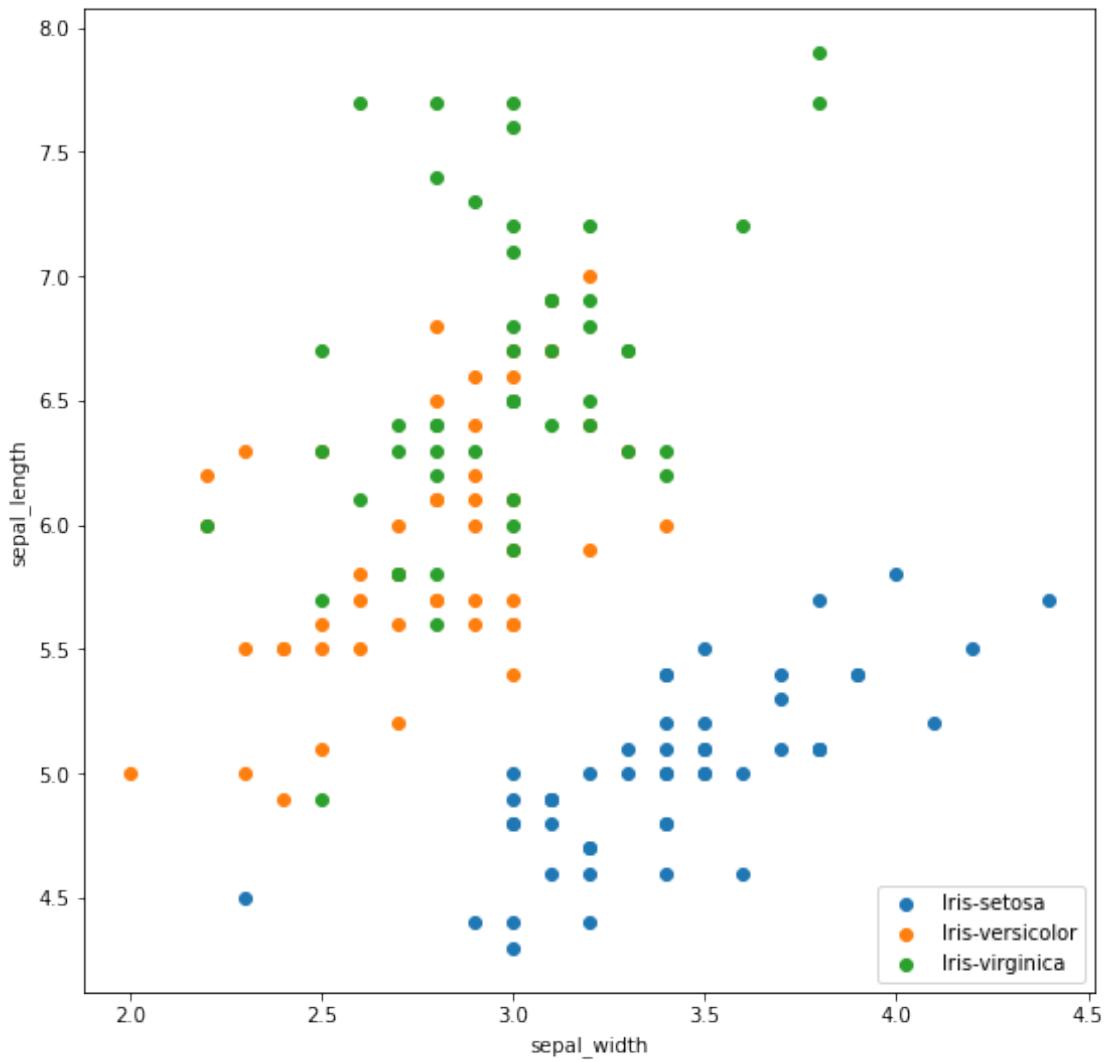
#####
DATASET NAME: iris

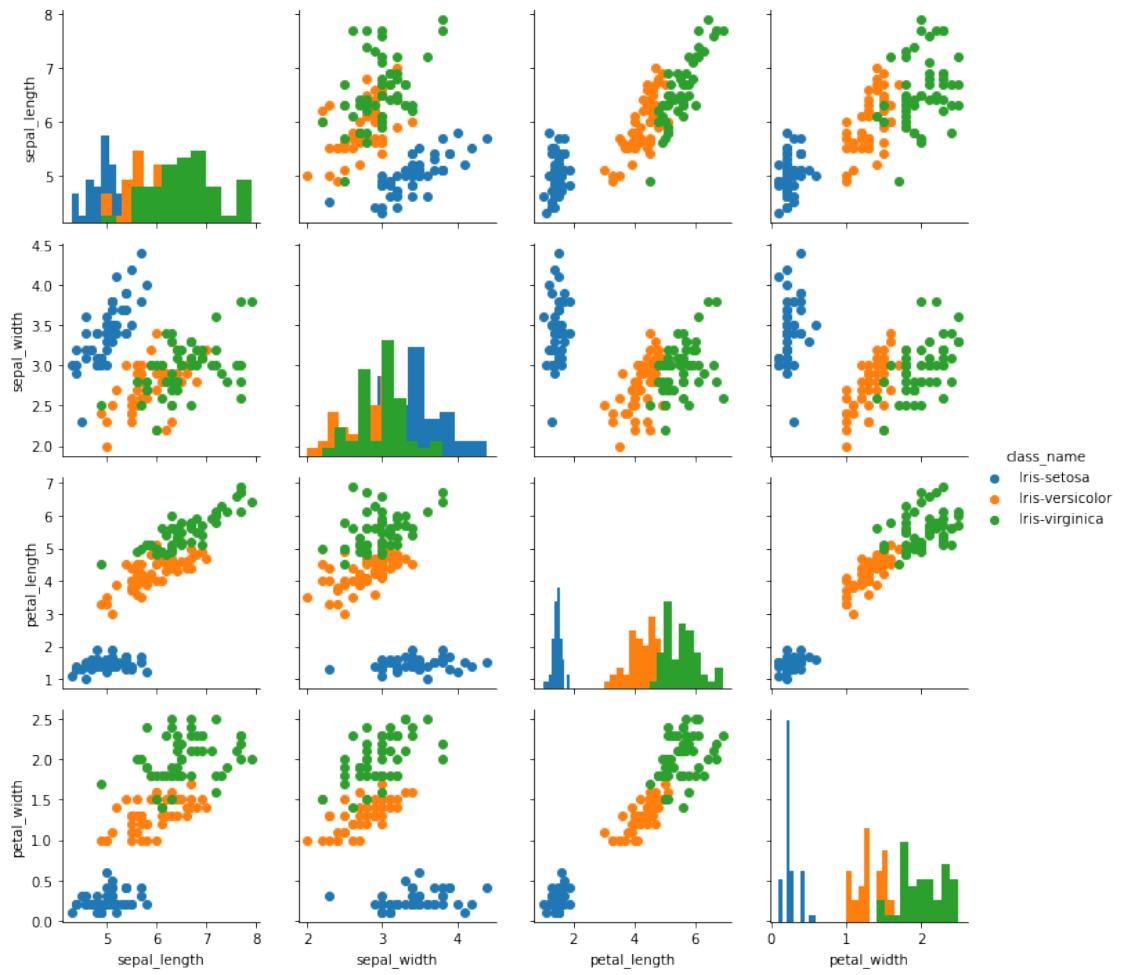
CLASS DISTRIBUTION ANALYSIS:
   class_name  class_idx  class_count  class_perc
0  Iris-virginica        2          50    33.333333
1  Iris-versicolor       1          50    33.333333
2    Iris-setosa         0          50    33.333333

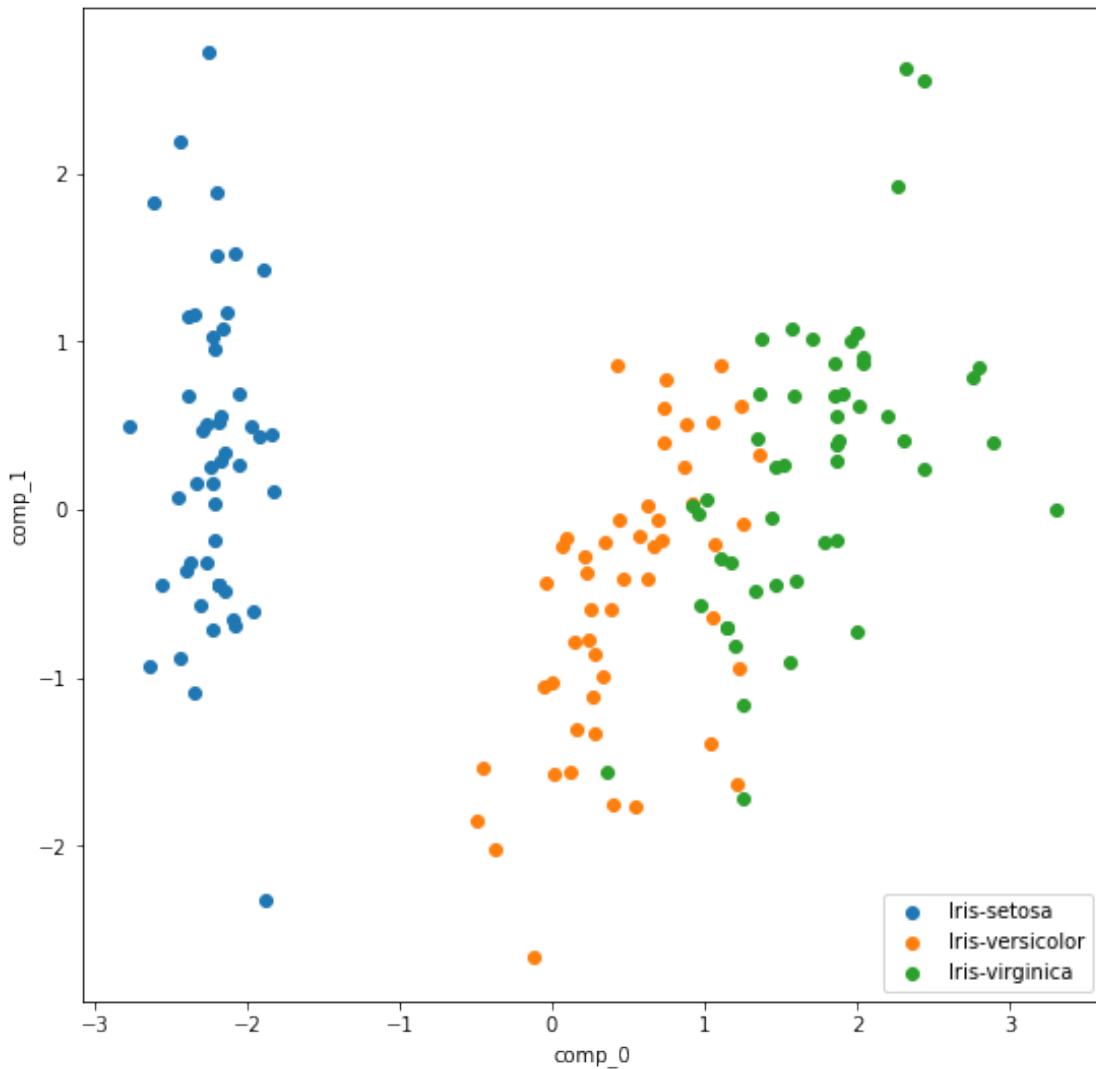
DATASET ATTRIBUTES:
['sepal_length', 'sepal_width', 'petal_length', 'petal_width']

DATASET ANALYSIS:
   sepal_length  sepal_width  petal_length  petal_width
count    150.000000    150.000000    150.000000    150.000000
mean      5.843333     3.054000     3.758667     1.198667
std       0.828066     0.433594     1.764420     0.763161
min       4.300000     2.000000     1.000000     0.100000
25%      5.100000     2.800000     1.600000     0.300000
50%      5.800000     3.000000     4.350000     1.300000
75%      6.400000     3.300000     5.100000     1.800000
max      7.900000     4.400000     6.900000     2.500000

```







```
[8]: perform_analysis(dataset_name='glass')
```

```
#####
DATASET NAME: glass
```

CLASS DISTRIBUTION ANALYSIS:

		class_name	class_idx	class_count	class_perc
0	building_windows_non_float_processed		2	76	35.514019
1	building_windows_float_processed		1	70	32.710280
2	headlamps		7	29	13.551402
3	vehicle_windows_float_processed		3	17	7.943925
4	containers		5	13	6.074766
5	tableware		6	9	4.205607

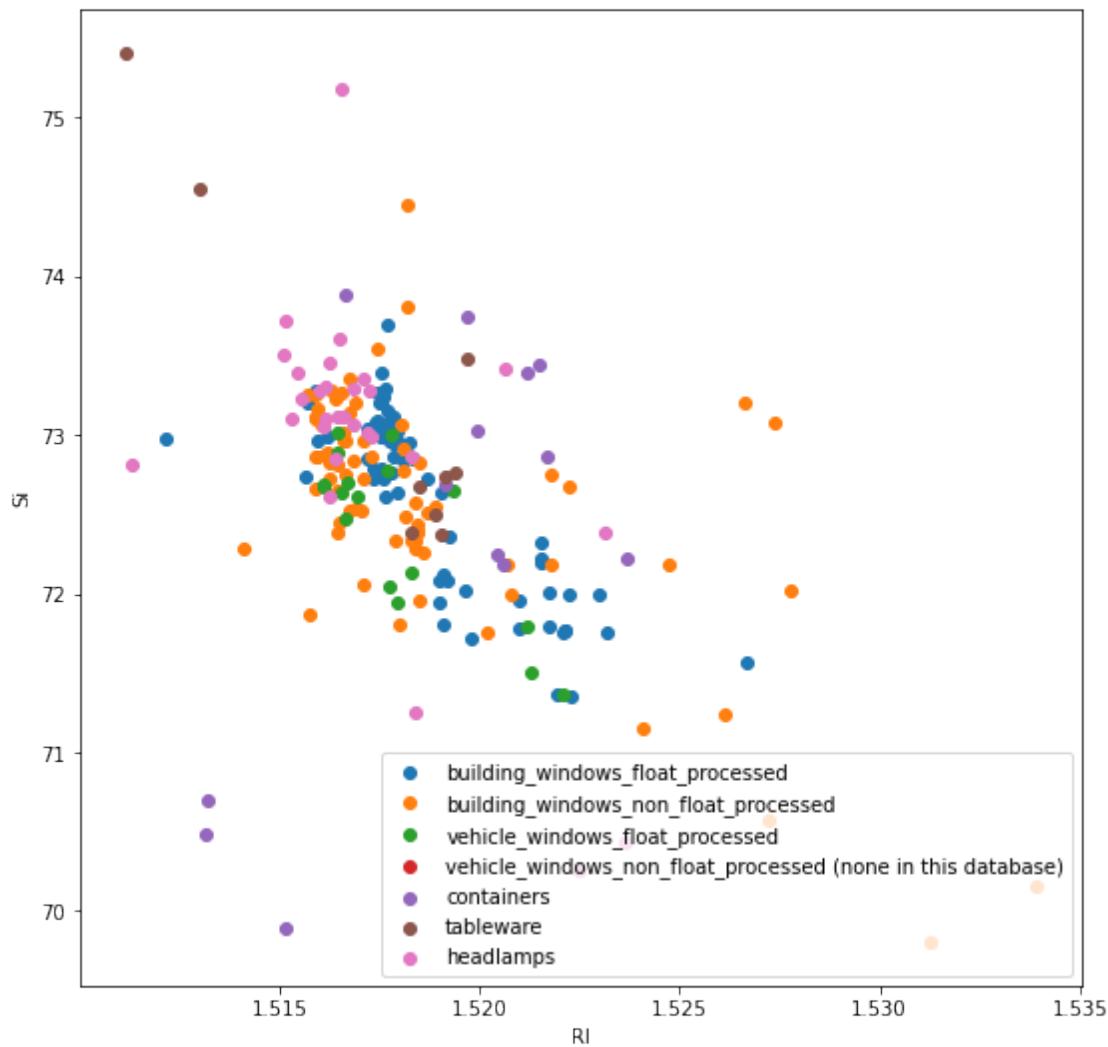
DATASET ATTRIBUTES:

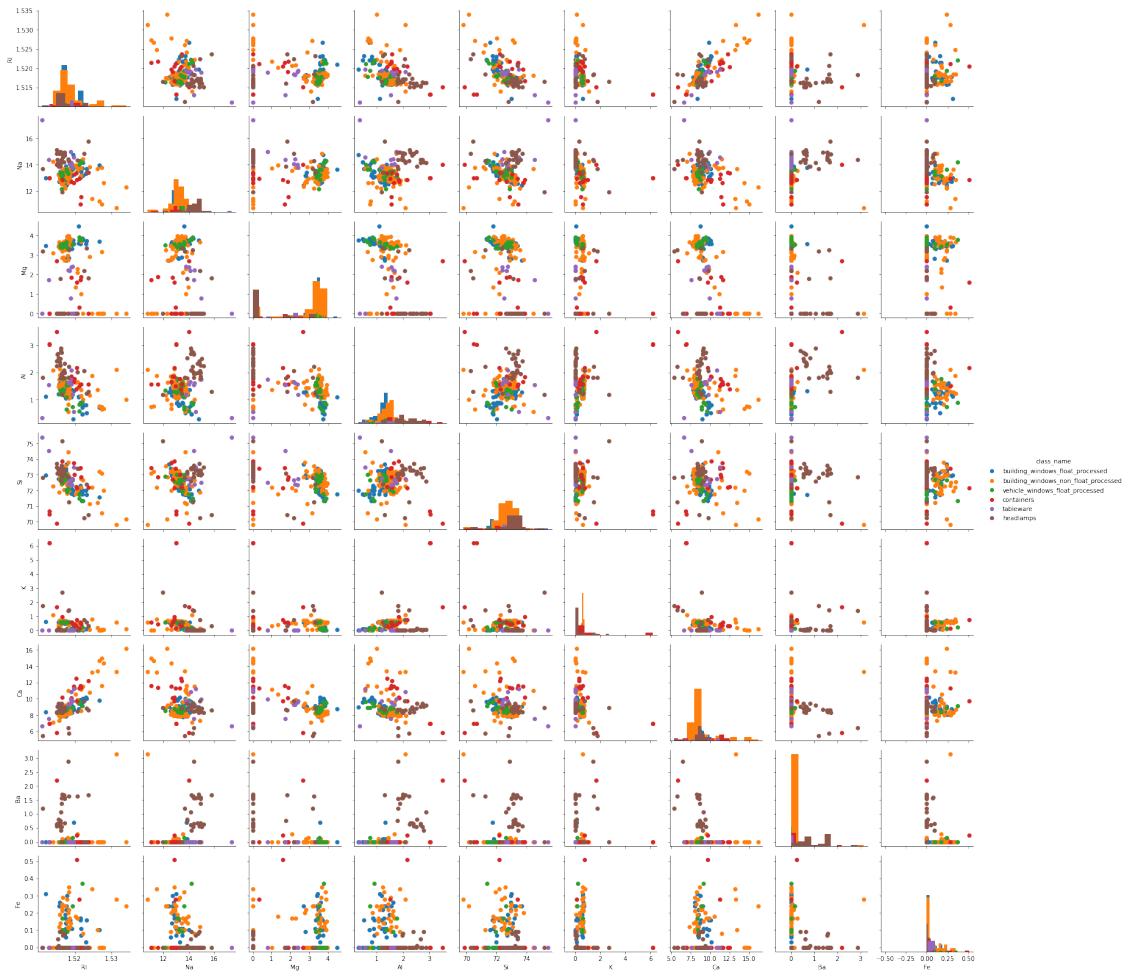
```
['RI', 'Na', 'Mg', 'Al', 'Si', 'K', 'Ca', 'Ba', 'Fe']
```

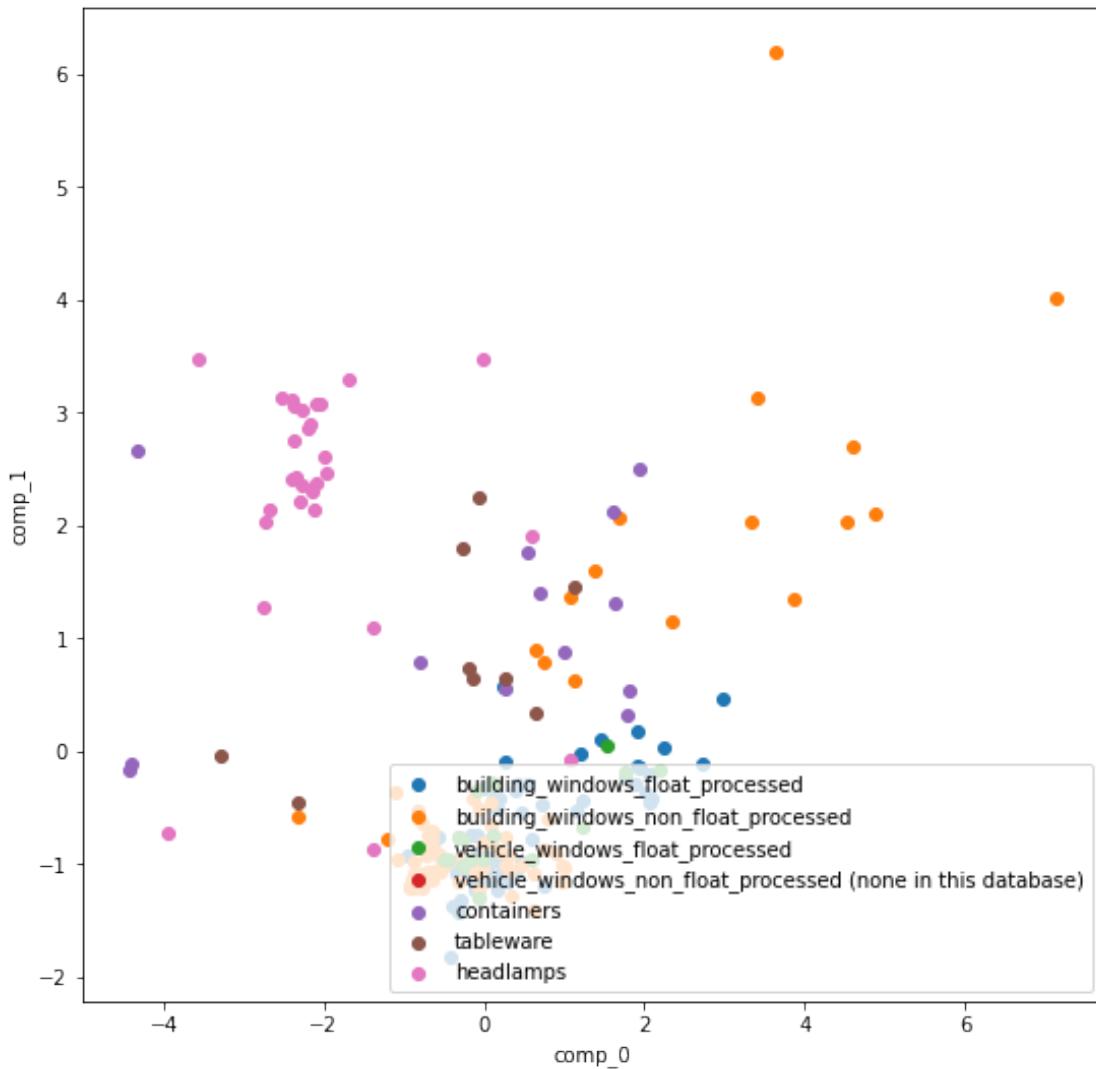
DATASET ANALYSIS:

	RI	Na	Mg	Al	Si	K	\
count	214.000000	214.000000	214.000000	214.000000	214.000000	214.000000	
mean	1.518365	13.407850	2.684533	1.444907	72.650935	0.497056	
std	0.003037	0.816604	1.442408	0.499270	0.774546	0.652192	
min	1.511150	10.730000	0.000000	0.290000	69.810000	0.000000	
25%	1.516523	12.907500	2.115000	1.190000	72.280000	0.122500	
50%	1.517680	13.300000	3.480000	1.360000	72.790000	0.555000	
75%	1.519157	13.825000	3.600000	1.630000	73.087500	0.610000	
max	1.533930	17.380000	4.490000	3.500000	75.410000	6.210000	

	Ca	Ba	Fe
count	214.000000	214.000000	214.000000
mean	8.956963	0.175047	0.057009
std	1.423153	0.497219	0.097439
min	5.430000	0.000000	0.000000
25%	8.240000	0.000000	0.000000
50%	8.600000	0.000000	0.000000
75%	9.172500	0.000000	0.100000
max	16.190000	3.150000	0.510000







```
[9]: perform_analysis(dataset_name='wine')
```

```
#####
DATASET NAME: wine
```

#### CLASS DISTRIBUTION ANALYSIS:

	class_name	class_idx	class_count	class_perc
0	class_2	2	71	39.887640
1	class_1	1	59	33.146067
2	class_3	3	48	26.966292

#### DATASET ATTRIBUTES:

```
['alcohol', 'malic_acid', 'ash', 'alcalinity_of_ash', 'magnesium',
'total_phenols', 'flavanoids', 'nonflavanoid_phenols', 'proanthocyanins',
```

```
'color_intensity', 'hue', 'OD280/OD315_of_diluted_wines', 'proline']
```

DATASET ANALYSIS:

	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	\
count	178.000000	178.000000	178.000000	178.000000	178.000000	
mean	13.000618	2.336348	2.366517	19.494944	99.741573	
std	0.811827	1.117146	0.274344	3.339564	14.282484	
min	11.030000	0.740000	1.360000	10.600000	70.000000	
25%	12.362500	1.602500	2.210000	17.200000	88.000000	
50%	13.050000	1.865000	2.360000	19.500000	98.000000	
75%	13.677500	3.082500	2.557500	21.500000	107.000000	
max	14.830000	5.800000	3.230000	30.000000	162.000000	
	total_phenols	flavanoids	nonflavanoid_phenols	proanthocyanins	\	
count	178.000000	178.000000	178.000000	178.000000	178.000000	
mean	2.295112	2.029270	0.361854	1.590899		
std	0.625851	0.998859	0.124453	0.572359		
min	0.980000	0.340000	0.130000	0.410000		
25%	1.742500	1.205000	0.270000	1.250000		
50%	2.355000	2.135000	0.340000	1.555000		
75%	2.800000	2.875000	0.437500	1.950000		
max	3.880000	5.080000	0.660000	3.580000		
	color_intensity	hue	OD280/OD315_of_diluted_wines	proline		
count	178.000000	178.000000	178.000000	178.000000	178.000000	
mean	5.058090	0.957449	2.611685	746.893258		
std	2.318286	0.228572	0.709990	314.907474		
min	1.280000	0.480000	1.270000	278.000000		
25%	3.220000	0.782500	1.937500	500.500000		
50%	4.690000	0.965000	2.780000	673.500000		
75%	6.200000	1.120000	3.170000	985.000000		
max	13.000000	1.710000	4.000000	1680.000000		

