

Лабораторная работа №5
по дисциплине
«Технологии машинного обучения»
на тему
«Линейные модели, SVM и деревья решений»

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1. Лабораторная работа №5. Линейные модели, SVM и деревья решений.

```
[0]: import numpy as np
import pandas as pd
from sklearn.datasets import load_wine
from sklearn.model_selection import train_test_split
```

1.1. Загрузка набора данных. Разбиение на тестовую и обучающую выборки

```
[0]: X, y = load_wine(return_X_y=True)
X_train, X_test, y_train, y_test = train_test_split(X, y,
↪ random_state=42)
```

```
[3]: print(X_train.shape)
print(X_test.shape)
print(y_train.shape)
print(y_test.shape)
```

```
(133, 13)
(45, 13)
(133,)
(45,)
```

```
[0]: from sklearn.svm import NuSVC
from sklearn.metrics import classification_report

# Отчёт о метриках модели
def test_model(model, x, y):
    results = model.predict(x)
    return classification_report(y, results)
```

1.2. Классификатор на методе опорных векторов

```
[5]: nu_svc = NuSVC(nu=0.5)
nu_svc.fit(X_train, y_train)
print(test_model(nu_svc, X_test, y_test))
```

	precision	recall	f1-score	support
0	1.00	1.00	1.00	15
1	1.00	1.00	1.00	18
2	1.00	1.00	1.00	12
accuracy			1.00	45

macro avg	1.00	1.00	1.00	45
weighted avg	1.00	1.00	1.00	45

1.3. Дерево решений

```
[0]: from sklearn.tree import DecisionTreeClassifier
```

```
[7]: dtc = DecisionTreeClassifier(max_depth=None)
dtc.fit(X_train, y_train)
print("Всего листьев: {}".format(dtc.get_n_leaves()))
print("Глубина дерева: {}".format(dtc.get_depth()))
print(test_model(dtc, X_test, y_test))
```

Всего листьев: 7
Глубина дерева: 4

	precision	recall	f1-score	support
0	0.93	0.93	0.93	15
1	0.95	1.00	0.97	18
2	1.00	0.92	0.96	12
accuracy			0.96	45
macro avg	0.96	0.95	0.95	45
weighted avg	0.96	0.96	0.96	45

1.4. Визуализация дерева решений

```
[0]: from sklearn.tree import plot_tree
from matplotlib import pyplot as plt
```

```
[10]: plt.figure(figsize=[15, 15])
plot_tree(dtc, feature_names=load_wine()["feature_names"], filled=False)

[10]: [Text(418.5, 733.86, 'color_intensity <= 3.82\ngini = 0.658\nsamples =
133\nvalue = [44, 53, 36]'),
Text(251.10000000000002, 570.78, 'proline <= 1010.0\ngini = 0.
↪08\nsamples =
48\nvalue = [2, 46, 0]'),
Text(167.4, 407.70000000000005, 'ash <= 3.07\ngini = 0.042\nsamples =
↪47\nvalue
= [1, 46, 0]'),
Text(83.7, 244.62, 'gini = 0.0\nsamples = 46\nvalue = [0, 46, 0]'),
Text(251.10000000000002, 244.62, 'gini = 0.0\nsamples = 1\nvalue = [1,
↪0, 0]'),
Text(334.8, 407.70000000000005, 'gini = 0.0\nsamples = 1\nvalue = [1,
↪0, 0]'),
```

```

Text(585.9, 570.78, 'flavanoids <= 1.4\ngini = 0.57\nsamples = 85\nvalue = [42,
7, 36]'),
Text(502.20000000000005, 407.70000000000005, 'gini = 0.0\nsamples = 36\nvalue =
[0, 0, 36]'),
Text(669.6, 407.70000000000005, 'proline <= 724.5\ngini = 0.
245\nsamples =
49\nvalue = [42, 7, 0]'),
Text(585.9, 244.62, 'alcohol <= 13.145\ngini = 0.219\nsamples = 8\nvalue = [1,
7, 0]'),
Text(502.20000000000005, 81.54000000000008, 'gini = 0.0\nsamples = 7\nvalue =
[0, 7, 0]'),
Text(669.6, 81.54000000000008, 'gini = 0.0\nsamples = 1\nvalue = [1, 0, 0]'),
Text(753.3000000000001, 244.62, 'gini = 0.0\nsamples = 41\nvalue = [41, 0,
0]')

```

