

ALGORITMA NAIVE BAYES PREDIKSI DIABETES

Margareta Valencia (A11.2022.14704)

1. Import Library

```
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
from sklearn import metrics
```

2. Import Dataset

```
dataset = pd.read_csv('diabetes.csv')
X = dataset.iloc[:, [2, 3]].values
y = dataset.iloc[:, -1].values
```

```
print(X)
```

```
[[72 35]
 [66 29]
 [64  0]
 ...
 [72 23]
 [60  0]
 [70 31]]
```

```
print(y)
```

```
[1 0 1 0 1 0 1 0 1 1 0 1 0 1 1 1 1 0 1 0 0 1 1 1 1 1 0 0 0 0 1 0 0 0
0 0
 1 1 1 0 0 0 1 0 1 0 0 1 0 0 0 0 1 0 0 1 0 0 0 0 1 0 0 1 0 1 0 0 0 1 0
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```

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0 1
1 0 0 1 0 0 1 0 1 1 1 0 0 1 1 1 0 1 0 1 0 1 0 0 0 0 0 1 0]

```

```
dataset.info()
```

```
<class 'pandas.core.frame.DataFrame'>
```

```
RangeIndex: 768 entries, 0 to 767
```

```
Data columns (total 9 columns):
```

#	Column	Non-Null Count	Dtype
0	Pregnancies	768 non-null	int64
1	Glucose	768 non-null	int64
2	BloodPressure	768 non-null	int64
3	SkinThickness	768 non-null	int64
4	Insulin	768 non-null	int64
5	BMI	768 non-null	float64
6	DiabetesPedigreeFunction	768 non-null	float64
7	Age	768 non-null	int64
8	Outcome	768 non-null	int64

```
dtypes: float64(2), int64(7)
```

```
memory usage: 54.1 KB
```

```
dataset.isna().sum()
```

Pregnancies	0
Glucose	0
BloodPressure	0
SkinThickness	0
Insulin	0
BMI	0

```
DiabetesPedigreeFunction    0
Age                         0
Outcome                     0
dtype: int64
```

#untuk model Gaussian hilangkan kolom diskrit

```
dataset = pd.DataFrame(dataset.drop('Pregnancies',axis=1))
dataset
```

	Glucose	BloodPressure	SkinThickness	Insulin	BMI	\
0	148	72	35	0	33.6	
1	85	66	29	0	26.6	
2	183	64	0	0	23.3	
3	89	66	23	94	28.1	
4	137	40	35	168	43.1	
..	
763	101	76	48	180	32.9	
764	122	70	27	0	36.8	
765	121	72	23	112	26.2	
766	126	60	0	0	30.1	
767	93	70	31	0	30.4	

	DiabetesPedigreeFunction	Age	Outcome
0	0.627	50	1
1	0.351	31	0
2	0.672	32	1
3	0.167	21	0
4	2.288	33	1
..
763	0.171	63	0
764	0.340	27	0
765	0.245	30	0
766	0.349	47	1
767	0.315	23	0

```
[768 rows x 8 columns]
```

```
dataset.describe()
```

	Glucose	BloodPressure	SkinThickness	Insulin
BMI				
\				
count	768.000000	768.000000	768.000000	768.000000
mean	120.894531	69.105469	20.536458	79.799479
std	31.972618	19.355807	15.952218	115.244002
min	0.000000	0.000000	0.000000	0.000000
25%	99.000000	62.000000	0.000000	0.000000

27.300000				
50%	117.000000	72.000000	23.000000	30.500000
32.000000				
75%	140.250000	80.000000	32.000000	127.250000
36.600000				
max	199.000000	122.000000	99.000000	846.000000
67.100000				

	DiabetesPedigreeFunction	Age	Outcome
count	768.000000	768.000000	768.000000
mean	0.471876	33.240885	0.348958
std	0.331329	11.760232	0.476951
min	0.078000	21.000000	0.000000
25%	0.243750	24.000000	0.000000
50%	0.372500	29.000000	0.000000
75%	0.626250	41.000000	1.000000
max	2.420000	81.000000	1.000000

3. Pembagian Dataset

```
# Membagi dataset menjadi data pelatihan dan data pengujian
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size =
0.35, random_state = 0)
```

4. Feature Scaling

```
from sklearn.preprocessing import StandardScaler
sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

print(X_train)
```

```
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 [ 0.01459831  0.69644524]
 [-0.24556436 -1.30803161]
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 [ 0.22272844 -0.11787348]
 [ 0.22272844  1.19756445]
 [ 0.11866337  0.44588563]]
```

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```

```
print(X_test)
```

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```

5. Pelatihan Model Naive Bayes Pada Data Pelatihan

```
from sklearn.naive_bayes import GaussianNB
classifier = GaussianNB()
classifier.fit(X_train, y_train)

GaussianNB()
```

6. Prediksi Hasil Data Pengujian

```
y_pred = classifier.predict(X_test)
print("Accuracy: ", metrics.accuracy_score(y_test, y_pred))

Accuracy:  0.6765799256505576
```

7. Evaluasi Model

```
# Menghitung dan menampilkan confusion matrix dan akurasi
from sklearn.metrics import confusion_matrix, accuracy_score
cm = confusion_matrix(y_test, y_pred)
print("\nConfusion Matrix:")
print(cm)
print("\nAkurasi:", accuracy_score(y_test, y_pred))

Confusion Matrix:
[[176   4]
```

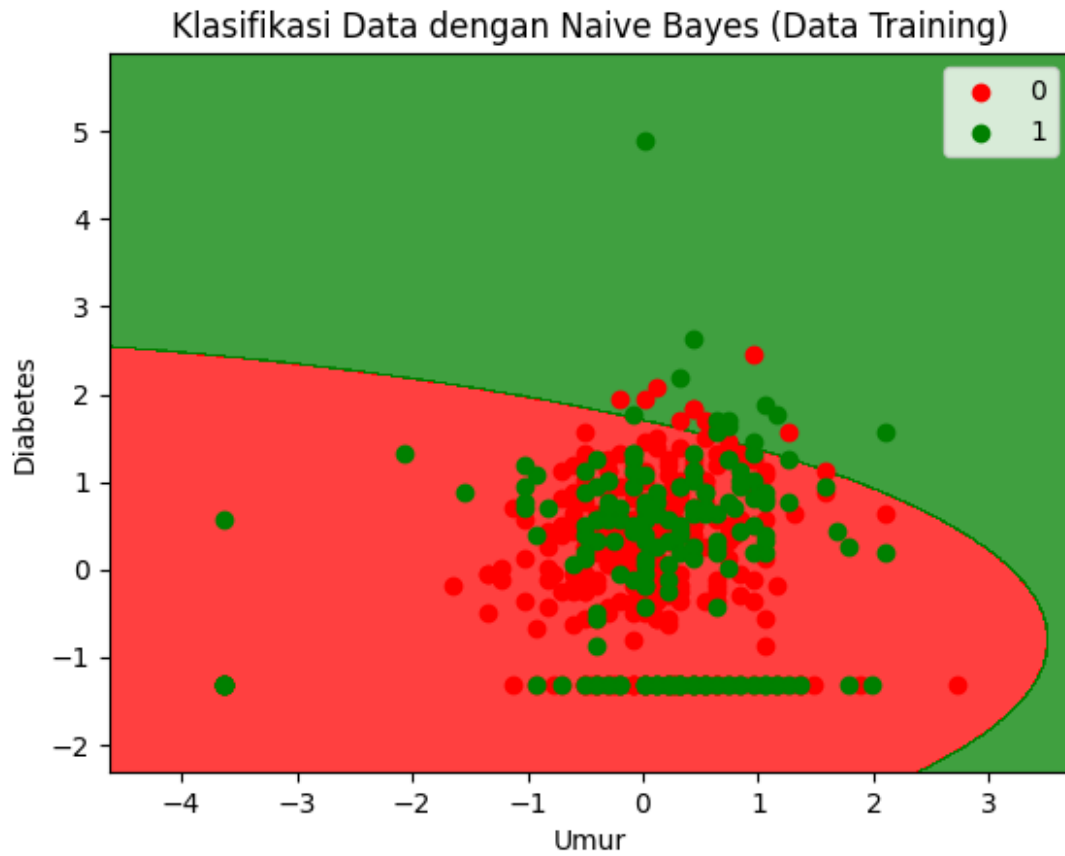
```
[ 83  6]]
```

Akurasi: 0.6765799256505576

8. Visualisasi Hasil Data Pelatihan

```
from matplotlib.colors import ListedColormap
X_set, y_set = X_train, y_train
X1, X2 = np.meshgrid(np.arange(start = X_set[:, 0].min() - 1, stop =
X_set[:, 0].max() + 1, step = 0.01),
                     np.arange(start = X_set[:, 1].min() - 1, stop =
X_set[:, 1].max() + 1, step = 0.01))
plt.contourf(X1, X2, classifier.predict(np.array([X1.ravel(),
X2.ravel()]).T).reshape(X1.shape),
             alpha = 0.75, cmap = ListedColormap(('red', 'green')))
plt.xlim(X1.min(), X1.max())
plt.ylim(X2.min(), X2.max())
for i, j in enumerate(np.unique(y_set)):
    plt.scatter(X_set[y_set == j, 0], X_set[y_set == j, 1],
               c = ListedColormap(('red', 'green'))(i), label = j)
plt.title('Klasifikasi Data dengan Naive Bayes (Data Training)')
plt.xlabel('Umur')
plt.ylabel('Diabetes')
plt.legend()
plt.show()
```

```
C:\Users\Acer\AppData\Local\Temp\ipykernel_53388\324515528.py:10:
UserWarning: *c* argument looks like a single numeric RGB or RGBA
sequence, which should be avoided as value-mapping will have
precedence in case its length matches with *x* & *y*. Please use the
*color* keyword-argument or provide a 2D array with a single row if
you intend to specify the same RGB or RGBA value for all points.
plt.scatter(X_set[y_set == j, 0], X_set[y_set == j, 1],
```



9. Visualisasi Hasil Data Pengujian

```
from matplotlib.colors import ListedColormap
X_set, y_set = X_test, y_test
X1, X2 = np.meshgrid(np.arange(start = X_set[:, 0].min() - 1, stop =
X_set[:, 0].max() + 1, step = 0.01),
                     np.arange(start = X_set[:, 1].min() - 1, stop =
X_set[:, 1].max() + 1, step = 0.01))
plt.contourf(X1, X2, classifier.predict(np.array([X1.ravel(),
X2.ravel()]).T).reshape(X1.shape),
             alpha = 0.75, cmap = ListedColormap(('red', 'green')))
plt.xlim(X1.min(), X1.max())
plt.ylim(X2.min(), X2.max())
for i, j in enumerate(np.unique(y_set)):
    plt.scatter(X_set[y_set == j, 0], X_set[y_set == j, 1],
                c = ListedColormap(('red', 'green'))(i), label = j)
plt.title('Klasifikasi Data dengan Naive Bayes (Data Testing)')
plt.xlabel('Umur')
plt.ylabel('Diabetes')
plt.legend()
plt.show()
```

```
C:\Users\Acer\AppData\Local\Temp\ipykernel_53388\369509195.py:10:  
UserWarning: *c* argument looks like a single numeric RGB or RGBA  
sequence, which should be avoided as value-mapping will have  
precedence in case its length matches with *x* & *y*. Please use the  
*color* keyword-argument or provide a 2D array with a single row if  
you intend to specify the same RGB or RGBA value for all points.  
plt.scatter(X_set[y_set == j, 0], X_set[y_set == j, 1],
```

