Medical Diagnosis with Naive Bayes

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Introduction

- Medical diagnosis is a critical domain where accurate and timely assessments can significantly impact patient outcomes.
- In this case study, we explore the application of the Naive Bayes algorithm to develop a diagnostic system for a medical research institute.
- Our objective is to create a classification model that aids in the medical diagnosis process by leveraging a dataset containing various medical test results, patient information, and corresponding diagnoses.

Data Exploration:

- Begin with dataset loading using Pandas.
- Features include medical test results and patient info, labels are diagnoses.
- Distribution of diagnoses: Analyze counts of each diagnosis for class balance.

```
data = pd.read_csv('data/heartdisease.csv')
data.head()
```

	age	sex	ср	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	ca	thal	targ
0	63	1	3	145	233	1	0	150	0	2.3	0	0	1	
1	37	1	2	130	250	0	1	187	0	3.5	0	0	2	
2	41	0	1	130	204	0	0	172	0	1.4	2	0	2	
3	56	1	1	120	236	0	1	178	0	0.8	2	0	2	
4	57	0	0	120	354	0	1	163	1	0.6	2	0	2	
•														-

x = data.iloc[:,:-1].values
y = data.iloc[:,-1:].values

Data Preprocessing

- Handle missing values: Impute or remove data points with missing values.
- Normalize or scale features if required for consistent magnitude.
- Encode categorical variables into numerical format.
- Calculate prior probabilities: P(Condition) and P(No Condition) based on class distribution.

Implementing Naive Bayes

- Implement Gaussian, Multinomial and Bernoulli Naive Bayes using scikit-learn.
- Split dataset into training and testing sets to evaluate the models.
- Feature Engineering:
 - Convert test results and patient info into features: Use one-hot encoding or numerical representation.
 - Importance of feature selection: Reducing dimensionality can improve model performance and interpretability.

```
# Gaussian Naive Baye
gauss nb = GaussianNB()
gauss nb.fit(xtrain,ytrain)
ypred = gauss nb.predict(xtest)
# Multinomial Naive Baye
from sklearn.naive bayes import MultinomialNB
multi nb = MultinomialNB()
multi nb.fit(xtrain,ytrain)
ypred = multi nb.predict(xtest)
#Bernoulli Naive Baye
from sklearn.naive bayes import BernoulliNB
ber nb = BernoulliNB()
ber nb.fit(xtrain,ytrain)
ypred = ber nb.predict(xtest)
```

Model Training and Model Evaluation

- Train Naive Bayes model on feature-engineered dataset.
- Probability estimation in Naive Bayes: Use conditional probabilities to calculate likelihood and apply Bayes' theorem.
- Assess model using evaluation metrics: Accuracy, precision, recall, F1-score.
- Interpret results: Discuss the model's ability to correctly classify medical conditions.

```
Accuracy score :
                        0.8947368421052632
Confusion Martix:
 [[32 4]
 [ 4 36]]
classification Report :
               precision
                            recall f1-score
                                               support
           0
                   0.89
                             0.89
                                       0.89
                                                   36
                   0.90
                             0.90
                                       0.90
                                                   40
                                       0.89
                                                   76
    accuracy
   macro avg
                   0.89
                             0.89
                                       0.89
                                                   76
weighted avg
                   0.89
                             0.89
                                       0.89
                                                   76
```

Gaussian Naive Bayes has Higher accuracy than other naive bayes models

Laplace Smoothing

```
Accuracy score :
                         0.8157894736842105
Confusion Martix:
 [[27 9]
 [ 5 35]]
classification Report :
               precision
                            recall f1-score
                                                support
           0
                   0.84
                             0.75
                                       0.79
                                                    36
                   0.80
                             0.88
                                       0.83
                                                    40
                                       0.82
                                                    76
    accuracy
                   0.82
                             0.81
                                       0.81
                                                    76
   macro avg
weighted avg
                                                    76
                   0.82
                             0.82
                                       0.81
```



Real-World Application

Importance of accurate medical diagnosis: Crucial for patient treatment and outcomes.

Practical implications of Naive Bayes: Faster, interpretable, and useful for initial screening or decision support in healthcare.

Model Limitations

Potential limitations of Naive Bayes: Assumes independence, may not capture complex relationships.

Scenarios where it may not perform well: Complex, high-dimensional data with strong dependencies.

Conclusion

- The application of the Naive Bayes algorithm in the context of medical diagnosis offers a valuable tool for healthcare and research.
- Through this case study, we have seen the importance of thorough data exploration, preprocessing, and feature engineering to prepare medical data for modeling.
- The use of Multinomial and Bernoulli Naive Bayes variants, as implemented in scikit-learn, allows for effective classification.