

decision-tree

December 18, 2023

0.1 This is a project of drug prediction using decision tree algorithm

0.1.1 Importing necessary libraries

```
[25]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
```

```
[26]: df = pd.read_csv("drug.csv")
```

```
[27]: df.head(5)
```

```
[27]:
```

| | Age | Sex | BP | Cholesterol | Na_to_K | Drug |
|---|-----|-----|--------|-------------|---------|-------|
| 0 | 23 | F | HIGH | HIGH | 25.355 | drugY |
| 1 | 47 | M | LOW | HIGH | 13.093 | drugC |
| 2 | 47 | M | LOW | HIGH | 10.114 | drugC |
| 3 | 28 | F | NORMAL | HIGH | 7.798 | drugX |
| 4 | 61 | F | LOW | HIGH | 18.043 | drugY |

0.1.2 Converting string values to numeric values for model fitting using LabelEncoder()

```
[28]: from sklearn.preprocessing import LabelEncoder
```

```
[29]: le = LabelEncoder()
```

```
[30]: df.Sex = le.fit_transform(df.Sex)
df.BP = le.fit_transform(df.BP)
df.Cholesterol = le.fit_transform(df.Cholesterol)
df.Drug = le.fit_transform(df.Drug)
```

```
[31]: df.head(5)
```

```
[31]:
```

| | Age | Sex | BP | Cholesterol | Na_to_K | Drug |
|---|-----|-----|----|-------------|---------|------|
| 0 | 23 | 0 | 0 | 0 | 25.355 | 4 |
| 1 | 47 | 1 | 1 | 0 | 13.093 | 2 |
| 2 | 47 | 1 | 1 | 0 | 10.114 | 2 |
| 3 | 28 | 0 | 2 | 0 | 7.798 | 3 |

```
4    61    0    1          0    18.043    4
```

0.1.3 Separating data into input and target variables

```
[36]: inputs = df.drop(["Drug"],axis = 'columns')
      inputs
```

```
[36]:
```

| | Age | Sex | BP | Cholesterol | Na_to_K |
|-----|-----|-----|----|-------------|---------|
| 0 | 23 | 0 | 0 | 0 | 25.355 |
| 1 | 47 | 1 | 1 | 0 | 13.093 |
| 2 | 47 | 1 | 1 | 0 | 10.114 |
| 3 | 28 | 0 | 2 | 0 | 7.798 |
| 4 | 61 | 0 | 1 | 0 | 18.043 |
| .. | ... | ... | .. | ... | ... |
| 195 | 56 | 0 | 1 | 0 | 11.567 |
| 196 | 16 | 1 | 1 | 0 | 12.006 |
| 197 | 52 | 1 | 2 | 0 | 9.894 |
| 198 | 23 | 1 | 2 | 1 | 14.020 |
| 199 | 40 | 0 | 1 | 1 | 11.349 |

```
[200 rows x 5 columns]
```

```
[38]: targets = df["Drug"]
      targets
```

```
[38]:
```

| | |
|-----|----|
| 0 | 4 |
| 1 | 2 |
| 2 | 2 |
| 3 | 3 |
| 4 | 4 |
| .. | .. |
| 195 | 2 |
| 196 | 2 |
| 197 | 3 |
| 198 | 3 |
| 199 | 3 |

```
Name: Drug, Length: 200, dtype: int32
```

0.1.4 Model fitting

```
[39]: from sklearn import tree
```

```
[40]: model = tree.DecisionTreeClassifier()
```

```
[41]: model.fit(inputs,targets)
```

```
[41]: DecisionTreeClassifier()
```

```
[42]: model.score(inputs,targets)
```

```
[42]: 1.0
```

```
[43]: model.predict([[23,0,0,0,25.355]])
```

```
C:\Users\luhar\anaconda3\lib\site-packages\sklearn\base.py:465: UserWarning: X
does not have valid feature names, but DecisionTreeClassifier was fitted with
feature names
```

```
warnings.warn(
```

```
[43]: array([4])
```

0.1.5 Here we got model score 1 which is excellent and we predicted the drugY(4) for 25 year old female with high BP, high cholesterol and 25.355 unit amount of sodium_pottasium

0.1.6 Prediction by splitting data into testing and training

```
[44]: from sklearn.model_selection import train_test_split
```

```
[48]: xtrain, xtest, ytrain, ytest = train_test_split(inputs, targets, test_size=0.3)
```

```
[49]: len(xtrain)
```

```
[49]: 140
```

```
[50]: len(xtest)
```

```
[50]: 60
```

```
[51]: model.fit(xtrain,ytrain)
```

```
[51]: DecisionTreeClassifier()
```

```
[52]: model.score(xtrain,ytrain)
```

```
[52]: 1.0
```

```
[54]: model.predict(xtest)
```

```
[54]: array([4, 4, 0, 3, 2, 0, 4, 1, 0, 2, 1, 0, 3, 4, 4, 2, 4, 3, 4, 4, 4, 4,
         4, 3, 3, 3, 3, 3, 3, 1, 0, 3, 3, 1, 2, 4, 0, 4, 3, 3, 4, 4, 4, 4,
         2, 2, 4, 0, 3, 4, 1, 3, 4, 2, 4, 0, 3, 4, 3, 2])
```

0.1.7 After splitting also we got the model score 1. We predicted the drug using our input testing data and it matches 100% with target testing data

[]: