```
In [11]: import pandas as pd
          from sklearn.model_selection import train_test_split
          from sklearn.preprocessing import OrdinalEncoder
          from sklearn.preprocessing import LabelEncoder
          from sklearn.feature_selection import SelectKBest
          from sklearn.feature_selection import chi2
          import matplotlib.pyplot as plt
          from sklearn.feature_selection import mutual_info_classif
          from sklearn.feature_selection import f_classif
In [3]: data = pd.read csv("drug200.csv")
 In [4]: dataset = data.values
          dataset
Out[4]: array([[23, 'F', 'HIGH', 'HIGH', 25.355, 'DrugY'],
                 [47, 'M', 'LOW', 'HIGH', 13.093, 'drugC'], [47, 'M', 'LOW', 'HIGH', 10.114, 'drugC'],
                 . . . ,
                 [52, 'M', 'NORMAL', 'HIGH', 9.894, 'drugX'],
                 [23, 'M', 'NORMAL', 'NORMAL', 14.02, 'drugX'],
                 [40, 'F', 'LOW', 'NORMAL', 11.349, 'drugX']], dtype=object)
 In [5]: X = dataset[:, :-1]
          Х
```

```
Out[5]: array([[23, 'F', 'HIGH', 'HIGH', 25.355],
                [47, 'M', 'LOW', 'HIGH', 13.093],
                [47, 'M', 'LOW', 'HIGH', 10.114],
                [28, 'F', 'NORMAL', 'HIGH', 7.798],
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                [22,
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                [43, 'M', 'LOW', 'NORMAL', 19.368],
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                [16, 'F', 'HIGH', 'NORMAL', 15.516],
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                     'F', 'LOW', 'NORMAL', 22.697],
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                [67, 'M', 'LOW', 'NORMAL', 20.693],
                [45, 'M', 'LOW', 'NORMAL', 8.37],
```

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          'NORMAL', 'HIGH', 13.884],
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```

```
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                                                                                            [46, 'F', 'HIGH', 'HIGH', 34.686],
                                                                                            [56, 'F', 'LOW', 'HIGH', 11.567],
                                                                                            [16, 'M', 'LOW', 'HIGH', 12.006],
                                                                                            [52, 'M', 'NORMAL', 'HIGH', 9.894],
                                                                                            [23, 'M', 'NORMAL', 'NORMAL', 14.02],
                                                                                            [40, 'F', 'LOW', 'NORMAL', 11.349]], dtype=object)
In [6]: Y = dataset[:, -1]
Out[6]: array(['DrugY', 'drugC', 'drugC', 'drugX', 'DrugY', 'drugX', 'DrugY',
                                                                                              'drugC', 'DrugY', 'DrugY', 'drugC', 'DrugY', 'DrugY', 'DrugY',
                                                                                            'drugX', 'DrugY', 'drugX', 'drugA', 'drugC', 'DrugY', 'DrugY', 'DrugY', 'DrugY', 'DrugY', 'DrugY', 'DrugY', 'DrugY', 'DrugY', 'drugX', 'DrugY', 'DrugY', 'drugX', 'dr
                                                                                            'DrugY', 'drugX', 'drugX', 'drugA', 'drugC', 'DrugY',
                                                                                              'DrugY', 'DrugY', 'drugX', 'DrugY', 'DrugY', 'drugB', 'drugC',
                                                                                              'drugB', 'DrugY', 'drugX', 'DrugY', 'DrugY', 'drugA', 'DrugY', 'drugX', 'drugX', 'DrugY', 'drugX', 'DrugY', 'Dr
                                                                                             'drugB', 'DrugY', 'drugX', 'DrugY', 'DrugY', 'DrugY', 'drugA',
                                                                                             'DrugY', 'drugA', 'drugX', 'drugB', 'drugX', 'drugC', 'drugA',
                                                                                              'drugC', 'drugB', 'drugX', 'DrugY', 'DrugY', 'DrugY', 'DrugY',
                                                                                              'DrugY', 'DrugY', 'DrugY', 'drugX', 'DrugY', 'DrugY',
                                                                                            'DrugY', 'DrugY', 'drugA', 'drugA', 'drugC', 'drugX', 'DrugY', 'drugX', 'DrugY', 'drugX', 'DrugY', 'DrugY', 'DrugY', 'DrugY', 'drugB', 'DrugY', 'DrugY', 'DrugY', 'DrugY', 'DrugY', 'DrugY', 'drugB', 'DrugY', 'Dr
                                                                                              'DrugY', 'drugX', 'DrugY', 'drugX', 'DrugY', 'DrugY', 'drugX',
                                                                                              'DrugY', 'DrugY', 'drugX', 'drugB', 'drugA', 'drugB', 'drugX',
                                                                                              'drugA', 'DrugY', 'drugB', 'DrugY', 'drugA', 'drugX', 'drugX', 'drugX', 'drugA', 'drugX', 'dr
                                                                                             'DrugY', 'drugC', 'drugA', 'DrugY', 'drugC', 'drugX', 'drugX',
                                                                                              'drugB', 'drugX', 'DrugY', 'DrugY', 'DrugY', 'DrugY', 'drugX',
                                                                                              'DrugY', 'drugA', 'drugX', 'drugX', 'DrugY', 'DrugY', 'drugA',
                                                                                              'DrugY', 'drugA', 'DrugY', 'DrugY', 'DrugY', 'DrugY', 'drugX',
                                                                                            'drugX', 'DrugY', 'DrugY', 'drugB', 'drugA', 'DrugY', 'DrugY', 'drugC', 'DrugY', 'drugC', 'drugC', 'drugC', 'drugC', 'drugC', 'drugC', 'drugX', 'drugX'], dtype=object)
In [7]: X_new = pd.DataFrame(X)
                                                  y new = pd.DataFrame(Y)
In [8]: X_{new} = X_{new.drop([0, 4], axis=1)}
                                                   X new
```

```
Out[8]:
                   HIGH
                           HIGH
                   LOW
                           HIGH
           1 M
           2 M
                   LOW
                           HIGH
              F NORMAL
                           HIGH
                    LOW
                           HIGH
         195
             F
                   LOW
                           HIGH
         196 M
                    LOW
                           HIGH
                NORMAL
                           HIGH
         197 M
         198 M NORMAL NORMAL
         199 F
                   LOW NORMAL
        200 rows × 3 columns
 In [9]:
        X_train, X_test, y_train, y_test = train_test_split(X_new, y_new, test_size=
In [10]: print(f"The Train shape of X : {X_train.shape} and y: {y_train.shape}")
         print(f"The Test shape of X: {X_test.shape} and y: {y_test.shape}")
         The Train shape of X : (134, 3) and y: (134, 1)
         The Test shape of X: (66, 3) and y: (66, 1)
In [17]: def prepare_inputs(X_train, X_test):
             ordinal_encoder = OrdinalEncoder()
             ordinal_encoder.fit(X_train)
             X_train_enc = ordinal_encoder.transform(X_train)
             X_test_enc = ordinal_encoder.transform(X_test)
             return X_train_enc, X_test_enc
In [18]:
         def prepare_target(y_train, y_test):
             label_encoder = LabelEncoder()
             label_encoder.fit(y_train)
             y_train_enc = label_encoder.transform(y_train)
             y_test_enc = label_encoder.transform(y_test)
             return y_train_enc, y_test_enc
In [19]: X_train_enc, X_test_enc = prepare_inputs(X_train, X_test)
         y_train_enc, y_test_enc = prepare_inputs(y_train, y_test)
In [20]:
         ANOVA F_statistic score based feature selection
In [23]:
         def select_features(X_train,y_train,X_test):
             fs=SelectKBest(score_func=f_classif,k=3)
             fs.fit(X_train,y_train)
             X_train_fs=fs.transform(X_train)
             X_test_fs=fs.transform(X_test)
             return X_train_fs,X_test_fs,fs
In [24]: | #Selecting best features and plotting
```

X train fs, X test fs, fs=select features (X train enc, y train enc, X test enc)

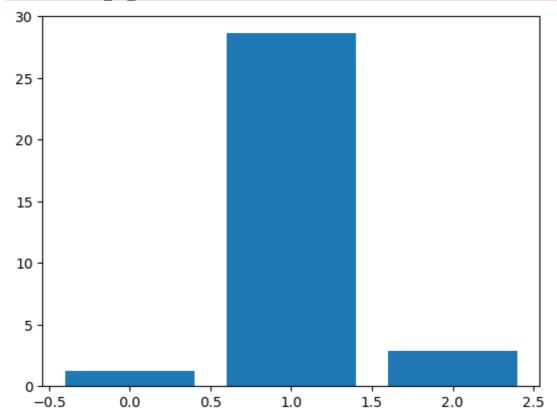
3

1

```
plt.bar([i for i in range(len(fs.scores_))],fs.scores_)
plt.show()
```

/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/utils/val idation.py:1143: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please change the shape of y to (n\_samples, ), for ex ample using ravel().

```
y = column_or_1d(y, warn=True)
```



```
In [25]: #feature scores fs.scores_
```

Out[25]: array([ 1.2711341 , 28.61641077, 2.8343434 ])

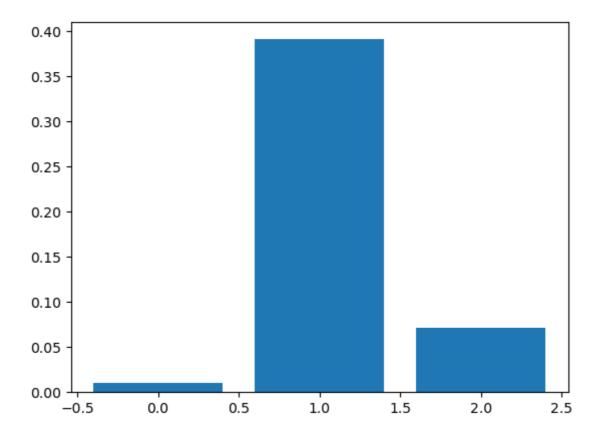
## Mutual information based feature selection

```
In [28]: def select_features_2(X_train,y_train,X_test):
    fs=SelectKBest(score_func=mutual_info_classif,k=3)
    fs.fit(X_train,y_train)
    X_train_fs=fs.transform(X_train)
    X_test_fs=fs.transform(X_test)
    return X_train_fs,X_test_fs,fs
```

```
In [30]: #Selecting best features using MI and plotting
X_train_fs2,X_test_fs2,fs2=select_features_2(X_train_enc, y_train_enc, X_test_plt.bar([i for i in range(len(fs2.scores_))],fs2.scores_)
    plt.show();
```

/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/utils/val idation.py:1143: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please change the shape of y to (n\_samples, ), for ex ample using ravel().

```
y = column_or_1d(y, warn=True)
```



```
In [31]: #feature scores fs2.scores_
```

Out[31]: array([0.00995744, 0.39129274, 0.07174999])

```
In [39]: from sklearn.ensemble import RandomForestClassifier
    from sklearn.metrics import accuracy_score

model1=RandomForestClassifier()
    model1.fit(X_train_enc,y_train_enc)
    yhat=model1.predict(X_test_enc)
    accuracy=accuracy_score(y_test_enc,yhat)
    print("Accuracy: %.2f" %(accuracy*100))
```

Accuracy: 51.52

/var/folders/nd/q79n0hd51vqcq6swmwgkc95w0000gn/T/ipykernel\_76679/3588750476. py:5: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please change the shape of y to (n\_samples,), for example using ravel().

model1.fit(X\_train\_enc,y\_train\_enc)

```
In [40]: model2=RandomForestClassifier()
    model2.fit(X_train_fs,y_train_enc)
    yhat=model1.predict(X_test_fs)
    accuracy=accuracy_score(y_test_enc,yhat)
    print("Accuracy: %.2f" %(accuracy*100))
```

Accuracy: 51.52

/var/folders/nd/q79n0hd51vqcq6swmwgkc95w0000gn/T/ipykernel\_76679/3067022059. py:2: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please change the shape of y to (n\_samples,), for example using ravel().

model2.fit(X\_train\_fs,y\_train\_enc)

```
In [42]: model3=RandomForestClassifier()
  model3.fit(X_train_fs2,y_train_enc)
  yhat=model1.predict(X_test_fs2)
```

```
accuracy_score(y_test_enc,yhat)
         print("Accuracy: %.2f" %(accuracy*100))
         Accuracy: 51.52
         /var/folders/nd/q79n0hd51vqcq6swmwgkc95w0000gn/T/ipykernel_76679/4190157872.
         py:2: DataConversionWarning: A column-vector y was passed when a 1d array wa
         s expected. Please change the shape of y to (n_samples,), for example using
         ravel().
           model3.fit(X_train_fs2,y_train_enc)
In [43]: from sklearn.pipeline import Pipeline
         from sklearn.model_selection import RepeatedStratifiedKFold
         from sklearn.model_selection import GridSearchCV
In [44]: def load_dataset(fname):
             data=pd.read_csv(fname, header=None)
             dataset=data.values
             X=dataset[:,:-1]
             y=dataset[:,-1]
             return X, y
In [45]: X, y = load_dataset("drug200.csv")
In [46]: #define the evaluation method - k-fold cross validation & k=10
         cv=RepeatedStratifiedKFold(n_splits=10,n_repeats=3,random_state=1)
In [47]: #define the pipeline to evaluate
         model=RandomForestClassifier()
         fs=SelectKBest(score_func=f_classif)
         pipeline=Pipeline(steps=[('anova',fs),('lr',model)])
In [48]: #define the grid
         grid=dict()
         grid['anova_k']=[i+1 for i in range(X.shape[1])]
In [50]: #define the grid search
         search=GridSearchCV(pipeline, grid, scoring='accuracy',n_jobs=-1, cv=cv)
         results=search.fit(X_train_enc, y_train_enc)
```

```
/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/utils/val
idation.py:1143: DataConversionWarning: A column-vector y was passed when a
1d array was expected. Please change the shape of y to (n_samples, ), for ex
ample using ravel().
 y = column or 1d(y, warn=True)
/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/pipeline.
py:405: DataConversionWarning: A column-vector y was passed when a 1d array
was expected. Please change the shape of y to (n_samples,), for example usin
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/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/model sel
ection/_validation.py:378: FitFailedWarning:
60 fits failed out of a total of 150.
The score on these train-test partitions for these parameters will be set to
If these failures are not expected, you can try to debug them by setting err
or_score='raise'.
Below are more details about the failures:
30 fits failed with the following error:
Traceback (most recent call last):
 File "/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/m
odel_selection/_validation.py", line 686, in _fit_and_score
    estimator.fit(X_train, y_train, **fit_params)
  File "/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/p
ipeline.py", line 401, in fit
    Xt = self. fit(X, y, **fit params steps)
                     ^^^^^
  File "/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/p
ipeline.py", line 359, in _fit
    X, fitted_transformer = fit_transform_one_cached(
  File "/Users/selva/enter/envs/torch/lib/python3.11/site-packages/joblib/me
mory.py", line 349, in __call_
    return self.func(*args, **kwargs)
           ^^^^^
 File "/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/p
ipeline.py", line 893, in _fit_transform_one
    res = transformer.fit_transform(X, y, **fit_params)
          ______
 File "/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/u
tils/_set_output.py", line 140, in wrapped
    data_to_wrap = f(self, X, *args, **kwargs)
 File "/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/b
ase.py", line 881, in fit_transform
    return self.fit(X, y, **fit_params).transform(X)
  File "/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/f
eature_selection/_univariate_selection.py", line 471, in fit
    self._check_params(X, y)
  File "/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/f
eature_selection/_univariate_selection.py", line 672, in _check_params
    raise ValueError(
ValueError: k should be <= n_features = 3; got 4. Use k='all' to return all
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    Xt = self._fit(X, y, **fit_params_steps)
         ^^^^^
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             self. check params(X, y)
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         ValueError: k should be <= n_features = 3; got 5. Use k='all' to return all
         features.
           warnings.warn(some_fits_failed_message, FitFailedWarning)
         /Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/model sel
         ection/ search.py:952: UserWarning: One or more of the test scores are non-f
         inite: [0.51703297 0.45677656 0.43644689
                                                         nan
                                                                    nan 1
           warnings.warn(
         /Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/utils/val
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          self._final_estimator.fit(Xt, y, **fit_params_last_step)
In [51]: print('Best Mean Accuracy: %.3f' % results.best_score_)
         print('Best config %s' %results.best_params_)
         Best Mean Accuracy: 0.517
         Best config {'anova_k': 1}
In [56]: #comparing different no. of features selected using ANOVA f-test
         from numpy import mean
         from numpy import std
         from sklearn.model_selection import cross_val_score
         def evaluate model(model):
             cv=RepeatedStratifiedKFold(n_repeats=3,n_splits=10, random_state=1)
             scores=cross_val_score(model, X_test_enc, y_test_enc, scoring='accuracy', cv
             return scores
In [53]: #define the no. of features
         num_features=[i+1 for i in range(X.shape[1])]
```

File "/Users/selva/enter/envs/torch/lib/python3.11/site-packages/sklearn/p

```
In [59]: #enumerate each no. of feature
    results=list()
    for k in num_features:
    #create pipeline
        model=RandomForestClassifier()
        fs=SelectKBest(score_func=f_classif,k=3)
        pipeline=Pipeline(steps=[('anova',fs),('lr',model)])
In [60]: #evaluate the model
    scores=evaluate_model(pipeline)
    results.append(scores)
```

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ection/split.py:700: UserWarning: The least populated class in y has only 4
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ection/_split.py:700: UserWarning: The least populated class in y has only 4
members, which is less than n_splits=10.
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In [61]: print('#%d %.3f (%.3f)' %(k,mean(scores),std(scores)))