as g Acc per	sture filtering method that is univarient method does not give ganrantee to get the better accuracy. Since in univarient method cortants are calculated indivisualy instaed of in a group. So, sometimes what happen top best performing features doesn't performing of some other method. Sturacy methods features importance are calculated individually instead of in a group so sometimes what happened at top 10 beforming features doesn't perform well as grouping of some other methods so what happens in the filtering methods sometime er into by selecting suboptimal features but filtering are univariate methods are quite fast and that those can be used as a screen
and afte less me	er into by selecting suboptimal features but filtering are univariate methods are quite fast and that those can be used as a screet sometimes you might get a better accuracy but most of the time you would get a better the training time that introduced trailer selecting some particular feature set so by keeping those things in mind you should read this lessons and one more thing in sons I was talking about the wrapper method as well as the imported methods which are better than the filter methods but who thods are very costly go ahead and I'll tell you to take you through this lessons for the first will be talking about are the classificablem in which shall be working on sent Android data set then first I'll remove the constant and duplicate features and then after remove all calculate the mutual information and then based on the mutual information Sal select top 10 top 10 percentage and
pro info staf in t	centile of the features which is the 23 features and then I build model and then finally I'll train it and after that I will work on regulate which is the best and data regression problem using Boston data set and using mutual information Sal calculate the in mormation gain on the individual features and then finally I select top 9 features which are to be stopped 9 features after selecting top 9 features and I calculate the performance compare the performance of the selected features and the overall features so his lesson. assification Problem Using Mutual Information Gain.
in in in %1	<pre>mport numpy as np mport matplotlib.pyplot as plt mport pandas as pd mport seaborn as sns matplotlib inline rom sklearn.feature_selection import mutual_info_classif,mutual_info_regression,SelectPercentile,Sel rom sklearn.model selection import train test split</pre>
f: f: f: d:	<pre>rom sklearn.ensemble import RandomForestClassifier,RandomForestRegressor rom sklearn.feature_selection import VarianceThreshold rom sklearn.preprocessing import LabelEncoder rom sklearn.metrics import accuracy_score,mean_squared_error,r2_score ata = pd.read_csv('titanic.csv') ata.head()</pre>
0 1 2 3	survivedpclasssexagesibspparchfareembarkedclasswhoadult_maledeckembark_townalivealone03male22.0107.2500SThirdmanTrueNaNSouthamptonnoFalse11female38.01071.2833CFirstwomanFalseCCherbourgyesFalse13female26.007.9250SThirdwomanFalseNaNSouthamptonyesTrue11female35.01053.1000SFirstwomanFalseCSouthamptonyesFalse03male35.008.0500SThirdmanTrueNaNSouthamptonnoTrue
do do do do	ata.dropna(inplace=True) abel = LabelEncoder() ata['sex'] = label.fit_transform(data['sex']) ata['embarked'] = label.fit_transform(data['embarked']) ata['class'] = label.fit_transform(data['embarked'])
da da da	<pre>ata['class'] = label.fit_transform(data['embarked']) ata['who']=label.fit_transform(data['adult_male']) ata['adult_male']=label.fit_transform(data['embark_town']) ata['embark_town']=label.fit_transform(data['embark_town']) ata['alive']=label.fit_transform(data['alive']) ata['alone'] = label.fit_transform(data['alone'])</pre> el data['deck']
x X:	<pre>= data.drop(labels='alive',axis=1) = data['alive'] .shape,y.shape 182, 13), (182,)) train,x test,y train,y test = train test split(x,y,train size=0.8,random state=0,stratify=y)</pre>
x ((train.shape,x_test.shape _train.shape,x_test.shape 145, 13), (37, 13)) move the constant,quaisi constant,duplicated onstant filter = VarianceThreshold(threshold=0.01)
X X (((<pre>onstant_filter.fit(x_train) train_filter=constant_filter.transform(x_train) test_filter= constant_filter.transform(x_test) _train_filter.shape,x_test_filter.shape 145, 13), (37, 13)) pollicated features Removal</pre>
x x x	_train_T = x_train_filter.T _test_T = x_test_filter.T _train_T = pd.DataFrame(x_train_T) _test_T = pd.DataFrame(x_test_T) uplicated_features = x_train_T.duplicated()
X X	<pre>eep_them = [not index for index in duplicated_features] _train_unique = x_train_T[keep_them].T _test_unique = x_test_T[keep_them].T _train_unique.shape,x_test_unique.shape 145, 11), (37, 11))</pre>
calc	<pre>culated mutual information i = mutual_info_classif(x_train_unique,y_train) en(mi)</pre>
If wance whi	i = pd.Series(mi) i.index = x_train_unique.columns i.sort_values(ascending= False, inplace = True) we see the graph here we can see that the few of the features are contributing the more information gain(individualy) than the column of the contribution of the features are added the contribution in mutual information gain and those who are contributing the more information are very important in this kind of classification. We are going to remove those features which minutly contributing any information of the features are added to the contribution of the features which minutly contributing any information of the features are added to the contribution of the features which minutly contributing any information of the features are contribution of the features which minutly contributing any information of the features are contribution of the features are contributed as a contribution of the features are contributed as a contribute of the features
info	ividually so for that we are going to select some k best or the Selectpercentile to get the those who are contributing the more ormation than the others. i.plot.bar(figsize = (15,6)) xesSubplot:>
0.6	5 -
0.3	
X	el = SelectPercentile(mutual_info_classif,percentile=90).fit(x_train_unique,y_train) _train_unique.columns[sel.get_support()]
9 x	<pre>t64Index([0, 1, 2, 3, 5, 6, 9, 10, 12], dtype='int64') en(x_train_unique.columns[sel.get_support()]) _train_mi = sel.transform(x_train_unique) _test_mi = sel.transform(x_test_unique)</pre>
((x	train_mi.shape,x_test_mi.shape 145, 9), (37, 9)) _train_mi.shape,x_test_mi.shape 145, 9), (37, 9))
#.	<pre>build a classifier ef randomforest(x_train, x_test, y_train, y_test): clf = RandomForestClassifier(random_state=0, n_estimators=1000, n_jobs=-1) clf.fit(x_train, y_train) y_pred = clf.predict(x_test) print('Accuracy :',accuracy_score(y_test, y_pred))</pre>
Ac Wa	<pre>print('Accuracy :',accuracy_score(y_test,y_pred)) %time andomforest(x_train_mi,x_test_mi,y_train,y_test) curacy : 1.0 ll time: 1.95 s %time andomforest(x_train,x_test,y_train,y_test)</pre>
Ac Wa Fro tim	
he #	elp(sel) this is how we can get the whole about it lp on SelectPercentile in module sklearn.feature_selectionunivariate_selection object: ass SelectPercentile(_BaseFilter) SelectPercentile(score_func= <function 0x000002d3ea398550="" at="" f_classif="">, *, percentile=10)</function>
	Select features according to a percentile of the highest scores. Read more in the :ref:`User Guide <univariate_feature_selection>`. Parameters score_func : callable, default=f_classif Function taking two arrays X and y, and returning a pair of arrays (scores, pvalues) or a single array with scores.</univariate_feature_selection>
	Default is f_classif (see below "See Also"). The default function only works with classification tasks. versionadded:: 0.18 percentile: int, default=10 Percent of features to keep. Attributes
	scores_: array-like of shape (n_features,) Scores of features. pvalues_: array-like of shape (n_features,) p-values of feature scores, None if `score_func` returned only scores. Examples
 	<pre>>>> from sklearn.datasets import load_digits >>> from sklearn.feature_selection import SelectPercentile, chi2 >>> X, y = load_digits(return_X_y=True) >>> X.shape (1797, 64) >>> X_new = SelectPercentile(chi2, percentile=10).fit_transform(X, y) >>> X_new.shape (1797, 7)</pre>
 	Notes Ties between features with equal scores will be broken in an unspecified way. See Also f_classif: ANOVA F-value between label/feature for classification tasks. mutual_info_classif: Mutual information for a discrete target.
 	<pre>chi2 : Chi-squared stats of non-negative features for classification tasks. f_regression : F-value between label/feature for regression tasks. mutual_info_regression : Mutual information for a continuous target. SelectKBest : Select features based on the k highest scores. SelectFpr : Select features based on a false positive rate test. SelectFdr : Select features based on an estimated false discovery rate. SelectFwe : Select features based on family-wise error rate. GenericUnivariateSelect : Univariate feature selector with configurable mode.</pre>
 	Method resolution order: SelectPercentile _BaseFilter sklearn.feature_selectionbase.SelectorMixin sklearn.base.TransformerMixin sklearn.base.BaseEstimator builtins.object Methods defined here:
	init(self, score_func= <function 0x000002d3ea398550="" at="" f_classif="">, *, percentile=10) Initialize self. See help(type(self)) for accurate signature. Data and other attributes defined here: abstractmethods = frozenset()</function>
 	Methods inherited from _BaseFilter: fit(self, X, y) Run score function on (X, y) and get the appropriate features. Parameters X: array-like of shape (n samples, n features)
	The training input samples. y: array-like of shape (n_samples,) The target values (class labels in classification, real numbers in regression). Returns self: object
 	Methods inherited from sklearn.feature_selectionbase.SelectorMixin: get_support(self, indices=False) Get a mask, or integer index, of the features selected Parameters
	<pre>indices : bool, default=False If True, the return value will be an array of integers, rather than a boolean mask. Returns support : array An index that selects the retained features from a feature vector. If `indices` is False, this is a boolean array of shape [# input features], in which an element is True iff its</pre>
 	corresponding feature is selected for retention. If `indices` is True, this is an integer array of shape [# output features] whose values are indices into the input feature vector. inverse_transform(self, X) Reverse the transformation operation Parameters
	<pre>X: array of shape [n_samples, n_selected_features] The input samples. Returns X_r: array of shape [n_samples, n_original_features] `X` with columns of zeros inserted where features would have been removed by :meth:`transform`. transform(self, X)</pre>
 	Reduce X to the selected features. Parameters X: array of shape [n_samples, n_features] The input samples. Returns
	<pre>X_r : array of shape [n_samples, n_selected_features]</pre>
	Fits transformer to `X` and `y` with optional parameters `fit_params` and returns a transformed version of `X`. Parameters X: array-like of shape (n_samples, n_features) Input samples. y: array-like of shape (n samples,) or (n samples, n outputs), default=None
 	Target values (None for unsupervised transformations). **fit_params : dict Additional fit parameters. Returns X_new : ndarray array of shape (n_samples, n_features_new) Transformed array.
 	Data descriptors inherited from sklearn.base.TransformerMixin: dict dictionary for instance variables (if defined) weakref list of weak references to the object (if defined)
	Methods inherited from sklearn.base.BaseEstimator: getstate(self) repr(self, N_CHAR_MAX=700) Return repr(self). setstate(self, state)
 	setstate(self, state) get_params(self, deep=True) Get parameters for this estimator. Parameters deep: bool, default=True If True, will return the parameters for this estimator and contained subobjects that are estimators.
	Returns params: dict Parameter names mapped to their values. set_params(self, **params) Set the parameters of this estimator.
 	The method works on simple estimators as well as on nested objects (such as :class: `~sklearn.pipeline.Pipeline`). The latter have parameters of the form `` <component><parameter>`` so that it's possible to update each component of a nested object. Parameters **params: dict Estimator parameters.</parameter></component>
	Returns self : estimator instance Estimator instance.
f:	<pre>utual Information Gain In Regression. rom sklearn import datasets rom sklearn.linear_model import LinearRegression rom sklearn.metrics import mean_squared_error, mean_absolute_error, r2_score oston = datasets.load_boston()</pre>
во	rint (boston.DESCR) _boston_dataset: ston house prices dataset
t.	:Attribute Information (in order): - CRIM per capita crime rate by town - ZN proportion of residential land zoned for lots over 25,000 sq.ft INDUS proportion of non-retail business acres per town
	- INDUS proportion of non-retail business acres per town - CHAS Charles River dummy variable (= 1 if tract bounds river; 0 otherwise) - NOX nitric oxides concentration (parts per 10 million) - RM average number of rooms per dwelling - AGE proportion of owner-occupied units built prior to 1940 - DIS weighted distances to five Boston employment centres - RAD index of accessibility to radial highways - TAX full-value property-tax rate per \$10,000 - PTRATIO pupil-teacher ratio by town - B 1000(Bk - 0.63)^2 where Bk is the proportion of black people by town
Th Th pr vo	
Th pr	e Boston house-price data has been used in many machine learning papers that address regression oblems. topic:: References - Belsley, Kuh & Welsch, 'Regression diagnostics: Identifying Influential Data and Sources of Colli ley, 1980. 244-261. - Quinlan, R. (1993). Combining Instance-Based and Model-Based Learning. In Proceedings on the Tenth
Her	- Quinlan, R. (1993). Combining Instance-Based and Model-Based Learning. In Proceedings on the Tenth nal Conference of Machine Learning, 236-243, University of Massachusetts, Amherst. Morgan Kaufmann. The we can see that there is 13 attributes (columns) and 506 instances (rows) - pd. DataFrame (data=boston.data, columns=boston.feature_names) . head() CRIM ZN INDUS CHAS NOX RM AGE DIS RAD TAX PTRATIO B LSTAT
0 1 2 3	CRIM ZN INDUS CHAS NOX RM AGE DIS RAD TAX PTRATIO B LSTAT 0.00632 18.0 2.31 0.0 0.538 6.575 65.2 4.0900 1.0 296.0 15.3 396.90 4.98 0.02731 0.0 7.07 0.0 0.469 6.421 78.9 4.9671 2.0 242.0 17.8 396.90 9.14 0.02729 0.0 7.07 0.0 0.469 7.185 61.1 4.9671 2.0 242.0 17.8 392.83 4.03 0.03237 0.0 2.18 0.0 0.458 6.998 45.8 6.0622 3.0 222.0 18.7 394.63 2.94 0.06905 0.0 2.18 0.0 0.458 7.147 54.2 6.0622 3.0 222.0 18.7 396.90 5.33
X.	0.06905
(((y	404, 13), (404,)) _train.shape,y_test.shape 404,), (37,)) i=mutual_info_regression(x_train,y_train)
m. m. m. LS	<pre>i =pd.Series(mi) i.index = x_train.columns i.sort_values(ascending =False,inplace = True) i TAT 0.680806 0.563681</pre>
IN PT NO TA	DUS 0.511407 RATIO 0.490240 X 0.453311 X 0.393179 IM 0.360620 E 0.338810 S 0.327494 ID 0.212547 O 0.204083
B CH dt	0.149616 AS 0.027637 ype: float64 i.plot.bar() xesSubplot:>
0.6 0.5 0.4	
0.2	LSTAT - RM - NOX - NOX - DIS - CHAS -
0.1 0.0	ve see the above plot in that the execpt one bar the other contributing more information and for that we can thses features are portant and the we are selecting the top 9 featues those who really contributing the imformation as expected.
If w imp	el = SelectKBest(mutual_info_regression, k=9).fit(x_train, y_train) _train.columns[sel.get_support()] dex(['CRIM', 'INDUS', 'NOX', 'RM', 'AGE', 'DIS', 'TAX', 'PTRATIO', 'LSTAT'], dtype='object')
If we imp	<pre>el = SelectKBest(mutual_info_regression, k=9).fit(x_train, y_train) _train.columns[sel.get_support()]</pre>
If we imp	<pre>el = SelectKBest(mutual_info_regression, k=9).fit(x_train, y_train) _train.columns[sel.get_support()] dex(['CRIM', 'INDUS', 'NOX', 'RM', 'AGE', 'DIS', 'TAX', 'PTRATIO', 'LSTAT'], dtype='object') en(x_train.columns[sel.get_support()]) _train = np.array(x_train) _test = np.array(x_test) _train = np.array(y_train)</pre>