

```
In [ ]: This document shows the Julia programming language code for the following medical data
```

```
In [ ]: Tinnitus data classification prediction accuracy results comparison on the testing data
```

```
In [1]: import Pkg; Pkg.add("DataFrames")
import Pkg; Pkg.add("CSV")
using CSV, DataFrames, Plots
```

```
Updating registry at `C:\Users\zizhe\.julia\registries\General.toml`
Resolving package versions...
Installed OpenJpeg_jll — v2.4.0+0
Installed ImageMagick_jll — v6.9.12+4
Installed LittleCMS_jll — v2.12.0+0
No Changes to `C:\Users\zizhe\.julia\environments\v1.7\Project.toml`
Updating `C:\Users\zizhe\.julia\environments\v1.7\Manifest.toml`
[c73af94c] ↑ ImageMagick_jll v6.9.12+3 ⇒ v6.9.12+4
[d3a379c0] + LittleCMS_jll v2.12.0+0
[643b3616] + OpenJpeg_jll v2.4.0+0
Precompiling project...
✓ LittleCMS_jll
✓ OpenJpeg_jll
✓ ImageMagick_jll
✓ ImageMagick
✓ Images
5 dependencies successfully precompiled in 27 seconds (370 already precompiled)
Resolving package versions...
No Changes to `C:\Users\zizhe\.julia\environments\v1.7\Project.toml`
No Changes to `C:\Users\zizhe\.julia\environments\v1.7\Manifest.toml`
```

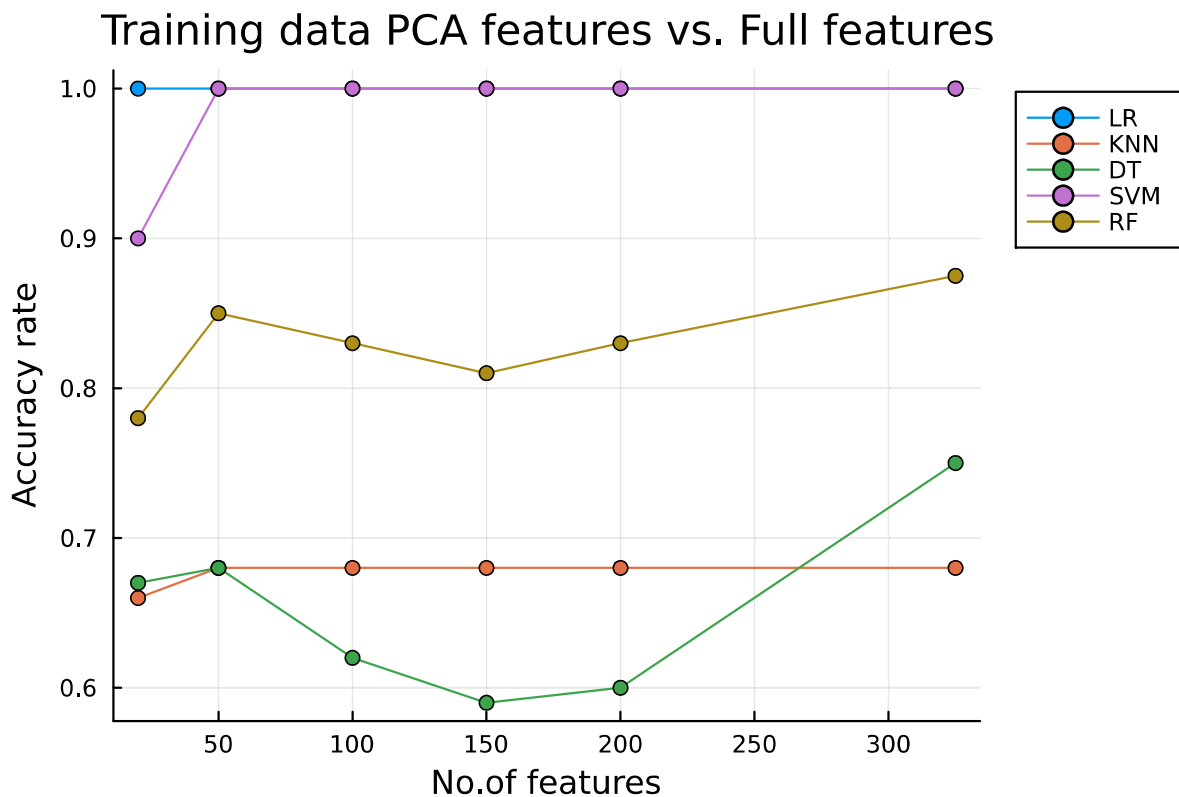
```
In [2]: # Load the PCA training data result
trPCA = CSV.read("C:/Users/Leo/Downloads/Training data PCA result in csv.csv", header=
```

```
Out[2]: 6 rows × 7 columns
```

	Connectivity Features	Number of Features	LR	KNN	DT	SVM	RF
	String	Int64	Int64	Float64	Float64	Float64	Float64
1	Full features	325	1	0.68	0.75	1.0	0.875
2	PCA feature extraction	200	1	0.68	0.6	1.0	0.83
3	PCA feature extraction	150	1	0.68	0.59	1.0	0.81
4	PCA feature extraction	100	1	0.68	0.62	1.0	0.83
5	PCA feature extraction	50	1	0.68	0.68	1.0	0.85
6	PCA feature extraction	20	1	0.66	0.67	0.9	0.78

```
In [3]: plot(trPCA[:,2],Matrix(trPCA[:,3:7]),xlabel="No.of features",ylabel="Accuracy rate",ma
```

Out[3]:



In [4]:

```
# Load the extra tree training data result
trET = CSV.read("C:/Users/Leo/Downloads/Training data extra tree result in csv.csv", k
```

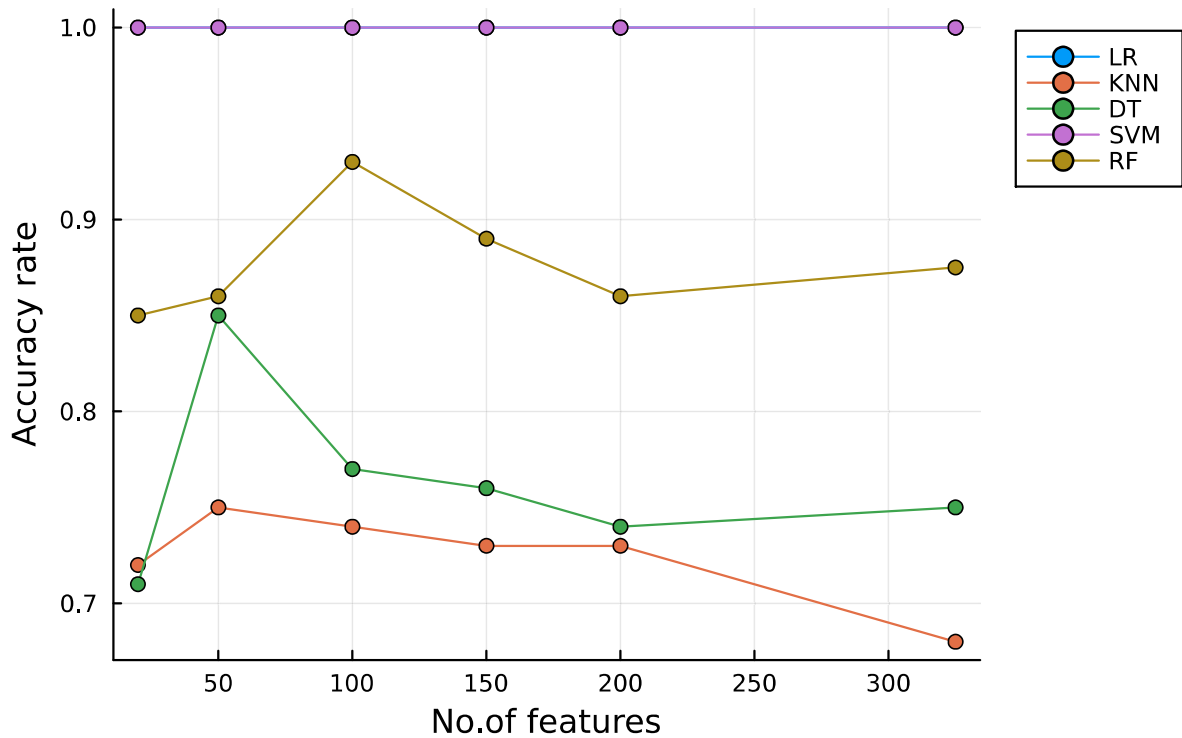
Out[4]: 6 rows × 7 columns

	Connectivity Features	Number of Features	LR	KNN	DT	SVM	RF
	String	Int64	Int64	Float64	Float64	Int64	Float64
1	Full features	325	1	0.68	0.75	1	0.875
2	Extra tree top features	200	1	0.73	0.74	1	0.86
3	Extra tree top features	150	1	0.73	0.76	1	0.89
4	Extra tree top features	100	1	0.74	0.77	1	0.93
5	Extra tree top features	50	1	0.75	0.85	1	0.86
6	Extra tree top features	20	1	0.72	0.71	1	0.85

In [5]:

```
plot(trET[:,2],Matrix(trET[:,3:7]),xlabel="No.of features",ylabel="Accuracy rate",mark
```

Out[5]: Training data Extra tree features vs. Full features



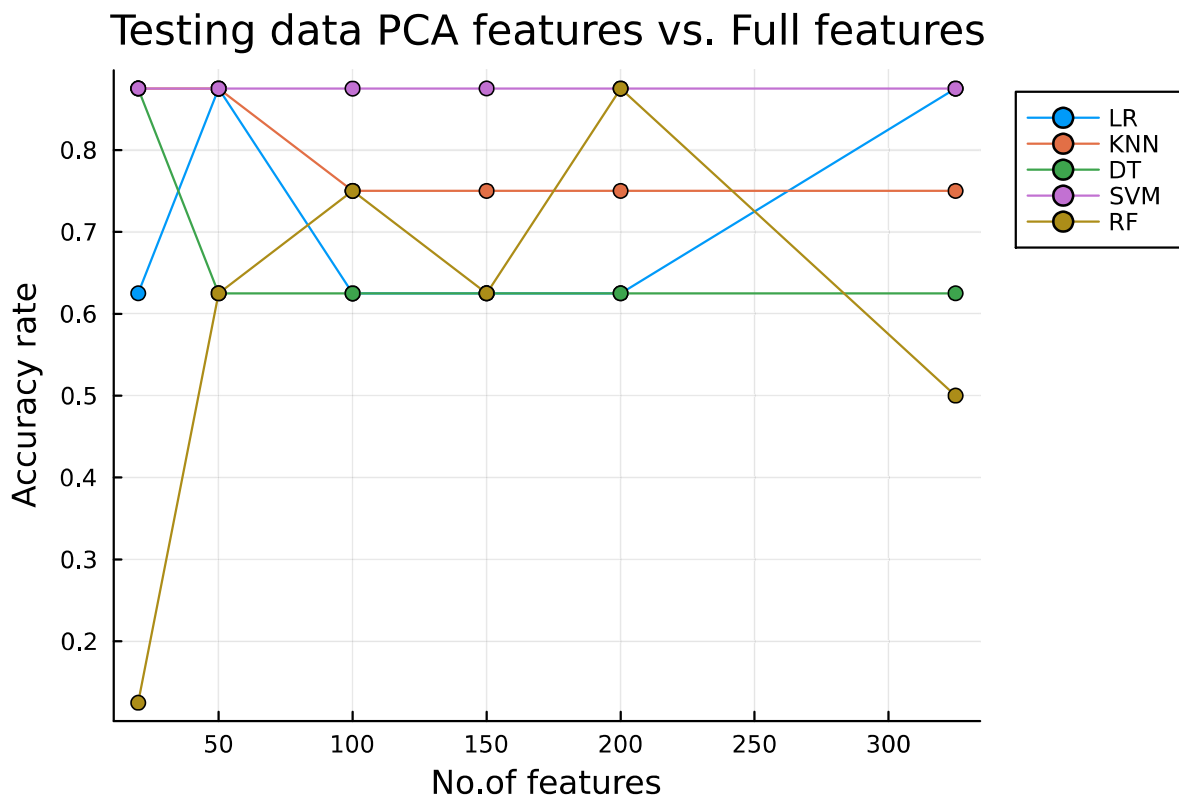
```
In [6]: # Load the PCA testing data result
tePCA = CSV.read("C:/Users/Leo/Downloads/Testing data PCA result in csv.csv", header=
```

Out[6]: 6 rows × 7 columns (omitted printing of 1 columns)

	Connectivity Features	Number of Features	LR	KNN	DT	SVM
	String	Int64	Float64	Float64	Float64	Float64
1	Full features	325	0.875	0.75	0.625	0.875
2	PCA feature extraction	200	0.625	0.75	0.625	0.875
3	PCA feature extraction	150	0.625	0.75	0.625	0.875
4	PCA feature extraction	100	0.625	0.75	0.625	0.875
5	PCA feature extraction	50	0.875	0.875	0.625	0.875
6	PCA feature extraction	20	0.625	0.875	0.875	0.875

```
In [7]: plot(tePCA[:,2],Matrix(tePCA[:,3:7]),xlabel="No. of features",ylabel="Accuracy rate",ma
```

Out[7]:



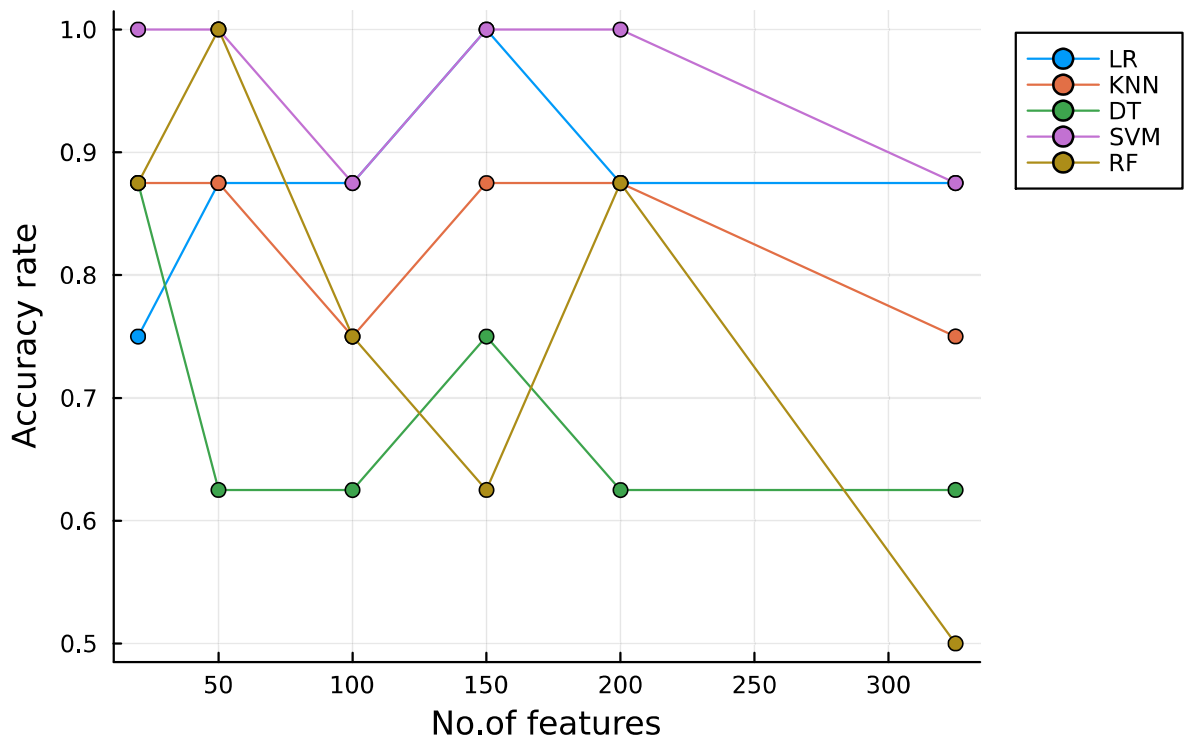
In [8]: `# Load the extra tree training data result`
`teET = CSV.read("C:/Users/Leo/Downloads/Testing data extra tree result in csv.csv", header=1)`

Out[8]: 6 rows × 7 columns (omitted printing of 1 columns)

	Connectivity Features	Number of Features	LR	KNN	DT	SVM
	String	Int64	Float64	Float64	Float64	Float64
1	Full features	325	0.875	0.75	0.625	0.875
2	Extra tree top features	200	0.875	0.875	0.625	1.0
3	Extra tree top features	150	1.0	0.875	0.75	1.0
4	Extra tree top features	100	0.875	0.75	0.625	0.875
5	Extra tree top features	50	0.875	0.875	0.625	1.0
6	Extra tree top features	20	0.75	0.875	0.875	1.0

In [9]: `plot(teET[:,2],Matrix(teET[:,3:7]),xlabel="No. of features",ylabel="Accuracy rate",mark`

Out[9]: Testing data Extra tree features vs. Full features



In []:

In []:

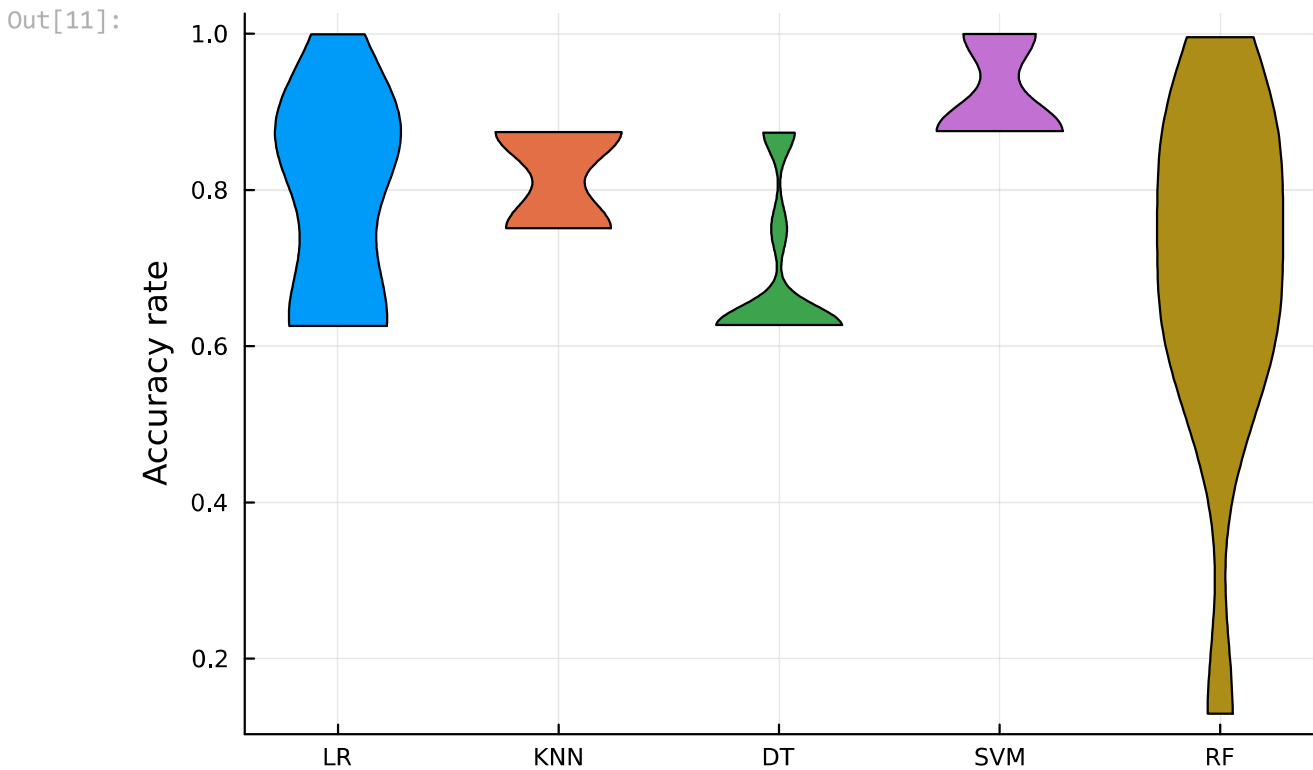
```
In [10]: # Load the full testing data result
te = CSV.read("C:/Users/Leo/Downloads/Testing data result in csv.csv", header=true, Dat
```

Out[10]: 11 rows × 7 columns (omitted printing of 1 columns)

	Connectivity Features	Number of Features	LR	KNN	DT	SVM
	String	Int64	Float64	Float64	Float64	Float64
1	Full features	325	0.875	0.75	0.625	0.875
2	PCA feature extraction	200	0.625	0.75	0.625	0.875
3	PCA feature extraction	150	0.625	0.75	0.625	0.875
4	PCA feature extraction	100	0.625	0.75	0.625	0.875
5	PCA feature extraction	50	0.875	0.875	0.625	0.875
6	PCA feature extraction	20	0.625	0.875	0.875	0.875
7	Extra tree top features	200	0.875	0.875	0.625	1.0
8	Extra tree top features	150	1.0	0.875	0.75	1.0
9	Extra tree top features	100	0.875	0.75	0.625	0.875
10	Extra tree top features	50	0.875	0.875	0.625	1.0
11	Extra tree top features	20	0.75	0.875	0.875	1.0

```
In [11]: Pkg.add("StatsPlots"); using StatsPlots
violin(["LR"], te[:,3], label=nothing, ylabel="Accuracy rate")
violin!(["KNN"], te[:,4], label=nothing)
violin!(["DT"], te[:,5], label=nothing)
violin!(["SVM"], te[:,6], label=nothing)
violin!(["RF"], te[:,7], label=nothing)
```

```
Resolving package versions...
No Changes to `C:\Users\zizhe\.julia\environments\v1.7\Project.toml`
No Changes to `C:\Users\zizhe\.julia\environments\v1.7\Manifest.toml`
```



```
In [12]: using Statistics
mean(te[:,3]),std(te[:,3])
```

Out[12]: (0.7840909090909091, 0.13796409282523808)

```
In [13]: mean(te[:,4]),std(te[:,4])
```

Out[13]: (0.8181818181818182, 0.06527912098338667)

```
In [14]: mean(te[:,5]),std(te[:,5])
```

Out[14]: (0.6818181818181818, 0.10252494153309054)

```
In [15]: mean(te[:,6]),std(te[:,6])
```

Out[15]: (0.9204545454545454, 0.06306562238868912)

```
In [16]: mean(te[:,7]),std(te[:,7])
```

Out[16]: (0.6931818181818182, 0.239554510782752)

```
In [17]: # as their variance are different, so use UnequalVarianceTTest
import Pkg; Pkg.add("HypothesisTests")
using HypothesisTests

UnequalVarianceTTest(te[:,3], te[:,4])
```

```
Resolving package versions...
No Changes to `C:\Users\zizhe\.julia\environments\v1.7\Project.toml`
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```

```
Out[17]: Two sample t-test (unequal variance)
-----
Population details:
  parameter of interest:  Mean difference
  value under h_0:       0
  point estimate:        -0.0340909
  95% confidence interval: (-0.1326, 0.06444)

Test summary:
  outcome with 95% confidence: fail to reject h_0
  two-sided p-value:       0.4708

Details:
  number of observations:  [11,11]
  t-statistic:             -0.740797197487194
  degrees of freedom:      14.263894781501907
  empirical standard error: 0.04601921984390133
```

```
In [18]: # statistical test between LR and KNN
MannWhitneyUTest(te[:,3], te[:,4])
```

```
Out[18]: Approximate Mann-Whitney U test
-----
Population details:
  parameter of interest:  Location parameter (pseudomedian)
  value under h_0:       0
  point estimate:        0.0

Test summary:
  outcome with 95% confidence: fail to reject h_0
  two-sided p-value:       0.6435

Details:
  number of observations in each group: [11, 11]
  Mann-Whitney-U statistic:             53.5
  rank sums:                           [119.5, 133.5]
  adjustment for ties:                  1590.0
  normal approximation ( $\mu$ ,  $\sigma$ ):      (-7.0, 14.0433)
```

```
In [19]: #above test all got "fail to reject h_0", since the null hypothesis is the means of the t
#the result fail to reject h_0" means there is at least 95% confidence that accuracy r
#logistic regression classifier and KNN are statistically identical! Both classifier c
```

```
In [20]: # statistical test between decision tree and KNN
UnequalVarianceTTest(te[:,4], te[:,5])
```

```
Out[20]: Two sample t-test (unequal variance)
-----
Population details:
  parameter of interest: Mean difference
  value under h_0:      0
  point estimate:       0.136364
  95% confidence interval: (0.05903, 0.2137)

Test summary:
  outcome with 95% confidence: reject h_0
  two-sided p-value:         0.0017

Details:
  number of observations: [11,11]
  t-statistic:            3.721042037676257
  degrees of freedom:     16.963613550815555
  empirical standard error: 0.03664662612862977
```

```
In [21]: MannWhitneyUTest(te[:,4], te[:,5])
```

```
Out[21]: Approximate Mann-Whitney U test
-----
Population details:
  parameter of interest: Location parameter (pseudomedian)
  value under h_0:      0
  point estimate:       0.25

Test summary:
  outcome with 95% confidence: reject h_0
  two-sided p-value:         0.0038

Details:
  number of observations in each group: [11, 11]
  Mann-Whitney-U statistic:            102.5
  rank sums:                          [168.5, 84.5]
  adjustment for ties:                 1218.0
  normal approximation ( $\mu$ ,  $\sigma$ ): (42.0, 14.3295)
```

```
In [22]: # above test all got "reject h_0", since the null hypothesis is the means of the two groups
# the result reject h_0 means there is at least 95% confidence that accuracy rates of
# from the plot, we can see the accuracy rate of KNN is better
```

```
In [23]: # statistical test between SVM and KNN
UnequalVarianceTTest(te[:,4], te[:,6])
```



```
Out[23]: Two sample t-test (unequal variance)
```

```
-----  
Population details:
```

```
parameter of interest: Mean difference  
value under h_0: 0  
point estimate: -0.102273  
95% confidence interval: (-0.1594, -0.04518)
```

```
Test summary:
```

```
outcome with 95% confidence: reject h_0  
two-sided p-value: 0.0013
```

```
Details:
```

```
number of observations: [11,11]  
t-statistic: -3.7370465934182957  
degrees of freedom: 19.976247030878863  
empirical standard error: 0.027367260406346124
```

```
In [24]: MannWhitneyUTest(te[:,4], te[:,6])
```

```
Out[24]: Approximate Mann-Whitney U test
```

```
-----  
Population details:
```

```
parameter of interest: Location parameter (pseudomedian)  
value under h_0: 0  
point estimate: 0.0
```

```
Test summary:
```

```
outcome with 95% confidence: reject h_0  
two-sided p-value: 0.0037
```

```
Details:
```

```
number of observations in each group: [11, 11]  
Mann-Whitney-U statistic: 21.0  
rank sums: [87.0, 166.0]  
adjustment for ties: 2364.0  
normal approximation ( $\mu$ ,  $\sigma$ ): (-39.5, 13.4284)
```

```
In [25]: # above test all got "reject h_0", since the null hypothesis is the means of the two groups are equal  
# the result reject h_0 means there is at least 95% confidence that accuracy rates of SVM are better than random  
# from the plot, we can see the accuracy rate of SVM is better
```

```
In [26]: # statistical test between SVM and random forest  
UnequalVarianceTTest(te[:,6], te[:,7])
```

```
Out[26]: Two sample t-test (unequal variance)
-----
Population details:
  parameter of interest: Mean difference
  value under h_0:      0
  point estimate:       0.227273
  95% confidence interval: (0.06355, 0.391)

Test summary:
  outcome with 95% confidence: reject h_0
  two-sided p-value:         0.0108

Details:
  number of observations: [11,11]
  t-statistic:            3.042903097250921
  degrees of freedom:     11.379512195121952
  empirical standard error: 0.0746894396597954
```

```
In [27]: MannWhitneyUTest(te[:,6], te[:,7])
```

```
Out[27]: Approximate Mann-Whitney U test
-----
Population details:
  parameter of interest: Location parameter (pseudomedian)
  value under h_0:      0
  point estimate:       0.125

Test summary:
  outcome with 95% confidence: reject h_0
  two-sided p-value:         0.0049

Details:
  number of observations in each group: [11, 11]
  Mann-Whitney-U statistic:            101.5
  rank sums:                          [167.5, 85.5]
  adjustment for ties:                 1140.0
  normal approximation ( $\mu$ ,  $\sigma$ ): (41.0, 14.3887)
```

```
In [28]: # above test all got "reject h_0", since the null hypothesis is the means of the two groups
# the result reject h_0" means there is at least 95% confidence that accuracy rates of
# classifier and random forest are statistically different!
# from the plot, we can see the accuracy rate of SVM is better
```

```
In [ ]: # After a few T test and MannWhitneyUTest, we can see the accuracy rate of SVM classifier
# statistically different from the other classifiers
# from the plot, we can see SVM is the classifier that has the highest prediction accuracy
```