```
In [ ]: This document shows the Julia programming language code
         for the following medical data analysis research project.
In [ ]: | Tinnitus data classification prediction accuracy results comparison
         on the testing dataset.
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] \uparrow ImageMagick jll v6.9.12+3 \Rightarrow 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`
```

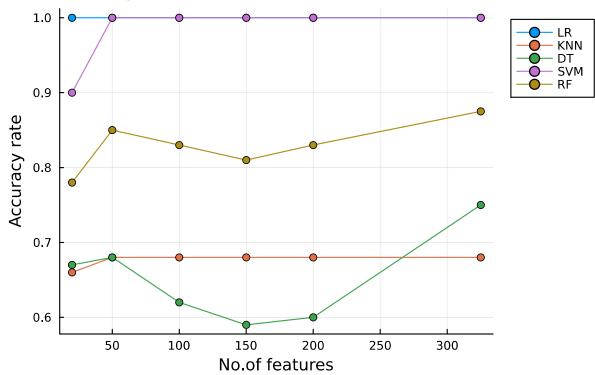
In [2]: # Load the PCA training data result
 trPCA = CSV.read("C:/Users/Leo/Downloads/Training data PCA result in csv.csv",
 header=true,DataFrame)

## Out[2]: 6 rows × 7 columns

	<b>Connectivity Features</b>	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",marker=:c,title="Training data PCA features vs. Full features",
   legend=:outertopright,label=["LR" "KNN" "DT" "SVM" "RF"])
```

## out[3]: Training data PCA features vs. Full features



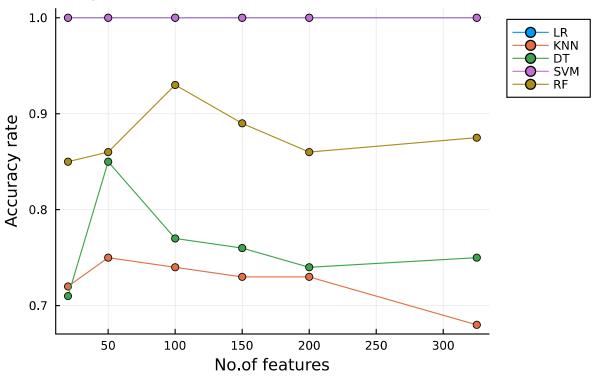
In [4]: # Load the extra tree training data result
trET = CSV.read("C:/Users/Leo/Downloads/Training data extra tree result in csv.csv",
header=true,DataFrame)

Out[4]: 6 rows × 7 columns

	<b>Connectivity Features</b>	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",
    marker=:c,title="Training data Extra tree features vs. Full features",
    legend=:outertopright,label=["LR" "KNN" "DT" "SVM" "RF"])
```

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



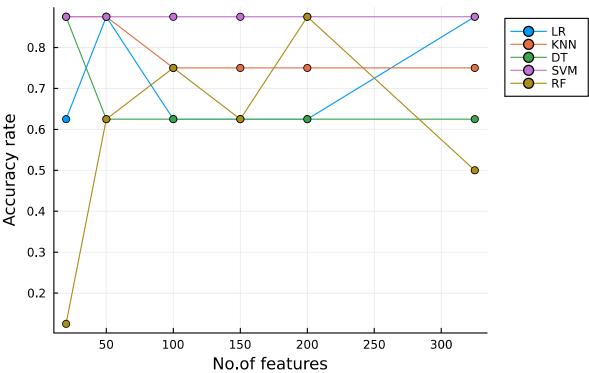
In [6]: # Load the PCA testing data result
tePCA = CSV.read("C:/Users/Leo/Downloads/Testing data PCA result in csv.csv",
header=true,DataFrame)

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

	<b>Connectivity Features</b>	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",
    marker=:c,title="Testing data PCA features vs. Full features",
    legend=:outertopright,label=["LR" "KNN" "DT" "SVM" "RF"])
```

## out[7]: Testing data PCA features vs. Full features



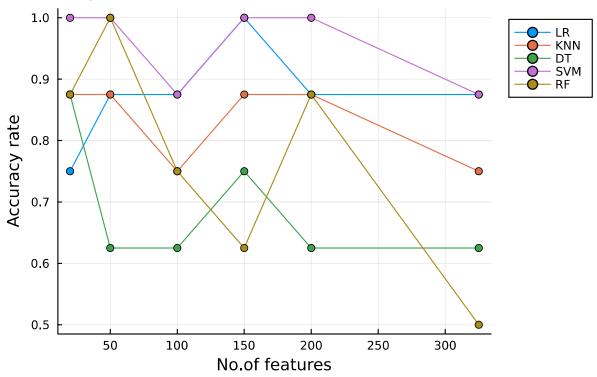
In [8]: # Load the extra tree training data result
teET = CSV.read("C:/Users/Leo/Downloads/Testing data extra tree result in csv.csv",
header=true,DataFrame)

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

	<b>Connectivity Features</b>	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",
    marker=:c,title="Testing data Extra tree features vs. Full features",
    legend=:outertopright,label=["LR" "KNN" "DT" "SVM" "RF"])
```

<code>out[9]: Testing data Extra tree features vs. Full features</code>



```
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,DataFrame)
```

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

	<b>Connectivity Features</b>	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

```
Pkg.add("StatsPlots"); using StatsPlots
In [11]:
          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`
Out[11]:
             1.0
             0.8
         Accuracy rate
             0.6
             0.4
             0.2
                      LR
                                      KNN
                                                      DT
                                                                      SVM
                                                                                       RF
          using Statistics
In [12]:
          mean(te[:,3]),std(te[:,3])
          (0.7840909090909091, 0.13796409282523808)
Out[12]:
          mean(te[:,4]),std(te[:,4])
In [13]:
          (0.8181818181818182, 0.06527912098338667)
Out[13]:
          mean(te[:,5]),std(te[:,5])
In [14]:
          (0.6818181818181818, 0.10252494153309054)
Out[14]:
          mean(te[:,6]),std(te[:,6])
In [15]:
          (0.9204545454545454, 0.06306562238868912)
Out[15]:
          mean(te[:,7]),std(te[:,7])
In [16]:
          (0.6931818181818182, 0.239554510782752)
Out[16]:
```

```
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`
           No Changes to `C:\Users\zizhe\.julia\environments\v1.7\Manifest.toml`
         Two sample t-test (unequal variance)
Out[17]:
         -----
         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])
        Approximate Mann-Whitney U test
Out[18]:
         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
                                                (-7.0, 14.0433)
             normal approximation (\mu, \sigma):
In [19]: Above test all got "fail to reject h_0", since the null hyposis
         is the means of the two groups are statistically identical,
         the result fail to reject h_0" means there is at least 95%
         confidence that accuracy rates of
         logistic regression classifier and KNN are statistically identical!
         Both classifier are equally good/bad
In [20]: # statistical test between decision tree and KNN
         UnequalVarianceTTest(te[:,4], te[:,5])
```

```
Two sample t-test (unequal variance)
Out[20]:
         Population details:
             parameter of interest: Mean difference
             value under h 0:
                                      0.136364
             point estimate:
             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])
         Approximate Mann-Whitney U test
Out[21]:
         Population details:
                                      Location parameter (pseudomedian)
             parameter of interest:
             value under h_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 hyposis is
         the means of the two groups are statistically identical,
         the result reject h 0" means there is at least 95% confidence
         that accuracy rates of decision tree classifier and KNN are statistically different!
         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])

```
Two sample t-test (unequal variance)
Out[23]:
         Population details:
             parameter of interest: Mean difference
             value under h 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])
         Approximate Mann-Whitney U test
Out[24]:
         Population details:
                                      Location parameter (pseudomedian)
             parameter of interest:
             value under h_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 hyposis is
         the means of the two groups are statistically identical,
         the result reject h 0" means there is at least 95% confidence
         that accuracy rates of SVM classifier and KNN are statistically different!
         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])

```
Two sample t-test (unequal variance)
Out[26]:
         Population details:
             parameter of interest: Mean difference
             value under h 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])
         Approximate Mann-Whitney U test
Out[27]:
         Population details:
             parameter of interest:
                                      Location parameter (pseudomedian)
             value under h_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 hyposis is
         the means of the two groups are statistically identical,
         the result reject h 0" means there is at least
         95% confidence that accuracy rates of SVM
         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 is
         statistically different from the other classifiers
         from the plot, we can see SVM is the classifier
         that has the highest prediction accuracy rate
```