

# Package ‘Dforest’

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**Type** Package

**Title** Decision Forest

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**Depends** R (>= 3.0)

**Imports** rpart, ggplot2, methods, stats

## Description

Provides R-implementation of Decision forest algorithm, which combines the predictions of multiple independent decision tree models for a consensus decision. In particular, Decision Forest is a novel pattern-recognition method which can be used to analyze: (1) DNA microarray data; (2) Surface-Enhanced Laser Desorption/Ionization Time-of-Flight Mass Spectrometry (SELDI-TOF-MS) data; and (3) Structure-Activity Relation (SAR) data. In this package, three fundamental functions are provided, as (1)DF\_train, (2)DF\_pred, and (3)DF\_CV. run Dforest() to see more instructions. Weida Tong (2003) <[doi:10.1021/ci020058s](https://doi.org/10.1021/ci020058s)>.

**License** GPL-2

**LazyLoad** yes

**LazyData** yes

**Encoding** UTF-8

**NeedsCompilation** no

**RoxygenNote** 6.0.1

**Repository** CRAN

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## Contents

cal_MCC . . . . .	2
Con_DT . . . . .	3
data_dili . . . . .	3
Dforest . . . . .	4
DF_acc . . . . .	4
DF_calp . . . . .	5
DF_ConfPlot . . . . .	5
DF_ConfPlot_accu . . . . .	6
DF_CV . . . . .	6
DF_CVsummary . . . . .	8
DF_dataFs . . . . .	8
DF_dataPre . . . . .	9
DF_easy . . . . .	9
DF_perf . . . . .	10
DF_pred . . . . .	11
DF_train . . . . .	12
DF_Trainsummary . . . . .	13
multiplot . . . . .	13
Pred_DT . . . . .	14
<b>Index</b>	<b>15</b>

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cal_MCC	<i>Performance evaluation from other modeling algorithm Result</i>
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### Description

Performance evaluation from other modeling algorithm Result

### Usage

```
cal_MCC(pred, label)
```

### Arguments

pred	Predictions
label	Known-endpoint

### Value

result\$ACC: Predicting Accuracy  
 result\$MIS: MisClassification Counts  
 result\$MCC: Matthew's Correlation Coefficients  
 result\$bACC: balanced Accuracy

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Con_DT	<i>Construct Decision Tree model with pruning</i>
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**Description**

Construct Decision Tree model with pruning

**Usage**

```
Con_DT(X, Y, min_split = 10, cp = 0.01)
```

**Arguments**

X	dataset
Y	data_Labels
min_split	minimum number of node in each leaf
cp	pre-defined Complexity Parameter (CP) rpart program

**Value**

Decision Tree Model with pruning Implemented by rpart

**See Also**

rpart

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data_dili	<i>QSAR dataset with DILI endpoint for demo</i>
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**Description**

This data set gives the DILI endpoint of various compounds (Most or No DILI-concern) with QSAR descriptors generated by MOLD2

**Usage**

```
rivers
```

**Format**

A List containing two vectors: X contains 958 observations and 777 variables. Y contains DILI endpoints of 958 observations

**Source**

In-house data

**References**

Minjun Chen (2011) *FDA-approved drug labeling for the study of drug-induced liver injury*. Drug discovery today

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Dforest	<i>Demo script to lean Decision Forest package Demo data are located in data/ folder</i>
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**Description**

Demo script to lean Decision Forest package Demo data are located in data/ folder

**Usage**

Dforest()

**Author(s)**

Leihong.Wu

**Examples**

Dforest()

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DF_acc	<i>Performance evaluation from Decision Tree Predictions</i>
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**Description**

Performance evaluation from Decision Tree Predictions

**Usage**

DF\_acc(pred, label)

**Arguments**

pred	Predictions
label	Known-endpoint

**Value**

result\$ACC: Predicting Accuracy  
 result\$MIS: MisClassification Counts  
 result\$MCC: Matthew's Correlation Coefficients  
 result\$bACC: balanced Accuracy

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DF_calp	<i>T-test for feature selection</i>
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**Description**

T-test for feature selection

**Usage**

DF\_calp(X, Y)

**Arguments**

X	X variable matrix
Y	Y label

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DF_ConfPlot	<i>Decision Forest algorithm: confidence level accumulated plot</i>
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**Description**

Draw accuracy curve according to the confidence level of predictions

**Usage**

DF\_ConfPlot(Pred\_result, Label, bin = 20, plot = T, smooth = F)

**Arguments**

Pred_result	Predictions
Label	known label for Test Dataset
bin	How many bins occurred in Conf Plot (Default is 20)
plot	Draw Plot if True, otherwise output the datamatrix
smooth	if TRUE, Fit the performance curve with smooth function (by ggplot2)

**Value**

ACC\_Conf: return data Matrix ("ConfidenceLevel", "Accuracy", "Matched Samples") for confidence plot (no plot)

ConfPlot: Draw Confidence Plot if True, need install ggplot2

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DF_ConfPlot_accu	<i>Decision Forest algorithm: confidence level accumulated plot (accumulated version)</i>
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---

### Description

Draw accuracy curve according to the confidence level of predictions

### Usage

```
DF_ConfPlot_accu(Pred_result, Label, bin = 20, plot = T, smooth = F)
```

### Arguments

Pred_result	Predictions
Label	known label for Test Dataset
bin	How many bins occurred in Conf Plot (Default is 20)
plot	Draw Plot if True, otherwise output the datamatrix
smooth	if TRUE, Fit the performance curve with smooth function (by ggplot2)

### Value

ACC\_Conf: return data Matrix ("ConfidenceLevel", "Accuracy", "Matched Samples") for confidence plot (no plot)

ConfPlot: Draw Confidence Plot if True, need install ggplot2

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DF_CV	<i>Decision Forest algorithm: Model training with Cross-validation</i>
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### Description

Decision Forest algorithm: Model training with Cross-validation Default is 5-fold cross-validation

### Usage

```
DF_CV(X, Y, stop_step = 10, CV_fold = 5, Max_tree = 20, min_split = 10,
      cp = 0.1, Filter = F, p_val = 0.05, Method = "bACC", Quiet = T,
      Grace_val = 0.05, imp_accu_val = 0.01, imp_accu_criteria = F)
```

**Arguments**

X	Training Dataset
Y	Training data endpoint
stop_step	How many extra step would be processed when performance not improved, 1 means one extra step
CV_fold	Fold of cross-validation (Default = 5)
Max_tree	Maximum tree number in Forest
min_split	minimum leaves in tree nodes
cp	parameters to pruning decision tree, default is 0.1
Filter	doing feature selection before training
p_val	P-value threshold measured by t-test used in feature selection, default is 0.05
Method	Which is used for evaluating training process. MIS: Misclassification rate; ACC: accuracy
Quiet	if TRUE (default), don't show any message during the process
Grace_val	Grace Value in evaluation: the next model should have a performance (Accuracy, bACC, MCC) not bad than previous model with threshold
imp_accu_val	improvement in evaluation: adding new tree should improve the overall model performance (Accuracy, bACC, MCC) by threshold
imp_accu_criteria	if TRUE, model must have improvement in accumulated accuracy

**Value**

.\$performance: Overall training accuracy (Cross-validation)  
 .\$pred: Detailed training prediction (Cross-validation)  
 .\$detail: Detailed usage of Decision tree Features/Models and their performances in all CVs  
 .\$Method: pass evaluating Methods used in training  
 .\$cp: pass cp value used in training decision trees

**Examples**

```

##data(iris)
X = iris[,1:4]
Y = iris[,5]
names(Y)=rownames(X)

random_seq=sample(nrow(X))
split_rate=3
split_sample = suppressWarnings(split(random_seq,1:split_rate))
Train_X = X[-random_seq[split_sample[[1]]],]
Train_Y = Y[-random_seq[split_sample[[1]]]]

CV_result = DF_CV(Train_X, Train_Y)

```

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DF_CVsummary	<i>output summary for Dforest Cross-validation results</i>
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**Description**

Draw plot for Dforest Cross-validation results

**Usage**

```
DF_CVsummary(CV_result, plot = T)
```

**Arguments**

CV_result	Training Dataset
plot	if TRUE (default), draw plot

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DF_dataFs	<i>Decision Forest algorithm: Feature Selection in pre-processing</i>
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**Description**

Decision Forest algorithm: feature selection for two-class predictions, kept statistical significant features pass the t-test

**Usage**

```
DF_dataFs(X, Y, p_val = 0.05)
```

**Arguments**

X	Training Dataset
Y	Training Labels
p_val	Correlation Coefficient threshold to filter out high correlated features; default is 0.95

**Value**

Keep\_feat: qualified features in data matrix after filtering

**Examples**

```
##data(iris)
X = iris[iris[,5]!="setosa",1:4]
Y = iris[iris[,5]!="setosa",5]
used_feat = DF_dataFs(X, Y)
```

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DF_dataPre	<i>Decision Forest algorithm: Data pre-processing</i>
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**Description**

Decision Forest algorithm: Data pre-processing, remove All-Zero columns/features and high correlated features

**Usage**

```
DF_dataPre(X, thres = 0.95)
```

**Arguments**

X	Training Dataset
thres	Correlation Coefficient threshold to filter out high correlated features; default is 0.95

**Value**

Keep\_feat: qualified features in data matrix after filtering

**Examples**

```
##data(iris)
X = iris[,1:4]
Keep_feat = DF_dataPre(X)
```

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DF_easy	<i>Simple pre-defined pipeline for Decision forest</i>
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**Description**

This is a script of decision forest for easy use t

**Usage**

```
DF_easy(Train_X, Train_Y, Test_X, Test_Y, mode = "default")
```

**Arguments**

Train_X	Training Dataset
Train_Y	Training data endpoint
Test_X	Testing Dataset
Test_Y	Testing data endpoint
mode	pre-defined modeling

**Value**

data\_matrix training and testing result

**Examples**

```
# data(demo_simple)
X = iris[,1:4]
Y = iris[,5]
names(Y)=rownames(X)

random_seq=sample(nrow(X))
split_rate=3
split_sample = suppressWarnings(split(random_seq,1:split_rate))
Train_X = X[-random_seq[split_sample[[1]]],]
Train_Y = Y[-random_seq[split_sample[[1]]]]
Test_X = X[random_seq[split_sample[[1]]],]
Test_Y = Y[random_seq[split_sample[[1]]]]

Result = DF_easy(Train_X, Train_Y, Test_X, Test_Y)
```

---

DF\_perf

*performance evaluation between two factors*


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**Description**

performance evaluation between two factors

**Usage**

```
DF_perf(pred, label)
```

**Arguments**

pred	Predictions
label	Known-endpoint

**Value**

result\$ACC: Predicting Accuracy  
result\$MIS: MisClassification Counts  
result\$MCC: Matthew's Correlation Coefficients  
result\$bACC: balanced Accuracy

DF\_pred

*Decision Forest algorithm: Model prediction***Description**

Decision Forest algorithm: Model prediction with constructed DF models. DT\_models is a list of Decision Tree models (rpart.objects) generated by DF\_train() DT\_train\_CV() is only designed for Cross-validation and won't generate models

**Usage**

```
DF_pred(DT_models, X, Y = NULL)
```

**Arguments**

DT_models	Constructed DF models
X	Test Dataset
Y	Test data endpoint

**Value**

.\$accuracy: Overall test accuracy  
 .\$predictions: Detailed test prediction

**Examples**

```
# data(demo_simple)
X = data_dili$X
Y = data_dili$Y
names(Y)=rownames(X)

random_seq=sample(nrow(X))
split_rate=3
split_sample = suppressWarnings(split(random_seq,1:split_rate))
Train_X = X[-random_seq[split_sample[[1]]],]
Train_Y = Y[-random_seq[split_sample[[1]]]]
Test_X = X[random_seq[split_sample[[1]]],]
Test_Y = Y[random_seq[split_sample[[1]]]]

used_model = DF_train(Train_X, Train_Y)
Pred_result = DF_pred(used_model,Test_X,Test_Y)
```

DF\_train

*Decision Forest algorithm: Model training***Description**

Decision Forest algorithm: Model training

**Usage**

```
DF_train(X, Y, stop_step = 5, Max_tree = 20, min_split = 10, cp = 0.1,
        Filter = F, p_val = 0.05, Method = "bACC", Quiet = T,
        Grace_val = 0.05, imp_accu_val = 0.01, imp_accu_criteria = F)
```

**Arguments**

X	Training Dataset
Y	Training data endpoint
stop_step	How many extra step would be processed when performance not improved, 1 means one extra step
Max_tree	Maximum tree number in Forest
min_split	minimum leaves in tree nodes
cp	parameters to pruning decision tree, default is 0.1
Filter	doing feature selection before training
p_val	P-value threshold measured by t-test used in feature selection, default is 0.05
Method	Which is used for evaluating training process. MIS: Misclassification rate; ACC: accuracy
Quiet	if TRUE (default), don't show any message during the process
Grace_val	Grace Value in evaluation: the next model should have a performance (Accuracy, bACC, MCC) not bad than previous model with threshold
imp_accu_val	improvement in evaluation: adding new tree should improve the overall model performance (Accuracy, bACC, MCC) by threshold
imp_accu_criteria	if TRUE, model must have improvement in accumulated accuracy

**Value**

.\$accuracy: Overall training accuracy  
 .\$pred: Detailed training prediction (fitting)  
 .\$detail: Detailed usage of Decision tree Features/Models and their performances  
 .\$models: Constructed (list of) Decision tree models  
 .\$Method: pass evaluating Methods used in training  
 .\$cp: pass cp value used in training decision trees

**Examples**

```
##data(iris)
X = iris[,1:4]
Y = iris[,5]
names(Y)=rownames(X)
used_model = DF_train(X, factor(Y))
```

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DF_Trainsummary	<i>output summary for Dforest test results</i>
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---

**Description**

Draw plot for Dforest test results

**Usage**

```
DF_Trainsummary(used_model, plot = T)
```

**Arguments**

used_model	Training result
plot	if TRUE (default), draw plot

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multiplot	<i>multiplot</i>
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**Description**

Multiple plot function

If the layout is something like `matrix(c(1,2,3,3), nrow=2, byrow=TRUE)`, then plot 1 will go in the upper left, 2 will go in the upper right, and 3 will go all the way across the bottom.

**Usage**

```
multiplot(..., plotlist = NULL, cols = 1, layout = NULL)
```

**Arguments**

...	ggplot objects
plotlist	a list of ggplot objects
cols	Number of columns in layout
layout	A matrix specifying the layout. If present, 'cols' is ignored.

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Pred\_DT

*Doing Prediction with Decision Tree model*

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**Description**

Doing Prediction with Decision Tree model

**Usage**

Pred\_DT(model, X)

**Arguments**

model	Decision Tree Model
X	dataset

**Value**

Decision Tree Predictions Different endpoints presented in multiple columns

**Source**

rpart

**See Also**

rpart

# Index

## \* datasets

- [data\\_dili, 3](#)
  
- [cal\\_MCC, 2](#)
- [Con\\_DT, 3](#)
  
- [data\\_dili, 3](#)
- [DF\\_acc, 4](#)
- [DF\\_calp, 5](#)
- [DF\\_ConfPlot, 5](#)
- [DF\\_ConfPlot\\_accu, 6](#)
- [DF\\_CV, 6](#)
- [DF\\_CVsummary, 8](#)
- [DF\\_dataFs, 8](#)
- [DF\\_dataPre, 9](#)
- [DF\\_easy, 9](#)
- [DF\\_perf, 10](#)
- [DF\\_pred, 11](#)
- [DF\\_train, 12](#)
- [DF\\_Trainsummary, 13](#)
- [Dforest, 4](#)
  
- [multiplot, 13](#)
  
- [Pred\\_DT, 14](#)