

Package ‘TailClassifier’

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Type Package

Title Tail Classifier

Version 0.1.1

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Description The function TailClassifier() suggests one of the following types of tail for your discrete data: 1) Power decaying tail; 2) Sub-exponential decaying tail; and 3) Near-exponential decaying tail. The function also provides an estimate of the parameter for the classified-distribution as a reference.

License GPL-3

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Imports ggplot2, cowplot, scales, stats, utils

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Description

The function TailClassifier() suggests one of the following types of tail for your discrete data: 1) Power decaying tail; 2) Sub-exponential decaying tail; and 3) Near-exponential decaying tail. The function also provides an estimate of the parameter for the classified-distribution as a reference.

Usage

```

TailClassifier(
  sample_frequencies,
  v_left = 20,
  v_right = min(floor(sum(sample_frequencies)/20),
    sum(sample_frequencies[sample_frequencies > 1]) - 1),
  plot_lower = v_left,
  plot_upper = v_right,
  Plot0_title = "Plot 0 of Heavy Tail Detection \n \n",
  Plot1_title = "Plot 1 of Heavy Tail Detection",
  Plot2_title = "Plot 2 of Heavy Tail Detection",
  Plot3_title = "Plot 3 of Heavy Tail Detection",
  C_Level = 0.95,
  ConfidenceBand = T,
  Plot_0_y_limit_lower_extend = 1.5,
  Plot_0_y_limit_upper_extend = 1.5,
  Plot_1_y_limit_lower_extend = 0.25,
  Plot_1_y_limit_upper_extend = 0.25,
  Plot_2_y_limit_lower_extend = 0.25,
  Plot_2_y_limit_upper_extend = 0.25,
  Plot_3_y_limit_lower_extend = 0.25,
  Plot_3_y_limit_upper_extend = 0.25
)

```

Arguments

sample_frequencies	The frequency counts for your discrete sample data.
v_left	The starting point of tail profile. 20 is recommended. A smaller v_left may lead to unreliable results. A larger v_left might be adopted if the sample size is extremely large.
v_right	The ending point of tail profile. Recommendation is no more than 100 regardless of sample size.
plot_lower	The lower range of v-axis.
plot_upper	The upper range of v-axis.
Plot0_title	The title for Plot0. The default is "Plot 0 of Heavy Tail Detection".
Plot1_title	The title for Plot1. The default is "Plot 1 of Heavy Tail Detection".
Plot2_title	The title for Plot2. The default is "Plot 2 of Heavy Tail Detection".
Plot3_title	The title for Plot3. The default is "Plot 3 of Heavy Tail Detection".
C_Level	The confidence level of confidence intervals in results. The default is 0.95.
ConfidenceBand	TRUE if a confidence band is requested. FALSE otherwise.
Plot_0_y_limit_lower_extend	Modify the y limit in Plot 0 to allow the confidence band to correctly display in different scenarios.

Plot_0_y_limit_upper_extend
 Modify the y limit in Plot 1 to allow the confidence band to correctly display in different scenarios.

Plot_1_y_limit_lower_extend
 Modify the y limit in Plot 2 to allow the confidence band to correctly display in different scenarios.

Plot_1_y_limit_upper_extend
 Modify the y limit in Plot 3 to allow the confidence band to correctly display in different scenarios.

Plot_2_y_limit_lower_extend
 Modify the y limit in Plot 0 to allow the confidence band to correctly display in different scenarios.

Plot_2_y_limit_upper_extend
 Modify the y limit in Plot 1 to allow the confidence band to correctly display in different scenarios.

Plot_3_y_limit_lower_extend
 Modify the y limit in Plot 2 to allow the confidence band to correctly display in different scenarios.

Plot_3_y_limit_upper_extend
 Modify the y limit in Plot 3 to allow the confidence band to correctly display in different scenarios.

Value

A statement on the type of tail.

Examples

```
## Power Example
# Generate data from power decaying distribution with parameter 1.5
rpar <- function(n, a, xm = 1) {
  v <- runif(n)
  xm / v^(1.0/a)
}
dpar <- function(x, a, xm = 1){
  return(a*xm^a/(x^(a+1)))
}
set.seed(2023)
data <- floor(rpar(1000, 0.5)) # lambda = 1.5
Result <- TailClassifier(table(data), plot_lower = 5, plot_upper = 400, v_left = 20, v_right = 54,
  Plot_0_y_limit_upper_extend = 8)
## display the results
Result
## call the classification decision
Result$Type
## call the confidence intervals for the parameters
data.frame(Result$Results[3])[,c(1,3:4)]
## call a specific plot
Result$Results[[1]][1]
Result$Results[[1]][2]
Result$Results[[1]][3]
```

```
Result$Results[[1]][4]  
## check the rank of possible type of tails  
Result$Rank
```

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