

Package ‘SURVFIT’

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Type Package

Title Rule Ensemble Learning and Analysis for Survival Data

Version 0.1.0

Author@R

`c(person("Ameer", "Hamza"), email = <ameerh.shakur@gmail.com>, role = c("aut", "cre"))`

Author Ameer Hamza [aut, cre]

Maintainer Ameer Hamza <ameerh.shakur@gmail.com>

Description This package derives doubly sparse predictive ensemble of rules from survival data using a second order cone formulation solved using Cplex. Double sparsity induces sparsity in both number of rules extracted as well as variables involved in the rules. This package also includes methods for comprehensive analysis of extracted rules such as statistical testing, decomposition analysis and sensitivity analysis.

URL <https://github.com/hamzameer/SURVFIT>

Depends R (>= 3.5.3)

License GPL-2 | GPL-3

Encoding UTF-8

LazyData true

Imports osqp,

dplyr,
ranger,
survival,
ggplot2,
Formula,
Rcplex

RoxygenNote 7.0.2

Suggests knitr,
rmarkdown

VignetteBuilder knitr

R topics documented:

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Description

SURVFIT extracts a "doubly sparse" (sparse in both number of rules and in number of variables in the rules) survival rule ensemble from survival data

Usage

```
SURVFIT(
  formula = formula,
  data = data,
  rulelength = 3,
  doubly.sparse = FALSE,
  gamma = NULL,
  lambda1 = NULL,
  lambda2 = NULL,
  crossvalidate = TRUE,
  nfolds = 4,
  num_toprules = 16,
  num_totalrules = 2000,
  input_rule_list = FALSE,
  rule_list = NULL,
  ntree = 200,
  digit = 10,
  seed = NULL,
  nodesize = NULL,
  trace = 1,
  max.grid = 25,
  ...
)
```

Arguments

| | |
|----------------|--|
| formula | formula. The model formula specifying time, status and dependent variables of the form <code>Surv(time, status)~ x1 + x2 + ..</code> |
| data | data.frame. Training data. |
| rulelength | Integer. Maximum length of the rule. (Default = 3) |
| doubly.sparse | Logical for whether double sparsity required. (Default = FALSE) |
| gamma | Numeric or list. Hyperparameter (Default = NULL) |
| lambda1 | Numeric or list. Hyperparameter (Default = NULL) |
| lambda2 | Numeric or list. Hyperparameter (Default = NULL) |
| crossvalidate | Logical. Whether crossvalidation to be done to find hyperparameters. (Default = TRUE) |
| nfolds | Integer. Number of cross validation folds. (Default = 5) |
| num_toprules | Integer. Number of rules extracted. (Default = 16) |
| num_totalrules | Integer. Number of rules considered. (Default = 2000) |

| | |
|------------------------------|--|
| <code>input_rule_list</code> | Logical Whether rule list supplied. (Default = FALSE) |
| <code>rule_list</code> | List. List of supplied rules. (Default = NULL) |
| <code>ntree</code> | Integer .Number of trees built |
| <code>digit</code> | Integer. Decimal points. |
| <code>seed</code> | Numeric. Seed for reproducible experiments. |
| <code>nodesize</code> | Integer. (Default = NULL) |
| <code>trace</code> | 0 or 1. : Turn CPLEX output on (1) or off(0). Default 1. |
| <code>...</code> | Other inputs |

Value

Object of class `list` with elements

| | |
|------------------------|---|
| <code>rules</code> | List of top <code>num_toprules</code> rules |
| <code>all_rules</code> | List of all <code>num_totalrules</code> rules |
| <code>rule_data</code> | Data.frame of rules evaluated over data |
| <code>beta</code> | Coefficients of <code>all_rules</code> in the model |

Examples

```
## For ovarian data from "survival" package.  
SURVFIT(Surv(futime, fustat) ~ ., data = ovarian)
```

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