

# Package ‘LogisticCopula’

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**Type** Package

**Title** A Copula Based Extension of Logistic Regression

**Version** 0.1.0

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**Description** An implementation of a method of extending a logistic regression model beyond linear effects of the co-variates. The extension is constructed by first equating the logistic regression model to a naive Bayes model where all the margins are specified to follow natural exponential distributions conditional on Y, that is, a model for Y given X that is specified through the distribution of X given Y, where the columns of X are assumed to be mutually independent conditional on Y. Subsequently, the model is expanded by adding vine - copulas to relax the assumption of mutual independence, where pair-copulas are added in a stage-wise, forward selection manner. Some heuristics are employed during the process of selecting edges, as well as the families of pair-copula models. After each component is added, the parameters are updated by a (smaller) number of gradient steps to maximise the likelihood. When the algorithm has stopped adding edges, based the criterion that a new edge should improve the likelihood more than k times the number new parameters, the parameters are updated with a larger number of gradient steps, or until convergence.

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**Imports** VineCopula (>= 2.5.0), rvinecopulib (>= 0.6.3.1.1), igraph (>= 2.0.3), numDeriv (>= 8-1.1), stringr (>= 1.5.1)

**Depends** R (>= 3.5.0), brglm2 (>= 0.9)

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.3.1

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**NeedsCompilation** no

**Repository** CRAN

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fit\_copula\_interactions  
*fit\_copula\_interactions*

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## Description

This is the main function of the package, which starting from an initial logistic regression model with only main effects of each covariate, selects and fits interaction terms in the form of two R-vine models with identical graphical structure, one for each class.

## Usage

```
fit_copula_interactions(
  y,
  x,
  xtype,
  family_set = c("gaussian", "clayton", "gumbel"),
  oos_validation = FALSE,
  tau = 2,
  which_include = NULL,
  reg.method = "glm",
  maxit_final = 1000,
  maxit_intermediate = 50,
  verbose = FALSE,
  adjust_intercept = TRUE,
  max_t = Inf,
  test_x = NULL,
  test_y = NULL,
  set_nonsig_zero = FALSE,
  reltol = sqrt(.Machine$double.eps)
)
```

## Arguments

**y** A vector of  $n$  observations of the (univariate) binary outcome variable  $y$

**x** A  $(n \times p)$  matrix of  $n$  observations of  $p$  covariates

xtype	A vector of p characters that have to take the value "c_a", "c_p", "d_b" or "d_b", to indicate whether each margin of the is continuous with full support, continuous with support on the positive real line, discrete (binary) or a counting variable.
family_set	A vector of strings that specifies the set of pair-copula families that the fitting algorithm chooses from. For an overview of which values that can be specified, see the documentation for <a href="#">bicop</a> .
oos_validation	Whether to use an external sample for validation instead of an in-sample likelihood based criteria. Would require that both test_x and test_y are provided if set to TRUE.
tau	Parameter used when selecting the structure, where the the criteria is (new_likelihood - previous_likelihood - tau), so that an additional edge in the copulas is only accepted if it leads to an increase in the likelihood that exceeds tau. Setting tau to NULL, has the same effect as -Inf.
which_include	The column indices of the covariates that could be included in the copula effects.
reg.method	The method by which the initial regression coefficients are fitted.
maxit_final	The maximum number of gradient optimisation iterations to use when the full structure has been selected to refit all the parameters. Defaults to 1000.
maxit_intermediate	The maximum number of gradient optimisation iterations to use when adding a newly selected component to refit the parameters. Defaults to 10.
verbose	Whether information about the progress should be printed to the console.
adjust_intercept	Whether to intermediately refit the intercept during the model/structure selection procedure. Defaults to true.
max_t	The maximum number of trees in the copula models. Defaults to Inf, i.e., no maximum.
test_x	Part of the optional validation set, see @oos_validation.
test_y	Part of the optional validation set, see @oos_validation.
set_nonsig_zero	If true, non-significant regression coefficients (in the initial glm model) will be set to zero
reltol	Relative convergence tolerance, see the documentation for <a href="#">optim</a> .

### Value

A logistic\_copula object, which contains the regression coefficients of the model, the parameters of the chosen conditional covariate distribution that corresponds to the regression coefficients, and the pair of vine-models that extend the logistic regression model.

### Examples

```
data("Ionosphere")
dset <- Ionosphere[, -(1:2)]
```

```

set.seed(20)
rowss <- sample(nrow(dset), round(nrow(dset) * 0.75))
colss <- sample(ncol(dset) - 1, 5)
x <- as.matrix(dset[rowss, colss])
xte <- as.matrix(dset[-rowss, colss])
y <- dset[rowss, ncol(dset)] == "bad"
yte <- dset[-rowss, ncol(dset)] == "bad"

xtype <- apply(x, 2, function(x) if(length(unique(x)) > 2) "c_a" else "d")

# Model with selection penalty tau=log(n)
md <- LogisticCopula::fit_copula_interactions(
  y, as.matrix(x), xtype, tau = log(nrow(x))
)
# Model with selection penalty tau=Inf, returns just the logistic
# regression model
mdglm <- LogisticCopula::fit_copula_interactions(
  y, as.matrix(x), xtype, tau = Inf
)

plot(predict(mdglm, xte), predict(md, xte), col = 3 + yte)

```

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fit\_model

*fit\_model*


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## Description

This function updates the parameters of a LogisticCopula model by maximum likelihood.

## Usage

```

fit_model(
  y,
  x,
  m_obj,
  maxit = 5,
  num_grad = FALSE,
  verbose = FALSE,
  hessian = FALSE,
  reltol = sqrt(.Machine$double.eps)
)

```

## Arguments

y	A vector of n observations of the (univariate) binary outcome variable y
x	A (n x p) matrix of n observations of p covariates
m_obj	The model object as returned from fit_copula_interactions

<code>maxit</code>	The maximum number of gradient steps
<code>num_grad</code>	Whether to compute gradients numerically.
<code>verbose</code>	Whether information about the progress should be printed to the console.
<code>hessian</code>	Whether to numerically compute the hessian matrix, see the documentation for <a href="#">optim</a> .
<code>reltol</code>	Relative convergence tolerance, see the documentation for <a href="#">optim</a> .

### Value

A `logistic_copula` object, which contains the regression coefficients of the model, the parameters of the chosen conditional covariate distribution that corresponds to the regression coefficients, and the pair of vine-models that extend the logistic regression model.

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Ionosphere	<i>Example data set</i>
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### Description

This radar data was collected by a system in Goose Bay, Labrador. This system consists of a phased array of 16 high-frequency antennas with a total transmitted power on the order of 6.4 kilowatts. See Sigillito, V. G., Wing, S. P., Hutton, L. V., & Baker, K. B. (1989) for more details. The targets were free electrons in the ionosphere. "Good" radar returns are those showing evidence of some type of structure in the ionosphere. "Bad" returns are those that do not; their signals pass through the ionosphere.

### Usage

```
data(Ionosphere)
```

### Format

List containing the following elements:

**x** 351 by 34 matrix of numeric values.

**Class** Character vector of length 351 containing 126 entries labeled "bad" and 225 labeled "good".

---

```
predict.logistic_copula  
  predict.logistic_copula
```

---

**Description**

Computes predicted probability of  $Y=1$  for a logistic regression model with a vine extension.

**Usage**

```
## S3 method for class 'logistic_copula'  
predict(object, new_x, ...)
```

**Arguments**

<code>object</code>	The model object as returned by <code>fit_copula_interactions</code>
<code>new_x</code>	A matrix of covariate values to compute predictions for.
<code>...</code>	Not used.

**Value**

A numeric vector of estimates of the conditional probability of  $Y=1 \mid x$ , computed for each row of `new_x`.

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