# Package: irboost (via r-universe)

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```
Type Package
Title Iteratively Reweighted Boosting for Robust Analysis
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Description Fit a predictive model using iteratively reweighted
      boosting (IRBoost) to minimize robust loss functions within the
      CC-family (concave-convex). This constitutes an application of
      iteratively reweighted convex optimization (IRCO), where convex
      optimization is performed using the functional descent boosting
      algorithm. IRBoost assigns weights to facilitate outlier
      identification. Applications include robust generalized linear
      models and robust accelerated failure time models. Wang (2021)
      <doi:10.48550/arXiv.2101.07718>.
Depends R (>= 3.5.0)
Imports mpath (>= 0.4-2.21), xgboost
```

Suggests R.rsp, DiagrammeR, survival, Hmisc

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License GPL (>= 3)

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NeedsCompilation no

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#### **Description**

generate random data for classification as in Long and Servedio (2010)

#### Usage

```
dataLS(ntr, ntu = ntr, nte, percon)
```

# Arguments

ntr number of training data

ntu number of tuning data, default is the same as ntr

nte number of test data

percon proportion of contamination, must between 0 and 1. If percon > 0, the labels

of the corresponding percenrage of response variable in the training and tuning

data are flipped.

#### Value

a list with elements xtr, xtu, xte, ytr, ytu, yte for predictors of disjoint training, tuning and test data, and response variable -1/1 of training, tuning and test data.

# Author(s)

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#### References

P. Long and R. Servedio (2010), *Random classification noise defeats all convex potential boosters*, *Machine Learning Journal*, 78(3), 287–304.

```
dat <- dataLS(ntr=100, nte=100, percon=0)</pre>
```

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irb.train

fit a robust predictive model with iteratively reweighted boosting algorithm

#### Description

Fit a predictive model with the iteratively reweighted convex optimization (IRCO) that minimizes the robust loss functions in the CC-family (concave-convex). The convex optimization is conducted by functional descent boosting algorithm in the R package **xgboost**. The iteratively reweighted boosting (IRBoost) algorithm reduces the weight of the observation that leads to a large loss; it also provides weights to help identify outliers. Applications include the robust generalized linear models and extensions, where the mean is related to the predictors by boosting, and robust accelerated failure time models. irb.train is an advanced interface for training an irboost model. The irboost function is a simpler wrapper for irb.train. See xgboost::xgb.train.

#### Usage

```
irb.train(
   params = list(),
   data,
   z_init = NULL,
   cfun = "ccave",
   s = 1,
   delta = 0.1,
   iter = 10,
   nrounds = 100,
   del = 1e-10,
   trace = FALSE,
   ...
)
```

#### Arguments

params

the list of parameters, params is passed to function xgboost::xgb.train which requires the same argument. The list must include objective, a convex component in the CC-family, the second C, or convex down. It is the same as objective in the xgboost::xgb.train. The following objective functions are currently implemented:

- reg: squarederror Regression with squared loss.
- binary:logitraw logistic regression for binary classification, predict linear predictor, not probabilies.
- binary: hinge hinge loss for binary classification. This makes predictions of -1 or 1, rather than producing probabilities.
- multi:softprob softmax loss function for multiclass problems. The result contains predicted probabilities of each data point in each class, say p\_k, k=0, ..., nclass-1. Note, label is coded as in [0, ..., nclass-1]. The loss

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function cross-entropy for the i-th observation is computed as  $-log(p_k)$  with  $k=lable_i$ , i=1,...,n.

- count:poisson: Poisson regression for count data, predict mean of poisson distribution.
- reg:gamma: gamma regression with log-link, predict mean of gamma distribution. The implementation in xgboost::xgb.train takes a parameterization in the exponential family:

xgboost/src/src/metric/elementwise\_metric.cu.

In particularly, there is only one parameter psi and set to 1. The implementation of the IRCO algorithm follows this parameterization. See Table 2.1, McCullagh and Nelder, Generalized linear models, Chapman & Hall, 1989, second edition.

- reg: tweedie: Tweedie regression with log-link. See also tweedie\_variance\_power in range: (1,2). A value close to 2 is like a gamma distribution. A value close to 1 is like a Poisson distribution.
- survival: aft: Accelerated failure time model for censored survival time data. irb.train invokes irb.train\_aft.

data	training dataset. irb.train accepts only an xgboost::xgb.DMatrix as the
	input. irboost, in addition, also accepts matrix, dgCMatrix, or name of a
	local data file. See xgboost::xgb.train.

z\_init vector of nobs with initial convex component values, must be non-negative with default values = weights if data has provided, otherwise z\_init = vector of 1s

cfun concave component of CC-family, can be "hacve", "acave", "bcave", "ccave", "dcave", "ecave", "foave", "hcave".

See Table 2 https://arxiv.org/pdf/2010.02848.pdf

s tuning parameter of cfun. s > 0 and can be equal to 0 for cfun="tcave". If s is too close to 0 for cfun="acave", "bcave", "ccave", the calculated weights

can become 0 for all observations, thus crash the program

delta a small positive number provided by user only if cfun="gcave" and 0 < s <1

iter number of iteration in the IRCO algorithm

nrounds boosting iterations within each IRCO iteration

del convergency criteria in the IRCO algorithm, no relation to delta

trace if TRUE, fitting progress is reported

... other arguments passing to xgb.train

#### Value

An object with S3 class xgb. train with the additional elments:

- weight\_update\_log a matrix of nobs row by iter column of observation weights in each iteration of the IRCO algorithm
- weight\_update a vector of observation weights in the last IRCO iteration that produces the final model fit

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• loss\_log sum of loss value of the composite function in each IRCO iteration. Note, cfun requires objective non-negative in some cases. Thus care must be taken. For instance, with objective="reg:gamma", the loss value is defined by gamma-nloglik - (1+log(min(y))), where y=label. The second term is introduced such that the loss value is non-negative. In fact, gamma-nloglik=y/ypre + log(ypre) in the xgboost::xgb.train, where ypre is the mean prediction value, can be negative. It can be derived that for fixed y, the minimum value of gamma-nloglik is achived at ypre=y, or 1+log(y). Thus, among all label values, the minimum of gamma-nloglik is 1+log(min(y)).

#### Author(s)

```
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```

#### References

Wang, Zhu (2021), Unified Robust Boosting, arXiv eprint, https://arxiv.org/abs/2101.07718

```
# logistic boosting
data(agaricus.train, package='xgboost')
data(agaricus.test, package='xgboost')
dtrain <- with(agaricus.train, xgboost::xgb.DMatrix(data, label = label))</pre>
dtest <- with(agaricus.test, xgboost::xgb.DMatrix(data, label = label))</pre>
watchlist <- list(train = dtrain, eval = dtest)</pre>
# A simple irb.train example:
param <- list(max_depth = 2, eta = 1, nthread = 2,</pre>
objective = "binary:logitraw", eval_metric = "auc")
bst <- xgboost::xgb.train(params=param, data=dtrain, nrounds = 2,</pre>
                           watchlist=watchlist, verbose=2)
bst <- irb.train(params=param, data=dtrain, nrounds = 2)</pre>
summary(bst$weight_update)
# a bug in xgboost::xgb.train
#bst <- irb.train(params=param, data=dtrain, nrounds = 2,</pre>
                   watchlist=watchlist, trace=TRUE, verbose=2)
# time-to-event analysis
X <- matrix(1:5, ncol=1)</pre>
# Associate ranged labels with the data matrix.
# This example shows each kind of censored labels.
# uncensored right left interval
y_{lower} = c(10, 15, -Inf, 30, 100)
y_{upper} = c(Inf, Inf, 20, 50, Inf)
dtrain <- xgboost::xgb.DMatrix(data=X, label_lower_bound=y_lower,</pre>
                                 label_upper_bound=y_upper)
param <- list(objective="survival:aft", aft_loss_distribution="normal",</pre>
               aft_loss_distribution_scale=1, max_depth=3, min_child_weight=0)
watchlist <- list(train = dtrain)</pre>
bst <- xgboost::xgb.train(params=param, data=dtrain, nrounds=15,</pre>
```

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```
watchlist=watchlist)
predict(bst, dtrain)
bst_cc <- irb.train(params=param, data=dtrain, nrounds=15, cfun="hcave",</pre>
                     s=1.5, trace=TRUE, verbose=0)
bst_cc$weight_update
```

irb.train\_aft

fit a robust accelerated failure time model with iteratively reweighted boosting algorithm

# **Description**

Fit an accelerated failure time model with the iteratively reweighted convex optimization (IRCO) that minimizes the robust loss functions in the CC-family (concave-convex). The convex optimization is conducted by functional descent boosting algorithm in the R package xgboost. The iteratively reweighted boosting (IRBoost) algorithm reduces the weight of the observation that leads to a large loss; it also provides weights to help identify outliers. For time-to-event data, an accelerated failure time model (AFT model) provides an alternative to the commonly used proportional hazards models. Note, function irboost\_aft was developed to facilitate a data input format used with function xgb.train for objective=survival:aft in package xgboost. In other ojective functions, the input format is different with function xgboost at the time.

#### **Usage**

```
irb.train_aft(
  params = list(),
  data,
  z_init = NULL,
  cfun = "ccave",
  s = 1,
  delta = 0.1,
  iter = 10,
  nrounds = 100,
  del = 1e-10,
  trace = FALSE,
)
```

# **Arguments**

params

the list of parameters used in xgb. train of **xgboost**.

Must include aft\_loss\_distribution, aft\_loss\_distribution\_scale, but there is no need to include objective. The complete list of parameters is available in the online documentation.

data

training dataset. irboost\_aft accepts only an xgb.DMatrix as the input.

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z_init	vector of nobs with initial convex component values, must be non-negative with default values = weights if provided, otherwise z_init = vector of 1s
cfun	concave component of CC-family, can be "hacve", "acave", "bcave", "ccave", "dcave", "ecave", "gcave", "hcave". See Table 2 at https://arxiv.org/pdf/2010.02848.pdf
S	tuning parameter of cfun. $s > 0$ and can be equal to 0 for cfun="tcave". If s is too close to 0 for cfun="acave", "bcave", "ccave", the calculated weights can become 0 for all observations, thus crash the program
delta	a small positive number provided by user only if cfun="gcave" and 0 < s <1
iter	number of iteration in the IRCO algorithm
nrounds	boosting iterations in xgb.train within each IRCO iteration
del	convergency criteria in the IRCO algorithm, no relation to delta
trace	if TRUE, fitting progress is reported
	other arguments passing to xgb.train

#### Value

An object of class xgb.Booster with additional elements:

- weight\_update\_log a matrix of nobs row by iter column of observation weights in each iteration of the IRCO algorithm
- weight\_update a vector of observation weights in the last IRCO iteration that produces the final model fit
- loss\_log sum of loss value of the composite function cfun(survival\_aft\_distribution) in each IRCO iteration

# Author(s)

Zhu Wang

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# References

Wang, Zhu (2021), Unified Robust Boosting, arXiv eprint, https://arxiv.org/abs/2101.07718

#### See Also

irboost

```
library("xgboost")
X <- matrix(1:5, ncol=1)

# Associate ranged labels with the data matrix.
# This example shows each kind of censored labels.
# uncensored right left interval</pre>
```

irboost

fit a robust predictive model with iteratively reweighted boosting algorithm

#### **Description**

Fit a predictive model with the iteratively reweighted convex optimization (IRCO) that minimizes the robust loss functions in the CC-family (concave-convex). The convex optimization is conducted by functional descent boosting algorithm in the R package **xgboost**. The iteratively reweighted boosting (IRBoost) algorithm reduces the weight of the observation that leads to a large loss; it also provides weights to help identify outliers. Applications include the robust generalized linear models and extensions, where the mean is related to the predictors by boosting, and robust accelerated failure time models.

# Usage

```
irboost(
  data,
  label,
  weights,
  params = list(),
  z_init = NULL,
  cfun = "ccave",
  s = 1,
  delta = 0.1,
  iter = 10,
  nrounds = 100,
  del = 1e-10,
  trace = FALSE,
  ...
)
```

#### **Arguments**

data

input data, if objective="survival:aft", it must be an xgb.DMatrix; otherwise, it can be a matrix of dimension nobs x nvars; each row is an observation vector. Can accept dgCMatrix

label

response variable. Quantitative for objective="reg:squarederror", objective="count:poisson" (non-negative counts) or objective="reg:gamma" (positive). For objective="binary:logitraw" or "binary:hinge", label should be a factor with two levels

weights

vector of nobs with non-negative weights

params

the list of parameters, params is passed to function xgboost::xgboost which requires the same argument. The list must include objective, a convex component in the CC-family, the second C, or convex down. It is the same as objective in the xgboost::xgboost. The following objective functions are currently implemented:

- reg: squarederror Regression with squared loss.
- binary:logitraw logistic regression for binary classification, predict linear predictor, not probabilies.
- binary:hinge hinge loss for binary classification. This makes predictions of -1 or 1, rather than producing probabilities.
- multi:softprob softmax loss function for multiclass problems. The result contains predicted probabilities of each data point in each class, say p\_k, k=0, ..., nclass-1. Note, label is coded as in [0, ..., nclass-1]. The loss function cross-entropy for the i-th observation is computed as -log(p\_k) with k=lable\_i, i=1, ..., n.
- count:poisson: Poisson regression for count data, predict mean of poisson distribution.
- reg:gamma: gamma regression with log-link, predict mean of gamma distribution. The implementation in xgboost takes a parameterization in the exponential family:
  - xgboost/src/src/metric/elementwise\_metric.cu.
  - In particularly, there is only one parameter psi and set to 1. The implementation of the IRCO algorithm follows this parameterization. See Table 2.1, McCullagh and Nelder, Generalized linear models, Chapman & Hall, 1989, second edition.
- reg: tweedie: Tweedie regression with log-link. See also tweedie\_variance\_power in range: (1,2). A value close to 2 is like a gamma distribution. A value close to 1 is like a Poisson distribution.
- survival:aft: Accelerated failure time model for censored survival time data. irboost invokes irb.train\_aft.

z\_init

vector of nobs with initial convex component values, must be non-negative with default values = weights if provided, otherwise z\_init = vector of 1s

cfun

concave component of CC-family, can be "hacve", "acave", "bcave", "ccave", "dcave", "ecave", "gcave", "hcave".

See Table 2 at https://arxiv.org/pdf/2010.02848.pdf

S	tuning parameter of cfun. $s > 0$ and can be equal to 0 for cfun="tcave". If s is too close to 0 for cfun="acave", "bcave", "ccave", the calculated weights can become 0 for all observations, thus crash the program
delta	a small positive number provided by user only if cfun="gcave" and 0 < s <1
iter	number of iteration in the IRCO algorithm
nrounds	boosting iterations within each IRCO iteration
del	convergency criteria in the IRCO algorithm, no relation to delta
trace	if TRUE, fitting progress is reported
	other arguments passing to xgboost

#### Value

An object with S3 class xgboost with the additional elments:

- weight\_update\_log a matrix of nobs row by iter column of observation weights in each iteration of the IRCO algorithm
- weight\_update a vector of observation weights in the last IRCO iteration that produces the final model fit
- loss\_log sum of loss value of the composite function in each IRCO iteration. Note, cfun requires objective non-negative in some cases. Thus care must be taken. For instance, with objective="reg:gamma", the loss value is defined by gamma-nloglik (1+log(min(y))), where y=label. The second term is introduced such that the loss value is non-negative. In fact, gamma-nloglik=y/ypre + log(ypre) in the xgboost, where ypre is the mean prediction value, can be negative. It can be derived that for fixed y, the minimum value of gamma-nloglik is achived at ypre=y, or 1+log(y). Thus, among all label values, the minimum of gamma-nloglik is 1+log(min(y)).

#### Author(s)

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#### References

Wang, Zhu (2021), Unified Robust Boosting, arXiv eprint, https://arxiv.org/abs/2101.07718

```
params=list(objective="binary:hinge", max_depth=1), trace=TRUE,
                 verbose=0, nrounds=50)
fit4 <- irboost(data=x, label=g2, cfun="acave",s=0.5,</pre>
                 params=list(objective="count:poisson", max_depth=1), trace=TRUE,
                 verbose=0, nrounds=50)
# Gamma regression
x <- matrix(rnorm(100*2), 100, 2)
g2 <- sample(rgamma(100, 1))</pre>
library("xgboost")
param <- list(objective="reg:gamma", max_depth=1)</pre>
fit5 <- xgboost(data=x, label=g2, params=param, nrounds=50)</pre>
fit6 <- irboost(data=x, label=g2, cfun="acave",s=5, params=param, trace=TRUE,</pre>
                 verbose=0, nrounds=50)
plot(predict(fit5, newdata=x), predict(fit6, newdata=x))
hist(fit6$weight_update)
plot(fit6$loss_log)
summary(fit6$weight_update)
# Tweedie regression
param <- list(objective="reg:tweedie", max_depth=1)</pre>
fit6t <- irboost(data=x, label=g2, cfun="acave",s=5, params=param,</pre>
                  trace=TRUE, verbose=0, nrounds=50)
# Gamma vs Tweedie regression
hist(fit6$weight_update)
hist(fit6t$weight_update)
plot(predict(fit6, newdata=x), predict(fit6t, newdata=x))
# multiclass classification in iris dataset:
lb <- as.numeric(iris$Species)-1</pre>
num_class <- 3</pre>
set.seed(11)
param <- list(objective="multi:softprob", max_depth=4, eta=0.5, nthread=2,</pre>
subsample=0.5, num_class=num_class)
fit7 <- irboost(data=as.matrix(iris[, -5]), label=lb, cfun="acave", s=50,</pre>
                 params=param, trace=TRUE, verbose=0, nrounds=10)
# predict for softmax returns num_class probability numbers per case:
pred7 <- predict(fit7, newdata=as.matrix(iris[, -5]))</pre>
# reshape it to a num_class-columns matrix
pred7 <- matrix(pred7, ncol=num_class, byrow=TRUE)</pre>
# convert the probabilities to softmax labels
pred7_labels <- max.col(pred7) - 1</pre>
# classification error: 0!
sum(pred7_labels != lb)/length(lb)
table(lb, pred7_labels)
hist(fit7$weight_update)
```

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