

# Package ‘noisemodel’

May 9, 2026

**Version** 1.0.2

**Title** Noise Models for Classification Datasets

**Description** Implementation of models for the controlled introduction of errors in classification datasets. This package contains the noise models described in Saez (2022) <[doi:10.3390/math10203736](https://doi.org/10.3390/math10203736)> that allow corrupting class labels, attributes and both simultaneously.

**License** GPL (>= 3)

**Depends** R (>= 3.5.0)

**Imports** caret, nnet, e1071, FNN, classInt, ggplot2, ExtDist, lsr, stringr, RColorBrewer, RSNNS, C50

**Suggests** knitr, rmarkdown, testthat (>= 3.0.0)

**VignetteBuilder** knitr

**Encoding** UTF-8

**RoxygenNote** 7.2.1

**NeedsCompilation** no

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**Repository** CRAN

**Date/Publication** 2022-10-17 06:20:02 UTC

**Config/testthat/edition** 3

## Contents

asy_def_ln . . . . .	3
asy_int_an . . . . .	5
asy_spa_ln . . . . .	7
asy_uni_an . . . . .	9
asy_uni_ln . . . . .	11
atm_uni_ln . . . . .	12
boud_gau_an . . . . .	14
clu_vot_ln . . . . .	16

diris2D . . . . .	18
exps_cuni_ln . . . . .	19
exp_bor_ln . . . . .	20
fra_bdir_ln . . . . .	22
gam_bor_ln . . . . .	24
gaum_bor_ln . . . . .	26
gau_bor_ln . . . . .	28
glev_uni_ln . . . . .	30
hubp_uni_ln . . . . .	32
imp_int_an . . . . .	33
iris2D . . . . .	35
irs_bdir_ln . . . . .	36
lap_bor_ln . . . . .	38
larm_uni_ln . . . . .	40
maj_udir_ln . . . . .	42
mind_bdir_ln . . . . .	44
minp_uni_ln . . . . .	46
mis_pre_ln . . . . .	47
mulc_udir_ln . . . . .	49
nei_bor_ln . . . . .	51
nlin_bor_ln . . . . .	53
oned_uni_ln . . . . .	54
opes_idnn_ln . . . . .	57
opes_idu_ln . . . . .	59
pai_bdir_ln . . . . .	61
plot.ndmodel . . . . .	63
pmd_con_ln . . . . .	64
print.ndmodel . . . . .	66
qua_uni_ln . . . . .	67
sco_con_ln . . . . .	69
sigb_uni_ln . . . . .	71
smam_bor_ln . . . . .	72
smu_cuni_ln . . . . .	74
summary.ndmodel . . . . .	76
symd_gau_an . . . . .	77
symd_gimg_an . . . . .	79
symd_rpix_an . . . . .	81
symd_uni_an . . . . .	83
sym_adj_ln . . . . .	84
sym_cen_ln . . . . .	86
sym_con_ln . . . . .	88
sym_cuni_an . . . . .	90
sym_cuni_cn . . . . .	91
sym_cuni_ln . . . . .	93
sym_ddef_ln . . . . .	95
sym_def_ln . . . . .	97
sym_dia_ln . . . . .	99
sym_dran_ln . . . . .	101

sym_end_an . . . . .	102
sym_exc_ln . . . . .	104
sym_gau_an . . . . .	106
sym_hienc_ln . . . . .	108
sym_hie_ln . . . . .	110
sym_int_an . . . . .	112
sym_natd_ln . . . . .	113
sym_nean_ln . . . . .	115
sym_nexc_ln . . . . .	117
sym_nuni_ln . . . . .	119
sym_opt_ln . . . . .	121
sym_pes_ln . . . . .	123
sym_sgau_an . . . . .	125
sym_uni_an . . . . .	126
sym_uni_ln . . . . .	128
sym_usim_ln . . . . .	130
ugau_bor_ln . . . . .	132
ulap_bor_ln . . . . .	134
uncs_guni_cn . . . . .	136
unc_fixw_an . . . . .	138
unc_vgau_an . . . . .	140

**Index****142**


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asy_def_ln	<i>Asymmetric default label noise</i>
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---

**Description**

Introduction of *Asymmetric default label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
asy_def_ln(x, y, level, def = 1, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
asy_def_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double vector with the noise levels in [0,1] to be introduced into each class.
def	an integer with the index of the default class (default: 1).
order	a character vector indicating the order of the classes (default: levels(y)).

sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

*Asymmetric default label noise* randomly selects  $(level[i] \cdot 100)\%$  of the samples of each class  $C[i]$  in the dataset -the order of the class labels is determined by order. Then, the labels of these samples are replaced by a fixed label ( $C[def]$ ) within the set of class labels.

### Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References.

### References

R. C. Prati, J. Luengo, and F. Herrera. **Emerging topics and challenges of learning from noisy data in nonstandard classification: a survey beyond binary class noise.** *Knowledge and Information Systems*, 60(1):63–97, 2019. doi:[10.1007/s1011501812444](https://doi.org/10.1007/s1011501812444).

### See Also

[sym\\_nean\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- asy_def_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = c(0.1, 0.2, 0.3), order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- asy_def_ln(formula = Species ~ ., data = iris2D,
                    level = c(0.1, 0.2, 0.3), order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

asy\_int\_an

*Asymmetric interval-based attribute noise*


---

**Description**

Introduction of *Asymmetric interval-based attribute noise* into a classification dataset.

**Usage**

```

## Default S3 method:
asy_int_an(x, y, level, nbins = 10, sortid = TRUE, ...)

## S3 method for class 'formula'
asy_int_an(formula, data, ...)

```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double vector with the noise levels in [0,1] to be introduced into each attribute.
nbins	an integer with the number of bins to create (default: 10).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.



```
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- asy_int_an(formula = Species ~ ., data = iris2D,
                    level = c(0.1, 0.2))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

asy\_spa\_ln

*Asymmetric sparse label noise*


---

## Description

Introduction of *Asymmetric sparse label noise* into a classification dataset.

## Usage

```
## Default S3 method:
asy_spa_ln(x, y, level0, levelE, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
asy_spa_ln(formula, data, ...)
```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level0	a double with the noise level in [0,1] to be introduced into each odd class.
levelE	a double with the noise level in [0,1] to be introduced into each even class.
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Asymmetric sparse label noise* randomly selects  $(level0 \cdot 100)\%$  of the samples in each odd class and  $(levelE \cdot 100)\%$  of the samples in each even class -the order of the class labels is determined by order. Then, each odd class is flipped to the next class, whereas each even class is flipped to the previous class. If the dataset has an odd number of classes, the last class is not corrupted.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

J. Wei and Y. Liu. **When optimizing f-divergence is robust with label noise**. In *Proc. 9th International Conference on Learning Representations*, pages 1-11, 2021. url:<https://openreview.net/forum?id=WesiCoRVQ15>.

**See Also**

[mind\\_bdir\\_ln](#), [fra\\_bdir\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- asy_spa_ln(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level0 = 0.1, levelE = 0.3, order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- asy_spa_ln(formula = Species ~ ., data = iris2D,
                    level0 = 0.1, levelE = 0.3, order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
```

```
identical(outdef$idnoise, outfrm$idnoise)
```

---

```
asy_uni_an
```

```
Asymmetric uniform attribute noise
```

---

## Description

Introduction of *Asymmetric uniform attribute noise* into a classification dataset.

## Usage

```
## Default S3 method:
asy_uni_an(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
asy_uni_an(formula, data, ...)
```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double vector with the noise levels in [0,1] to be introduced into each attribute.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Asymmetric uniform attribute noise* corrupts  $(level[i] \cdot 100)\%$  of the values for each attribute  $A[i]$  in the dataset. In order to corrupt an attribute  $A[i]$ ,  $(level[i] \cdot 100)\%$  of the samples in the dataset are chosen. Then, their values for  $A[i]$  are replaced by random different ones between the minimum and maximum of the domain of the attribute following a uniform distribution (for numerical attributes) or choosing a random value (for nominal attributes).

## Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per attribute.
idnoise	an integer vector list with the indices of noisy samples per attribute.
numclean	an integer vector with the amount of clean samples per attribute.

idclean	an integer vector list with the indices of clean samples per attribute.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References.

### References

A. Petety, S. Tripathi, and N. Hemachandra. **Attribute noise robust binary classification**. In *Proc. 34th AAAI Conference on Artificial Intelligence*, pages 13897-13898, 2020.

### See Also

[synd\\_gimg\\_an](#), [unc\\_vgau\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- asy_uni_an(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = c(0.1, 0.2))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- asy_uni_an(formula = Species ~ ., data = iris2D,
                    level = c(0.1, 0.2))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

asy_uni_ln	<i>Asymmetric uniform label noise</i>
------------	---------------------------------------

---

**Description**

Introduction of *Asymmetric uniform label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
asy_uni_ln(x, y, level, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
asy_uni_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double vector with the noise levels in [0,1] to be introduced into each class.
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Asymmetric uniform label noise* randomly selects (level[i]·100)% of the samples of each class C[i] in the dataset -the order of the class labels is determined by order. Finally, the labels of these samples are randomly replaced by other different ones within the set of class labels.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

Z. Zhao, L. Chu, D. Tao, and J. Pei. **Classification with label noise: a Markov chain sampling framework**. *Data Mining and Knowledge Discovery*, 33(5):1468-1504, 2019. doi:[10.1007/s10618-01805928](https://doi.org/10.1007/s10618-01805928).

**See Also**

[maj\\_udir\\_ln](#), [asy\\_def\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- asy_uni_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                   level = c(0.1, 0.2, 0.3), order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- asy_uni_ln(formula = Species ~ ., data = iris2D,
                   level = c(0.1, 0.2, 0.3), order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

attm\_uni\_ln

*Attribute-mean uniform label noise*


---

**Description**

Introduction of *Attribute-mean uniform label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
attm_uni_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
attm_uni_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

For each sample, its distance to the mean of each attribute is computed. Then,  $(level \cdot 100)\%$  of the samples in the dataset are randomly selected to be mislabeled, more likely choosing samples whose features are generally close to the mean. The labels of these samples are randomly replaced by other different ones within the set of class labels.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References

**References**

B. Nicholson, V. S. Sheng, and J. Zhang. **Label noise correction and application in crowdsourcing**. *Expert Systems with Applications*, 66:149-162, 2016. doi:10.1016/j.eswa.2016.09.003.

**See Also**

[qua\\_uni\\_ln](#), [exps\\_cuni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- attm_uni_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- attm_uni_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

boud\_gau\_an

*Boundary/dependent Gaussian attribute noise*


---

**Description**

Introduction of *Boundary/dependent Gaussian attribute noise* into a classification dataset.

**Usage**

```

## Default S3 method:
boud_gau_an(x, y, level, k = 0.2, sortid = TRUE, ...)

## S3 method for class 'formula'
boud_gau_an(formula, data, ...)

```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
k	a double in [0,1] with the scale used for the standard deviation (default: 0.2).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Boundary/dependent Gaussian attribute noise* corrupts  $(level \cdot 100)\%$  samples among the  $((level + 0.1) \cdot 100)\%$  of samples closest to the decision boundary. Their attribute values are corrupted by adding a random number that follows a Gaussian distribution of *mean* = 0 and *standard deviation* =  $(max - min) \cdot k$ , being *max* and *min* the limits of the attribute domain. For nominal attributes, a random value is chosen.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per attribute.
<code>idnoise</code>	an integer vector list with the indices of noisy samples per attribute.
<code>numclean</code>	an integer vector with the amount of clean samples per attribute.
<code>idclean</code>	an integer vector list with the indices of clean samples per attribute.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

J. Bi and T. Zhang. **Support vector classification with input data uncertainty**. In *Advances in Neural Information Processing Systems*, volume 17, pages 161-168, 2004. url:<https://proceedings.neurips.cc/paper/2004/hash/22b1f2e0983160db6f7bb9f62f4dbb39-Abstract.html>.

**See Also**

[imp\\_int\\_an](#), [asy\\_int\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- boud_gau_an(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)
```

```
# usage of the method for class formula
set.seed(9)
outfrm <- boud_gau_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

clu\_vot\_ln

*Clustering-based voting label noise*


---

## Description

Introduction of *Clustering-based voting label noise* into a classification dataset.

## Usage

```
## Default S3 method:
clu_vot_ln(x, y, k = nlevels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
clu_vot_ln(formula, data, ...)
```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
k	an integer with the number of clusters (default: nlevels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Clustering-based voting label noise* divides the dataset into k clusters. Then, the labels of each cluster are relabeled with the majority class among its samples.

## Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.

idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References, which considers  $k$ -means as unsupervised clustering method.

### References

Q. Wang, B. Han, T. Liu, G. Niu, J. Yang, and C. Gong. **Tackling instance-dependent label noise via a universal probabilistic model**. In *Proc. 35th AAAI Conference on Artificial Intelligence*, pages 10183-10191, 2021. url:<https://ojs.aaai.org/index.php/AAAI/article/view/17221>.

### See Also

[sco\\_con\\_ln](#), [mis\\_pre\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- clu_vot_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)])

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- clu_vot_ln(formula = Species ~ ., data = iris2D)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

`diris2D`*diris2D dataset*

---

**Description**

Discretized version of the `iris2D` dataset.

**Usage**

```
data(diris2D)
```

**Format**

A `data.frame` with 103 samples (rows) and 3 variables (columns) named `Petal.Length`, `Petal.Width` and `Species`.

**Source**

Data collected by E. Anderson (1935).

**References**

R. A. Fisher. **The use of multiple measurements in taxonomic problems.** *Annals of Eugenics*, 7:179-188, 1936.

E. Anderson. **The irises of the Gaspé Peninsula.** *Bulletin of the American Iris Society*, 59:2-5, 1935.

**See Also**

[iris2D](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(diris2D)

# noise introduction
set.seed(9)
outdef <- sym_uni_ln(x = diris2D[,-ncol(diris2D)], y = diris2D[,ncol(diris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
```

---

exps_cuni_ln	<i>Exponential/smudge completely-uniform label noise</i>
--------------	--

---

### Description

Introduction of *Exponential/smudge completely-uniform label noise* into a classification dataset.

### Usage

```
## Default S3 method:
exps_cuni_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
exps_cuni_ln(formula, data, ...)
```

### Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the lambda value.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

*Exponential/smudge completely-uniform label noise* includes an additional attribute (*smudge*) in the dataset with random values in [0,1]. This attribute is used to compute the mislabeling probability for each sample based on an exponential function (in which *level* is used as lambda). It selects samples in the dataset based on these probabilities. Finally, the labels of these samples are randomly replaced by others within the set of class labels (this model can choose the original label of a sample as noisy).

### Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.

distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References.

### References

B. Denham, R. Pears, and M. A. Naeem. **Null-labelling: A generic approach for learning in the presence of class noise**. In *Proc. 20th IEEE International Conference on Data Mining*, pages 990–995, 2020. doi:10.1109/ICDM50108.2020.00114.

### See Also

[opes\\_idu\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- exps_cuni_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.8)

# show results
summary(outdef, showid = TRUE)
plot(outdef, pca = TRUE)

# usage of the method for class formula
set.seed(9)
outfrm <- exps_cuni_ln(formula = Species ~ ., data = iris2D, level = 0.8)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

exp\_bor\_ln

*Exponential borderline label noise*

---

### Description

Introduction of *Exponential borderline label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
exp_bor_ln(x, y, level, rate = 1, k = 1, sortid = TRUE, ...)

## S3 method for class 'formula'
exp_bor_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
rate	a double with the rate for the exponential distribution (default: 1).
k	an integer with the number of nearest neighbors to be used (default: 1).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Exponential borderline label noise* uses an SVM to induce the decision border in the dataset. For each sample, its distance to the decision border is computed. Then, an exponential distribution with parameter `rate` is used to compute the value for the probability density function associated to each distance. Finally,  $(level \cdot 100)\%$  of the samples in the dataset are randomly selected to be mislabeled according to their values of the probability density function. For each noisy sample, the majority class among its `k`-nearest neighbors of a different class is chosen as the new label.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References, considering SVM with linear kernel as classifier, a mislabeling process using the neighborhood of noisy samples and a noise level to control the number of errors in the data.

**References**

J. Bootkrajang. **A generalised label noise model for classification in the presence of annotation errors.** *Neurocomputing*, 192:61–71, 2016. doi:10.1016/j.neucom.2015.12.106.

**See Also**

[pmd\\_con\\_ln](#), [clu\\_vot\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- exp_bor_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- exp_bor_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

fra\_bdir\_ln

*Fraud bidirectional label noise*


---

**Description**

Introduction of *Fraud bidirectional label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
fra_bdir_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
fra_bdir_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Fraud bidirectional label noise* randomly selects  $(level \cdot 100)\%$  of the samples from the minority class in the dataset and  $level \cdot 10$  samples from the majority class. Then, minority class samples are mislabeled as belonging to the majority class and majority class samples are mislabeled as belonging to the minority class. In case of ties determining minority and majority classes, a random class is chosen among them.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

Z. Salekshahrezaee, J. L. Leevy, and T. M. Khoshgoftaar. **A reconstruction error-based framework for label noise detection.** *Journal of Big Data*, 8(1):1-16, 2021. doi:10.1186/s40537021-004475.

**See Also**

[irs\\_bdir\\_ln](#), [pai\\_bdir\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- fra_bdir_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- fra_bdir_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

gam\_bor\_ln

*Gamma borderline label noise***Description**

Introduction of *Gamma borderline label noise* into a classification dataset.

**Usage**

```

## Default S3 method:
gam_bor_ln(x, y, level, shape = 1, rate = 0.5, k = 1, sortid = TRUE, ...)

## S3 method for class 'formula'
gam_bor_ln(formula, data, ...)

```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
shape	a double with the shape for the gamma distribution (default: 1)
rate	a double with the rate for the gamma distribution (default: 0.5).
k	an integer with the number of nearest neighbors to be used (default: 1).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Gamma borderline label noise* uses an SVM to induce the decision border in the dataset. For each sample, its distance to the decision border is computed. Then, a gamma distribution with parameters (shape, rate) is used to compute the value for the probability density function associated to each distance. Finally, (level·100)% of the samples in the dataset are randomly selected to be mislabeled according to their values of the probability density function. For each noisy sample, the majority class among its k-nearest neighbors of a different class is chosen as the new label.

## Value

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

## Note

Noise model adapted from the papers in References, considering SVM with linear kernel as classifier, a mislabeling process using the neighborhood of noisy samples and a noise level to control the number of errors in the data.

## References

J. Bootkrajang. **A generalised label noise model for classification**. In *Proc. 23rd European Symposium on Artificial Neural Networks*, pages 349-354, 2015. url:<https://dblp.org/rec/conf/esann/Bootkrajang15.html?view=bibtex>.

## See Also

[exp\\_bor\\_ln](#), [pmd\\_con\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- gam_bor_ln(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)
```

```

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- gam_bor_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

gaum\_bor\_ln

*Gaussian-mixture borderline label noise*


---

## Description

Introduction of *Gaussian-mixture borderline label noise* into a classification dataset.

## Usage

```

## Default S3 method:
gaum_bor_ln(
  x,
  y,
  level,
  mean = c(0, 2),
  sd = c(sqrt(0.5), sqrt(0.5)),
  w = c(0.5, 0.5),
  k = 1,
  sortid = TRUE,
  ...
)

## S3 method for class 'formula'
gaum_bor_ln(formula, data, ...)

```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
mean	a double vector with the mean for each Gaussian distribution (default: c(0,2)).
sd	a double vector with the standard deviation for each Gaussian distribution (default: c(sqrt(0.5),sqrt(0.5))).
w	a double vector with the weight for each Gaussian distribution (default: c(0.5,0.5)).

k	an integer with the number of nearest neighbors to be used (default: 1).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

*Gaussian-mixture borderline label noise* uses an SVM to induce the decision border in the dataset. For each sample, its distance to the decision border is computed. Then, a Gaussian mixture distribution with parameters (mean, sd) and weights  $w$  is used to compute the value for the probability density function associated to each distance. Finally,  $(level \cdot 100)\%$  of the samples in the dataset are randomly selected to be mislabeled according to their values of the probability density function. For each noisy sample, the majority class among its  $k$ -nearest neighbors of a different class is chosen as the new label.

### Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References, considering SVM with linear kernel as classifier, a mislabeling process using the neighborhood of noisy samples and a noise level to control the number of errors in the data.

### References

J. Bootkrajang and J. Chaijaruwanich. **Towards instance-dependent label noise-tolerant classification: a probabilistic approach.** *Pattern Analysis and Applications*, 23(1):95-111, 2020. [doi:10.1007/s100440180750z](https://doi.org/10.1007/s100440180750z).

### See Also

[gau\\_bor\\_ln](#), [sigb\\_uni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- gaum_bor_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- gaum_bor_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

gau\_bor\_ln

*Gaussian borderline label noise*


---

**Description**

Introduction of *Gaussian borderline label noise* into a classification dataset.

**Usage**

```

## Default S3 method:
gau_bor_ln(x, y, level, mean = 0, sd = 1, k = 1, sortid = TRUE, ...)

## S3 method for class 'formula'
gau_bor_ln(formula, data, ...)

```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
mean	a double with the mean for the Gaussian distribution (default: 0).
sd	a double with the standard deviation for the Gaussian distribution (default: 1).
k	an integer with the number of nearest neighbors to be used (default: 1).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Gaussian borderline label noise* uses an SVM to induce the decision border in the dataset. For each sample, its distance to the decision border is computed. Then, a Gaussian distribution with parameters (mean, sd) is used to compute the value for the probability density function associated to each distance. Finally,  $(level \cdot 100)\%$  of the samples in the dataset are randomly selected to be mislabeled according to their values of the probability density function. For each noisy sample, the majority class among its k-nearest neighbors of a different class is chosen as the new label.

## Value

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

## Note

Noise model adapted from the papers in References to multiclass data, considering SVM with linear kernel as classifier, a mislabeling process using the neighborhood of noisy samples and a noise level to control the number of errors in the data.

## References

J. Bootkrajang and J. Chaijaruwanich. **Towards instance-dependent label noise-tolerant classification: a probabilistic approach.** *Pattern Analysis and Applications*, 23(1):95-111, 2020. [doi:10.1007/s100440180750z](https://doi.org/10.1007/s100440180750z).

## See Also

[sigb\\_uni\\_ln](#), [larm\\_uni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- gau_bor_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1)
```

```

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- gau_bor_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

glev\_uni\_ln

*Gaussian-level uniform label noise*


---

## Description

Introduction of *Gaussian-level uniform label noise* into a classification dataset.

## Usage

```

## Default S3 method:
glev_uni_ln(x, y, level, sd = 0.01, sortid = TRUE, ...)

## S3 method for class 'formula'
glev_uni_ln(formula, data, ...)

```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sd	a double with the standard deviation for the Gaussian distribution (default: 0.01).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

For each sample, *Gaussian-level uniform label noise* assigns a random probability following a Gaussian distribution of mean = level and standard deviation sd. Noisy samples are chosen according to these probabilities. The labels of these samples are randomly replaced by other different ones within the set of class labels.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

D. Liu, G. Yang, J. Wu, J. Zhao, and F. Lv. **Robust binary loss for multi-category classification with label noise**. In *Proc. 2021 IEEE International Conference on Acoustics, Speech and Signal Processing*, pages 1700-1704, 2021. doi:[10.1109/ICASSP39728.2021.9414493](https://doi.org/10.1109/ICASSP39728.2021.9414493).

**See Also**

[sym\\_hienc\\_ln](#), [sym\\_nexc\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- glev_uni_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- glev_uni_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

 hubp\_uni\_ln

*Hubness-proportional uniform label noise*


---

### Description

Introduction of *Hubness-proportional uniform label noise* into a classification dataset.

### Usage

```
## Default S3 method:
hubp_uni_ln(x, y, level, k = 3, sortid = TRUE, ...)

## S3 method for class 'formula'
hubp_uni_ln(formula, data, ...)
```

### Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
k	an integer with the number of neighbors to compute the hubness of each sample (default: 3).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

*Hubness-proportional uniform label noise* is based on the presence of hubs in the dataset. It selects  $(level \cdot 100)\%$  of the samples in the dataset using a discrete probability distribution based on the concept of hubness, which is computed using the nearest neighbors of each sample. Then, the class labels of these samples are randomly replaced by different ones from the  $c$  classes.

### Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.

distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

N. Tomasev and K. Buza. **Hubness-aware kNN classification of high-dimensional data in presence of label noise.** *Neurocomputing*, 160:157-172, 2015. doi:10.1016/j.neucom.2014.10.084.

**See Also**

[smu\\_cuni\\_ln](#), [oned\\_uni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- hubp_uni_ln(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- hubp_uni_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

imp\_int\_an

---

*Importance interval-based attribute noise*


---

**Description**

Introduction of *Importance interval-based attribute noise* into a classification dataset.

**Usage**

```
## Default S3 method:
imp_int_an(x, y, level, nbins = 10, ascending = TRUE, sortid = TRUE, ...)

## S3 method for class 'formula'
imp_int_an(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double vector with the noise levels in [0,1] to be introduced into each attribute.
nbins	an integer with the number of bins to create (default: 10).
ascending	a boolean indicating how noise levels are assigned to attributes: <ul style="list-style-type: none"> <li>• TRUE: the lowest noise level is assigned to the most important attribute (default value).</li> <li>• FALSE: the highest noise level is assigned to the most important attribute.</li> </ul>
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

The values in `level` are ordered and assigned to attributes according to their information gain (using the ordering given by `ascending`). Then, *Importance interval-based attribute noise* corrupts  $(level[i] \cdot 100)\%$  of the values for each attribute  $A[i]$  in the dataset. In order to corrupt each attribute  $A[i]$ ,  $(level[i] \cdot 100)\%$  of the samples in the dataset are chosen. To corrupt a value in numeric attributes, the attribute is split into equal-frequency intervals, one of its closest intervals is picked out and a random value within the interval is chosen as noisy. For nominal attributes, a random value within the domain is chosen.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per attribute.
idnoise	an integer vector list with the indices of noisy samples per attribute.
numclean	an integer vector with the amount of clean samples per attribute.
idclean	an integer vector list with the indices of clean samples per attribute.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

M. V. Mannino, Y. Yang, and Y. Ryu. **Classification algorithm sensitivity to training data with non representative attribute noise**. *Decision Support Systems*, 46(3):743-751, 2009. doi:[10.1016/j.dss.2008.11.021](https://doi.org/10.1016/j.dss.2008.11.021).

**See Also**

[asy\\_int\\_an](#), [asy\\_uni\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- imp_int_an(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = c(0.1, 0.2))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- imp_int_an(formula = Species ~ ., data = iris2D,
                    level = c(0.1, 0.2))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

iris2D

*iris2D dataset*

---

**Description**

A 2-dimensional version of the well-known *iris* dataset. It maintains the attributes `Petal.Length` and `Petal.Width`, which give the measurements in centimeters of the petal length and width of iris flowers belonging to three different species (*setosa*, *versicolor* and *virginica*). Duplicate and contradictory samples are removed from the dataset, resulting in a total of 103 samples.

**Usage**

```
data(iris2D)
```

**Format**

A data.frame with 103 samples (rows) and 3 variables (columns) named Petal.Length, Petal.Width and Species.

**Source**

Data collected by E. Anderson (1935).

**References**

R. A. Fisher. **The use of multiple measurements in taxonomic problems.** *Annals of Eugenics*, 7:179-188, 1936.

E. Anderson. **The irises of the Gaspé Peninsula.** *Bulletin of the American Iris Society*, 59:2-5, 1935.

**See Also**

[sym\\_uni\\_ln](#), [sym\\_uni\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
library(ggplot2)
library(RColorBrewer)

data(iris2D)

ggplot(data = iris2D, aes(x = iris2D[,1], y = iris2D[,2], color = iris2D[,3])) +
  geom_point(stroke = 0.5) +
  xlim(min(iris2D[,1]), max(iris2D[,1])) +
  ylim(min(iris2D[,2]), max(iris2D[,2])) +
  xlab(names(iris2D)[1]) +
  ylab(names(iris2D)[2]) +
  labs(color='Species') +
  scale_color_manual(values = brewer.pal(3, "Dark2")) +
  theme(panel.border = element_rect(colour = "black", fill=NA),
        aspect.ratio = 1,
        axis.text = element_text(colour = 1, size = 12),
        legend.background = element_blank(),
        legend.box.background = element_rect(colour = "black"))
```

---

 irs\_bdir\_ln

*IR-stable bidirectional label noise*


---

**Description**

Introduction of *IR-stable bidirectional label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
irs_bdir_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
irs_bdir_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*IR-stable bidirectional label noise* randomly selects  $(level \cdot 100)\%$  of the samples from the minority class in the dataset and the same amount of samples from the majority class. Then, minority class samples are mislabeled as belonging to the majority class and majority class samples are mislabeled as belonging to the minority class. In case of ties determining minority and majority classes, a random class is chosen among them.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

## References

B. Chen, S. Xia, Z. Chen, B. Wang, and G. Wang. **RSMOTE: A self-adaptive robust SMOTE for imbalanced problems with label noise.** *Information Sciences*, 553:397-428, 2021. doi:10.1016/j.ins.2020.10.013.

## See Also

[pai\\_bdir\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- irs_bdir_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- irs_bdir_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

lap\_bor\_ln

*Laplace borderline label noise*

---

## Description

Introduction of *Laplace borderline label noise* into a classification dataset.

## Usage

```
## Default S3 method:
lap_bor_ln(x, y, level, mu = 0, b = 1, k = 1, sortid = TRUE, ...)

## S3 method for class 'formula'
lap_bor_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
mu	a double with the location for the Laplace distribution (default: 0).
b	a double with the scale for the Laplace distribution (default: 1).
k	an integer with the number of nearest neighbors to be used (default: 1).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Laplace borderline label noise* uses an SVM to induce the decision border in the dataset. For each sample, its distance to the decision border is computed. Then, a Laplace distribution with parameters ( $\mu$ ,  $b$ ) is used to compute the value for the probability density function associated to each distance. Finally,  $(level \cdot 100)\%$  of the samples in the dataset are randomly selected to be mislabeled according to their values of the probability density function. For each noisy sample, the majority class among its  $k$ -nearest neighbors of a different class is chosen as the new label.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References to multiclass data, considering SVM with linear kernel as classifier, a mislabeling process using the neighborhood of noisy samples and a noise level to control the number of errors in the data.

## References

J. Du and Z. Cai. **Modelling class noise with symmetric and asymmetric distributions**. In *Proc. 29th AAAI Conference on Artificial Intelligence*, pages 2589-2595, 2015. url:<https://dl.acm.org/doi/10.5555/2886521.2886681>.

## See Also

[ugau\\_bor\\_ln](#), [gaum\\_bor\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- lap_bor_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- lap_bor_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

larm\_uni\_ln

*Large-margin uniform label noise*

---

## Description

Introduction of *Large-margin uniform label noise* into a classification dataset.

## Usage

```
## Default S3 method:
larm_uni_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
larm_uni_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Large-margin uniform label noise* uses an SVM to induce the decision border in the dataset. For each sample, its distance to the decision border is computed. Then, the samples are ordered according to their distance and  $(level \cdot 100)\%$  of the most distant correctly classified samples to the decision boundary are selected to be mislabeled with a random different class.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References to multiclass data, considering SVM with linear kernel as classifier.

**References**

E. Amid, M. K. Warmuth, and S. Srinivasan. **Two-temperature logistic regression based on the Tsallis divergence**. In *Proc. 22nd International Conference on Artificial Intelligence and Statistics*, volume 89 of PMLR, pages 2388-2396, 2019. url:<http://proceedings.mlr.press/v89/amid19a.html>.

**See Also**

[hubp\\_uni\\_ln](#), [smu\\_cuni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- larm_uni_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.3)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- larm_uni_ln(formula = Species ~ ., data = iris2D, level = 0.3)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

maj\_udir\_ln

*Majority-class unidirectional label noise*


---

**Description**

Introduction of *Majority-class unidirectional label noise* into a classification dataset.

**Usage**

```

## Default S3 method:
maj_udir_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
maj_udir_ln(formula, data, ...)

```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

Let  $A$  be the majority class and  $B$  be the second majority class in the dataset. The *Majority-class unidirectional label noise* introduction model randomly selects  $(level \cdot 100)\%$  of the samples of  $A$  and labels them as  $B$ .

## Value

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

## Note

Noise model adapted from the papers in References to multiclass data.

## References

J. Li, Q. Zhu, Q. Wu, Z. Zhang, Y. Gong, Z. He, and F. Zhu. **SMOTE- NaN-DE: Addressing the noisy and borderline examples problem in imbalanced classification by natural neighbors and differential evolution.** *Knowledge-Based Systems*, 223:107056, 2021. doi:10.1016/j.knosys.2021.107056.

## See Also

[asy\\_def\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- maj_udir_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)
```

```
# usage of the method for class formula
set.seed(9)
outfrm <- maj_udir_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

mind\_bdir\_ln

*Minority-driven bidirectional label noise*


---

### Description

Introduction of *Minority-driven bidirectional label noise* into a classification dataset.

### Usage

```
## Default S3 method:
mind_bdir_ln(x, y, level, pos = 0.1, sortid = TRUE, ...)

## S3 method for class 'formula'
mind_bdir_ln(formula, data, ...)
```

### Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
pos	a double in [0,1] with the proportion of samples from the positive class (default: 0.1).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

*Minority-driven bidirectional label noise* randomly selects  $n = 2m \cdot \text{level}$  samples in the dataset (with  $m$  the number of samples in the minority class), making sure that  $n \cdot \text{pos}$  samples belong to the minority class and the rest to the majority class. Then, minority class samples are mislabeled as belonging to the majority class and majority class samples are mislabeled as belonging to the minority class. In case of ties determining minority and majority classes, a random class is chosen among them.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References to multiclass data.

**References**

A. Folleco, T. M. Khoshgoftaar, J. V. Hulse, and L. A. Bullard. **Software quality modeling: The impact of class noise on the random forest classifier**. In *Proc. 2008 IEEE Congress on Evolutionary Computation*, pages 3853–3859, 2008. doi:10.1109/CEC.2008.4631321.

**See Also**

[fra\\_bdir\\_ln](#), [irs\\_bdir\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- mind_bdir_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.5)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- mind_bdir_ln(formula = Species ~ ., data = iris2D, level = 0.5)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

minp\_uni\_In

*Minority-proportional uniform label noise*


---

### Description

Introduction of *Minority-proportional uniform label noise* into a classification dataset.

### Usage

```
## Default S3 method:
minp_uni_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
minp_uni_ln(formula, data, ...)
```

### Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

Given a dataset, assume the original class distribution of class  $i$  is  $p_i$  and the distribution of the minority class is  $p_m$ . Let `level` be the noise level, *Minority-proportional uniform label noise* introduces noise proportionally to different classes, where a sample with its label  $i$  has a probability  $(p_m/p_i) \cdot \text{level}$  to be corrupted as another random class. That is, the least common class is used as the baseline for noise introduction.

### Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.

model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References.

### References

X. Zhu and X. Wu. **Cost-guided class noise handling for effective cost-sensitive learning**. In *Proc. 4th IEEE International Conference on Data Mining*, pages 297–304, 2004. doi:10.1109/ICDM.2004.10108.

### See Also

[asy\\_uni\\_ln](#), [maj\\_udir\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- minp_uni_ln(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- minp_uni_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

mis\_pre\_ln

*Misclassification prediction label noise*

---

### Description

Introduction of *Misclassification prediction label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
mis_pre_ln(x, y, sortid = TRUE, ...)

## S3 method for class 'formula'
mis_pre_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Misclassification prediction label noise* creates a Multi-Layer Perceptron (MLP) model from the dataset and relabels each sample with the class predicted by the classifier.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

Q. Wang, B. Han, T. Liu, G. Niu, J. Yang, and C. Gong. **Tackling instance-dependent label noise via a universal probabilistic model**. In *Proc. 35th AAAI Conference on Artificial Intelligence*, pages 10183-10191, 2021. url:<https://ojs.aaai.org/index.php/AAAI/article/view/17221>.

**See Also**

[smam\\_bor\\_ln](#), [nlin\\_bor\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- mis_pre_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)])

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- mis_pre_ln(formula = Species ~ ., data = iris2D)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

mulc\_udir\_ln

---

*Multiple-class unidirectional label noise*


---

**Description**

Introduction of *Multiple-class unidirectional label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
mulc_udir_ln(x, y, level, goal, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
mulc_udir_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
goal	an integer vector with the indices of noisy classes for each class.
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).

... other options to pass to the function.  
 formula a formula with the output class and, at least, one input attribute.  
 data a data frame in which to interpret the variables in the formula.

### Details

*Multiple-class unidirectional label noise* introduction model randomly selects  $(level \cdot 100)\%$  of the samples of each class  $c$  with  $goal[c] \neq NA$ . Then, the labels  $c$  of these samples are replaced by the class indicated in  $goal[c]$ . The order of indices in  $goal$  is determined by order.

### Value

An object of class `ndmodel` with elements:

`xnoise` a data frame with the noisy input attributes.  
`ynoise` a factor vector with the noisy output class.  
`numnoise` an integer vector with the amount of noisy samples per class.  
`idnoise` an integer vector list with the indices of noisy samples.  
`numclean` an integer vector with the amount of clean samples per class.  
`idclean` an integer vector list with the indices of clean samples.  
`distr` an integer vector with the samples per class in the original data.  
`model` the full name of the noise introduction model used.  
`param` a list of the argument values.  
`call` the function call.

### Note

Noise model adapted from the papers in References.

### References

Q. Wang, B. Han, T. Liu, G. Niu, J. Yang, and C. Gong. **Tackling instance-dependent label noise via a universal probabilistic model**. In *Proc. 35th AAAI Conference on Artificial Intelligence*, pages 10183-10191, 2021. url:<https://ojs.aaai.org/index.php/AAAI/article/view/17221>.

### See Also

[minp\\_uni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- mulc_udir_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1,
```

```

goal = c(NA, 1, 2), order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- mulc_udir_ln(formula = Species ~ ., data = iris2D, level = 0.1,
                      goal = c(NA, 1, 2), order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

nei\_bor\_ln

*Neighborwise borderline label noise*


---

## Description

Introduction of *Neighborwise borderline label noise* into a classification dataset.

## Usage

```

## Default S3 method:
nei_bor_ln(x, y, level, k = 1, sortid = TRUE, ...)

## S3 method for class 'formula'
nei_bor_ln(formula, data, ...)

```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
k	an integer with the number of nearest neighbors to be used (default: 1).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

For each sample in the dataset, *Neighborwise borderline label noise* computes the ratio of two distances: the distance to its nearest neighbor from the same class and the distance to its nearest neighbor from another class. Then, these values are ordered in descending order and the first  $(level \cdot 100)\%$  of them are used to determine the noisy samples. For each noisy sample, the majority class among its k-nearest neighbors of a different class is chosen as the new label.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References, considering a mislabeling process using the neighborhood of noisy samples.

**References**

L. P. F. Garcia, J. Lehmann, A. C. P. L. F. de Carvalho, and A. C. Lorena. **New label noise injection methods for the evaluation of noise filters.** *Knowledge-Based Systems*, 163:693–704, 2019. [doi:10.1016/j.knosys.2018.09.031](https://doi.org/10.1016/j.knosys.2018.09.031).

**See Also**

[ulap\\_bor\\_ln](#), [lap\\_bor\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- nei_bor_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- nei_bor_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

nlin\_bor\_ln

*Non-linearwise borderline label noise*


---

**Description**

Introduction of *Non-linearwise borderline label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
nlin_bor_ln(x, y, level, k = 1, sortid = TRUE, ...)

## S3 method for class 'formula'
nlin_bor_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
k	an integer with the number of nearest neighbors to be used (default: 1).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Non-linearwise borderline label noise* uses an SVM to induce the decision border in the dataset. Then, for each sample, its distance to the decision border is computed. Finally, the distances obtained are ordered in ascending order and the first  $(level \cdot 100)\%$  of them are used to determine the noisy samples. For each noisy sample, the majority class among its  $k$ -nearest neighbors of a different class is chosen as the new label.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.

distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References, considering a mislabeling process using the neighborhood of noisy samples.

**References**

L. P. F. Garcia, J. Lehmann, A. C. P. L. F. de Carvalho, and A. C. Lorena. **New label noise injection methods for the evaluation of noise filters.** *Knowledge-Based Systems*, 163:693–704, 2019. doi:10.1016/j.knosys.2018.09.031.

**See Also**

[nei\\_bor\\_ln](#), [ulap\\_bor\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- nlin_bor_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- nlin_bor_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

oned\_uni\_ln

*One-dimensional uniform label noise*

---

**Description**

Introduction of *One-dimensional uniform label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
oned_uni_ln(
  x,
  y,
  level,
  att,
  lower,
  upper,
  order = levels(y),
  sortid = TRUE,
  ...
)

## S3 method for class 'formula'
oned_uni_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
att	an integer with the index of the attribute determining noisy samples.
lower	a vector with the lower bound to determine the noisy region of each class.
upper	a vector with the upper bound to determine the noisy region of each class.
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*One-dimensional uniform label noise* is based on the introduction of noise according to the values of the attribute `att`. Samples of class  $i$  with the attribute `att` falling between `lower[i]` and `upper[i]` have a probability `level` of being mislabeled. The labels of these samples are randomly replaced by other different ones within the set of class labels. The order of the class labels is determined by `order`.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.

numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References to multiclass data, considering a noise level to control the number of errors in the data

### References

N. Gornitz, A. Porbadnigk, A. Binder, C. Sannelli, M. L. Braun, K. Muller, and M. Kloft. **Learning and evaluation in presence of non-i.i.d. label noise**. In *Proc. 17th International Conference on Artificial Intelligence and Statistics*, volume 33 of PMLR, pages 293–302, 2014. url:<https://proceedings.mlr.press/v33/gornitz14.html>.

### See Also

[attn\\_uni\\_ln](#), [qua\\_uni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- oned_uni_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)],
                    level = 0.5, att = 1, lower = c(1.5, 2, 6), upper = c(2, 4, 7))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- oned_uni_ln(formula = Species ~ ., data = iris2D,
                    level = 0.5, att = 1, lower = c(1.5, 2, 6), upper = c(2, 4, 7))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

opes\_idnn\_ln

*Open-set ID/nearest-neighbor label noise***Description**

Introduction of *Open-set ID/nearest-neighbor label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
opes_idnn_ln(
  x,
  y,
  level,
  openset = c(1),
  order = levels(y),
  sortid = TRUE,
  ...
)

## S3 method for class 'formula'
opes_idnn_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double with the noise level in [0,1] to be introduced.
openset	an integer vector with the indices of classes in the open set (default: c(1)).
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Open-set ID/nearest-neighbor label noise* corrupts  $(level \cdot 100)\%$  of the samples with classes in openset. Then, the labels of these samples are replaced by the label of the nearest sample of a different in-distribution class. The order of the class labels for the indices in openset is determined by order.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

P. H. Seo, G. Kim, and B. Han. **Combinatorial inference against label noise**. In *Advances in Neural Information Processing Systems*, volume 32, pages 1171-1181, 2019. url:<https://proceedings.neurips.cc/paper/2019/hash/0cb929eae7a499e50248a3a78f7acfc7-Abstract.html>.

**See Also**

[opes\\_idu\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- opes_idnn_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                      level = 0.4, order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- opes_idnn_ln(formula = Species ~ ., data = iris2D,
                      level = 0.4, order = c("virginica", "setosa", "versicolor"))
```

```
# check the match of noisy indices
identical(outdef$idnoise, outfmr$idnoise)
```

---

opes_idu_ln	<i>Open-set ID/uniform label noise</i>
-------------	--

---

## Description

Introduction of *Open-set ID/uniform label noise* into a classification dataset.

## Usage

```
## Default S3 method:
opes_idu_ln(x, y, level, openset = c(1), order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
opes_idu_ln(formula, data, ...)
```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double with the noise level in [0,1] to be introduced.
openset	an integer vector with the indices of classes in the open set (default: c(1)).
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Open-set ID/uniform label noise* corrupts (level·100)% of the samples with classes in openset. For each sample selected, a label from in-distribution classes is randomly chosen. The order of the class labels for the indices in openset is determined by order.

## Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.

numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References.

### References

P. H. Seo, G. Kim, and B. Han. **Combinatorial inference against label noise**. In *Advances in Neural Information Processing Systems*, volume 32, pages 1171-1181, 2019. url:<https://proceedings.neurips.cc/paper/2019/hash/0cb929eae7a499e50248a3a78f7acfc7-Abstract.html>.

### See Also

[asy\\_spa\\_ln](#), [mind\\_bdir\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- opes_idu_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = 0.4, order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- opes_idu_ln(formula = Species ~ ., data = iris2D,
                    level = 0.4, order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

pai_bdir_ln	<i>Pairwise bidirectional label noise</i>
-------------	---

---

**Description**

Introduction of *Pairwise bidirectional label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
pai_bdir_ln(x, y, level, pairs, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
pai_bdir_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
pairs	a list of integer vectors with the indices of classes to corrupt.
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

For each vector ( $c1$ ,  $c2$ ) in *pairs*, *Pairwise bidirectional label noise* randomly selects  $(level \cdot 100)\%$  of the samples from class  $c1$  in the dataset and  $(level \cdot 100)\%$  of the samples from class  $c2$ . Then,  $c1$  samples are mislabeled as belonging to  $c2$  and  $c2$  samples are mislabeled as belonging to  $c1$ . The order of the class labels is determined by *order*.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.

distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References.

### References

S. Fefilyayev, M. Shreve, K. Kramer, L. O. Hall, D. B. Goldgof, R. Kasturi, K. Daly, A. Remsen, and H. Bunke. **Label-noise reduction with support vector machines**. In *Proc. 21st International Conference on Pattern Recognition*, pages 3504-3508, 2012. url:<https://ieeexplore.ieee.org/document/6460920/>.

### See Also

[print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# create new class with some samples
class <- as.character(iris2D$Species)
class[iris2D$Petal.Length > 6] <- "newclass"
iris2D$Species <- as.factor(class)

# usage of the default method
set.seed(9)
outdef <- pai_bdir_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = 0.1, pairs = list(c(1,2), c(3,4)),
                    order = c("virginica", "setosa", "newclass", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- pai_bdir_ln(formula = Species ~ ., data = iris2D,
                    level = 0.1, pairs = list(c(1,2), c(3,4)),
                    order = c("virginica", "setosa", "newclass", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

plot.ndmodel

*Plot function for class ndmodel*


---

### Description

Representation of the dataset contained in an object of class `ndmodel` after the application of a noise introduction model.

### Usage

```
## S3 method for class 'ndmodel'
plot(x, ..., noise = NA, xvar = 1, yvar = 2, pca = FALSE)
```

### Arguments

<code>x</code>	an object of class <code>ndmodel</code> .
<code>...</code>	other options to pass to the function.
<code>noise</code>	a logical indicating which samples to show. The valid options are: <ul style="list-style-type: none"> <li>• <code>TRUE</code>: to show only the noisy samples.</li> <li>• <code>FALSE</code>: to show only the clean samples.</li> <li>• <code>NA</code>: to show both the clean and noisy samples (default value).</li> </ul>
<code>xvar</code>	an integer with the index of the input attribute (if <code>pca = FALSE</code> ) or the principal component (if <code>pca = TRUE</code> ) to represent in the $x$ axis (default: 1).
<code>yvar</code>	an integer with the index of the input attribute (if <code>pca = FALSE</code> ) or the principal component (if <code>pca = TRUE</code> ) to represent in the $y$ axis (default: 2).
<code>pca</code>	a logical indicating if PCA must be used (default: <code>FALSE</code> ).

### Details

This function performs a two-dimensional representation using the `ggplot2` package of the dataset contained in the object `x` of class `ndmodel`. Each of the classes in the dataset (available in `x$ynoise`) is represented by a different color. There are two options to represent the input attributes of the samples on the  $x$  and  $y$  axes of the graph:

- If `pca = FALSE`, the values in the graph are taken from the current attribute values found in `x$ynoise`. In this case, `xvar` and `yvar` indicate the indices of the attributes to show in the  $x$  and  $y$  axes, respectively.
- If `pca = TRUE`, the values in the graph are taken after performing a PCA over `x$ynoise`. In this case, `xvar` and `yvar` indicate the index of the principal component according to the variance explained to show in the  $x$  and  $y$  axes, respectively.

Finally, the parameter `noise` is used to indicate which samples (noisy, clean or all) to show. Clean samples are represented by circles in the graph, while noisy samples are represented by crosses.

**Value**

An object of class `ggplot` and `gg` with the graph created using the `ggplot2` package.

**See Also**

[print.ndmodel](#), [summary.ndmodel](#), [sym\\_uni\\_ln](#), [sym\\_cuni\\_ln](#), [sym\\_uni\\_an](#)

**Examples**

```
# load the dataset
data(iris)

# apply the noise introduction model
set.seed(9)
output <- sym_uni_ln(x = iris[,-ncol(iris)], y = iris[,ncol(iris)], level = 0.1)

# plots for all the samples, the clean samples and the noisy samples using PCA
plot(output, pca = TRUE)
plot(output, noise = FALSE, pca = TRUE)
plot(output, noise = TRUE, pca = TRUE)

# plots using the Petal.Length and Petal.Width variables
plot(output, xvar = 3, yvar = 4)
plot(output, noise = FALSE, xvar = 3, yvar = 4)
plot(output, noise = TRUE, xvar = 3, yvar = 4)
```

---

pmd\_con\_ln

*PMD-based confidence label noise*

---

**Description**

Introduction of *PMD-based confidence label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
pmd_con_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
pmd_con_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).

...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

*PMD-based confidence label noise* approximates the probability of noise using the confidence prediction of a neural network. These predictions are used to estimate the mislabeling probability and the most possible noisy class label for each sample. Finally,  $(level \cdot 100)\%$  of the samples in the dataset are randomly selected to be mislabeled according to their values of probability computed.

### Value

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

### Note

Noise model adapted from the papers in References.

### References

Y. Zhang, S. Zheng, P. Wu, M. Goswami, and C. Chen. **Learning with feature-dependent label noise: A progressive approach**. In *Proc. 9th International Conference on Learning Representations*, pages 1-13, 2021. url:<https://openreview.net/forum?id=ZPa2SyGcbwh>.

### See Also

[clu\\_vot\\_ln](#), [sco\\_con\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
```

```

outdef <- pmd_con_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- pmd_con_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

```
print.ndmodel
```

```
Print function for class ndmodel
```

---

### **Description**

This method displays the basic information about the noise introduction process contained in an object of class `ndmodel`.

### **Usage**

```
## S3 method for class 'ndmodel'
print(x, ...)
```

### **Arguments**

<code>x</code>	an object of class <code>ndmodel</code> .
<code>...</code>	other options to pass to the function.

### **Details**

This function presents the basic information of the noise introduction process and the resulting noisy dataset contained in the object `x` of class `ndmodel`. The information offered is as follows:

- the name of the noise introduction model.
- the parameters associated with the noise model.
- the number of noisy and clean samples in the dataset.

### **Value**

This function does not return any value.

### **See Also**

[summary.ndmodel](#), [plot.ndmodel](#), [sym\\_uni\\_ln](#), [sym\\_cuni\\_ln](#), [sym\\_uni\\_an](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_uni_ln(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
print(outdef)
```

---

qua_uni_ln	<i>Quadrant-based uniform label noise</i>
------------	---

---

**Description**

Introduction of *Quadrant-based uniform label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
qua_uni_ln(x, y, level, att1 = 1, att2 = 2, sortid = TRUE, ...)

## S3 method for class 'formula'
qua_uni_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double vector with the noise levels in [0,1] in each quadrant.
att1	an integer with the index of the first attribute forming the quadrants (default: 1).
att2	an integer with the index of the second attribute forming the quadrants (default: 2).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

For each sample, the probability of flipping its label is based on which quadrant (with respect to the attributes `att1` and `att2`) the sample falls in. The probability of mislabeling for each quadrant is expressed with the argument `level`, whose length is equal to 4. Let  $m1$  and  $m2$  be the mean values of the domain of `att1` and `att2`, respectively. Each quadrant is defined as follows: values  $\leq m1$  and  $\leq m2$  (first quadrant); values  $\leq m1$  and  $> m2$  (second quadrant); values  $> m1$  and  $\leq m2$  (third quadrant); and values  $> m1$  and  $> m2$  (fourth quadrant). Finally, the labels of these samples are randomly replaced by other different ones within the set of class labels.

## Value

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

## Note

Noise model adapted from the papers in References.

## References

A. Ghosh, N. Manwani, and P. S. Sastry. **Making risk minimization tolerant to label noise.** *Neurocomputing*, 160:93-107, 2015. doi:10.1016/j.neucom.2014.09.081.

## See Also

[exps\\_cuni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- qua_uni_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)],
                    level = c(0.05, 0.15, 0.20, 0.4))
```

```

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- qua_uni_ln(formula = Species ~ ., data = iris2D,
                    level = c(0.05, 0.15, 0.20, 0.4))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

sco\_con\_ln

*Score-based confidence label noise*


---

## Description

Introduction of *Score-based confidence label noise* into a classification dataset.

## Usage

```

## Default S3 method:
sco_con_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sco_con_ln(formula, data, ...)

```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Score-based confidence label noise* follows the intuition that hard samples are more likely to be mislabeled. Given the confidence per class of each sample, if it is predicted with a different class with a high probability, it means that it is hard to clearly distinguish the sample from this class. The confidence information is used to compute a mislabeling score for each sample and its potential noisy label. Finally,  $(level \cdot 100)\%$  of the samples with the highest mislabeling scores are chosen as noisy.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

P. Chen, J. Ye, G. Chen, J. Zhao, and P. Heng. **Beyond class-conditional assumption: A primary attempt to combat instance-dependent label noise.** In *Proc. 35th AAAI Conference on Artificial Intelligence*, pages 11442-11450, 2021. url:<https://ojs.aaai.org/index.php/AAAI/article/view/17363>.

**See Also**

[mis\\_pre\\_ln](#), [smam\\_bor\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sco_con_ln(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sco_con_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

sigb\_uni\_ln

*Sigmoid-bounded uniform label noise***Description**

Introduction of *Sigmoid-bounded uniform label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
sigb_uni_ln(x, y, level, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
sigb_uni_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double vector with the noise levels in [0,1] to be introduced into each class.
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Sigmoid-bounded uniform label noise* generates bounded instance-dependent and label-dependent label noise at random using a weight for each sample in the dataset to compute its noise probability through a sigmoid function. Note that this noise model considers the maximum noise level per class given by level, so the current noise level in each class may be lower than that specified. The order of the class labels is determined by order.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.

distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References to multiclass data.

**References**

J. Cheng, T. Liu, K. Ramamohanarao, and D. Tao. **Learning with bounded instance and label-dependent label noise**. In *Proc. 37th International Conference on Machine Learning*, volume 119 of PMLR, pages 1789-1799, 2020. url:<http://proceedings.mlr.press/v119/cheng20c.html>.

**See Also**

[larm\\_uni\\_ln](#), [hubp\\_uni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sigb_uni_ln(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = c(0.1, 0.2, 0.3))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sigb_uni_ln(formula = Species ~ ., data = iris2D,
                    level = c(0.1, 0.2, 0.3))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

smam\_bor\_ln

*Small-margin borderline label noise*


---

**Description**

Introduction of *Small-margin borderline label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
smam_bor_ln(x, y, level, k = 1, sortid = TRUE, ...)

## S3 method for class 'formula'
smam_bor_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
k	an integer with the number of nearest neighbors to be used (default: 1).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Small-margin borderline label noise* uses an SVM to induce the decision border in the dataset. For each sample, its distance to the decision border is computed. Then, the samples are ordered according to their distance and  $(level \cdot 100)\%$  of the closest correctly classified samples to the decision boundary are selected to be mislabeled. For each noisy sample, the majority class among its k-nearest neighbors of a different class is chosen as the new label.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References to multiclass data, considering SVM with linear kernel as classifier and a mislabeling process using the neighborhood of noisy samples.

## References

E. Amid, M. K. Warmuth, and S. Srinivasan. **Two-temperature logistic regression based on the Tsallis divergence**. In *Proc. 22nd International Conference on Artificial Intelligence and Statistics*, volume 89 of PMLR, pages 2388-2396, 2019. url:<http://proceedings.mlr.press/v89/amid19a.html>.

## See Also

[nlin\\_bor\\_ln](#), [nei\\_bor\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- smam_bor_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- smam_bor_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

smu\_cuni\_ln

*Smudge-based completely-uniform label noise*

---

## Description

Introduction of *Smudge-based completely-uniform label noise* into a classification dataset.

## Usage

```
## Default S3 method:
smu_cuni_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
smu_cuni_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Smudge-based completely-uniform label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. Then, the labels of these samples are randomly replaced by others within the set of class labels. An additional attribute `smudge` is included in the dataset with value equal to 1 in mislabeled samples and equal to 0 in clean samples.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

S. Thulasidasan, T. Bhattacharya, J. A. Bilmes, G. Chennupati, and J. Mohd-Yusof. **Combating label noise in deep learning using abstention**. In *Proc. 36th International Conference on Machine Learning*, volume 97 of PMLR, pages 6234-6243, 2019. url:<http://proceedings.mlr.press/v97/thulasidasan19a.html>.

**See Also**

[oned\\_uni\\_ln](#), [attm\\_uni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- smu_cuni_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef, pca = TRUE)

# usage of the method for class formula
set.seed(9)
outfrm <- smu_cuni_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

summary.ndmodel

*Summary function for class ndmodel*


---

**Description**

This method displays a summary containing information about the noise introduction process contained in an object of class `ndmodel`.

**Usage**

```

## S3 method for class 'ndmodel'
summary(object, ..., showid = FALSE)

```

**Arguments**

<code>object</code>	an object of class <code>ndmodel</code> .
<code>...</code>	other options to pass to the function.
<code>showid</code>	a logical indicating if the indices of noisy samples must be displayed (default: <code>FALSE</code> ).

**Details**

This function presents a summary containing information of the noise introduction process and the resulting noisy dataset contained in the object `object` of class `ndmodel`. The information offered is as follows:

- the function call.
- the name of the noise introduction model.

- the parameters associated with the noise model.
- the number of noisy and clean samples in the dataset.
- the number of noisy samples per class/attribute.
- the number of clean samples per class/attribute.
- the indices of the noisy samples (if showid = TRUE).

### Value

A list with the elements of object, including the showid argument.

### See Also

[print.ndmodel](#), [plot.ndmodel](#), [sym\\_uni\\_ln](#), [sym\\_cuni\\_ln](#), [sym\\_uni\\_an](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_uni_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
```

---

symd\_gau\_an

*Symmetric/dependent Gaussian attribute noise*

---

### Description

Introduction of *Symmetric/dependent Gaussian attribute noise* into a classification dataset.

### Usage

```
## Default S3 method:
symd_gau_an(x, y, level, k = 0.2, sortid = TRUE, ...)

## S3 method for class 'formula'
symd_gau_an(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
k	a double in [0,1] with the scale used for the standard deviation (default: 0.2).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric/dependent Gaussian attribute noise* corrupts  $(level \cdot 100)\%$  of the samples in the dataset. Their attribute values are modified adding a random value that follows a Gaussian distribution of  $mean = 0$  and  $standard\ deviation = (max - min) \cdot k$ , being  $max$  and  $min$  the limits of the attribute domain. For nominal attributes, a random value is chosen.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per attribute.
idnoise	an integer vector list with the indices of noisy samples per attribute.
numclean	an integer vector with the amount of clean samples per attribute.
idclean	an integer vector list with the indices of clean samples per attribute.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

X. Huang, L. Shi, and J. A. K. Suykens. **Support vector machine classifier with pinball loss.** *IEEE Transactions on Pattern Analysis and Machine Intelligence*, 36(5):984-997, 2014. doi:10.1109/TPAMI.2013.178.

**See Also**

[sym\\_gau\\_an](#), [sym\\_int\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- symd_gau_an(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- symd_gau_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

symd\_gimg\_an

*Symmetric/dependent Gaussian-image attribute noise*


---

**Description**

Introduction of *Symmetric/dependent Gaussian-image attribute noise* into a classification dataset.

**Usage**

```

## Default S3 method:
symd_gimg_an(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
symd_gimg_an(formula, data, ...)

```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric/dependent Gaussian-image attribute noise* corrupts (level·100)% of the samples in the dataset. For each sample, a Gaussian distribution (with matching mean and variance to the original sample) is used to generate random attribute values for that sample.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per attribute.
<code>idnoise</code>	an integer vector list with the indices of noisy samples per attribute.
<code>numclean</code>	an integer vector with the amount of clean samples per attribute.
<code>idclean</code>	an integer vector list with the indices of clean samples per attribute.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

L. Huang, C. Zhang, and H. Zhang. **Self-adaptive training: Beyond empirical risk minimization**. In *Proceedings of the Advances in Neural Information Processing Systems*, 2020, Vol. 33, pp. 19365–19376. <https://proceedings.neurips.cc/paper/2020/file/e0ab531ec312161511493b002f9be2ee-Paper.pdf>

**See Also**

[unc\\_vgau\\_an](#), [symd\\_rpix\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- symd_gimg_an(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)
```

```
# usage of the method for class formula
set.seed(9)
outfrm <- symd_gimg_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

 symd\_rpix\_an

*Symmetric/dependent random-pixel attribute noise*


---

## Description

Introduction of *Symmetric/dependent random-pixel attribute noise* into a classification dataset.

## Usage

```
## Default S3 method:
symd_rpix_an(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
symd_rpix_an(formula, data, ...)
```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Symmetric/dependent random-pixel attribute noise* corrupts  $(level \cdot 100)\%$  of the samples in the dataset. For each sample, its attribute values are shuffled using independent random permutations.

## Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per attribute.

idnoise	an integer vector list with the indices of noisy samples per attribute.
numclean	an integer vector with the amount of clean samples per attribute.
idclean	an integer vector list with the indices of clean samples per attribute.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References.

### References

L. Huang, C. Zhang, and H. Zhang. **Self-adaptive training: Beyond empirical risk minimization**. In *Proceedings of the Advances in Neural Information Processing Systems*, 2020, Vol. 33, pp. 19365–19376. <https://proceedings.neurips.cc/paper/2020/file/e0ab531ec312161511493b002f9be2ee-Paper.pdf>

### See Also

[unc\\_fixw\\_an](#), [sym\\_end\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- symd_rpix_an(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- symd_rpix_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

symd_uni_an	<i>Symmetric/dependent uniform attribute noise</i>
-------------	--

---

**Description**

Introduction of *Symmetric/dependent uniform attribute noise* into a classification dataset.

**Usage**

```
## Default S3 method:
symd_uni_an(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
symd_uni_an(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric/dependent uniform attribute noise* corrupts  $(level \cdot 100)\%$  of the samples in the dataset. Their attribute values are replaced by random different ones between the minimum and maximum of the domain of each attribute following a uniform distribution (for numerical attributes) or choosing a random value (for nominal attributes).

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per attribute.
idnoise	an integer vector list with the indices of noisy samples per attribute.
numclean	an integer vector with the amount of clean samples per attribute.
idclean	an integer vector list with the indices of clean samples per attribute.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

A. Petety, S. Tripathi, and N. Hemachandra. **Attribute noise robust binary classification**. In *Proc. 34th AAAI Conference on Artificial Intelligence*, pages 13897-13898, 2020.

**See Also**

[sym\\_uni\\_an](#), [sym\\_cuni\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- symd_uni_an(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- symd_uni_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_adj\_ln

*Symmetric adjacent label noise*

---

**Description**

Introduction of *Symmetric adjacent label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
sym_adj_ln(x, y, level, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
sym_adj_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric adjacent label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. Then, the labels of these samples are replaced by a random adjacent class label according to order.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

J. R. Cano, J. Luengo, and S. Garcia. **Label noise filtering techniques to improve monotonic classification**. *Neurocomputing*, 353:83-95, 2019. doi:10.1016/j.neucom.2018.05.131.

**See Also**

[sym\\_dran\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_adj_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_adj_ln(formula = Species ~ ., data = iris2D,
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

sym\_cen\_ln

*Symmetric center-based label noise*


---

**Description**

Introduction of *Symmetric center-based label noise* into a classification dataset.

**Usage**

```

## Default S3 method:
sym_cen_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_cen_ln(formula, data, ...)

```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Symmetric center-based label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. The probability for choosing the noisy label is determined based on the distance between class centers. Thus, the mislabeling probability between classes increases as the distance between their centers decreases. This model is consistent with the intuition that samples in similar classes are more likely to be mislabeled. Besides, the model also allows mislabeling data in dissimilar classes with a relatively small probability, which corresponds to label noise caused by random errors.

## Value

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

## Note

Noise model adapted from the papers in References.

## References

X. Pu and C. Li. **Probabilistic information-theoretic discriminant analysis for industrial label-noise fault diagnosis**. *IEEE Transactions on Industrial Informatics*, 17(4):2664-2674, 2021. doi:10.1109/TII.2020.3001335.

## See Also

[glev\\_uni\\_ln](#), [sym\\_hienc\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_cen_ln(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)
```

```
# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_cen_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym_con_ln	<i>Symmetric confusion label noise</i>
------------	--

---

## Description

Introduction of *Symmetric confusion label noise* into a classification dataset.

## Usage

```
## Default S3 method:
sym_con_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_con_ln(formula, data, ...)
```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Symmetric confusion label noise* considers that the mislabeling probability for each class is level. It obtains the confusion matrix from the dataset, which is row-normalized to estimate the transition matrix and get the probability of selecting each class when noise occurs.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References, considering C5.0 as classifier.

**References**

D. Ortego, E. Arazo, P. Albert, N. E. O'Connor, and K. McGuinness. **Towards robust learning with different label noise distributions**. In *Proc. 25th International Conference on Pattern Recognition*, pages 7020-7027, 2020. doi:10.1109/ICPR48806.2021.9412747.

**See Also**

[sym\\_cen\\_ln](#), [glev\\_uni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_con_ln(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_con_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

sym\_cuni\_an

*Symmetric completely-uniform attribute noise***Description**

Introduction of *Symmetric completely-uniform attribute noise* into a classification dataset.

**Usage**

```
## Default S3 method:
sym_cuni_an(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_cuni_an(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric completely-uniform attribute noise* corrupts  $(level \cdot 100)\%$  of the values of each attribute in the dataset. In order to corrupt an attribute  $A$ ,  $(level \cdot 100)\%$  of the samples in the dataset are randomly chosen. Then, their values for  $A$  are replaced by random ones from the domain of the attribute. Note that the original attribute value of a sample can be chosen as noisy and the actual percentage of noise in the dataset can be lower than the theoretical noise level.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per attribute.
idnoise	an integer vector list with the indices of noisy samples per attribute.
numclean	an integer vector with the amount of clean samples per attribute.
idclean	an integer vector list with the indices of clean samples per attribute.
distr	an integer vector with the samples per class in the original data.

model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References, only considering attribute noise introduction.

### References

C. Teng. **Polishing blemishes: Issues in data correction.** *IEEE Intelligent Systems*, 19(2):34-39, 2004. doi:10.1109/MIS.2004.1274909.

### See Also

[sym\\_uni\\_an](#), [sym\\_cuni\\_cn](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_cuni_an(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_cuni_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_cuni\_cn

*Symmetric completely-uniform combined noise*

---

### Description

Introduction of *Symmetric completely-uniform combined noise* into a classification dataset.

**Usage**

```
## Default S3 method:
sym_cuni_cn(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_cuni_cn(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric completely-uniform combined noise* corrupts  $(level \cdot 100)\%$  of the values of each attribute in the dataset. In order to corrupt an attribute  $A$ ,  $(level \cdot 100)\%$  of the samples in the dataset are randomly chosen. Then, their values for  $A$  are replaced by random ones from the domain of the attribute.

Additionally, this noise model also selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. The labels of these samples are randomly replaced by other ones within the set of class labels.

Note that, for both attributes and class labels, the original value of a sample can be chosen as noisy and the actual percentage of noise in the dataset can be lower than the theoretical noise level.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per variable.
idnoise	an integer vector list with the indices of noisy samples per variable.
numclean	an integer vector with the amount of clean samples per variable.
idclean	an integer vector list with the indices of clean samples per variable.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

C. Teng. **Polishing blemishes: Issues in data correction.** *IEEE Intelligent Systems*, 19(2):34-39, 2004. doi:10.1109/MIS.2004.1274909.

**See Also**

[uncs\\_guni\\_cn](#), [sym\\_cuni\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_cuni_cn(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_cuni_cn(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_cuni\_ln

*Symmetric completely-uniform label noise*

---

**Description**

Introduction of *Symmetric completely-uniform label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
sym_cuni_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_cuni_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric completely-uniform label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. Then, the labels of these samples are randomly replaced by others within the set of class labels. Note that this model can choose the original label of a sample as noisy.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

A. Ghosh and A. S. Lan. **Contrastive learning improves model robustness under label noise**. In *Proc. 2021 IEEE Conference on Computer Vision and Pattern Recognition Workshops*, pages 2703-2708, 2021. doi:[10.1109/CVPRW53098.2021.00304](https://doi.org/10.1109/CVPRW53098.2021.00304).

**See Also**

[sym\\_uni\\_ln](#), [sym\\_cuni\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_cuni_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_cuni_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

sym\_ddef\_ln

*Symmetric double-default label noise*


---

**Description**

Introduction of *Symmetric double-default label noise* into a classification dataset.

**Usage**

```

## Default S3 method:
sym_ddef_ln(
  x,
  y,
  level,
  def1 = 1,
  def2 = 2,
  order = levels(y),
  sortid = TRUE,
  ...
)

## S3 method for class 'formula'
sym_ddef_ln(formula, data, ...)

```

**Arguments**

x                    a data frame of input attributes.  
y                    a factor vector with the output class of each sample.

level	a double in [0,1] with the noise level to be introduced.
def1	an integer with the index of the first default class (default: 1).
def2	an integer with the index of the second default class (default: 2).
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

*Symmetric double-default label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. Then, the labels of these samples are replaced by one of two fixed labels (def1 or def2) within the set of class labels. The indices def1 and def2 are taken according to the order given by order.

### Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References.

### References

B. Han, J. Yao, G. Niu, M. Zhou, I. W. Tsang, Y. Zhang, and M. Sugiyama. **Masking: A new perspective of noisy supervision.** In *Advances in Neural Information Processing Systems*, volume 31, pages 5841-5851, 2018. url:<https://proceedings.neurips.cc/paper/2018/hash/aee92f16efd522b9326c25cc3237ac15-Abstract.html>.

### See Also

[sym\\_exc\\_ln](#), [sym\\_cuni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_ddef_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_ddef_ln(formula = Species ~ ., data = iris2D,
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

sym\_def\_ln

*Symmetric default label noise*


---

**Description**

Introduction of *Symmetric default label noise* into a classification dataset.

**Usage**

```

## Default S3 method:
sym_def_ln(x, y, level, def = 1, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
sym_def_ln(formula, data, ...)

```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
def	an integer with the index of the default class (default: 1).
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric default label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. Then, the labels of these samples are replaced by a fixed label (def) within the set of class labels. The index def is taken according to the order given by order.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

M. Ren, W. Zeng, B. Yang, and R. Urtasun. **Learning to reweight examples for robust deep learning**. In *Proc. 35th International Conference on Machine Learning*, volume 80 of PMLR, pages 4331-4340, 2018. url:<http://proceedings.mlr.press/v80/ren18a.html>.

**See Also**

[sym\\_ddef\\_ln](#), [sym\\_exc\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_def_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)
```

```
# usage of the method for class formula
set.seed(9)
outfrm <- sym_def_ln(formula = Species ~ ., data = iris2D,
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_dia\_ln

*Symmetric diametrical label noise*


---

## Description

Introduction of *Symmetric diametrical label noise* into a classification dataset.

## Usage

```
## Default S3 method:
sym_dia_ln(x, y, level, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
sym_dia_ln(formula, data, ...)
```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Symmetric diametrical label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. In this model, diametrical (opposite) classes are more likely to have their labels mixed. The probability of mislabel a sample of class  $i$  as belonging to class  $j$  is computed as  $d_{ij}/S$ , where  $d_{ij} = \text{abs}(i-j)$  and  $S$  is the sum of distances to class  $i$ . The order of the classes is determined by order.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

R. C. Prati, J. Luengo, and F. Herrera. **Emerging topics and challenges of learning from noisy data in nonstandard classification: a survey beyond binary class noise.** *Knowledge and Information Systems*, 60(1):63–97, 2019. doi:10.1007/s1011501812444.

**See Also**

[sym\\_pes\\_ln](#), [sym\\_opt\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_dia_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_dia_ln(formula = Species ~ ., data = iris2D,
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
```

```
identical(outdef$idnoise, outfrm$idnoise)
```

---

```
sym_dran_ln          Symmetric double-random label noise
```

---

## Description

Introduction of *Symmetric double-random label noise* into a classification dataset.

## Usage

```
## Default S3 method:
sym_dran_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_dran_ln(formula, data, ...)
```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Symmetric double-random label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. Then, each of the original class labels is flipped to one between two other random labels.

## Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.

distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

A. Ghosh and A. S. Lan. **Do we really need gold samples for sample weighting under label noise?** In *Proc. 2021 IEEE Winter Conference on Applications of Computer Vision*, pages 3921-3930, 2021. doi:[10.1109/WACV48630.2021.00397](https://doi.org/10.1109/WACV48630.2021.00397).

**See Also**

[sym\\_hie\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_dran_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_dran_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_end\_an

*Symmetric end-directed attribute noise*


---

**Description**

Introduction of *Symmetric end-directed attribute noise* into a classification dataset.

**Usage**

```
## Default S3 method:
sym_end_an(x, y, level, scale = 0.2, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_end_an(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
scale	a double in (0,1) with the scale to be used (default: 0.2).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

For each attribute  $A$ , *Symmetric end-directed attribute noise* computes a value  $k = \text{scale} \cdot \max(A)$ . Then, it chooses  $(\text{level} \cdot 100)\%$  of the values of that attribute. For each value, it applies the following procedure:

- If the value is less than the median of the attribute, the value transforms into adding  $k$  to the maximum of the attribute  $A$ .
- If the value is greater than the median of the attribute, the value transforms into subtracting  $k$  from the minimum of the attribute  $A$ .
- If the value matches the median, one of the two previous alternatives is chosen.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per attribute.
idnoise	an integer vector list with the indices of noisy samples per attribute.
numclean	an integer vector with the amount of clean samples per attribute.
idclean	an integer vector list with the indices of clean samples per attribute.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

T. M. Khoshgoftaar and J. V. Hulse. **Empirical case studies in attribute noise detection.** *IEEE Transactions on Systems, Man and Cybernetics Part C: Applications and Reviews*, 39(4):379-388, 2009. doi:[10.1109/TSMCC.2009.2013815](https://doi.org/10.1109/TSMCC.2009.2013815).

**See Also**

[sym\\_sgau\\_an](#), [symd\\_gau\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_end_an(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_end_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_exc\_ln

*Symmetric exchange label noise*

---

**Description**

Introduction of *Symmetric exchange label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
sym_exc_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_exc_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric exchange label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. These samples are divided into two groups: *A* and *B*. Then, each sample of group *A* is labeled with the label of a sample of group *B* and vice versa.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

J. Schneider, J. P. Handali, and J. vom Brocke. **Increasing trust in (big) data analytics**. In *Proc. 2018 Advanced Information Systems Engineering Workshops*, volume 316 of LNBIP, pages 70-84, 2018. doi:10.1007/9783319928982\_6.

**See Also**

[sym\\_cuni\\_ln](#), [sym\\_cuni\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_exc_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_exc_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

sym\_gau\_an

*Symmetric Gaussian attribute noise***Description**

Introduction of *Symmetric Gaussian attribute noise* into a classification dataset.

**Usage**

```

## Default S3 method:
sym_gau_an(x, y, level, k = 0.2, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_gau_an(formula, data, ...)

```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
k	a double in [0,1] with the scale used for the standard deviation (default: 0.2).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Symmetric Gaussian attribute noise* corrupts  $(level \cdot 100)\%$  of the values of each attribute in the dataset. In order to corrupt an attribute  $A$ ,  $(level \cdot 100)\%$  of the samples in the dataset are chosen. Then, their values for  $A$  are corrupted adding a random value that follows a Gaussian distribution of  $mean = 0$  and  $standard\ deviation = (max - min) \cdot k$ , being  $max$  and  $min$  the limits of the attribute domain. For nominal attributes, a random value is chosen.

## Value

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per attribute.
<code>idnoise</code>	an integer vector list with the indices of noisy samples per attribute.
<code>numclean</code>	an integer vector with the amount of clean samples per attribute.
<code>idclean</code>	an integer vector list with the indices of clean samples per attribute.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

## Note

Noise model adapted from the papers in References.

## References

J. A. Sáez, M. Galar, J. Luengo, and F. Herrera. **Analyzing the presence of noise in multi-class problems: alleviating its influence with the one-vs-one decomposition.** *Knowledge and Information Systems*, 38(1):179-206, 2014. doi:[10.1007/s1011501205701](https://doi.org/10.1007/s1011501205701).

## See Also

[sym\\_int\\_an](#), [symd\\_uni\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_gau_an(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
```

```

plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_gau_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

sym\_hienc\_In

*Symmetric hierarchical/next-class label noise*


---

### Description

Introduction of *Symmetric hierarchical/next-class label noise* into a classification dataset.

### Usage

```

## Default S3 method:
sym_hienc_In(x, y, level, group, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
sym_hienc_In(formula, data, ...)

```

### Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
group	a list of integer vectors with the indices of classes in each superclass.
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

*Symmetric hierarchical/next-class label noise* randomly selects (level·100)% of the samples in the dataset with independence of their class. Then, the labels of these samples are replaced by the next class within the set of class labels related to them (given by the argument group). The indices in group are taken according to the order given by order. Note that if a class does not belong to any superclass, it may be mislabeled as any other class.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

T. Kaneko, Y. Ushiku, and T. Harada. **Label-noise robust generative adversarial networks**. In *Proc. 2019 IEEE Conference on Computer Vision and Pattern Recognition*, pages 2462-2471, 2019. [doi:10.1109/CVPR.2019.00257](https://doi.org/10.1109/CVPR.2019.00257).

**See Also**

[sym\\_nexc\\_ln](#), [sym\\_dia\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_hienc_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1,
                      group = list(c(1,2)), order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_hienc_ln(formula = Species ~ ., data = iris2D, level = 0.1,
                      group = list(c(1,2)), order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
```

```
identical(outdef$idnoise, outfrm$idnoise)
```

---

```
sym_hie_ln
```

```
Symmetric hierarchical label noise
```

---

## Description

Introduction of *Symmetric hierarchical label noise* into a classification dataset.

## Usage

```
## Default S3 method:
sym_hie_ln(x, y, level, group, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
sym_hie_ln(formula, data, ...)
```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
group	a list of integer vectors with the indices of classes in each superclass.
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Symmetric hierarchical label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. Then, the labels of these samples are randomly replaced by other ones within the set of class labels related to them (given by the argument `group`). The indices in `group` are taken according to the order given by `order`. Note that if a class does not belong to any superclass, it may be mislabeled as any other class.

## Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.

idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References.

### References

D. Hendrycks, M. Mazeika, D. Wilson, and K. Gimpel. **Using trusted data to train deep networks on labels corrupted by severe noise**. In *Advances in Neural Information Processing Systems*, volume 31, pages 10477-10486, 2018. url:<https://proceedings.neurips.cc/paper/2018/hash/ad554d8c3b06d6b97ee76a2448bd7913-Abstract.html>.

### See Also

[sym\\_uni\\_ln](#), [sym\\_def\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method: a superclass with labels of indices 1 and 2
set.seed(9)
outdef <- sym_hie_ln(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1,
                    group = list(c(1,2)), order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_hie_ln(formula = Species ~ ., data = iris2D, level = 0.1,
                    group = list(c(1,2)), order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

 sym\_int\_an

*Symmetric interval-based attribute noise*


---

### Description

Introduction of *Symmetric interval-based attribute noise* into a classification dataset.

### Usage

```
## Default S3 method:
sym_int_an(x, y, level, nbins = 10, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_int_an(formula, data, ...)
```

### Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
nbins	an integer with the number of bins to create (default: 10).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

*Symmetric interval-based attribute noise* corrupts  $(level \cdot 100)\%$  of the values of each attribute in the dataset. In order to corrupt an attribute  $A$ ,  $(level \cdot 100)\%$  of the samples in the dataset are selected. To corrupt numeric attributes, the attribute is split into `nbins` equal-frequency intervals, one of its closest intervals is chosen and a random value within the interval is picked out as noisy. For nominal attributes, a random value within the domain is chosen.

### Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per attribute.
idnoise	an integer vector list with the indices of noisy samples per attribute.
numclean	an integer vector with the amount of clean samples per attribute.
idclean	an integer vector list with the indices of clean samples per attribute.

distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References.

### References

M. V. Mannino, Y. Yang, and Y. Ryu. **Classification algorithm sensitivity to training data with non representative attribute noise**. *Decision Support Systems*, 46(3):743-751, 2009. doi:[10.1016/j.dss.2008.11.021](https://doi.org/10.1016/j.dss.2008.11.021).

### See Also

[symd\\_uni\\_an](#), [sym\\_uni\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_int_an(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_int_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_natd\_ln

*Symmetric natural-distribution label noise*

---

### Description

Introduction of *Symmetric natural-distribution label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
sym_natd_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_natd_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric natural-distribution label noise* randomly selects (level·100)% of the samples in the dataset with independence of their class. Then, the labels of these samples are randomly replaced by other different ones within the set of class labels. When noise for a certain class occurs, another class with a probability proportional to the natural class distribution replaces it.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

## References

R. C. Prati, J. Luengo, and F. Herrera. **Emerging topics and challenges of learning from noisy data in nonstandard classification: a survey beyond binary class noise.** *Knowledge and Information Systems*, 60(1):63–97, 2019. doi:10.1007/s1011501812444.

## See Also

[sym\\_nuni\\_ln](#), [sym\\_adj\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_natd_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_natd_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_nean\_ln

*Symmetric nearest-neighbor label noise*

---

## Description

Introduction of *Symmetric nearest-neighbor label noise* into a classification dataset.

## Usage

```
## Default S3 method:
sym_nean_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_nean_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric nearest-neighbor label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. Then, the labels of these samples are replaced by the label of the nearest sample of a different class.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

P. H. Seo, G. Kim, and B. Han. **Combinatorial inference against label noise**. In *Advances in Neural Information Processing Systems*, volume 32, pages 1171-1181, 2019. url:<https://proceedings.neurips.cc/paper/2019/hash/0cb929eae7a499e50248a3a78f7acfc7-Abstract.html>.

**See Also**

[sym\\_con\\_ln](#), [sym\\_cen\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_nexc_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_nexc_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_nexc\_ln

*Symmetric next-class label noise*


---

**Description**

Introduction of *Symmetric next-class label noise* into a classification dataset.

**Usage**

```
## Default S3 method:
sym_nexc_ln(x, y, level, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
sym_nexc_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

The *Symmetric next-class label noise* introduction model randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. Then, the labels of these samples are replaced by the next class label according to order.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References

**References**

S. Gehlot, A. Gupta, and R. Gupta. **A CNN-based unified framework utilizing projection loss in unison with label noise handling for multiple Myeloma cancer diagnosis.** *Medical Image Analysis*, 72:102099, 2021. doi:10.1016/j.media.2021.102099.

**See Also**

[sym\\_dia\\_ln](#), [sym\\_pes\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_nexc_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)
```

```
# usage of the method for class formula
set.seed(9)
outfrm <- sym_nexc_ln(formula = Species ~ ., data = iris2D,
                     level = 0.1, order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_nuni\_ln

*Symmetric non-uniform label noise*


---

## Description

Introduction of *Symmetric non-uniform label noise* into a classification dataset.

## Usage

```
## Default S3 method:
sym_nuni_ln(x, y, level, tramat, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_nuni_ln(formula, data, ...)
```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
tramat	a double matrix with the values of the transition matrix.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Symmetric non-uniform label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. Then, the labels of these samples are randomly replaced by other different ones according to the probabilities given in the transition matrix `tramat`. For details about the structure of the transition matrix, see Kang *et al.* (2021).

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

J. Kang, R. Fernandez-Beltran, P. Duan, X. Kang, and A. J. Plaza. **Robust normalized softmax loss for deep metric learning-based characterization of remote sensing images with label noise.** *IEEE Transactions on Geoscience and Remote Sensing*, 59(10):8798-8811, 2021. doi:10.1109/TGRS.2020.3042607.

**See Also**

[sym\\_adj\\_ln](#), [sym\\_dran\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
trammat <- matrix(data = c(0.9, 0.03, 0.07, 0.03, 0.9, 0.07, 0.03, 0.07, 0.9),
                  nrow = 3, ncol = 3, byrow = TRUE)
outdef <- sym_nuni_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)],
                    level = 0.1, trammat = trammat)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_nuni_ln(formula = Species ~ ., data = iris2D, level = 0.1, trammat = trammat)
```

```
# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym_opt_ln	<i>Symmetric optimistic label noise</i>
------------	---

---

## Description

Introduction of *Symmetric optimistic label noise* into a classification dataset.

## Usage

```
## Default S3 method:
sym_opt_ln(x, y, level, levelH = 0.9, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
sym_opt_ln(formula, data, ...)
```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
levelH	a double in (0.5, 1] with the noise level for higher classes (default: 0.9).
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Symmetric optimistic label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. In the optimistic case, the probability of a class  $i$  of being mislabeled as class  $j$  is higher for  $j > i$  in comparison to  $j < i$ . Thus, when noise for a certain class occurs, it is assigned to a random higher class with probability  $levelH$  and to a random lower class with probability  $1 - levelH$ . The order of the classes is determined by `order`.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

R. C. Prati, J. Luengo, and F. Herrera. **Emerging topics and challenges of learning from noisy data in nonstandard classification: a survey beyond binary class noise.** *Knowledge and Information Systems*, 60(1):63–97, 2019. doi:10.1007/s1011501812444.

**See Also**

[sym\\_usim\\_ln](#), [sym\\_natd\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_opt_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_opt_ln(formula = Species ~ ., data = iris2D,
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
```

```
identical(outdef$idnoise, outfrm$idnoise)
```

---

```
sym_pes_ln          Symmetric pessimistic label noise
```

---

## Description

Introduction of *Symmetric pessimistic label noise* into a classification dataset.

## Usage

```
## Default S3 method:
sym_pes_ln(x, y, level, levelL = 0.9, order = levels(y), sortid = TRUE, ...)

## S3 method for class 'formula'
sym_pes_ln(formula, data, ...)
```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
levelL	a double in (0.5, 1] with the noise level for lower classes (default: 0.9).
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Symmetric pessimistic label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. In the pessimistic case, the probability of a class  $i$  of being mislabeled as class  $j$  is higher for  $j < i$  in comparison to  $j > i$ . Thus, when noise for a certain class occurs, it is assigned to a random lower class with probability  $levelL$  and to a random higher class with probability  $1 - levelL$ . The order of the classes is determined by `order`.

## Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.

idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References.

### References

R. C. Prati, J. Luengo, and F. Herrera. **Emerging topics and challenges of learning from noisy data in nonstandard classification: a survey beyond binary class noise.** *Knowledge and Information Systems*, 60(1):63–97, 2019. doi:[10.1007/s1011501812444](https://doi.org/10.1007/s1011501812444).

### See Also

[sym\\_opt\\_ln](#), [sym\\_usim\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

### Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_pes_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)],
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_pes_ln(formula = Species ~ ., data = iris2D,
                    level = 0.1, order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_sgau\_an

*Symmetric scaled-Gaussian attribute noise*


---

### Description

Introduction of *Symmetric scaled-Gaussian attribute noise* into a classification dataset.

### Usage

```
## Default S3 method:
sym_sgau_an(x, y, level, k = 0.2, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_sgau_an(formula, data, ...)
```

### Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
k	a double in [0,1] with the scale used for the standard deviation (default: 0.2).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

*Symmetric scaled-Gaussian attribute noise* corrupts  $(level \cdot 100)\%$  of the values of each attribute in the dataset. In order to corrupt an attribute  $A$ ,  $(level \cdot 100)\%$  of the samples in the dataset are chosen. Then, their values for  $A$  are modified adding a random value that follows a Gaussian distribution of  $mean = 0$  and  $standard\ deviation = (max - min) \cdot k \cdot level$ , being  $max$  and  $min$  the limits of the attribute domain. For nominal attributes, a random value is chosen.

### Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per attribute.
idnoise	an integer vector list with the indices of noisy samples per attribute.
numclean	an integer vector with the amount of clean samples per attribute.
idclean	an integer vector list with the indices of clean samples per attribute.

distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

M. Kozierski, B. Krawczyk, and M. Wozniak. **Radial-based oversampling for noisy imbalanced data classification.** *Neurocomputing*, 343:19–33, 2019. doi:10.1016/j.neucom.2018.04.089.

**See Also**

[sym\\_sgau\\_an](#), [sym\\_gau\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_sgau_an(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_sgau_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_uni\_an

*Symmetric uniform attribute noise*


---

**Description**

Introduction of *Symmetric uniform attribute noise* into a classification dataset.

**Usage**

```
## Default S3 method:
sym_uni_an(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_uni_an(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric uniform attribute noise* corrupts  $(level \cdot 100)\%$  of the values of each attribute in the dataset. In order to corrupt an attribute  $A$ ,  $(level \cdot 100)\%$  of the samples in the dataset are randomly chosen. Then, their values for  $A$  are replaced by random different ones from the domain of the attribute.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per attribute.
idnoise	an integer vector list with the indices of noisy samples per attribute.
numclean	an integer vector with the amount of clean samples per attribute.
idclean	an integer vector list with the indices of clean samples per attribute.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

## References

J. A. Sáez, M. Galar, J. Luengo, and F. Herrera. **Tackling the problem of classification with noisy data using Multiple Classifier Systems: Analysis of the performance and robustness.** *Information Sciences*, 247:1-20, 2013. doi:10.1016/j.ins.2013.06.002.

## See Also

[sym\\_cuni\\_an](#), [sym\\_cuni\\_cn](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_uni_an(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_uni_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

sym\_uni\_ln

*Symmetric uniform label noise*

---

## Description

Introduction of *Symmetric uniform label noise* into a classification dataset.

## Usage

```
## Default S3 method:
sym_uni_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_uni_ln(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Symmetric uniform label noise* randomly selects  $(\text{level} \cdot 100)\%$  of the samples in the dataset with independence of their class. Then, the labels of these samples are randomly replaced by other different ones within the set of class labels.

**Value**

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

**Note**

Noise model adapted from the papers in References.

**References**

Y. Wei, C. Gong, S. Chen, T. Liu, J. Yang, and D. Tao. **Harnessing side information for classification under label noise**. *IEEE Transactions on Neural Networks and Learning Systems*, 31(9):3178–3192, 2020. doi:10.1109/TNNLS.2019.2938782.

**See Also**

[sym\\_def\\_ln](#), [sym\\_ddef\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_uni_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_uni_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

sym\_usim\_ln

*Symmetric unit-simplex label noise*


---

**Description**

Introduction of *Symmetric unit-simplex label noise* into a classification dataset.

**Usage**

```

## Default S3 method:
sym_usim_ln(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
sym_usim_ln(formula, data, ...)

```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Symmetric unit-simplex label noise* randomly selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. Then, the labels of these samples are randomly replaced by other different ones within the set of class labels. The probability for each noisy class is drawn uniformly and independently from the  $M-1$ -dimensional unit simplex (with  $M$  the number of classes).

## Value

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

## Note

Noise model adapted from the papers in References.

## References

I. Jindal, D. Pressel, B. Lester, and M. S. Nokleby. **An effective label noise model for DNN text classification**. In *Proc. 2019 Conference of the North American Chapter of the Association for Computational Linguistics: Human Language Technologies*, pages 3246-3256, 2019. [doi:10.18653/v1/n191328](https://doi.org/10.18653/v1/n191328).

## See Also

[sym\\_natd\\_ln](#), [sym\\_nuni\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- sym_usim_ln(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
```

```

plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- sym_usim_ln(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

ugau\_bor\_ln

*Uneven-Gaussian borderline label noise*


---

## Description

Introduction of *Uneven-Gaussian borderline label noise* into a classification dataset.

## Usage

```

## Default S3 method:
ugau_bor_ln(
  x,
  y,
  level,
  mean = 0,
  sd = 1,
  k = 1,
  order = levels(y),
  sortid = TRUE,
  ...
)

## S3 method for class 'formula'
ugau_bor_ln(formula, data, ...)

```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double vector with the noise levels in [0,1] to be introduced into each class.
mean	a double with the mean for the Gaussian distribution (default: 0).
sd	a double with the standard deviation for the Gaussian distribution (default: 1).
k	an integer with the number of nearest neighbors to be used (default: 1).
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

## Details

*Uneven-Gaussian borderline label noise* uses an SVM to induce the decision border in the dataset. For each sample, its distance to the decision border is computed. Then, a Gaussian distribution with parameters (mean, sd) is used to compute the value for the probability density function associated to each distance. For each class  $c[i]$ , it randomly selects  $(level[i] \cdot 100)\%$  of the samples in the dataset based on their values of the probability density function -the order of the class labels is determined by order. For each noisy sample, the majority class among its  $k$ -nearest neighbors of a different class is chosen as the new label.

## Value

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per class.
<code>idnoise</code>	an integer vector list with the indices of noisy samples.
<code>numclean</code>	an integer vector with the amount of clean samples per class.
<code>idclean</code>	an integer vector list with the indices of clean samples.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

## Note

Noise model adapted from the papers in References to multiclass data, considering SVM with linear kernel as classifier, a mislabeling process using the neighborhood of noisy samples and a noise level to control the number of errors in the data.

## References

J. Du and Z. Cai. **Modelling class noise with symmetric and asymmetric distributions**. In *Proc. 29th AAAI Conference on Artificial Intelligence*, pages 2589-2595, 2015. url:<https://dl.acm.org/doi/10.5555/2886521.2886681>.

## See Also

[gaum\\_bor\\_ln](#), [gau\\_bor\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

## Examples

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
```

```

outdef <- ugau_bor_ln(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = c(0.1, 0.2, 0.3), order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- ugau_bor_ln(formula = Species ~ ., data = iris2D,
                    level = c(0.1, 0.2, 0.3), order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

---

ulap\_bor\_ln

*Uneven-Laplace borderline noise*


---

## Description

Introduction of *Uneven-Laplace borderline noise* into a classification dataset.

## Usage

```

## Default S3 method:
ulap_bor_ln(
  x,
  y,
  level,
  mu = 0,
  b = 1,
  k = 1,
  order = levels(y),
  sortid = TRUE,
  ...
)

## S3 method for class 'formula'
ulap_bor_ln(formula, data, ...)

```

## Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double vector with the noise levels in [0,1] to be introduced into each class.
mu	a double with the location for the Laplace distribution (default: 0).

b	a double with the scale for the Laplace distribution (default: 1).
k	an integer with the number of nearest neighbors to be used (default: 1).
order	a character vector indicating the order of the classes (default: levels(y)).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

*Uneven-Laplace borderline noise* uses an SVM to induce the decision border in the dataset. For each sample, its distance to the decision border is computed. Then, a Laplace distribution with parameters ( $\mu$ ,  $b$ ) is used to compute the value for the probability density function associated to each distance. For each class  $c[i]$ , it randomly selects  $(level[i] \cdot 100)\%$  of the samples in the dataset based on their values of the probability density function -the order of the class labels is determined by order. For each noisy sample, the majority class among its  $k$ -nearest neighbors of a different class is chosen as the new label.

### Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per class.
idnoise	an integer vector list with the indices of noisy samples.
numclean	an integer vector with the amount of clean samples per class.
idclean	an integer vector list with the indices of clean samples.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References to multiclass data, considering SVM with linear kernel as classifier, a mislabeling process using the neighborhood of noisy samples and a noise level to control the number of errors in the data.

### References

J. Du and Z. Cai. **Modelling class noise with symmetric and asymmetric distributions**. In *Proc. 29th AAAI Conference on Artificial Intelligence*, pages 2589-2595, 2015. url:<https://dl.acm.org/doi/10.5555/2886521.2886681>.

**See Also**

[lap\\_bor\\_ln](#), [ugau\\_bor\\_ln](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- ulap_bor_ln(x = iris2D[,ncol(iris2D)], y = iris2D[,ncol(iris2D)],
                    level = c(0.1, 0.2, 0.3), order = c("virginica", "setosa", "versicolor"))

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- ulap_bor_ln(formula = Species ~ ., data = iris2D,
                    level = c(0.1, 0.2, 0.3), order = c("virginica", "setosa", "versicolor"))

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

uncs\_guni\_cn

*Unconditional/symmetric Gaussian/uniform combined noise*

---

**Description**

Introduction of *Unconditional/symmetric Gaussian/uniform combined noise* into a classification dataset.

**Usage**

```
## Default S3 method:
uncs_guni_cn(x, y, level, k = 0.2, sortid = TRUE, ...)

## S3 method for class 'formula'
uncs_guni_cn(formula, data, ...)
```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
k	a double in [0,1] with the scale used for the standard deviation (default: 0.2).

sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

*Unconditional/symmetric Gaussian/uniform combined noise* corrupts all the samples for each attribute in the dataset. Their values are corrupted by adding a random value following a Gaussian distribution of  $mean = 0$  and  $standard\ deviation = (max-min) \cdot k$ , being  $max$  and  $min$  the limits of the attribute domain. For nominal attributes, a random value is chosen. Additionally, this noise model also selects  $(level \cdot 100)\%$  of the samples in the dataset with independence of their class. The labels of these samples are randomly replaced by different ones within the set of class labels.

### Value

An object of class `ndmodel` with elements:

xnoise	a data frame with the noisy input attributes.
ynoise	a factor vector with the noisy output class.
numnoise	an integer vector with the amount of noisy samples per variable.
idnoise	an integer vector list with the indices of noisy samples per variable.
numclean	an integer vector with the amount of clean samples per variable.
idclean	an integer vector list with the indices of clean samples per variable.
distr	an integer vector with the samples per class in the original data.
model	the full name of the noise introduction model used.
param	a list of the argument values.
call	the function call.

### Note

Noise model adapted from the papers in References.

### References

S. Kazmierczak and J. Mandziuk. **A committee of convolutional neural networks for image classification in the concurrent presence of feature and label noise**. In *Proc. 16th International Conference on Parallel Problem Solving from Nature*, volume 12269 of LNCS, pages 498-511, 2020. doi:10.1007/9783030581121\_34.

### See Also

[sym\\_cuni\\_cn](#), [sym\\_cuni\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```

# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- uncs_guni_cn(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- uncs_guni_cn(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)

```

unc\_fixw\_an

*Unconditional fixed-width attribute noise***Description**

Introduction of *Unconditional fixed-width attribute noise* into a classification dataset.

**Usage**

```

## Default S3 method:
unc_fixw_an(x, y, level, k = 0.1, sortid = TRUE, ...)

## S3 method for class 'formula'
unc_fixw_an(formula, data, ...)

```

**Arguments**

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced in nominal attributes.
k	a double in [0,1] with the domain proportion of the noise width (default: 0.1).
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

**Details**

*Unconditional fixed-width attribute noise* corrupts all the samples in the dataset. For each attribute  $A$ , all the original values are corrupted by adding a random number in the interval  $[-width, width]$ , being  $width = (max(A)-min(A)) \cdot k$ . For nominal attributes,  $(level \cdot 100)\%$  of the samples in the dataset are chosen and a random value is selected as noisy.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per attribute.
<code>idnoise</code>	an integer vector list with the indices of noisy samples per attribute.
<code>numclean</code>	an integer vector with the amount of clean samples per attribute.
<code>idclean</code>	an integer vector list with the indices of clean samples per attribute.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References, corrupting all samples and allowing nominal attributes.

**References**

A. Ramdas, B. Póczos, A. Singh, and L. A. Wasserman. **An analysis of active learning with uniform feature noise**. In *Proc. 17th International Conference on Artificial Intelligence and Statistics*, volume 33 of JMLR, pages 805-813, 2014. url:<http://proceedings.mlr.press/v33/ramdas14.html>.

**See Also**

[sym\\_end\\_an](#), [sym\\_sgau\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- unc_fixw_an(x = iris2D[, -ncol(iris2D)], y = iris2D[, ncol(iris2D)], level = 0.1)

# show results
```

```
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- unc_fixw_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

---

unc\_vgau\_an

*Unconditional vp-Gaussian attribute noise*


---

### Description

Introduction of *Unconditional vp-Gaussian attribute noise* into a classification dataset.

### Usage

```
## Default S3 method:
unc_vgau_an(x, y, level, sortid = TRUE, ...)

## S3 method for class 'formula'
unc_vgau_an(formula, data, ...)
```

### Arguments

x	a data frame of input attributes.
y	a factor vector with the output class of each sample.
level	a double in [0,1] with the noise level to be introduced.
sortid	a logical indicating if the indices must be sorted at the output (default: TRUE).
...	other options to pass to the function.
formula	a formula with the output class and, at least, one input attribute.
data	a data frame in which to interpret the variables in the formula.

### Details

In *Unconditional vp-Gaussian attribute noise*, the noise level for numeric attributes indicates the magnitude of the errors introduced. For each attribute  $A$ , all the original values are corrupted by adding a random number that follows a Gaussian distribution with  $mean = 0$  and  $variance = level\%$  of the variance of  $A$ . For nominal attributes,  $(level \cdot 100)\%$  of the samples in the dataset are chosen and a random value is selected as noisy.

**Value**

An object of class `ndmodel` with elements:

<code>xnoise</code>	a data frame with the noisy input attributes.
<code>ynoise</code>	a factor vector with the noisy output class.
<code>numnoise</code>	an integer vector with the amount of noisy samples per attribute.
<code>idnoise</code>	an integer vector list with the indices of noisy samples per attribute.
<code>numclean</code>	an integer vector with the amount of clean samples per attribute.
<code>idclean</code>	an integer vector list with the indices of clean samples per attribute.
<code>distr</code>	an integer vector with the samples per class in the original data.
<code>model</code>	the full name of the noise introduction model used.
<code>param</code>	a list of the argument values.
<code>call</code>	the function call.

**Note**

Noise model adapted from the papers in References, corrupting all samples and allowing nominal attributes.

**References**

X. Huang, L. Shi, and J. A. K. Suykens. **Support vector machine classifier with pinball loss.** *IEEE Transactions on Pattern Analysis and Machine Intelligence*, 36(5):984-997, 2014. doi:10.1109/TPAMI.2013.178.

**See Also**

[synd\\_rpix\\_an](#), [unc\\_fixw\\_an](#), [print.ndmodel](#), [summary.ndmodel](#), [plot.ndmodel](#)

**Examples**

```
# load the dataset
data(iris2D)

# usage of the default method
set.seed(9)
outdef <- unc_vgau_an(x = iris2D[,-ncol(iris2D)], y = iris2D[,ncol(iris2D)], level = 0.1)

# show results
summary(outdef, showid = TRUE)
plot(outdef)

# usage of the method for class formula
set.seed(9)
outfrm <- unc_vgau_an(formula = Species ~ ., data = iris2D, level = 0.1)

# check the match of noisy indices
identical(outdef$idnoise, outfrm$idnoise)
```

# Index

## \* datasets

diris2D, 18  
iris2D, 35

asy\_def\_ln, 3, 12, 43  
asy\_int\_an, 5, 15, 35  
asy\_spa\_ln, 7, 60  
asy\_uni\_an, 6, 9, 35  
asy\_uni\_ln, 11, 47  
attn\_uni\_ln, 12, 56, 75

boud\_gau\_an, 14

clu\_vot\_ln, 16, 22, 65

diris2D, 18

exp\_bor\_ln, 20, 25  
exps\_cuni\_ln, 13, 19, 68

fra\_bdir\_ln, 8, 22, 45

gam\_bor\_ln, 24  
gau\_bor\_ln, 27, 28, 133  
gaum\_bor\_ln, 26, 40, 133  
glev\_uni\_ln, 30, 87, 89

hubp\_uni\_ln, 32, 41, 72

imp\_int\_an, 15, 33  
iris, 35  
iris2D, 18, 35  
irs\_bdir\_ln, 23, 36, 45

lap\_bor\_ln, 38, 52, 136  
larm\_uni\_ln, 29, 40, 72

maj\_udir\_ln, 12, 42, 47  
mind\_bdir\_ln, 8, 44, 60  
minp\_uni\_ln, 46, 50  
mis\_pre\_ln, 17, 47, 70  
mulc\_udir\_ln, 49

nei\_bor\_ln, 51, 54, 74  
nlin\_bor\_ln, 49, 53, 74

oned\_uni\_ln, 33, 54, 75  
opes\_idnn\_ln, 57  
opes\_idu\_ln, 20, 58, 59

pai\_bdir\_ln, 23, 38, 61  
plot.ndmodel, 4, 6, 8, 10, 12, 13, 15, 17, 18,  
20, 22, 23, 25, 27, 29, 31, 33, 35, 36,  
38, 40, 41, 43, 45, 47, 49, 50, 52, 54,  
56, 58, 60, 62, 63, 65, 66, 68, 70, 72,  
74, 75, 77, 78, 80, 82, 84, 85, 87, 89,  
91, 93, 94, 96, 98, 100, 102, 104,  
105, 107, 109, 111, 113, 115, 116,  
118, 120, 122, 124, 126, 128, 129,  
131, 133, 136, 137, 139, 141

pmd\_con\_ln, 22, 25, 64  
print.ndmodel, 4, 6, 8, 10, 12, 13, 15, 17, 18,  
20, 22, 23, 25, 27, 29, 31, 33, 35, 36,  
38, 40, 41, 43, 45, 47, 49, 50, 52, 54,  
56, 58, 60, 62, 64, 65, 66, 68, 70, 72,  
74, 75, 77, 78, 80, 82, 84, 85, 87, 89,  
91, 93, 94, 96, 98, 100, 102, 104,  
105, 107, 109, 111, 113, 115, 116,  
118, 120, 122, 124, 126, 128, 129,  
131, 133, 136, 137, 139, 141

qua\_uni\_ln, 13, 56, 67

sco\_con\_ln, 17, 65, 69  
sigb\_uni\_ln, 27, 29, 71  
smam\_bor\_ln, 49, 70, 72  
smu\_cuni\_ln, 33, 41, 74  
summary.ndmodel, 4, 6, 8, 10, 12, 13, 15, 17,  
18, 20, 22, 23, 25, 27, 29, 31, 33, 35,  
36, 38, 40, 41, 43, 45, 47, 49, 50, 52,  
54, 56, 58, 60, 62, 64–66, 68, 70, 72,  
74, 75, 76, 78, 80, 82, 84, 85, 87, 89,  
91, 93, 94, 96, 98, 100, 102, 104,

*105, 107, 109, 111, 113, 115, 116,  
118, 120, 122, 124, 126, 128, 129,  
131, 133, 136, 137, 139, 141*

*sym\_adj\_ln, 84, 115, 120*  
*sym\_cen\_ln, 86, 89, 116*  
*sym\_con\_ln, 88, 116*  
*sym\_cuni\_an, 84, 90, 93, 94, 105, 128, 137*  
*sym\_cuni\_cn, 91, 91, 128, 137*  
*sym\_cuni\_ln, 64, 66, 77, 93, 96, 105*  
*sym\_ddef\_ln, 95, 98, 129*  
*sym\_def\_ln, 97, 111, 129*  
*sym\_dia\_ln, 99, 109, 118*  
*sym\_dran\_ln, 85, 101, 120*  
*sym\_end\_an, 82, 102, 139*  
*sym\_exc\_ln, 96, 98, 104*  
*sym\_gau\_an, 78, 106, 126*  
*sym\_hie\_ln, 102, 110*  
*sym\_hienc\_ln, 31, 87, 108*  
*sym\_int\_an, 78, 107, 112*  
*sym\_natd\_ln, 113, 122, 131*  
*sym\_nean\_ln, 4, 115*  
*sym\_nexc\_ln, 31, 109, 117*  
*sym\_nuni\_ln, 115, 119, 131*  
*sym\_opt\_ln, 100, 121, 124*  
*sym\_pes\_ln, 100, 118, 123*  
*sym\_sgau\_an, 104, 125, 126, 139*  
*sym\_uni\_an, 36, 64, 66, 77, 84, 91, 113, 126*  
*sym\_uni\_ln, 36, 64, 66, 77, 94, 111, 128*  
*sym\_usim\_ln, 122, 124, 130*  
*symd\_gau\_an, 77, 104*  
*symd\_gimg\_an, 6, 10, 79*  
*symd\_rpix\_an, 80, 81, 141*  
*symd\_uni\_an, 83, 107, 113*

*ugau\_bor\_ln, 40, 132, 136*  
*ulap\_bor\_ln, 52, 54, 134*  
*unc\_fixw\_an, 82, 138, 141*  
*unc\_vgau\_an, 10, 80, 140*  
*uncs\_guni\_cn, 93, 136*