

# Package ‘saens’

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**Type** Package

**Title** Small Area Estimation on the Fay-Herriot Model for Non-Sampled Area Estimation with Cluster Information.

**Version** 0.1.0

**Description** This package provides several methods for small area estimation on the Fay-Herriot model for non-sampled area estimation with cluster information.

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**URL** <https://github.com/Alfrzlp/sae-ns>

**BugReports** <https://github.com/Alfrzlp/sae-ns/issues>

**Encoding** UTF-8

**LazyData** true

**Depends** R (>= 4.00)

**RoxygenNote** 7.2.0

**Imports** cli, dplyr, ggplot2, methods, rlang, stats, tidyr

**NeedsCompilation** no

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AIC.eblupres	<i>Akaike's An Information Criterion</i>
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**Description**

Generic function calculating Akaike's "An Information Criterion" for EBLUP model

**Usage**

```
## S3 method for class 'eblupres'
AIC(object, ...)

## S3 method for class 'eblupres'
BIC(object, ...)
```

**Arguments**

object	EBLUP model
...	further arguments passed to or from other methods.

**Value**

AIC value

**Examples**

```
m1 <- eblupfh_cluster(y ~ x1 + x2 + x3, data = mys, vardir = "var", cluster = "clust")
AIC(m1)
```

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autoplot	<i>Create a complete ggplot appropriate to a particular data type</i>
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**Description**

autoplot() uses ggplot2 to draw a particular plot for an object of a particular class in a single command. This defines the S3 generic that other classes and packages can extend.

**Usage**

```
autoplot(object, ...)
```

**Arguments**

object	an object, whose class will determine the behaviour of autoplot
...	other arguments passed to specific methods

**Value**

a ggplot object

**See Also**

`autolayer()`, `ggplot()` and `fortify()`

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autoplot.eblupres	<i>Autoplot EBLUP results</i>
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**Description**

Autoplot EBLUP results

**Usage**

```
## S3 method for class 'eblupres'
autoplot(object, variable = "RSE", ...)
```

**Arguments**

object	EBLUP model
variable	variable to plot
...	further arguments passed to or from other methods.

**Value**

plot

**Examples**

```
library(saens)

m1 <- eblupfh_cluster(y ~ x1 + x2 + x3, data = mys, vardir = "var", cluster = "clust")
autoplot(m1)
```

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coef.eblupres	<i>Extract Model Coefficients</i>
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**Description**

Extract Model Coefficients

**Usage**

```
## S3 method for class 'eblupres'
coef(object, ...)
```

**Arguments**

object	EBLUP model
...	further arguments passed to or from other methods.

**Value**

model coefficients

**Examples**

```
m1 <- eblupfh_cluster(y ~ x1 + x2 + x3, data = mys, vardir = "var", cluster = "clust")
coef(m1)
```

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eblupfh

*EBLUPs based on a Fay-Herriot Model.*


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**Description**

This function gives the EBLUP (or EB predictor under normality) based on a Fay-Herriot model.

**Usage**

```
eblupfh(
  formula,
  data,
  vardir,
  method = "REML",
  maxiter = 100,
  precision = 1e-04,
  scale = FALSE,
  print_result = TRUE
)
```

**Arguments**

formula	an object of class formula that contains a description of the model to be fitted. The variables included in the formula must be contained in the data.
data	a data frame or a data frame extension (e.g. a tibble)
vardir	vector or column names from data that contain variance sampling from the direct estimator for each area
method	Fitting method can be chosen between 'ML' and 'REML'
maxiter	maximum number of iterations allowed in the Fisher-scoring algorithm. Default is 100 iterations.
precision	convergence tolerance limit for the Fisher-scoring algorithm. Default value is 0.0001.
scale	scaling auxiliary variable or not, default value is FALSE
print_result	print coefficient or not, default value is TRUE

**Details**

The model has a form that is response ~ auxiliary variables. where numeric type response variables can contain NA. When the response variable contains NA it will be estimated with cluster information.

**Value**

The function returns a list with the following objects (`df_res` and `fit`): `df_res` a data frame that contains the following columns:

- `y` variable response
- `eblup` estimated results for each area
- `random_effect` random effect for each area
- `vardir` variance sampling from the direct estimator for each area
- `mse` Mean Square Error
- `cluster` cluster information for each area
- `rse` Relative Standart Error (%)

`fit` a list containing the following objects:

- `estcoef` a data frame with the estimated model coefficients in the first column (`beta`), their asymptotic standard errors in the second column (`std.error`), the t-statistics in the third column (`tvalue`) and the p-values of the significance of each coefficient in last column (`pvalue`)
- `model_formula` model formula applied
- `method` type of fitting method applied (ML or REML)
- `random_effect_var` estimated random effect variance
- `convergence` logical value that indicates the Fisher-scoring algorithm has converged or not
- `n_iter` number of iterations performed by the Fisher-scoring algorithm.
- `goodness` vector containing several goodness-of-fit measures: loglikelihood, AIC, and BIC

**References**

1. Rao, J. N., & Molina, I. (2015). Small area estimation. John Wiley & Sons.
2. Anisa, R., Kurnia, A., & Indahwati, I. (2013). Cluster information of non-sampled area in small area estimation. E-Prosiding Internasional Departemen Statistika FMIPA Universitas Padjadjaran, 1(1), 69-76.

## Examples

```
library(saens)

m1 <- eblupfh(y ~ x1 + x2 + x3, data = na.omit(mys), vardir = "var")
m1 <- eblupfh(y ~ x1 + x2 + x3, data = na.omit(mys), vardir = ~var)
```

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eblupfh_cluster	<i>EBLUPs based on a Fay-Herriot Model with Cluster Information.</i>
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## Description

This function gives the EBLUP (or EB predictor under normality) based on a Fay-Herriot model with cluster information for non-sampled areas.

## Usage

```
eblupfh_cluster(
  formula,
  data,
  vardir,
  cluster,
  method = "REML",
  maxiter = 100,
  precision = 1e-04,
  scale = FALSE,
  print_result = TRUE
)
```

## Arguments

formula	an object of class formula that contains a description of the model to be fitted. The variables included in the formula must be contained in the data.
data	a data frame or a data frame extension (e.g. a tibble)
vardir	vector or column names from data that contain variance sampling from the direct estimator for each area
cluster	vector or column name from data that contain cluster information.
method	Fitting method can be chosen between 'ML' and 'REML'
maxiter	maximum number of iterations allowed in the Fisher-scoring algorithm. Default is 100 iterations.
precision	convergence tolerance limit for the Fisher-scoring algorithm. Default value is 0.0001.
scale	scaling auxiliary variable or not, default value is FALSE
print_result	print coefficient or not, default value is TRUE

## Details

The model has a form that is response ~ auxiliary variables. where numeric type response variables can contain NA. When the response variable contains NA it will be estimated with cluster information.

**Value**

The function returns a list with the following objects `df_res` and `fit`: `df_res` a data frame that contains the following columns:

- `y` variable response
- `eblup` estimated results for each area
- `random_effect` random effect for each area
- `vardir` variance sampling from the direct estimator for each area
- `mse` Mean Square Error
- `cluster` cluster information for each area
- `rse` Relative Standart Error (%)

`fit` a list containing the following objects:

- `estcoef` a data frame with the estimated model coefficients in the first column (`beta`), their asymptotic standard errors in the second column (`std.error`), the t-statistics in the third column (`tvalue`) and the p-values of the significance of each coefficient in last column (`pvalue`)
- `model_formula` model formula applied
- `method` type of fitting method applied (ML or REML)
- `random_effect_var` estimated random effect variance
- `convergence` logical value that indicates the Fisher-scoring algorithm has converged or not
- `n_iter` number of iterations performed by the Fisher-scoring algorithm.
- `goodness` vector containing several goodness-of-fit measures: loglikelihood, AIC, and BIC

**References**

1. Rao, J. N., & Molina, I. (2015). Small area estimation. John Wiley & Sons.
2. Anisa, R., Kurnia, A., & Indahwati, I. (2013). Cluster information of non-sampled area in small area estimation. E-Prosiding Internasional Departemen Statistika FMIPA Universitas Padjadjaran, 1(1), 69-76.

**Examples**

```
library(saens)

m1 <- eblupfh_cluster(y ~ x1 + x2 + x3, data = mys, vardir = "var", cluster = "clust")
m1 <- eblupfh_cluster(y ~ x1 + x2 + x3, data = mys, vardir = ~var, cluster = ~clust)
```

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logLik.eblupres	<i>Extract Log-Likelihood</i>
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**Description**

Extract Log-Likelihood

**Usage**

```
## S3 method for class 'eblupres'
logLik(object, ...)
```

**Arguments**

object	EBLUP model
...	further arguments passed to or from other methods.

**Value**

Log-Likelihood value

**Examples**

```
library(saens)

model1 <- eblupfh_cluster(y ~ x1 + x2 + x3, data = mys, vardir = "var", cluster = "clust")
logLik(model1)
```

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milk	<i>milk: Data on fresh milk expenditure.</i>
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**Description**

Data on fresh milk expenditure, used by Arora and Lahiri (1997) and by You and Chapman (2006).

**Usage**

```
milk
```



**Format**

A data frame with 43 observations on the following 6 variables.

**SmallArea** areas of inferential interest.

**ni** sample sizes of small areas.

**yi** average expenditure on fresh milk for the year 1989 (direct estimates for the small areas).

**SD** estimated standard deviations of yi.

**var** variance sampling from the direct estimator (yi) for each area

**CV** estimated coefficients of variation of yi.

**MajorArea** major areas created by You and Chapman (2006). These areas have similar direct estimates and produce a large CV reduction when using a FH model.

**References**

1. Arora, V. and Lahiri, P. (1997). On the superiority of the Bayesian method over the BLUP in small area estimation problems. *Statistica Sinica* 7, 1053-1063.
2. You, Y. and Chapman, B. (2006). Small area estimation using area level models and estimated sampling variances. *Survey Methodology* 32, 97-103.

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mys

*mys: mean years of schooling people with disabilities in Papua Island, Indonesia.*

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**Description**

A dataset containing the mean years of schooling people with disabilities in Papua Island, Indonesia in 2021.

**Usage**

mys

**Format**

A data frame with 42 rows and 7 variables with 10 domains are non-sampled areas.

**area** regency municipality

**y** mean years of schooling people with disabilities

**var** variance sampling from the direct estimator for each area

**rse** relative standard error (%)

**x1** Number of Elementary Schools

**x2** Number of Junior High Schools

**x3** Number of Senior High Schools

**clust** Cluster

**Source**

<https://www.bps.go.id>

saens

*saens : Small Area Estimation (sae) for Non-sampled Areas.***Description**

saens package Provides several functions for area level of small area estimation for non sampled areas using cluster information.

**Author(s)**

Ridson Al Farizal P

**References**

1. Rao, J. N., & Molina, I. (2015). Small area estimation. John Wiley & Sons.
2. Anisa, R., Kurnia, A., & Indahwati, I. (2013). Cluster information of non-sampled area in small area estimation. E-Prosiding Internasional Departemen Statistika FMIPA Universitas Padjadjaran, 1(1), 69-76.

summary.eblupres

*Summarizing EBLUP Model Fits***Description**

'summary' method for class "eblupres"

**Usage**

```
## S3 method for class 'eblupres'
summary(object, ...)
```

**Arguments**

object            EBLUP model  
...                further arguments passed to or from other methods.

**Value**

The function returns a data frame that contains the following columns:

- \* y variable response
- \* eblup estimated results for each area
- \* random\_effect random effect for each area
- \* vardir variance sampling from the direct estimator for each area
- \* mse Mean Square Error
- \* cluster cluster information for each area
- \* rse Relative Standart Error (

**Examples**

```
library(saens)

model1 <- eblupfh_cluster(y ~ x1 + x2 + x3, data = mys, vardir = "var", cluster = "clust")
summary(model1)
```

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