

# Package ‘vmsae’

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**Title** Variational Multivariate Spatial Small Area Estimation

**Version** 0.1.0

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**Description** Variational Autoencoded Multivariate Spatial Fay-Herriot models are designed to efficiently estimate population parameters in small area estimation. This package implements the variational generalized multivariate spatial Fay-Herriot model (VGMSFH) using 'NumPyro' and 'PyTorch' backends, as demonstrated by Wang, Parker, and Holan (2025) <[doi:10.48550/arXiv.2503.14710](https://doi.org/10.48550/arXiv.2503.14710)>. The 'vmsae' package provides utility functions to load weights of the pretrained variational autoencoders (VAEs) as well as tools to train custom VAEs tailored to users specific applications.

**Depends** R (>= 3.5.0)

**Imports** dplyr, ggplot2, gridExtra, sf, tidyr, reticulate, methods,  
rlang

**URL** <https://github.com/zhenhua-wang/vmsae>

**BugReports** <https://github.com/zhenhua-wang/vmsae/issues>

**License** MIT + file LICENSE

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|                    |  |
|--------------------|--|
| coef,VGMSFH-method | <i>Extract Coefficients from a VGMSFH Object</i> |
|--------------------|--|

---

**Description**

This method extracts posterior mean estimates of model coefficients from a VGMSFH object. It can return either fixed effect coefficients or spatial random effects.

**Usage**

```
## S4 method for signature 'VGMSFH'
coef(object, var_idx = 1, type = "fixed")
```

**Arguments**

- |         |  |
|---------|--|
| object  | An object of class VGMSFH.   |
| var_idx | Integer. The index of the variable of interest (for multivariate models). Default is 1.  |
| type    | Character. The type of coefficient to extract. Options are: <ul style="list-style-type: none"><li>• "fixed" – extract the posterior mean of fixed effect coefficients (default).</li><li>• "spatial" – extract the posterior mean of spatial random effects.</li></ul> |

**Value**

A numeric vector of posterior means for the selected coefficient type.

**Examples**

```
library(vmsae)
example_model <- readRDS(system.file("example", "example_model.Rds", package = "vmsae"))
coef(example_model) # Get fixed effect coefficients
coef(example_model, type = "spatial") # Get spatial random effects
```

---

confint, VGMSFH-method *Compute Credible Intervals for VGMSFH Parameters*

---

## Description

This method computes 95\

## Usage

```
## S4 method for signature 'VGMSFH'
confint(object, var_idx = 1, field = "yhat_samples")
```

## Arguments

|         |  |
|---------|--|
| object  | An object of class VGMSFH.   |
| var_idx | Integer. The index of the variable of interest (for multivariate models). Default is 1.  |
| field   | Character. The name of the slot to summarize (e.g., "yhat_samples", "beta_samples", "spatial_samples"). Default is "yhat_samples". |

## Details

The function extracts posterior samples for the specified variable and then computes quantiles to form 95\

## Value

A data frame with two columns:

- lower: the 2.5\
- upper: the 97.5\

## Examples

```
library(vmsae)
example_model <- readRDS(system.file("example", "example_model.Rds", package = "vmsae"))
confint(example_model) # Get credible intervals for predicted values
confint(example_model, field = "beta_samples") # For fixed effects
```

---

|               |                         |
|---------------|-------------------------|
| Decoder-class | <i>Decoder S4 Class</i> |
|---------------|-------------------------|

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**Description**

An S4 class to represent a neural network decoder, used for emulating spatial priors. The class includes parameters for input and output weight matrices and biases, as well as region identifiers.

**Slots**

- GE0ID A character vector of region or area identifiers.
- W\_in An array representing the input weight matrix of the decoder.
- B\_in An array representing the input bias vector of the decoder.
- W\_out An array representing the output weight matrix of the decoder.
- B\_out An array representing the output bias vector of the decoder.

---

|                         |  |
|-------------------------|--|
| download_pretrained_vae | <i>Download and Extract a Pretrained VAE Model</i> |
|-------------------------|--|

---

**Description**

This function downloads a pretrained VAE model archive from Zenodo, extracts its contents into a specified directory, and removes the downloaded ZIP file after extraction.

**Usage**

download\_pretrained\_vae(model\_name, save\_dir, verbose = TRUE)

**Arguments**

- model\_name Character. The name of the model file (without extension) to download. This should correspond to a \*model\_name\*.zip file hosted on Zenodo (e.g., "ca\_county").
- save\_dir Character. The local directory where the model should be saved and extracted.
- verbose Logical; if TRUE (default), prints progress and error messages.

**Value**

No return value, called for side effects

**Examples**

```
## Not run:
library(vmsae)
# this function is time consuming for the first run
install_environment()
load_environment()
download_pretrained_vae("mo_county", tempdir())

## End(Not run)
```

---

|                     |                                    |
|---------------------|------------------------------------|
| install_environment | <i>Install python environment.</i> |
|---------------------|------------------------------------|

---

**Description**

This function creates the vmsae python environment and installs required packages.

**Usage**

```
install_environment(envname = "vmsae")
```

**Arguments**

|         |  |
|---------|--|
| envname | Character. The name of the Python environment to create or update. Default is "vmsae". |
|---------|--|

**Value**

No return value, called for side effects

**Examples**

```
## Not run:
library(vmsae)
# this function is time consuming for the first run
install_environment()      # Install into default "vmsae" environment

# this step is time consuming for the first run
install_environment("custom") # Install into a custom-named environment

## End(Not run)
```

---

|                  |   |
|------------------|---|
| load_environment | <i>Load Python Environment and Source Model Modules</i> |
|------------------|---|

---

### Description

This function activates a specified Python virtual environment and sources Python modules used by the **vmsae** package, including models and python scripts.

### Usage

```
load_environment(envname = "vmsae")
```

### Arguments

|         |  |
|---------|--|
| envname | Character. The name of the Python environment to create or update. Default is "vmsae". |
|---------|--|

### Details

The function loads four Python scripts located in the package's `py/` directory:

- `vgmcar.py`
- `vae.py`
- `train_vae.py`
- `car_dataset.py`

The environment must be created beforehand (e.g., using `install_environment()`), and must include all Python dependencies required by these modules.

### Value

No return value, called for side effects

### Examples

```
## Not run:
library(vmsae)

# this function is time consuming for the first run
install_environment()
load_environment()      # Load default "vmsae" environment

# this function is time consuming for the first run
install_environment("custom")
load_environment("custom") # Load custom virtual environment

## End(Not run)
```

---

|          |                                    |
|----------|------------------------------------|
| load_vae | <i>Load Pretrained VAE Decoder</i> |
|----------|------------------------------------|

---

## Description

Load a pretrained Variational Autoencoder (VAE) decoder from disk. This function reads the saved PyTorch model weights and corresponding GEOID list, and constructs a Decoder S4 object with the loaded parameters.

## Usage

```
load_vae(model_name, save_dir = NULL)
```

## Arguments

|            |   |
|------------|---|
| model_name | Character. The name of the trained VAE model (without .zip extensions).                                   |
| save_dir   | Character. The directory where the trained VAE model is saved. Defaults to the current directory if NULL. |

## Details

This function assumes the model was trained and saved using `train_vae()`, and that the decoder weights are stored in a file compatible with `torch::load()` (via `reticulate`). It extracts the decoder input/output weights and biases, along with region GEOIDs, and returns them as an S4 object of class `Decoder`.

## Value

An object of class `Decoder`, containing the decoder weights and region identifiers.

## Examples

```
## Not run:
library(vmsae)
# this function is time consuming for the first run
install_environment()
load_environment()
decoder <- load_vae(model_name = "mo_county")

## End(Not run)
```

---

plot, VGMSFH-method      *Plot VGMSFH Result*


---

### Description

This method plots spatial summaries of results from a VGMSFH object, including model estimates and comparisons with direct estimates.

### Usage

```
## S4 method for signature 'VGMSFH'
plot(x, shp = NULL, var_idx = 1, type = "compare", verbose = TRUE)
```

### Arguments

|         |   |
|---------|---|
| x       | An object of class VGMSFH, containing posterior samples and direct estimates from the model.  |
| shp     | An sf object representing the spatial shapefile. If NULL, the function will automatically download a shapefile associated with the pretrained model.  |
| var_idx | Integer. The index of the variable of interest (for multivariate models).   |
| type    | Character. The type of plot to generate. Options are: <ul style="list-style-type: none"> <li>"compare" – compare direct estimates and model-based estimates.</li> <li>"estimate" – show the posterior mean and standard deviation of the model estimate.</li> </ul> |
| verbose | Logical; if TRUE (default), prints error messages.  |

### Details

The function provides spatial visualization of model results. It supports both univariate and multivariate response settings. When type = "compare", it generates side-by-side choropleth maps for the direct and model-based estimates. When type = "estimate", it plots the posterior mean and standard deviation of the VGMSFH model output.

If no shapefile is provided, a default geometry is loaded from the pretrained repository.

### Value

A ggplot object. The plot is rendered to the active device.

### Examples

```
library(vmsae)
library(sf)
example_model <- readRDS(system.file("example", "example_model.Rds", package = "vmsae"))
example_shp <- read_sf(system.file("example", "mo_county.shp", package = "vmsae"))
plot(example_model, shp = example_shp, type = "compare")
plot(example_model, shp = example_shp, type = "estimate", var_idx = 2)
```



---

summary, VGMSFH-method *Summarize VGMSFH Result*


---

## Description

This method provides a summary of posterior samples from a VGMSFH object, including posterior means and credible intervals for a specified parameter field.

## Usage

```
## S4 method for signature 'VGMSFH'
summary(object, var_idx = 1, field = "beta_samples")
```

## Arguments

|         |   |
|---------|---|
| object  | An object of class VGMSFH, containing posterior samples from the model.   |
| var_idx | Integer. The index of the variable of interest (for multivariate models). Default is 1.   |
| field   | Character. The name of the slot in the VGMSFH object to summarize (e.g., "beta_samples", "spatial_samples", "yhat_samples"). Default is "beta_samples". |

## Details

This function extracts the posterior samples for the specified variable index, and combines it with `confint()` to compute credible intervals. The result is a compact summary table of central tendency and uncertainty.

## Value

A data frame with columns:

- mean: Posterior mean,
- lower: Lower bound of the credible interval,
- upper: Upper bound of the credible interval.

## Examples

```
library(vmsae)
example_model <- readRDS(system.file("example", "example_model.Rds", package = "vmsae"))
summary(example_model) # Summary of beta_samples for variable 1
summary(example_model, var_idx = 2, field = "yhat_samples")
```

train\_vae

*Train VAE for CAR Prior***Description**

Trains a Variational Autoencoder (VAE) to learn the spatial structure implied by the Conditional Autoregressive (CAR) prior. The trained VAE parameters are saved and can later be used as a generator within Hamiltonian Monte Carlo (HMC) sampling.

**Usage**

```
train_vae(
    W,
    GEOID,
    model_name,
    save_dir,
    n_samples = 10000,
    batch_size = 256,
    epoch = 10000,
    lr_init = 0.001,
    lr_min = 1e-07,
    verbose = TRUE
)
```

**Arguments**

|            |  |
|------------|--|
| W          | Matrix. A proximity or adjacency matrix representing spatial relationships.  |
| GEOID      | Character vector. Identifiers for spatial units (e.g., region or area codes).  |
| model_name | Character. The name of the trained VAE model.  |
| save_dir   | Character. Directory to save the trained VAE model and associated metadata. Defaults to the current working directory. |
| n_samples  | Integer. Number of samples to draw from the prior for training. Default is 10000.                                      |
| batch_size | Integer. Batch size for VAE training. Default is 256.  |
| epoch      | Integer. Number of training epochs. Default is 10000.  |
| lr_init    | Numeric. Initial learning rate. Default is 0.001.  |
| lr_min     | Numeric. Minimum learning rate at the final epoch. Default is 1e-7.  |
| verbose    | Logical; if TRUE (default), prints progress.   |

**Details**

The function requires a configured Python environment via the **reticulate** interface, with VAE training implemented in Python. It uses `py$train_vae()` defined in the sourced Python modules (see [load\\_environment](#)).

**Value**

A named list containing:

|      |                             |
|------|-----------------------------|
| loss | Total training loss         |
| RCL  | Reconstruction error        |
| KLD  | Kullback–Leibler divergence |

**Examples**

```
## Not run:
library(vmsae)
library(sf)
# this function is time consuming for the first run
install_environment()
load_environment()

acs_data <- read_sf(system.file("example", "mo_county.shp", package = "vmsae"))
W <- readRDS(system.file("example", "W.Rds", package = "vmsae"))

loss <- train_vae(W = W,
  GEOID = acs_data$GEOID,
  model_name = "test",
  save_dir = tempdir(),
  n_samples = 1000, # set to larger values in practice, e.g. 10000.
  batch_size = 256,
  epoch = 1000)    # set to larger values in practice, e.g. 10000.

## End(Not run)
```

---

VGMSFH-class

VGMSFH S4 Class

---

**Description**

An S4 class to store results from the Variational Gaussian Markov Small Area Estimation with Fay-Herriot model (VGMSFH). This class holds the posterior samples for various model components as well as the original direct estimates.

**Slots**

`model_name` Character. The name of the trained VAE model.

`direct_estimate` Array. Direct estimates of parameters.

`yhat_samples` Array. Posterior samples of the estimated parameters.

`spatial_samples` Array. Posterior samples of the estimated spatial random effects.

`beta_samples` Array. Posterior samples of the fixed effect coefficients.

`all_samples` List. Posterior samples of all parameters in the VGMSFH model.

vgmsfh\_numpyro

*Run VGMSFH Using NumPyro***Description**

This function runs the Variational Generalized Multivariate Spatil Fay-Herriot model (VGMSFH) using NumPyro as the inference backend. It loads pretrained VAE decoder weights, prepares the data, and performs posterior sampling.

**Usage**

```
vgmsfh_numpyro(
    y,
    y_sigma,
    X,
    W,
    GEOID,
    model_name,
    save_dir = NULL,
    num_warmup = 1000,
    num_samples = 1000
)
```

**Arguments**

|                          |  |
|--------------------------|--|
| <code>y</code>           | Matrix. Response variables (direct estimates).   |
| <code>y_sigma</code>     | Matrix. Reported standard deviations of the responses.   |
| <code>X</code>           | Matrix. Covariate matrix.  |
| <code>W</code>           | Matrix. Proximity or adjacency matrix defining spatial structure.  |
| <code>GEOID</code>       | Character vector. FIPS codes or other region identifiers used to match with the pretrained VAE model.          |
| <code>model_name</code>  | Character. The name of the pretrained VAE model.   |
| <code>save_dir</code>    | Character. The directory where the VAE model is stored. If NULL, a default pretrained model directory is used. |
| <code>num_warmup</code>  | Integer. Number of warmup (burn-in) iterations. Default is 1000.   |
| <code>num_samples</code> | Integer. Number of posterior samples to draw. Default is 1000.   |

**Details**

This function uses a pretrained VAE decoder to parameterize the CAR prior and enables scalable inference through NumPyro. It is suitable for both univariate and multivariate response modeling in spatial domains.

**Value**

An object of class VGMSFH, which contains:

- `direct_estimate`: the observed response data,
- `yhat_samples`: posterior samples of the latent population process,
- `spatial_samples`: posterior samples of spatial random effects (CAR),
- `beta_samples`: posterior samples of fixed effect coefficients,
- `all_samples`: a list containing all sampled parameters, including  $\mu$ ,  $\delta$ , and other intermediate quantities.

**References**

Wang, Z., Parker, P. A., & Holan, S. H. (2025). Variational Autoencoded Multivariate Spatial Fay-Herriot Models. arXiv:2503.14710. <https://arxiv.org/abs/2503.14710>

**Examples**

```
## Not run:
library(sf)
library(vmsae)
# this function is time consuming for the first run
install_environment()
load_environment()

acs_data <- read_sf(system.file("example", "mo_county.shp", package = "vmsae"))
y <- readRDS(system.file("example", "y.Rds", package = "vmsae"))
y_sigma <- readRDS(system.file("example", "y_sigma.Rds", package = "vmsae"))
X <- readRDS(system.file("example", "X.Rds", package = "vmsae"))
W <- readRDS(system.file("example", "W.Rds", package = "vmsae"))

num_samples <- 1000 # set to larger values in practice, e.g. 10000.
model <- vgmsfh_numpyro(y, y_sigma, X, W,
  GEOID = acs_data$GEOID,
  model_name = "mo_county", save_dir = NULL,
  num_samples = num_samples, num_warmup = num_samples)
y_hat_np <- model@yhat_samples
y_hat_mean_np <- apply(y_hat_np, c(2, 3), mean)
y_hat_lower_np <- apply(y_hat_np, c(2, 3), quantile, 0.025)
y_hat_upper_np <- apply(y_hat_np, c(2, 3), quantile, 0.975)

plot(model, shp = acs_data, type = "compare", var_idx = 2)

## End(Not run)
```

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