

# Package: soilVAE (via r-universe)

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**Type** Package

**Title** Supervised Variational Autoencoder Regression via 'reticulate'

**Version** 0.1.9

**Description** Supervised latent-variable regression for high-dimensional predictors such as soil reflectance spectra. The model uses an encoder-decoder neural network with a stochastic Gaussian latent representation regularized by a Kullback-Leibler term, and a supervised prediction head trained jointly with the reconstruction objective. The implementation interfaces R with a 'Python' deep-learning backend and provides utilities for training, tuning, and prediction.

**License** MIT + file LICENSE

**Encoding** UTF-8

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.3.3

**Imports** reticulate, stats

**Suggests** knitr, rmarkdown, prospectr, pls

**VignetteBuilder** knitr

**SystemRequirements** Python (>= 3.9); TensorFlow (>= 2.13); Keras (>= 3)

**URL** <https://hugomachadorodrigues.github.io/soilVAE/>,  
<https://github.com/HugoMachadoRodrigues/soilVAE/>

**BugReports** <https://github.com/HugoMachadoRodrigues/soilVAE/issues>

**Config/pak/sysreqs** libpng-dev python3

**Repository** <https://hugomachadorodrigues.r-universe.dev>

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datsoilspc	<i>Soil spectroscopy example dataset used in the soilVAE vignettes</i>
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### Description

A small soil spectroscopy dataset packaged with **soilVAE** for demonstrating typical spectral pre-processing (reflectance  $\rightarrow$  absorbance, resampling, SNV, smoothing) and for comparing a classic PLS baseline model against supervised VAE regression via **soilVAE**.

### Format

A data.frame or list containing at minimum:

**spc** Numeric matrix/data.frame of reflectance spectra (samples  $\times$  wavelengths).

**TotalCarbon** Numeric vector of total carbon values.

### Details

The object datsoilspc contains:

- spc: a numeric matrix (or data.frame) of reflectance spectra, with rows as samples and columns as wavelengths (nm). Column names should be interpretable as numeric wavelengths.
- TotalCarbon: a numeric vector with the soil total carbon content for each sample.

Depending on the original source, additional columns may be present (e.g., sample identifiers or other soil properties).

The dataset is intended for examples and unit-sized demonstrations. It is not meant to be a comprehensive soil spectral library.

**Examples**

```
data("datsoilspc", package = "soilVAE")
str(datsoilspc)

# basic plot of reflectance spectra
spc <- as.matrix(datsoilspc$spc)
wav <- as.numeric(colnames(spc))
matplot(wav, t(spc), type = "l", lty = 1,
        xlab = "Wavelength / nm", ylab = "Reflectance")
```

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select\_best\_from\_grid *Select the best configuration from a tuning table*

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**Description**

Select the best configuration from a tuning table

**Usage**

```
select_best_from_grid(
  tuning_df,
  selection_metric = c("euclid", "rmse", "r2", "rpiq")
)
```

**Arguments**

tuning\_df            Data frame containing RMSE\_val, R2\_val, and RPIQ\_val.  
selection\_metric    One of: "euclid", "rmse", "r2", "rpiq".

**Value**

List with best (one-row data frame) and best\_score.

**Examples**

```
tuning_df <- data.frame(
  cfg_id = 1:3,
  RMSE_val = c(0.5, 0.3, 0.4),
  R2_val = c(0.8, 0.9, 0.85),
  RPIQ_val = c(2.0, 3.0, 2.5)
)
best <- select_best_from_grid(tuning_df, selection_metric = "rmse")
best$best
```

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tune\_vae\_train\_val      *Tune VAEReg on a train/validation split*

---

### Description

Tune VAEReg on a train/validation split

### Usage

```
tune_vae_train_val(X_tr, y_tr, X_va, y_va, seed = 123, grid_vae)
```

### Arguments

X_tr	Train predictors matrix.
y_tr	Train response numeric.
X_va	Validation predictors matrix.
y_va	Validation response numeric.
seed	Integer seed.
grid_vae	Data frame with required columns: hidden_enc (list), hidden_dec (list), latent_dim, dropout, lr, beta_kl, alpha_y, epochs, batch_size, patience.

### Value

A list with grid (input grid) and tuning\_df (metrics per config).

### Examples

```
## Not run:
vae_configure()
X <- matrix(rnorm(500), nrow = 50, ncol = 10)
y <- rnorm(50)
grid <- data.frame(
  latent_dim = 4L, dropout = 0.1, lr = 0.001,
  beta_kl = 1, alpha_y = 1, epochs = 5L,
  batch_size = 16L, patience = 3L
)
grid$hidden_enc <- list(c(32L, 16L))
grid$hidden_dec <- list(c(16L, 32L))
result <- tune_vae_train_val(
  X_tr = X[1:40, ], y_tr = y[1:40],
  X_va = X[41:50, ], y_va = y[41:50],
  grid_vae = grid
)

## End(Not run)
```

---

`vae_build`*Build a supervised VAE regression model (VAEReg)*

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**Description**

Build a supervised VAE regression model (VAEReg)

**Usage**

```
vae_build(  
  input_dim,  
  hidden_enc = c(512L, 256L),  
  hidden_dec = c(256L, 512L),  
  latent_dim = 32L,  
  dropout = 0.1,  
  lr = 0.001,  
  beta_kl = 1,  
  alpha_y = 1  
)
```

**Arguments**

<code>input_dim</code>	integer
<code>hidden_enc</code>	integer vector
<code>hidden_dec</code>	integer vector
<code>latent_dim</code>	integer
<code>dropout</code>	numeric
<code>lr</code>	numeric learning rate
<code>beta_kl</code>	numeric
<code>alpha_y</code>	numeric

**Value**

A fitted 'Python' 'Keras' model object (VAEReg).

**Examples**

```
## Not run:  
vae_configure()  
model <- vae_build(input_dim = 100L)  
  
## End(Not run)
```

---

vae\_configure                    *Configure Python / reticulate for soilVAE*

---

**Description**

Configure Python / reticulate for soilVAE

**Usage**

```
vae_configure(python = NULL, venv = NULL, conda = NULL)
```

**Arguments**

python	Path to python executable (optional).
venv	Name of virtualenv to use (optional).
conda	Name of conda env to use (optional).

**Value**

TRUE invisibly.

**Examples**

```
## Not run:  
# Let reticulate auto-detect Python  
vae_configure()  
  
# Or point to an existing virtual environment  
vae_configure(venv = "r-soilvae")  
  
## End(Not run)
```

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vae\_encode                    *Extract latent embeddings (z) from VAEReg*

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**Description**

Extract latent embeddings (z) from VAEReg

**Usage**

```
vae_encode(model, X)
```

**Arguments**

model	Python VAEReg
X	matrix

**Value**

Matrix of latent embeddings (samples x latent\_dim).

**Examples**

```
## Not run:
vae_configure()
X <- matrix(rnorm(200), nrow = 20, ncol = 10)
y <- rnorm(20)
model <- vae_build(input_dim = ncol(X))
vae_fit(model, X, y, epochs = 5L)
z <- vae_encode(model, X)

## End(Not run)
```

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vae\_fit

*Fit VAEReg*


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**Description**

Fit VAEReg

**Usage**

```
vae_fit(
  model,
  X,
  y,
  X_val = NULL,
  y_val = NULL,
  epochs = 80L,
  batch_size = 64L,
  patience = 10L,
  verbose = 0L
)
```

**Arguments**

model	Python VAEReg object from vae_build()
X	matrix (n x p)
y	numeric vector (n)
X_val	optional matrix
y_val	optional numeric vector
epochs	integer
batch_size	integer
patience	integer for early stopping (only if validation provided)
verbose	0/1/2

**Value**

The fitted model, invisibly.

**Examples**

```
## Not run:
vae_configure()
X <- matrix(rnorm(200), nrow = 20, ncol = 10)
y <- rnorm(20)
model <- vae_build(input_dim = ncol(X))
vae_fit(model, X, y, epochs = 5L)

## End(Not run)
```

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vae_predict	<i>Predict y using VAEReg (via latent z -&gt; y_head)</i>
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**Description**

Predict y using VAEReg (via latent z -> y\_head)

**Usage**

```
vae_predict(model, X)
```

**Arguments**

model	Python VAEReg
X	matrix

**Value**

Numeric vector of predicted values.

**Examples**

```
## Not run:
vae_configure()
X <- matrix(rnorm(200), nrow = 20, ncol = 10)
y <- rnorm(20)
model <- vae_build(input_dim = ncol(X))
vae_fit(model, X, y, epochs = 5L)
preds <- vae_predict(model, X)

## End(Not run)
```

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