

Package ‘SVEMnet’

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Type Package

Title Self-Validated Ensemble Models with Elastic Net Regression

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Description Implements Self-Validated Ensemble Models (SVEM, Lemkus et al. (2021) <[doi:10.1016/j.chemolab.2021.104439](https://doi.org/10.1016/j.chemolab.2021.104439)>) using Elastic Net regression via 'glmnet' (Friedman et al. <[doi:10.18637/jss.v033.i01](https://doi.org/10.18637/jss.v033.i01)>). SVEM averages predictions from multiple models fitted to fractionally weighted bootstraps of the data, tuned with anti-correlated validation weights. Also implements the randomized permutation whole model test for SVEM (Karl (2024) <[doi:10.1016/j.chemolab.2024.105122](https://doi.org/10.1016/j.chemolab.2024.105122)>). Code for the whole model test was supplementary material of Karl (2024). Development of this package was assisted by 'GPT o1-preview' for code structure and documentation.

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Description

The SVEMnet package implements Self-Validated Ensemble Models (SVEM) using Elastic Net (including lasso and ridge) regression via `glmnet`. SVEM averages predictions from multiple models fitted to fractionally weighted bootstraps of the data, tuned with anti-correlated validation weights.

Functions

`SVEMnet` Fit an SVEMnet model using Elastic Net regression.
`svem_significance_test` Perform a whole-model significance test for SVEM models.
`predict.svem_model` Predict method for SVEM models.
`plot.svem_model` Plot method for SVEM models.
`coef.svem_model` Plot method for SVEM models.

Acknowledgments

Development of this package was assisted by GPT o1-preview, which helped in constructing the structure of some of the code and the roxygen documentation. The code for the significance test is taken from the supplementary material of Karl (2024) (it was handwritten by that author).

Author(s)

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References

- Gotwalt, C., & Ramsey, P. (2018). Model Validation Strategies for Designed Experiments Using Bootstrapping Techniques With Applications to Biopharmaceuticals. *JMP Discovery Conference*. <https://community.jmp.com/t5/Discovery-Summit-2018/Model-Validation-Strategies-for-Designed-Experiments-p/73730>
- Karl, A. T. (2024). A randomized permutation whole-model test heuristic for Self-Validated Ensemble Models (SVEM). *Chemometrics and Intelligent Laboratory Systems*, 249, 105122. doi:10.1016/j.chemolab.2024.105122
- Lemkus, T., Gotwalt, C., Ramsey, P., & Weese, M. L. (2021). Self-Validated Ensemble Models for Design of Experiments. *Chemometrics and Intelligent Laboratory Systems*, 219, 104439. doi:10.1016/j.chemolab.2021.104439
- Xu, L., Gotwalt, C., Hong, Y., King, C. B., & Meeker, W. Q. (2020). General Applications of the Fractional-Random-Weight Bootstrap. *Technometrics*, 62(3), 345–358. doi:10.1080/00031305.2020.1731599
- Ramsey, P., Gaudard, M., & Levin, W. (2021). Accelerating Innovation with Space Filling Mixture Designs, Neural Networks and SVEM. *JMP Discovery Conference*. <https://community.jmp.com>

[com/t5/Abstracts/Accelerating-Innovation-with-Space-Filling-Mixture-Designs/ev-p/756841](https://community.jmp.com/t5/Abstracts/Accelerating-Innovation-with-Space-Filling-Mixture-Designs/ev-p/756841)

Ramsey, P., & Gotwalt, C. (2018). Model Validation Strategies for Designed Experiments Using Bootstrapping Techniques With Applications to Biopharmaceuticals. *JMP Discovery Conference - Europe*. <https://community.jmp.com/t5/Discovery-Summit-Europe-2018/Model-Validation-Strategies-for-Design-Experiments-Using-Bootstrapping-Techniques-With-Applications-to-Biopharmaceuticals/ev-p/51286>

Ramsey, P., Levin, W., Lemkus, T., & Gotwalt, C. (2021). SVEM: A Paradigm Shift in Design and Analysis of Experiments. *JMP Discovery Conference - Europe*. <https://community.jmp.com/t5/Abstracts/SVEM-A-Paradigm-Shift-in-Design-and-Analysis-of-Experiments-2021/ev-p/756634>

Ramsey, P., & McNeill, P. (2023). CMC, SVEM, Neural Networks, DOE, and Complexity: It's All About Prediction. *JMP Discovery Conference*.

coef.svem_model

Plot Coefficient Nonzero Percentages from a SVEMnet Model

Description

This function calculates the percentage of bootstrap iterations in which each coefficient is nonzero.

Usage

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

Arguments

object An object of class svem_model returned by the SVEMnet function.
 ... other arguments to pass.

Value

Invisibly returns a data frame containing the percentage of bootstraps where each coefficient is nonzero.

Acknowledgments

Development of this package was assisted by GPT o1-preview, which helped in constructing the structure of some of the code and the roxygen documentation. The code for the significance test is taken from the supplementary material of Karl (2024) (it was handwritten by that author).

plot.svem_model *Plot Method for SVEM Models*

Description

Plots actual versus predicted values for an svem_model using ggplot2.

Usage

```
## S3 method for class 'svem_model'  
plot(x, plot_debiased = TRUE, ...)
```

Arguments

x An object of class svem_model.
plot_debiased Logical; if TRUE, includes debiased predictions if available (default is TRUE).
... Additional arguments passed to ggplot2 functions.

Details

This function creates an actual vs. predicted plot for the SVEM model. If plot_debiased is TRUE and debiased predictions are available, it includes them in the plot.

****Plot Features:****

- ****Actual vs. Predicted Points:**** Plots the actual response values against the predicted values from the SVEM model.
- ****Debiased Predictions:**** If available and plot_debiased is TRUE, debiased predictions are included.
- ****Ideal Fit Line:**** A dashed line representing perfect prediction (slope = 1, intercept = 0) is included for reference.

Value

A ggplot object showing actual versus predicted values.

Acknowledgments

Development of this package was assisted by GPT o1-preview, which helped in constructing the structure of some of the code and the roxygen documentation. The code for the significance test is taken from the supplementary material of Karl (2024) (it was handwritten by that author).

`plot.svem_significance_test`*Plot SVEM Significance Test Results for Multiple Responses*

Description

Plots the Mahalanobis distances for the original and permuted data from multiple SVEM significance test results.

Usage

```
## S3 method for class 'svem_significance_test'  
plot(..., labels = NULL)
```

Arguments

<code>...</code>	One or more objects of class <code>svem_significance_test</code> , which are the outputs from svem_significance_test .
<code>labels</code>	Optional character vector of labels for the responses. If not provided, the function uses the response variable names.

Details

This function creates a combined plot of the Mahalanobis distances (d_Y and d_{π_Y}) for the original and permuted data from multiple SVEM significance test results. It groups the data by response and source type, displaying original and permutation distances side by side for each response.

****Usage Notes:****

- Use this function to compare the significance test results across multiple responses.
- The plot shows original and permutation distances next to each other for each response.

Value

A `ggplot` object showing the distributions of Mahalanobis distances for all responses.

Acknowledgments

Development of this package was assisted by GPT o1-preview, which helped in constructing the structure of some of the code and the roxygen documentation. The code for the significance test is taken from the supplementary material of Karl (2024) (it was handwritten by that author).

predict.svem_model *Predict Method for SVEM Models*

Description

Generates predictions from a fitted svem_model.

Usage

```
## S3 method for class 'svem_model'  
predict(object, newdata, debias = TRUE, se.fit = FALSE, ...)
```

Arguments

object	An object of class svem_model.
newdata	A data frame of new predictor values.
debias	Logical; default is TRUE.
se.fit	Logical; if TRUE, returns standard errors (default is FALSE).
...	Additional arguments.

Details

A debiased fit is output (along with the standard fit). This is provided to allow the user to match the output of JMP. https://www.jmp.com/support/help/en/18.1/?utm_source=help&utm_medium=redirect#page/jmp/
The debiasing coefficients are always calculated by SVEMnet(), and the predict() function determines whether the raw or debiased predictions are returned via the debias argument.

Value

Predictions or a list containing predictions and standard errors.

Acknowledgments

Development of this package was assisted by GPT o1-preview, which helped in constructing the structure of some of the code and the roxygen documentation. The code for the significance test is taken from the supplementary material of Karl (2024) (it was handwritten by that author).

Description

Wrapper for 'glmnet' (Friedman et al. 2010) to fit an ensemble of Elastic Net models using the Self-Validated Ensemble Model method (SVEM, Lemkus et al. 2021). Allows searching over multiple alpha values in the Elastic Net penalty.

Usage

```
SVEMnet(
  formula,
  data,
  nBoot = 200,
  glmnet_alpha = c(0, 0.25, 0.5, 0.75, 1),
  weight_scheme = c("SVEM", "FWR", "Identity"),
  objective = c("wSSE", "wAIC"),
  ...
)
```

Arguments

formula	A formula specifying the model to be fitted.
data	A data frame containing the variables in the model.
nBoot	Number of bootstrap iterations (default is 200).
glmnet_alpha	Elastic Net mixing parameter(s) (default is <code>c(0, 0.25, 0.5, 0.75, 1)</code>). Can be a vector of alpha values, where <code>alpha = 1</code> corresponds to Lasso and <code>alpha = 0</code> corresponds to Ridge regression.
weight_scheme	Weighting scheme for SVEM (default is "SVEM"). Valid options are "SVEM", "FWR", and "Identity". "FWR" calculates the Fractional Weight Regression (Xu et al., 2020) and is included for demonstration; "SVEM" generally provides better performance. "Identity" simply sets the training and validation weights to 1. Use with <code>nBoot = 1</code> and <code>objective = "wAIC"</code> to get an elastic net fit on the training data using AIC.
objective	Objective function for selecting lambda (default is "wSSE"). Valid options are "wAIC" and "wSSE". The "w" refers to "weighted" validation.
...	Additional arguments passed to the underlying <code>glmnet()</code> function.

Details

The Self-Validated Ensemble Model (SVEM, Lemkus et al., 2021) framework provides a bootstrap approach to improve predictions from various base learning models, including Elastic Net regression as implemented in 'glmnet'. SVEM is particularly suited for situations where a complex response surface is modeled with relatively few experimental runs.

In each of the ‘nBoot’ iterations, SVEMnet applies random exponentially distributed weights to the observations. Anti-correlated weights are used for validation.

SVEMnet allows for the Elastic Net mixing parameter (‘glmnet_alpha’) to be a vector, enabling the function to search over multiple ‘alpha’ values within each bootstrap iteration. Within each iteration, the model is fit for each specified ‘alpha’, and the best ‘alpha’ is selected based on the specified ‘objective’.

objective options:

“wAIC” Weighted Akaike Information Criterion. Balances model fit with complexity by penalizing the number of parameters. It is calculated as $AIC = n \times \log(wSSE / n) + 2 \times k$, where wSSE is the weighted sum of squared errors, n is the number of observations, and k is the number of parameters. Typically used with `weight_scheme="FWR"` or `weight_scheme="Identity"`

“wSSE” Weighted Sum of Squared Errors. Selects the lambda that minimizes the weighted validation error without penalizing model complexity. While this may lead to models that overfit when the number of parameters is large relative to the number of observations, SVEM mitigates overfitting (high prediction variance) by averaging over multiple bootstrap models. Typically used with `weight_scheme="SVEM"`

weight_scheme options:

“SVEM” Uses anti-correlated fractional weights for training and validation sets, improving model generalization by effectively simulating multiple training-validation splits (Lemkus et al. (2021)). Typically used with `objective="wSSE"`.

“FWR” Fractional Weight Regression as described by Xu et al. (2020). Weights are the same for both training and validation sets. This method does not provide the self-validation benefits of SVEM but is included for comparison. Typically used with `objective="wAIC"`.

“Identity” Uses weights of 1 for both training and validation. This uses the full dataset for both training and validation, effectively disabling the self-validation mechanism. Use with `objective="wAIC"` and `nBoot=1` to fit the Elastic Net on the AIC of the training data.

A debiased fit is output (along with the standard fit). This is provided to allow the user to match the output of JMP, which returns a debiased fit whenever `nBoot >= 10`.? https://www.jmp.com/support/help/en/18.1/?utm_
The debiasing coefficients are always calculated by SVEMnet(), and the `predict()` function determines whether the raw or debiased predictions are returned via its `debias` argument.

The output includes: ****Model Output:**** The returned object is a list of class `svem_model`, containing the following components:

- `parms`: Averaged coefficients across all bootstrap iterations.
- `debias_fit`: The debiasing linear model fit (if applicable). This is a linear model of the form $y \sim y_{pred}$, used to adjust the predictions and reduce bias.
- `coef_matrix`: Matrix of coefficients from each bootstrap iteration. Each row corresponds to a bootstrap iteration, and each column corresponds to a model coefficient.
- `nBoot`: Number of bootstrap iterations performed.
- `glmnet_alpha`: The Elastic Net mixing parameter(s) used. This is the alpha parameter from `glmnet`.
- `best_alphas`: The best alpha values selected during the fitting process for each bootstrap iteration.

- `best_lambdas`: The best lambda values selected during the fitting process for each bootstrap iteration.
- `weight_scheme`: The weighting scheme used in SVEM. Indicates whether "SVEM", "FWR", or "Identity" weights were used.
- `actual_y`: The response vector used in the model.
- `training_X`: The predictor matrix used in the model.
- `y_pred`: The predicted response values from the ensemble model before debiasing.
- `y_pred_debiased`: The debiased predicted response values (if debiasing is applied). Adjusted predictions using the `debias_fit` model.
- `nobs`: The number of observations in the dataset.
- `nparam`: The number of parameters (including the intercept), calculated as $\text{ncol}(X) + 1$.
- `formula`: The formula used in the model fitting.
- `terms`: The terms object extracted from the model frame.

Value

An object of class `svem_model`.

Acknowledgments

Development of this package was assisted by GPT o1-preview, which helped in constructing the structure of some of the code and the roxygen documentation. The code for the significance test is taken from the supplementary material of Karl (2024) (it was handwritten by that author).

References

- Gotwalt, C., & Ramsey, P. (2018). Model Validation Strategies for Designed Experiments Using Bootstrapping Techniques With Applications to Biopharmaceuticals. *JMP Discovery Conference*. <https://community.jmp.com/t5/Discovery-Summit-2018/Model-Validation-Strategies-for-Designed-Experiments/p/73730>
- Karl, A. T. (2024). A randomized permutation whole-model test heuristic for Self-Validated Ensemble Models (SVEM). *Chemometrics and Intelligent Laboratory Systems*, 249, 105122. doi:10.1016/j.chemolab.2024.105122
- Lemkus, T., Gotwalt, C., Ramsey, P., & Weese, M. L. (2021). Self-Validated Ensemble Models for Design of Experiments. *Chemometrics and Intelligent Laboratory Systems*, 219, 104439. doi:10.1016/j.chemolab.2021.104439
- Xu, L., Gotwalt, C., Hong, Y., King, C. B., & Meeker, W. Q. (2020). General Applications of the Fractional-Random-Weight Bootstrap. *Technometrics*, 62(3), 345–358. doi:10.1080/00031305.2020.1731599
- Ramsey, P., Gaudard, M., & Levin, W. (2021). Accelerating Innovation with Space Filling Mixture Designs, Neural Networks and SVEM. *JMP Discovery Conference*. <https://community.jmp.com/t5/Abstracts/Accelerating-Innovation-with-Space-Filling-Mixture-Designs/ev-p/756841>
- Ramsey, P., & Gotwalt, C. (2018). Model Validation Strategies for Designed Experiments Using Bootstrapping Techniques With Applications to Biopharmaceuticals. *JMP Discovery Conference* -


```

nSVEM = 5,
nPerm = 125,
percent = 90,
nBoot = 200,
glmnet_alpha = c(1),
weight_scheme = c("SVEM", "FWR"),
debias = FALSE,
objective = c("wSSE", "wAIC"),
verbose = FALSE,
...
)

```

Arguments

formula	A formula specifying the model to be tested.
data	A data frame containing the variables in the model.
nPoint	The number of random points to generate in the factor space (default: 2000).
nSVEM	The number of SVEM models to fit to the original data (default: 5).
nPerm	The number of SVEM models to fit to permuted data for reference distribution (default: 125).
percent	The percentage of variance to capture in the SVD (default: 90).
nBoot	The number of bootstrap iterations within SVEM (default: 200).
glmnet_alpha	The alpha parameter(s) for glmnet (default: c(1)).
weight_scheme	The weight scheme to use in SVEM (default: "SVEM"). Valid options are "SVEM" and "FWR".
debias	Logical; debiasing option passed to SVEMnet (default: FALSE).
objective	Character; the objective function to use in SVEMnet . Options are "wAIC" or "wSSE" (default: "wSSE").
verbose	Logical; if TRUE, displays progress messages (default: FALSE).
...	Additional arguments passed to the underlying SVEMnet() and then glmnet() functions.

Details

The ‘svem_significance_test’ function implements a whole-model test designed to gauge the significance of a fitted SVEM model compared to the null hypothesis of a constant response surface. This method helps identify responses that have relatively stronger or weaker relationships with study factors.

The test constructs standardized predictions by centering the SVEM predictions by the response mean and scaling by the ensemble standard deviation. A reference distribution is created by fitting the SVEM model to multiple randomized permutations of the response vector. The Mahalanobis distances of the original and permuted models are calculated using a reduced-rank singular value decomposition.

The R code to perform this test (using matrices of nSVEM and nPerm predictions) is taken from the supplementary material of Karl (2024).

Value

A list containing the test results.

Acknowledgments

Development of this package was assisted by GPT o1-preview, which helped in constructing the structure of some of the code and the roxygen documentation. The code for the significance test is taken from the supplementary material of Karl (2024) (it was handwritten by that author).

References

Karl, A. T. (2024). A randomized permutation whole-model test heuristic for Self-Validated Ensemble Models (SVEM). *Chemometrics and Intelligent Laboratory Systems*, 249, 105122. doi:10.1016/j.chemolab.2024.105122

Examples

```
# Simulate data
set.seed(0)
n <- 21
X1 <- runif(n)
X2 <- runif(n)
X3 <- runif(n)
y <- 1 + X1 + X2 + X1 * X2 + X1^2 + rnorm(n)
data <- data.frame(y, X1, X2, X3)

# Perform the SVEM significance test
test_result <- svem_significance_test(
  y ~ (X1 + X2 + X3)^2 + I(X1^2) + I(X2^2) + I(X3^2),
  data = data,
  nPoint = 2000,
  nSVEM = 5,
  nPerm = 125,
  nBoot = 200
)

# View the p-value
test_result$p_value

test_result2 <- svem_significance_test(
  y ~ (X1 + X2 )^2 + I(X1^2) + I(X2^2),
  data = data,
  nPoint = 2000,
  nSVEM = 5,
  nPerm = 125,
  nBoot = 200
)

# View the p-value
test_result2$p_value
```

```
# Plot the Mahalanobis distances  
plot(test_result, test_result2)
```

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