# Package 'CVEK'

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Title Cross-Validated Kernel Ensemble

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**Description** Using a library of base kernels, it learns a proper generating function from data by directly minimizing the ensemble model's cross-validation error, therefore guaranteeing robust test for a wide range of data-generating functions.

**Depends** R (>= 3.0.1), mvtnorm, MASS, psych, limSolve

License GPL-2

**Encoding** UTF-8

LazyData true

RoxygenNote 6.0.1

Suggests knitr, rmarkdown

VignetteBuilder knitr

NeedsCompilation no

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# R topics documented:

base Estimate

**Estimating Projection Matrices** 

### **Description**

Calculate the estiamted projection matrices for every kernels in the kernel library.

# Usage

baseEstimate(size, magn, Y, X1, X2, Kernlist, mode, lambda)

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### **Arguments**

size	A numeric number specifying the number of observations.
magn	A numeric number specifying the number of kernels in the kernel library.
Υ	Reponses of the dataframe.
X1	The first type of factor in the dataframe (could contains several subfactors).
X2	The second type of factor in the dataframe (could contains several subfactors).

Kernlist The kernel library containing several kernels given by user.

mode A character string indicating which tuning parameter criteria is to be used.

1 ambda A numeric string specifying the range of noise to be chosen. The lower limit of

lambda must be above 0.

#### **Details**

For a given mode, this function return a list of projection matrices for every kernels in the kernel library and a size\*magn matrix indicating errors.

### Value

A_hat	A list of projection matrices for every kernels in the kernel library.
error_mat	A size*magn matrix indicating errors.

### Author(s)

Wenying Deng

#### References

Jeremiah Zhe Liu and Brent Coull. Robust Hypothesis Test for Nonlinear Effect with Gaussian Processes. October 2017.

# **Examples**

```
##baseEstimate(size = 50, magn = 3, Y, X1, X2, Kernlist = NULL,
##mode = "loocv", lambda = exp(seq(-5, 5)))
```

dataGenerate

Generating Original Data

### **Description**

Generate original data based on specific kernels.

# Usage

```
dataGenerate(size, label_names, method = "rbf", int_effect = 0, l = 1,
    p = 2, eps = 0.01)
```

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### **Arguments**

size A numeric number specifying the number of observations.

label\_names A character string indicating all the interior variables included in each predictor.

method A character string indicating which kernel is to be computed.

int\_effect A numeric number specifying the size of interaction.

1 A numeric number indicating the hyperparameter (flexibility) of a specific ker-

nel.

p For polynomial, p is the power; for matern, v = p + 1/2; for rational, alpha = p.

eps A numeric number indicating the size of noise.

#### **Details**

This function generates with a specific kernel. The argument int\_effect represents the strength of interaction relative to the main effect since all sampled functions have been standardized to have unit norm.

### Value

data A dataframe to be fitted.

### Author(s)

Wenying Deng

# **Examples**

```
##data <- dataGenerate(size = 50, label_names =
##list(X1 = c("x1", "x2"), X2 = c("x3", "x4")),
##method = "rbf", int_effect = 0, l = 1, p = 2, eps = 0.01)</pre>
```

defineModel

Defining the Model

### **Description**

Give the complete formula and generate the expected kernel library.

# Usage

```
defineModel(formula, label_names, data, Kern_par)
```

### Arguments

formula A symbolic description of the model to be fitted.

label\_names A character string indicating all the interior variables included in each predictor.

data A dataframe to be fitted.

Kern\_par A dataframe indicating the parameters of base kernels to be created.

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#### **Details**

It processes data based on formula and label\_names and creates a kernel library according to the parameters given in Kern\_par.

#### Value

Y Reponses of the dataframe.

X1 The first type of factor in the dataframe (could contains several subfactors).
X2 The second type of factor in the dataframe (could contains several subfactors).

Kernlist The kernel library containing several kernels given by user.

### Author(s)

Wenying Deng

#### See Also

method: kernelGenerate

### **Examples**

```
##Kern_par <- data.frame(method = c("rbf", "polynomial", "matern"),
##Sigma = rep(0, 3), l = c(.5, 1, 1.5), p = 1:3)
##Kern_par$method <- as.character(Kern_par$method)
##defineModel(formula = Y \sim X1 + X2,
##label_names = list(X1 = c("x1", "x2"), X2 = c("x3", "x4")),
##data, Kern_par)
```

ensemble

Estimating Ensemble Kernel Matrices

### **Description**

Give a list of estimated kernel matrices and their weights.

### Usage

```
ensemble(n, D, strategy, beta, error_mat, A_hat)
```

### **Arguments**

n A	A numeric number	specifying the	e number of o	bservations.

D A numeric number specifying the number of kernels in the kernel library.

Strategy A character string indicating which ensemble strategy is to be used.

A numeric value specifying the parameter when strategy = "exp".

error\_mat A n\*D matrix indicating errors.

A\_hat A list of projection matrices for every kernels in the kernel library.

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#### **Details**

There are three ensemble strategies available here:

### **Empirical Risk Minimization**

After obtaining the estimated errors  $\{\hat{\epsilon}_d\}_{d=1}^D$ , we estimate the ensemble weights  $u = \{u_d\}_{d=1}^D$  such that it minimizes the overall error

$$\hat{u} = u \in \Delta argmin \parallel \sum_{d=1}^{D} u_d \hat{\epsilon}_d \parallel^2 \quad where \ \Delta = \{u | u \ge 0, \parallel u \parallel_1 = 1\}$$

Then produce the final ensemble prediction:

$$\hat{h} = \sum_{d=1}^{D} \hat{u}_d h_d = \sum_{d=1}^{D} \hat{u}_d A_{d,\hat{\lambda}_d} y = \hat{A}y$$

where  $\hat{A} = \sum_{d=1}^{D} \hat{u}_d A_{d,\hat{\lambda}_d}$  is the ensemble matrix.

### **Simple Averaging**

Motivated by existing literature in omnibus kernel, we propose another way to obtain the ensemble matrix by simply choosing unsupervised weights  $u_d = 1/D$  for d = 1, 2, ...D.

### **Exponential Weighting**

Additionally, another scholar gives a new strategy to calculate weights based on the estimated errors  $\{\hat{\epsilon}_d\}_{d=1}^D$ .

$$u_d(\beta) = \frac{exp(-\parallel \hat{\epsilon}_d \parallel_2^2 / \beta)}{\sum_{d=1}^{D} exp(-\parallel \hat{\epsilon}_d \parallel_2^2 / \beta)}$$

### Value

A\_est A list of estimated kernel matrices.

u\_hat A vector of weights of the kernels in the library.

### Author(s)

Wenying Deng

### References

Jeremiah Zhe Liu and Brent Coull. Robust Hypothesis Test for Nonlinear Effect with Gaussian Processes. October 2017.

Xiang Zhan, Anna Plantinga, Ni Zhao, and Michael C. Wu. A fast small-sample kernel independence test for microbiome community-level association analysis. December 2017.

Arnak S. Dalalyan and Alexandre B. Tsybakov. Aggregation by Exponential Weighting and Sharp Oracle Inequalities. In Learning Theory, Lecture Notes in Computer Science, pages 97–111. Springer, Berlin, Heidelberg, June 2007.

### See Also

mode: tuning

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### **Examples**

```
\#\#ensemble(n = 50, D = 6, strategy = "erm", beta = 1, error_mat, A_hat)
```

estimation

Conducting Gaussian Process Regression

### **Description**

Conduct gaussian process regression based on the estimated ensemble kernel matrix.

# Usage

```
estimation(Y, X1, X2, Kernlist, mode = "loocv", strategy = "erm",
beta = 1, lambda = exp(seq(-5, 5)))
```

# Arguments

Υ	Reponses of the dataframe.
X1	The first type of factor in the dataframe (could contains several subfactors).
X2	The second type of factor in the dataframe (could contains several subfactors).
Kernlis	The kernel library containing several kernels given by user.
mode	A character string indicating which tuning parameter criteria is to be used.
strategy	A character string indicating which ensemble strategy is to be used.
beta	A numeric value specifying the parameter when strategy = "exp".
lambda	A numeric string specifying the range of noise to be chosen. The lower limit of lambda must be above 0.

### **Details**

After obtaining the ensemble kernel matrix, we can calculate the outpur of gaussian process regression, the solution is given by

$$\hat{\beta} = [1^T (K + \lambda I)^{-1} 1]^{-1} 1^T (K + \lambda I)^{-1} y$$
$$\hat{\alpha} = (K + \lambda I)^{-1} (y - \hat{\beta} 1)$$

where  $\beta = intercept$ .

# Value

lam	The selected tuning parameter based on the estimated ensemble kernel matrix.
intercept	Estimated bias of the model.
alpha	Estimated coefficients of the estimated ensemble kernel matrix.
K	Estimated ensemble kernel matrix.
u_hat	A vector of weights of the kernels in the library.

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### Author(s)

Wenying Deng

### See Also

```
strategy: ensemble
```

### **Examples**

```
##estimation(Y, X1, X2, Kernlist, mode = "loocv", strategy = "erm",
##beta = 1, lambda = exp(seq(-5, 5)))
```

genericFormula

From Vectors to Single Variables

# **Description**

Transform format of predictors from vectors to single variables.

### Usage

```
genericFormula(formula, label_names)
```

### **Arguments**

formula A symbolic description of the model to be fitted.

label\_names A character string indicating all the interior variables included in each predictor.

### Value

generic\_formula

A symbolic description of the model written in single variables format.

length\_main A numeric value indicating the length of main effects.

### Author(s)

Wenying Deng

```
##generic_formula0 <- genericFormula(formula = Y \sim X1 + X2, ##label_names = list(X1 = c("x1", "x2"), X2 = c("x3", "x4")))
```

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infoMat	Computing Information Matrices	

# Description

Compute information matrices based on block matrices.

# Usage

```
infoMat(P0_mat, mat_del = NULL, mat_sigma2 = NULL, mat_tau = NULL)
```

# **Arguments**

P0_mat	Scale projection matrix under REML.
mat_del	Derivative of the scale covariance matrix of Y with respect to delta.
mat_sigma2	Derivative of the scale covariance matrix of Y with respect to sigma2.
mat_tau	Derivative of the scale covariance matrix of Y with respect to tau

### **Details**

This function gives the information value of the interaction strength.

### Value

The computed information value.

# Author(s)

Wenying Deng

### References

Arnab Maity and Xihong Lin. Powerful tests for detecting a gene effect in the presence of possible gene-gene interactions using garrote kernel machines. December 2011.

```
##I0 <- infoMat(P0_mat, mat_del = drV0_del,
##mat_sigma2 = drV0_sigma2, mat_tau = drV0_tau)</pre>
```

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kernelGenerate

Generating A Single Kernel

### **Description**

Generate kernels for the kernel library.

# Usage

kernelGenerate(method = "rbf", Sigma = 0, l = 1, p = 2)

### **Arguments**

method A character string indicating which kernel is to be computed.

Sigma The covariance matrix for neural network kernel.

1 A numeric number indicating the hyperparameter (flexibility) of a specific ker-

nel.

p For polynomial, p is the power; for matern, v = p + 1/2; for rational, alpha = p.

### **Details**

There are seven kinds of kernel available here. For convenience, we define r = |x - x'|.

#### **Gaussian RBF Kernels**

$$k_{SE}(r) = exp\left(-\frac{r^2}{2l^2}\right)$$

### **Matern Kernels**

$$k_{Matern}(r) = \frac{2^{1-\nu}}{\Gamma(\nu)} \left(\frac{\sqrt{2\nu r}}{l}\right)^{\nu} K_{\nu} \left(\frac{\sqrt{2\nu r}}{l}\right)$$

### **Rational Quadratic Kernels**

$$k_{RQ}(r) = \left(1 + \frac{r^2}{2\alpha l^2}\right)^{-\alpha}$$

### **Polynomial Kernels**

$$k(x, x') = (x \cdot x')^p$$

We have intercept kernel when p = 0, and linear kernel when p = 1.

### **Neural Network Kernels**

$$k_{NN}(x,x') = \frac{2}{\pi} sin^{-1} \left( \frac{2\tilde{x}^T \Sigma \tilde{x}'}{\sqrt{(1+2\tilde{x}^T \Sigma \tilde{x})(1+2\tilde{x}'^T \Sigma \tilde{x}')}} \right)$$

### Value

Kerr

A function indicating the generated kernel.

### Author(s)

Wenying Deng

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### References

The MIT Press. Gaussian Processes for Machine Learning, 2006.

### **Examples**

```
##kernelGenerate(method = "rbf", Sigma = 0, 1 = 1, p = 2)
##Kernlist <- NULL
##Kernlist <- c(Kernlist, kernelGenerate('rbf', 1 = .6))
##Kernlist <- c(Kernlist, kernelGenerate('rbf', 1 = 1))
##Kernlist <- c(Kernlist, kernelGenerate('rbf', 1 = 2))</pre>
```

noise Estimate

Estimating Noise

# Description

An implementation of Gaussian processes for estimating noise.

# Usage

```
noiseEstimate(Y, lambda_hat, beta_hat, alpha_hat, K_hat)
```

### **Arguments**

Y Reponses of the dataframe.

beta\_hat Estimated bias of the model.

alpha\_hat Estimated coefficients of the estimated ensemble kernel matrix.

K\_hat Estimated ensemble kernel matrix.

# Value

sigma2\_hat The estimated noise of the fixed effects.

### Author(s)

Wenying Deng

### References

Jeremiah Zhe Liu and Brent Coull. Robust Hypothesis Test for Nonlinear Effect with Gaussian Processes. October 2017.

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### **Examples**

```
##sigma2_hat <- noiseEstimate(y, lam, beta0, alpha0, K_gpr)</pre>
```

scoreStat

Computing Score Test Statistics.

### **Description**

Compute score test statistics.

### Usage

```
scoreStat(n, Y, X12, beta0, sigma2_hat, tau_hat, K_gpr)
```

### **Arguments**

n A numeric number specifying the number of observations.

Y Reponses of the dataframe.

X12 The interaction items of first and second types of factors in the dataframe.

beta0 Estimated bias of the model.

sigma2\_hat The estimated noise of the fixed effects.
tau\_hat The estimated noise of the random effects.

K\_gpr Estimated ensemble kernel matrix.

# **Details**

The test statistic is distributed as a scaled Chi-squared distribution.

### Value

test\_stat The computed test statistic.

### Author(s)

Wenying Deng

#### References

Arnab Maity and Xihong Lin. Powerful tests for detecting a gene effect in the presence of possible gene-gene interactions using garrote kernel machines. December 2011.

```
##scoreStat(n, Y, X12, beta0, sigma2_hat, tau_hat, K_gpr)
```

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testing	Conducting Score Tests for Interaction	

### **Description**

Conduct score tests comparing a fitted model and a more general alternative model.

# Usage

```
testing(formula_int, label_names, Y, X1, X2, Kernlist, mode = "loocv",
   strategy = "erm", beta = 1, test = "boot", lambda = exp(seq(-5, 5)),
   B = 100)
```

# **Arguments**

formula_int	A symbolic description of the model with interaction.
label_names	A character string indicating all the interior variables included in each predictor.
Υ	Reponses of the dataframe.
X1	The first type of factor in the dataframe (could contains several subfactors).
X2	The second type of factor in the dataframe (could contains several subfactors).
Kernlist	The kernel library containing several kernels given by user.
mode	A character string indicating which tuning parameter criteria is to be used.
strategy	A character string indicating which ensemble strategy is to be used.
beta	A numeric value specifying the parameter when strategy = "exp".
test	A character string indicating which test is to be used.
lambda	A numeric string specifying the range of noise to be chosen. The lower limit of lambda must be above 0.
В	A numeric value indicating times of resampling when test = "boot".

### **Details**

There are two tests available here:

### **Asymptotic Test**

This is based on the classical variance component test to construct a testing procedure for the hypothesis about Gaussian process function.

### **Bootstrap Test**

When it comes to small sample size, we can use bootstrap test instead, which can give valid tests with moderate sample sizes and requires similar computational effort to a permutation test.

### Value

pvalue p-value of the test.

### Author(s)

Wenying Deng

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#### References

Xihong Lin. Variance component testing in generalised linear models with random effects. June 1997.

Arnab Maity and Xihong Lin. Powerful tests for detecting a gene effect in the presence of possible gene-gene interactions using garrote kernel machines. December 2011.

Petra Bu z'kova, Thomas Lumley, and Kenneth Rice. Permutation and parametric bootstrap tests for gene-gene and gene-environment interactions. January 2011.

### See Also

method: kernelGenerate

mode: tuning
strategy: ensemble

### **Examples**

```
##testing(formula_int = Y \sim X1 * X2,
##label_names = list(X1 = c("x1", "x2"), X2 = c("x3", "x4")),
##Y, X1, X2, Kernlist, mode = "loocv", strategy = "erm",
##beta = 1, test = "boot", lambda = exp(seq(-5, 5)), B = 100)
```

tuning

Calculating Tuning Parameters

### **Description**

Calculate tuning parameters based on given criteria.

### Usage

```
tuning(Y, K_mat, mode, lambda)
```

### **Arguments**

Y Reponses of the dataframe.

K\_mat Estimated ensemble kernel matrix.

mode A character string indicating which tuning parameter criteria is to be used.

lambda A numeric string specifying the range of noise to be chosen. The lower limit of

lambda must be above 0.

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#### **Details**

There are four tuning parameter selections here:

#### leave-one-out Cross Validation

$$\lambda_{n-CV} = \lambda \in \Lambda argmin \left\{ log \ y^{\star T} [I - diag(A_{\lambda}) - \frac{1}{n} I]^{-1} (I - A_{\lambda})^2 [I - diag(A_{\lambda}) - \frac{1}{n} I]^{-1} y^{\star} \right\}$$

#### **Akaike Information Criteria**

$$\lambda_{AICc} = \lambda \in \Lambda argmin \Big\{ log \ y^{\star T} (I - A_{\lambda})^2 y^{\star} + \frac{2[tr(A_{\lambda}) + 2]}{n - tr(A_{\lambda}) - 3} \Big\}$$

### **Generalized Cross Validation**

$$\lambda_{GCVc} = \lambda \in \Lambda argmin \Big\{ log \ y^{\star T} (I - A_{\lambda})^2 y^{\star} - 2log [1 - \frac{tr(A_{\lambda})}{n} - \frac{2}{n}]_+ \Big\}$$

### Generalized Maximum Profile Marginal Likelihood

$$\lambda_{GMPML} = \lambda \in \Lambda argmin \left\{ log \ y^{\star T} (I - A_{\lambda}) y^{\star} - \frac{1}{n-1} log \ | \ I - A_{\lambda} \ | \ \right\}$$

#### Value

lambda0 The estimated tuning parameter.

#### Author(s)

Wenying Deng

# References

Philip S. Boonstra, Bhramar Mukherjee, and Jeremy M. G. Taylor. A Small-Sample Choice of the Tuning Parameter in Ridge Regression. July 2015.

Trevor Hastie, Robert Tibshirani, and Jerome Friedman. The Elements of Statistical Learning: Data Mining, Inference, and Prediction, Second Edition. Springer Series in Statistics. Springer-Verlag, New York, 2 edition, 2009.

Hirotogu Akaike. Information Theory and an Extension of the Maximum Likelihood Princi- ple. In Selected Papers of Hirotugu Akaike, Springer Series in Statistics, pages 199–213. Springer, New York, NY, 1998.

Clifford M. Hurvich and Chih-Ling Tsai. Regression and time series model selection in small samples. June 1989.

Hurvich Clifford M., Simonoff Jeffrey S., and Tsai Chih-Ling. Smoothing parameter selection in nonparametric regression using an improved Akaike information criterion. January 2002.

```
##tuning(Y, K_mat = K, mode = "loocv", lambda = exp(seq(-5, 5)))
```