

# Package: SALES (via r-universe)

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**Title** The (Adaptive) Elastic Net and Lasso Penalized Sparse Asymmetric Least Squares (SALES) and Coupled Sparse Asymmetric Least Squares (COSALES) using Coordinate Descent and Proximal Gradient Algorithms

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**Imports** grDevices, graphics, stats, methods, Matrix

**Description** A coordinate descent algorithm for computing the solution paths of the sparse and coupled sparse asymmetric least squares, including the (adaptive) elastic net and Lasso penalized SALES and COSALES regressions.

**License** GPL (>= 2)

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coef	<i>Extract Model Coefficients</i>
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### Description

coef is a generic function which extracts model coefficients from objects returned by modeling functions. coefficients is an *alias* for it.

### Usage

```
coef(object, ...)
```

### Arguments

object	an object for which the extraction of model coefficients is meaningful.
...	other arguments.

### Value

Coefficients extracted from the model object object.

### See Also

[coef.ernet](#), [coef.cpernet](#)

---

`coef.cpernet`*Get coefficients from a cpernet object*

---

**Description**

Computes the coefficients or returns a list of the indices of the nonzero coefficients at the requested values for `lambda` from a fitted `cpernet` object.

**Usage**

```
## S3 method for class 'cpernet'  
coef(object, s = NULL, type = c("coefficients", "nonzero"), ...)
```

**Arguments**

<code>object</code>	fitted <a href="#">cpernet</a> model object.
<code>s</code>	value(s) of the penalty parameter <code>lambda</code> at which predictions are to be made. Default is the entire sequence used to create the model.
<code>type</code>	type "coefficients" computes coefficients at the requested values for <code>s</code> . Type "nonzero" returns a list of the indices of nonzero coefficients for each value of <code>s</code> . Default is "coefficients".
<code>...</code>	not used. Other arguments to <code>predict</code> .

**Details**

`s` is the new vector at which predictions are requested. If `s` is not in the `lambda` sequence used for fitting the model, the `coef` function will use linear interpolation to make predictions. The new values are interpolated using a fraction of coefficients from both left and right `lambda` indices.

**Value**

The object returned depends on `type`.

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**See Also**

[cpernet](#), [predict.cpernet](#), [print.cpernet](#), [plot.cpernet](#)

**Examples**

```

set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.30
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
w <- 2.0
lambda2 <- 1
m2 <- cpernet(y = y, x = x, w = w, tau = tau, eps = 1e-8,
              pf.mean = pf, pf.scale = pf2, intercept = TRUE,
              standardize = FALSE, lambda2 = lambda2)

mean.coef <- as.vector(coef(m2, s = m2$lambda[50])[[1]])
scale.coef <- as.vector(coef(m2, s = m2$lambda[50])[[2]])

```

coef.cv.cpernet

*Get coefficients from a cv.cpernet object***Description**

This function gets coefficients from a cross-validated cpernet model, using the fitted cv.cpernet object, and the optimal value chosen for lambda.

**Usage**

```

## S3 method for class 'cv.cpernet'
coef(object, s = c("lambda.1se", "lambda.min"), ...)

```

**Arguments**

object	fitted <a href="#">cv.cpernet</a> object.
s	value(s) of the penalty parameter lambda at which predictions are required. Default is the value s="lambda.1se" stored on the CV object, it is the largest value of lambda such that error is within 1 standard error of the minimum. Alternatively s="lambda.min" can be used, it is the optimal value of lambda that gives minimum cross validation error cvm. If s is numeric, it is taken as the value(s) of lambda to be used.
...	not used. Other arguments to predict.

**Details**

This function makes it easier to use the results of cross-validation to get coefficients or make coefficient predictions.

**Value**

The object returned depends the ... argument which is passed on to the `predict` method for `cpernet` objects.

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**See Also**

[cv.cpernet](#), [predict.cv.cpernet](#)

**Examples**

```
set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.30
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
w <- 2.0
lambda2 <- 1
m2.cv <- cv.cpernet(y = y, x = x, w = w, tau = tau, eps = 1e-8,
                   pf.mean = pf, pf.scale = pf2,
                   standardize = FALSE, lambda2 = lambda2)
as.vector(coef(m2.cv, s = "lambda.min")$beta)
as.vector(coef(m2.cv, s = "lambda.min")$theta)
```

---

coef.cv.ernet

*Get coefficients from a cv.ernet object*

---

**Description**

This function gets coefficients from a cross-validated ernet model, using the fitted `cv.ernet` object, and the optimal value chosen for `lambda`.

**Usage**

```
## S3 method for class 'cv.ernet'
coef(object, s = c("lambda.1se", "lambda.min"), ...)
```

**Arguments**

object	fitted <a href="#">cv.ernet</a> object.
s	value(s) of the penalty parameter lambda at which predictions are required. Default is the value <code>s="lambda.1se"</code> stored on the CV object, it is the largest value of lambda such that error is within 1 standard error of the minimum. Alternatively <code>s="lambda.min"</code> can be used, it is the optimal value of lambda that gives minimum cross validation error <code>cvm</code> . If s is numeric, it is taken as the value(s) of lambda to be used.
...	not used. Other arguments to predict.

**Details**

This function makes it easier to use the results of cross-validation to get coefficients or make coefficient predictions.

**Value**

The object returned depends the ... argument which is passed on to the [predict](#) method for [ernet](#) objects.

**Author(s)**

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**See Also**

[cv.ernet](#), [predict.cv.ernet](#)

**Examples**

```
set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.90
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
lambda2 <- 1
m1.cv <- cv.ernet(y = y, x = x, tau = tau, eps = 1e-8, pf = pf,
                 pf2 = pf2, standardize = FALSE, intercept = FALSE,
                 lambda2 = lambda2)
as.vector(coef(m1.cv, s = "lambda.min"))
```

---

`coef.ernet`*Get coefficients from an ernet object*

---

### Description

Computes the coefficients or returns a list of the indices of the nonzero coefficients at the requested values for `lambda` from a fitted ernet object.

### Usage

```
## S3 method for class 'ernet'  
coef(object, s = NULL, type = c("coefficients", "nonzero"), ...)
```

### Arguments

<code>object</code>	fitted <a href="#">ernet</a> model object.
<code>s</code>	value(s) of the penalty parameter <code>lambda</code> at which predictions are to be made. Default is the entire sequence used to create the model.
<code>type</code>	type "coefficients" computes coefficients at the requested values for <code>s</code> . Type "nonzero" returns a list of the indices of nonzero coefficients for each value of <code>s</code> . Default is "coefficients".
<code>...</code>	not used. Other arguments to <code>predict</code> .

### Details

`s` is the new vector at which predictions are requested. If `s` is not in the `lambda` sequence used for fitting the model, the `coef` function will use linear interpolation to make predictions. The new values are interpolated using a fraction of coefficients from both left and right `lambda` indices.

### Value

The object returned depends on `type`.

### Author(s)

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### See Also

[ernet](#), [predict.ernet](#), [print.ernet](#), [plot.ernet](#)

**Examples**

```

set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.90
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
lambda2 <- 1
m1 <- ernet(y = y, x = x, tau = tau, eps = 1e-8, pf = pf,
            pf2 = pf2, standardize = FALSE, intercept = FALSE,
            lambda2 = lambda2)
as.vector(coef(m1, s = m1$lambda[5]))

```

---

cpernet

*Regularization paths for the coupled sparse asymmetric least squares (COSALES) regression (or the coupled sparse expectile regression)*

---

**Description**

Fits regularization paths for coupled sparse asymmetric least squares regression at a sequence of regularization parameters.

**Usage**

```

cpernet(
  x,
  y,
  w = 1,
  nlambda = 100L,
  method = "cper",
  lambda.factor = ifelse(2 * nobs < nvars, 0.01, 1e-04),
  lambda = NULL,
  lambda2 = 0,
  pf.mean = rep(1, nvars),
  pf2.mean = rep(1, nvars),
  pf.scale = rep(1, nvars),
  pf2.scale = rep(1, nvars),
  exclude,
  dfmax = nvars + 1,
  pmax = min(dfmax * 1.2, nvars),
  standardize = TRUE,
  intercept = TRUE,
  eps = 1e-08,
  maxit = 100000L,
  tau = 0.8
)

```



**Arguments**

x	matrix of predictors, of dimension (nobs * nvars); each row is an observation.
y	response variable.
w	weight applied to the asymmetric squared error loss of the mean part. See details. Default is 1.0.
nlambda	the number of lambda values (default is 100).
method	a character string specifying the loss function to use. Only cper is available now.
lambda.factor	The factor for getting the minimal lambda in the lambda sequence, where we set $\min(\lambda) = \lambda.factor * \max(\lambda)$ with $\max(\lambda)$ being the smallest value of lambda that penalizes all coefficients to zero. The default value depends on the relationship between $N$ (the number of observations) and $p$ (the number of predictors). If $N < p$ , the default is 0.01. If $N > p$ , the default is 0.0001, closer to zero. A very small value of lambda.factor will lead to a saturated fit. The argument takes no effect if there is a user-supplied lambda sequence.
lambda	a user-supplied lambda sequence. Typically, by leaving this option unspecified users can have the program compute its own lambda sequence based on nlambda and lambda.factor. It is better to supply, if necessary, a decreasing sequence of lambda values than a single (small) value. The program will ensure that the user-supplied lambda sequence is sorted in decreasing order.
lambda2	regularization parameter lambda2 for the quadratic penalty of the coefficients. Default is 0, meaning no L2 penalization.
pf.mean, pf.scale	L1 penalty factor of length $p$ used for adaptive LASSO or adaptive elastic net. Separate L1 penalty weights can be applied to each mean or scale coefficient to allow different L1 shrinkage. Can be 0 for some variables, which imposes no shrinkage and results in that variable being always included in the model. Default is 1 for all variables (and implicitly infinity for variables listed in exclude).
pf2.mean, pf2.scale	L2 penalty factor of length $p$ used for adaptive elastic net. Separate L2 penalty weights can be applied to each mean or scale coefficient to allow different L2 shrinkage. Can be 0 for some variables, which imposes no shrinkage. Default is 1 for all variables.
exclude	indices of variables to be excluded from the model. Default is none. Equivalent to an infinite penalty factor.
dfmax	limit the maximum number of variables in the model. Useful for very large $p$ , if a partial path is desired. Default is $p + 1$ .
pmax	limit the maximum number of variables ever to be nonzero. For example once $\beta$ enters the model, no matter how many times it exits or re-enters the model through the path, it will be counted only once. Default is $\min(dfmax * 1.2, p)$ .
standardize	logical flag for variable standardization, prior to fitting the model sequence. The coefficients are always returned to the original scale. Default is TRUE.
intercept	Should intercept(s) be fitted (default=TRUE) or set to zero (FALSE).

eps	convergence threshold for coordinate descent. Each inner coordinate descent loop continues until the maximum change in any coefficient is less than eps. Defaults value is 1e-8.
maxit	maximum number of outer-loop iterations allowed at fixed lambda values. Default is 1e7. If the algorithm does not converge, consider increasing maxit.
tau	the parameter tau in the coupled ALS regression model. The value must be in (0,1) and cannot be 0.5. Default is 0.8.

### Details

Note that the objective function in cpernet is

$$w * 1' \Psi(y - X\beta, 0.5) / N + 1' \Psi(y - X\beta - X\theta, \tau) / N + \lambda_1 * \|\beta\|_1 + 0.5\lambda_2 \|\beta\|_2^2 + \mu_1 * \|\theta\| + 0.5\mu_2 \|\theta\|_2^2,$$

where  $\Psi(u, \tau) = |\tau - I(u < 0)| * u^2$  denotes the asymmetric squared error loss and the penalty is a combination of L1 and L2 terms for both the mean and scale coefficients.

For faster computation, if the algorithm is not converging or running slow, consider increasing eps, decreasing nlambda, or increasing lambda.factor before increasing maxit.

### Value

An object with S3 class `cpernet`.

call	the call that produced this object.
b0, t0	intercept sequences both of length length(lambda) for the mean and scale respectively.
beta, theta	p*length(lambda) matrices of coefficients for the mean and scale respectively, stored as sparse matrices (dgCMatrix class, the standard class for sparse numeric matrices in the Matrix package). To convert them into normal R matrices, use as.matrix().
lambda	the actual sequence of lambda values used
df.beta, df.theta	the number of nonzero mean and scale coefficients respectively for each value of lambda.
dim	dimensions of coefficient matrices.
npasses	total number of iterations summed over all lambda values.
jerr	error flag, for warnings and errors, 0 if no error.

### Author(s)

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### References

Gu, Y., and Zou, H. (2016). "High-dimensional generalizations of asymmetric least squares regression and their applications." *The Annals of Statistics*, 44(6), 2661–2694.

**See Also**

[plot.cpernet](#), [coef.cpernet](#), [predict.cpernet](#), [print.cpernet](#)

**Examples**

```
set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.30
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
w <- 2.0
lambda2 <- 1
m2 <- cpernet(y = y, x = x, w = w, tau = tau, eps = 1e-8,
             pf.mean = pf, pf.scale = pf2,
             standardize = FALSE, lambda2 = lambda2)
```

---

cv.cpernet

*Cross-validation for cpernet*

---

**Description**

Does k-fold cross-validation for cpernet, produces a plot, and returns a value for lambda. This function is based on the cv function from the glmnet package.

**Usage**

```
cv.cpernet(
  x,
  y,
  w = 1,
  lambda = NULL,
  pred.loss = "loss",
  nfolds = 5,
  foldid,
  tau = 0.8,
  ...
)
```

**Arguments**

x                    x matrix as in [cpernet](#).

y                    response variable y as in [cpernet](#).

w                    weight applied to the asymmetric squared error loss of the mean part. Default is 1.0.

lambda	optional user-supplied lambda sequence; default is NULL, and <code>cpernet</code> chooses its own sequence.
pred.loss	loss function used to calculate cross-validation error. The only option now is "loss", which is the asymmetric squared error loss (ASEL).
nfolds	number of folds. Default value is 5. Although <code>nfolds</code> can be as large as the sample size (leave-one-out CV), it is not recommended for large datasets. Smallest value allowed is 3.
foldid	an optional vector of values between 1 and <code>nfolds</code> , identifying what fold each observation is in. If supplied, <code>nfolds</code> will be suppressed.
tau	the asymmetry coefficient $\tau$ used in the asymmetric squared error loss.
...	other arguments that can be passed to <code>cpernet</code> .

### Details

The function runs `cpernet` `nfolds+1` times. The first gets the lambda sequence, and the remainder fits the model with each of the folds removed. The average error and standard deviation over the folds are computed.

### Value

an object of class `cv.cpernet` is returned, which is a list with the ingredients of the cross-validation fit.

lambda	the values of lambda used in the fits.
cvm	the mean cross-validated error - a vector of length <code>length(lambda)</code> .
cvsd	estimate of standard error of <code>cvm</code> .
cvupper	upper curve = <code>cvm+cvsd</code> .
cvlower	lower curve = <code>cvm-cvsd</code> .
nzero	a list of two components, each representing the number of non-zero coefficients at each lambda in the mean and scale part.
name	a text string indicating type of measure (for plotting purposes).
<code>cpernet.fit</code>	a fitted <code>cpernet</code> object for the full data.
lambda.min	The optimal value of lambda that gives minimum cross validation error <code>cvm</code> .
lambda.1se	The largest value of lambda such that error is within 1 standard error of the minimum.

### Author(s)

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### See Also

[cpernet](#)

**Examples**

```

set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.30
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
w <- 2.0
lambda2 <- 1
m2.cv <- cv.cpernet(y = y, x = x, w = w, tau = tau, eps = 1e-8,
                    pf.mean = pf, pf.scale = pf2,
                    standardize = FALSE, lambda2 = lambda2)

```

---

cv.ernet

*Cross-validation for ernet*


---

**Description**

Does k-fold cross-validation for ernet, produces a plot, and returns a value for lambda. This function is based on the cv function from the glmnet package.

**Usage**

```

cv.ernet(
  x,
  y,
  lambda = NULL,
  pred.loss = "loss",
  nfold = 5,
  foldid,
  tau = 0.5,
  ...
)

```

**Arguments**

x	x matrix as in <a href="#">ernet</a> .
y	response variable y as in <a href="#">ernet</a> .
lambda	optional user-supplied lambda sequence; default is NULL, and <a href="#">ernet</a> chooses its own sequence.
pred.loss	loss function used to calculate cross-validation error. The only option now is "loss", which is the asymmetric squared error loss (ASEL).

<code>nfolds</code>	number of folds. Default value is 5. Although <code>nfolds</code> can be as large as the sample size (leave-one-out CV), it is not recommended for large datasets. Smallest value allowed is 3.
<code>foldid</code>	an optional vector of values between 1 and <code>nfolds</code> , identifying what fold each observation is in. If supplied, <code>nfolds</code> will be suppressed.
<code>tau</code>	the asymmetry coefficient $\tau$ used in the asymmetric squared error loss.
<code>...</code>	other arguments that can be passed to <code>ernet</code> .

### Details

The function runs `ernet` `nfolds+1` times; the first to get the `lambda` sequence, and the remainder to compute the fit with each of the folds removed. The average error and standard deviation over the folds are computed.

### Value

an object of class `cv.ernet` is returned, which is a list with the ingredients of the cross-validation fit.

<code>lambda</code>	the values of <code>lambda</code> used in the fits.
<code>cvm</code>	the mean cross-validated error - a vector of length <code>length(lambda)</code> .
<code>cvsd</code>	estimate of standard error of <code>cvm</code> .
<code>cvupper</code>	upper curve = <code>cvm+cvsd</code> .
<code>cvlower</code>	lower curve = <code>cvm-cvsd</code> .
<code>nzero</code>	number of non-zero coefficients at each <code>lambda</code> .
<code>name</code>	a text string indicating type of measure (for plotting purposes).
<code>ernet.fit</code>	a fitted <code>ernet</code> object for the full data.
<code>lambda.min</code>	The optimal value of <code>lambda</code> that gives minimum cross validation error <code>cvm</code> .
<code>lambda.1se</code>	The largest value of <code>lambda</code> such that error is within 1 standard error of the minimum.

### Author(s)

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### See Also

[ernet](#)

**Examples**

```

set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.90
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
lambda2 <- 1
m1.cv <- cv.ernet(y = y, x = x, tau = tau, eps = 1e-8, pf = pf,
                 pf2 = pf2, standardize = FALSE, intercept = FALSE,
                 lambda2 = lambda2)

```

---

ermet	<i>Regularization paths for the sparse asymmetric least squares (SALES) regression (or the sparse expectile regression)</i>
-------	---

---

**Description**

Fits regularization paths for the Lasso or elastic net penalized asymmetric least squares regression at a sequence of regularization parameters.

**Usage**

```

ernet(
  x,
  y,
  nlambda = 100L,
  method = "er",
  lambda.factor = ifelse(nobs < nvars, 0.01, 1e-04),
  lambda = NULL,
  lambda2 = 0,
  pf = rep(1, nvars),
  pf2 = rep(1, nvars),
  exclude,
  dfmax = nvars + 1,
  pmax = min(dfmax * 1.2, nvars),
  standardize = TRUE,
  intercept = TRUE,
  eps = 1e-08,
  maxit = 100000L,
  tau = 0.5
)

```

**Arguments**

<code>x</code>	matrix of predictors, of dimension (nobs * nvars); each row is an observation.
<code>y</code>	response variable.
<code>nlambda</code>	the number of lambda values (default is 100).
<code>method</code>	a character string specifying the loss function to use. only er is available now.
<code>lambda.factor</code>	The factor for getting the minimal lambda in the lambda sequence, where we set $\min(\text{lambda}) = \text{lambda.factor} * \max(\text{lambda})$ with $\max(\text{lambda})$ being the smallest value of lambda that penalizes all coefficients to zero. The default depends on the relationship between $N$ (the number of rows in the matrix of predictors) and $p$ (the number of predictors). If $N < p$ , the default is 0.01. If $N > p$ , the default is 0.0001, closer to zero. A very small value of lambda.factor will lead to a saturated fit. It takes no effect if there is a user-supplied lambda sequence.
<code>lambda</code>	a user-supplied lambda sequence. Typically, by leaving this option unspecified users can have the program compute its own lambda sequence based on nlambda and lambda.factor. It is better to supply, if necessary, a decreasing sequence of lambda values than a single (small) value. The program will ensure that the user-supplied lambda sequence is sorted in decreasing order before fitting the model.
<code>lambda2</code>	regularization parameter lambda2 for the quadratic penalty of the coefficients.
<code>pf</code>	L1 penalty factor of length $p$ used for the adaptive LASSO or adaptive elastic net. Separate L1 penalty weights can be applied to each coefficient to allow different L1 shrinkage. Can be 0 for some variables, which imposes no shrinkage, and results in that variable always be included in the model. Default is 1 for all variables (and implicitly infinity for variables listed in exclude).
<code>pf2</code>	L2 penalty factor of length $p$ used for adaptive elastic net. Separate L2 penalty weights can be applied to each coefficient to allow different L2 shrinkage. Can be 0 for some variables, which imposes no shrinkage. Default is 1 for all variables.
<code>exclude</code>	indices of variables to be excluded from the model. Default is none. Equivalent to an infinite penalty factor.
<code>dfmax</code>	the maximum number of variables allowed in the model. Useful for very large $p$ when a partial path is desired. Default is $p + 1$ .
<code>pmax</code>	the maximum number of coefficients allowed ever to be nonzero. For example once $\beta$ enters the model, no matter how many times it exits or re-enters the model through the path, it will be counted only once. Default is $\min(\text{dfmax} * 1.2, p)$ .
<code>standardize</code>	logical flag for variable standardization, prior to fitting the model sequence. The coefficients are always returned to the original scale. Default is TRUE.
<code>intercept</code>	Should intercept(s) be fitted (default is TRUE) or set to zero (FALSE)?
<code>eps</code>	convergence threshold for coordinate descent. Each inner coordinate descent loop continues until the maximum change in any coefficient is less than eps. Defaults value is $1e-8$ .



<code>maxit</code>	maximum number of outer-loop iterations allowed at fixed <code>lambda</code> values. Default is <code>1e7</code> . If the algorithm does not converge, consider increasing <code>maxit</code> .
<code>tau</code>	the parameter $\tau$ in the ALS regression model. The value must be in $(0,1)$ . Default is <code>0.5</code> .

### Details

Note that the objective function in `ernet` is

$$1' \Psi_{\tau}(y - X\beta)/N + \lambda_1 * \|\beta\|_1 + 0.5\lambda_2 * \|\beta\|_2^2,$$

where  $\Psi_{\tau}$  denotes the asymmetric squared error loss and the penalty is a combination of weighted L1 and L2 terms.

For faster computation, if the algorithm is not converging or running slow, consider increasing `eps`, decreasing `nlambda`, or increasing `lambda.factor` before increasing `maxit`.

### Value

An object with S3 class `ernet`.

<code>call</code>	the call that produced this object
<code>b0</code>	intercept sequence of length <code>length(lambda)</code>
<code>beta</code>	a <code>p*length(lambda)</code> matrix of coefficients, stored as a sparse matrix ( <code>dgCMatrix</code> class, the standard class for sparse numeric matrices in the <code>Matrix</code> package.). To convert it into normal type matrix use <code>as.matrix()</code> .
<code>lambda</code>	the actual sequence of <code>lambda</code> values used
<code>df</code>	the number of nonzero coefficients for each value of <code>lambda</code> .
<code>dim</code>	dimension of coefficient matrix
<code>npasses</code>	total number of iterations summed over all <code>lambda</code> values
<code>jerr</code>	error flag, for warnings and errors, 0 if no error.

### Author(s)

Yuwen Gu and Hui Zou

Maintainer: Yuwen Gu <yuwen.gu@uconn.edu>

### References

Gu, Y., and Zou, H. (2016). "High-dimensional generalizations of asymmetric least squares regression and their applications." *The Annals of Statistics*, 44(6), 2661–2694.

### See Also

`plot.ernet`, `coef.ernet`, `predict.ernet`, `print.ernet`

**Examples**

```

set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.90
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
lambda2 <- 1
m1 <- ernet(y = y, x = x, tau = tau, eps = 1e-8, pf = pf,
            pf2 = pf2, standardize = FALSE, intercept = FALSE,
            lambda2 = lambda2)

```

---

plot.cpernet

*Plot coefficients from a cpernet object*


---

**Description**

Produces a coefficient profile plot of the coefficient paths for a fitted cpernet object. This function is modified based on the plot method in the glmnet package.

**Usage**

```

## S3 method for class 'cpernet'
plot(x, xvar = c("norm", "lambda"), color = FALSE, label = FALSE, ...)

```

**Arguments**

x	fitted <a href="#">cpernet</a> model
xvar	what is on the x-axis. "norm" plots against the L1-norm of the coefficients, "lambda" against the log-lambda sequence.
color	if TRUE, plot the curves with rainbow colors. Otherwise, plot the curves with gray colors. Default is FALSE.
label	if TRUE, label the curves with variable sequence numbers. Otherwise, do not put labels. Default is FALSE.
...	other graphical parameters to plot.

**Details**

Two coefficient profile plots are produced, one for the mean coefficients and the other for the scale coefficients.

**Author(s)**

Yuwen Gu and Hui Zou

Maintainer: Yuwen Gu &lt;yuwen.gu@uconn.edu&gt;

**See Also**[plot.cv.cpernet](#)**Examples**

```

set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.30
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
w <- 2.0
lambda2 <- 1
m2 <- cpernet(y = y, x = x, w = w, tau = tau, eps = 1e-8,
              pf.mean = pf, pf.scale = pf2, intercept = TRUE,
              standardize = FALSE, lambda2 = lambda2)
plot(m2)

```

plot.cv.cpernet

*Plot the cross-validated curve produced by cv.cpernet***Description**

Plots the cross-validated curve, and upper and lower standard deviation curves, as a function of the lambda values used. This function is modified based on the plot.cv.glmnet function from the glmnet package.

**Usage**

```

## S3 method for class 'cv.cpernet'
plot(x, sign.lambda = 1, ...)

```

**Arguments**

x	fitted <a href="#">cv.cpernet</a> object
sign.lambda	either plot against $\log(\lambda)$ (default) or its negative if sign.lambda=-1.
...	other graphical parameters to plot

**Details**

A plot is produced.

**Author(s)**

Yuwen Gu and Hui Zou

Maintainer: Yuwen Gu <yuwen.gu@uconn.edu>

**See Also**

[plot.cpernet](#)

**Examples**

```
set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.30
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
w <- 2.0
lambda2 <- 1
m2.cv <- cv.cpernet(y = y, x = x, w = w, tau = tau, eps = 1e-8,
                  pf.mean = pf, pf.scale = pf2,
                  standardize = FALSE, lambda2 = lambda2)
plot(m2.cv)
```

---

plot.cv.ernet

*Plot the cross-validated curve produced by cv.ernet*

---

**Description**

Plots the cross-validated curve, and upper and lower standard deviation curves, as a function of the lambda values used. This function is modified based on the plot.cv.glmnet function from the glmnet package.

**Usage**

```
## S3 method for class 'cv.ernet'
plot(x, sign.lambda = 1, ...)
```

**Arguments**

`x` fitted `cv.ernet` object  
`sign.lambda` either plot against  $\log(\lambda)$  (default) or its negative if `sign.lambda=-1`.  
`...` other graphical parameters to plot

**Details**

A plot is produced.

**Author(s)**

Yuwen Gu and Hui Zou

Maintainer: Yuwen Gu <yuwen.gu@uconn.edu>

**See Also**

[plot.ernet](#)

**Examples**

```
set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.90
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
lambda2 <- 1
m1.cv <- cv.ernet(y = y, x = x, tau = tau, eps = 1e-8, pf = pf,
                 pf2 = pf2, standardize = FALSE, intercept = FALSE,
                 lambda2 = lambda2)
plot(m1.cv)
```

---

plot.ernet

*Plot coefficients from an ernet object*

---

**Description**

Produces a coefficient profile plot of the coefficient paths for a fitted ernet object. This function is modified based on the `plot` method in the `glmnet` package.

**Usage**

```
## S3 method for class 'ernet'
plot(x, xvar = c("norm", "lambda"), color = FALSE, label = FALSE, ...)
```

**Arguments**

x	fitted <a href="#">ernet</a> model
xvar	what is on the x-axis. "norm" plots against the L1-norm of the coefficients, "lambda" against the log-lambda sequence.
color	if TRUE, plot the curves with rainbow colors. Otherwise, plot the curves with gray colors. Default is FALSE.
label	if TRUE, label the curves with variable sequence numbers. Otherwise, do not put labels. Default is FALSE.
...	other graphical parameters to plot.

**Details**

A coefficient profile plot is produced.

**Author(s)**

Yuwen Gu and Hui Zou

Maintainer: Yuwen Gu <yuwen.gu@uconn.edu>

**See Also**

[plot.cv.ernet](#)

**Examples**

```
set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.90
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
lambda2 <- 1
m1 <- ernet(y = y, x = x, tau = tau, eps = 1e-8, pf = pf,
           pf2 = pf2, standardize = FALSE, intercept = FALSE,
           lambda2 = lambda2)
plot(m1)
```

---

predict	<i>Model predictions</i>
---------	--------------------------

---

### Description

predict is a generic function for predictions from the results of various model fitting functions. The function invokes particular *methods* which depend on the `class` of the first argument.

### Usage

```
predict(object, ...)
```

### Arguments

object	a model object for which prediction is desired.
...	additional arguments affecting the predictions produced.

### Value

The form of the value returned by predict depends on the class of its argument. See the documentation of the particular methods for details of what is produced by that method.

### See Also

[predict.ernet](#), [predict.cpernet](#).

---

predict.cpernet	<i>Make predictions from a cpernet object</i>
-----------------	---

---

### Description

Similar to other predict methods, this function predicts fitted values from a cpernet object.

### Usage

```
## S3 method for class 'cpernet'  
predict(object, newx, s = NULL, type = "response", ...)
```

**Arguments**

object	fitted <a href="#">cpernet</a> model object.
newx	matrix of new values for x at which predictions are to be made. NOTE: newx must be a matrix, predict function does not accept a vector or other formats of newx.
s	value(s) of the penalty parameter lambda at which predictions are to be made. Default is the entire sequence used to create the model.
type	type of prediction required. Only response is available. Gives predicted response for regression problems.
...	Not used. Other arguments to predict.

**Details**

s is the new vector at which predictions are to be made. If s is not in the lambda sequence used for fitting the model, the predict function will use linear interpolation to make predictions. The new values are interpolated using a fraction of predicted values from both left and right lambda indices.

**Value**

The object returned depends on type.

**Author(s)**

Yuwen Gu and Hui Zou

Maintainer: Yuwen Gu <yuwen.gu@uconn.edu>

**See Also**

[cpernet](#), [coef.cpernet](#), [plot.cpernet](#), [print.cpernet](#)

**Examples**

```
set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.30
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
w <- 2.0
lambda2 <- 1
m2 <- cpernet(y = y, x = x, w = w, tau = tau, eps = 1e-8,
             pf.mean = pf, pf.scale = pf2,
             standardize = FALSE, lambda2 = lambda2)
predict(m2, newx = x, s = m2$lambda[50])
```



---

predict.cv.cpernet      *Make predictions from a cv.cpernet object*

---

### Description

This function makes predictions from a cross-validated cpernet model, using the fitted cv.cpernet object, and the optimal value chosen for lambda.

### Usage

```
## S3 method for class 'cv.cpernet'  
predict(object, newx, s = c("lambda.1se", "lambda.min"), ...)
```

### Arguments

object	fitted <a href="#">cv.cpernet</a> object.
newx	matrix of new values for x at which predictions are to be made. Must be a matrix. See documentation for <a href="#">predict.cpernet</a> .
s	value(s) of the penalty parameter lambda at which predictions are to be made. Default is the value <code>s = "lambda.1se"</code> stored on the CV object. Alternatively <code>s = "lambda.min"</code> can be used. If s is numeric, it is taken as the value(s) of lambda to be used.
...	not used. Other arguments to <a href="#">predict</a> .

### Details

This function makes it easier to use the results of cross-validation to make a prediction.

### Value

The object returned depends the ... argument which is passed on to the [predict](#) method for [cpernet](#) objects.

### Author(s)

Yuwen Gu and Hui Zou

Maintainer: Yuwen Gu <[yuwen.gu@uconn.edu](mailto:yuwen.gu@uconn.edu)>

### See Also

[cv.cpernet](#), [coef.cv.cpernet](#), [plot.cv.cpernet](#)

**Examples**

```

set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.30
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
w <- 2.0
lambda2 <- 1
m2.cv <- cv.cpernet(y = y, x = x, w = w, tau = tau, eps = 1e-8,
                   pf.mean = pf, pf.scale = pf2,
                   standardize = FALSE, lambda2 = lambda2)
as.vector(predict(m2.cv, newx = x, s = "lambda.min"))

```

---

predict.cv.ernet

*Make predictions from a cv.ernet object*

---

**Description**

This function makes predictions from a cross-validated ernet model, using the fitted `cv.ernet` object, and the optimal value chosen for `lambda`.

**Usage**

```

## S3 method for class 'cv.ernet'
predict(object, newx, s = c("lambda.1se", "lambda.min"), ...)

```

**Arguments**

<code>object</code>	fitted <code>cv.ernet</code> object.
<code>newx</code>	matrix of new values for <code>x</code> at which predictions are to be made. Must be a matrix. See documentation for <code>predict.ernet</code> .
<code>s</code>	value(s) of the penalty parameter <code>lambda</code> at which predictions are to be made. Default is the value <code>s = "lambda.1se"</code> stored on the CV object. Alternatively <code>s = "lambda.min"</code> can be used. If <code>s</code> is numeric, it is taken as the value(s) of <code>lambda</code> to be used.
<code>...</code>	not used. Other arguments to <code>predict</code> .

**Details**

This function makes it easier to use the results of cross-validation to make a prediction.

**Value**

The object returned depends the ... argument which is passed on to the `predict` method for `ernet` objects.

**Author(s)**

Yuwen Gu and Hui Zou

Maintainer: Yuwen Gu <yuwen.gu@uconn.edu>

**See Also**

[cv.ernet](#), [coef.cv.ernet](#), [plot.cv.ernet](#)

**Examples**

```
set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.90
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
lambda2 <- 1
m1.cv <- cv.ernet(y = y, x = x, tau = tau, eps = 1e-8, pf = pf,
                 pf2 = pf2, standardize = FALSE, intercept = FALSE,
                 lambda2 = lambda2)
as.vector(predict(m1.cv, newx = x, s = "lambda.min"))
```

---

predict.ernet

*Make predictions from an ernet object*

---

**Description**

Similar to other predict methods, this functions predicts fitted values from a fitted ernet object.

**Usage**

```
## S3 method for class 'ernet'
predict(object, newx, s = NULL, type = "response", ...)
```

**Arguments**

object	fitted <a href="#">ernet</a> model object.
newx	matrix of new values for x at which predictions are to be made. NOTE: newx must be a matrix, predict function does not accept a vector or other formats of newx.
s	value(s) of the penalty parameter lambda at which predictions are to be made. Default is the entire sequence used to create the model.
type	type of prediction required. Only response is available. Gives predicted response for regression problems.
...	Not used. Other arguments to predict.

**Details**

s is the new vector at which predictions are to be made. If s is not in the lambda sequence used for fitting the model, the predict function will use linear interpolation to make predictions. The new values are interpolated using a fraction of predicted values from both left and right lambda indices.

**Value**

The object returned depends on type.

**Author(s)**

Yuwen Gu and Hui Zou

Maintainer: Yuwen Gu <yuwen.gu@uconn.edu>

**See Also**

[ernet](#), [coef.ernet](#), [plot.ernet](#), [print.ernet](#)

---

`print.cpernet`

*Print a cpernet object*

---

**Description**

Print a summary of the cpernet path at each step along the path.

**Usage**

```
## S3 method for class 'cpernet'
print(x, digits = max(3, getOption("digits") - 3), ...)
```

**Arguments**

x	fitted <code>cpernet</code> object.
digits	significant digits in the output.
...	additional print arguments.

**Details**

The call that produced the `cpernet` object is printed, followed by a three-column matrix with columns Df1, Df2 and Lambda. The Df1 and Df2 columns are the number of nonzero mean and scale coefficients respectively.

**Value**

a three-column matrix, the first two columns are the number of nonzero mean and scale coefficients respectively and the third column is Lambda.

**Author(s)**

Yuwen Gu and Hui Zou

Maintainer: Yuwen Gu <yuwen.gu@uconn.edu>

**Examples**

```
set.seed(1)
n <- 100
p <- 400
x <- matrix(rnorm(n * p), n, p)
y <- rnorm(n)
tau <- 0.30
pf <- abs(rnorm(p))
pf2 <- abs(rnorm(p))
w <- 2.0
lambda2 <- 1
m2 <- cpernet(y = y, x = x, w = w, tau = tau, eps = 1e-8,
              pf.mean = pf, pf.scale = pf2,
              standardize = FALSE, lambda2 = lambda2)
print(m2)
```

---

print.ernet

*Print an ernet object*

---

**Description**

Print a summary of the ernet path at each step along the path.

**Usage**

```
## S3 method for class 'ernet'  
print(x, digits = max(3, getOption("digits") - 3), ...)
```

**Arguments**

x	fitted <a href="#">ernet</a> object.
digits	significant digits in the output.
...	additional print arguments.

**Details**

The call that produced the [ernet](#) object is printed, followed by a two-column matrix with columns Df and Lambda. The Df column is the number of nonzero coefficients.

**Value**

a two-column matrix, the first column is the number of nonzero coefficients and the second column is Lambda.

**Author(s)**

Yuwen Gu and Hui Zou

Maintainer: Yuwen Gu <yuwen.gu@uconn.edu>

**Examples**

```
set.seed(1)  
n <- 100  
p <- 400  
x <- matrix(rnorm(n * p), n, p)  
y <- rnorm(n)  
tau <- 0.90  
pf <- abs(rnorm(p))  
pf2 <- abs(rnorm(p))  
lambda2 <- 1  
m1 <- ernet(y = y, x = x, tau = tau, eps = 1e-8, pf = pf,  
            pf2 = pf2, standardize = FALSE, intercept = FALSE,  
            lambda2 = lambda2)  
print(m1)
```

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