

# Package ‘oem’

February 21, 2017

**Type** Package

**Title** Orthogonalizing EM

**Version** 2.0.4

**Date** 2017-02-15

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**Description** Solves penalized least squares problems using the orthogonalizing EM algorithm of Xiong et al. (2016) <doi:10.1080/00401706.2015.1054436>. The main fitting function is `oem()` and the functions `cv.oem()` and `xval.oem()` are for cross validation, the latter being an accelerated cross validation function for linear models. The `big.oem()` function allows for out of memory fitting.

**URL** <https://github.com/jaredhuling/oem>

**BugReports** <https://github.com/jaredhuling/oem/issues>

**License** GPL (>= 2)

**LazyData** TRUE

**Depends** R (>= 3.2.0), bigmemory

**Imports** Rcpp (>= 0.11.0), Matrix, foreach, methods

**LinkingTo** Rcpp, RcppEigen, BH, bigmemory, RcppArmadillo

**RoxygenNote** 5.0.1

**Suggests** knitr, rmarkdown

**VignetteBuilder** knitr

**NeedsCompilation** yes

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**Repository** CRAN

**Date/Publication** 2017-02-21 08:43:23

**R topics documented:**

big.oem	2
cv.oem	4
logLik.oem	6
oem	7
oem.txt	10
oemfit	12
plot.oem	14
predict.cv.oem	16
predict.oem	18
predict.xval.oem	19
print.summary.cv.oem	20
summary.cv.oem	21
xval.oem	21

<b>Index</b>	<b>24</b>
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big.oem	<i>Orthogonalizing EM</i>
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**Description**

Orthogonalizing EM

**Usage**

```
big.oem(x, y, family = c("gaussian", "binomial"), penalty = c("elastic.net",
  "lasso", "ols", "mcp", "scad", "grp.lasso"), weights = numeric(0),
  lambda = numeric(0), nlambda = 100L, lambda.min.ratio = NULL,
  alpha = 1, gamma = 3, groups = numeric(0), penalty.factor = NULL,
  group.weights = NULL, standardize = TRUE, intercept = TRUE,
  maxit = 500L, tol = 1e-07, irls.maxit = 100L, irls.tol = 0.001,
  compute.loss = FALSE, hessian.type = c("full", "upper.bound"))
```

**Arguments**

x	input big.matrix object pointing to design matrix Each row is an observation, each column corresponds to a covariate
y	numeric response vector of length nobs.
family	"gaussian" for least squares problems, "binomial" for binary response. "binomial" currently not available.
penalty	Specification of penalty type in lowercase letters. Choices include "lasso", "ols" (Ordinary least squares, no penalty), "elastic.net", "scad", "mcp", "grp.lasso"
weights	observation weights. Not implemented yet. Defaults to 1 for each observation (setting weight vector to length 0 will default all weights to 1)

<code>lambda</code>	A user supplied lambda sequence. By default, the program computes its own lambda sequence based on <code>nlambda</code> and <code>lambda.min.ratio</code> . Supplying a value of lambda overrides this.
<code>nlambda</code>	The number of lambda values - default is 100.
<code>lambda.min.ratio</code>	Smallest value for lambda, as a fraction of <code>lambda.max</code> , the (data derived) entry value (i.e. the smallest value for which all coefficients are zero). The default depends on the sample size <code>nobs</code> relative to the number of variables <code>nvars</code> . If <code>nobs &gt; nvars</code> , the default is 0.0001, close to zero. If <code>nobs &lt; nvars</code> , the default is 0.01. A very small value of <code>lambda.min.ratio</code> will lead to a saturated fit when <code>nobs &lt; nvars</code> .
<code>alpha</code>	mixing value for <code>elastic.net</code> . penalty applied is $(1 - \text{alpha}) * (\text{ridge penalty}) + \text{alpha} * (\text{lasso penalty})$
<code>gamma</code>	tuning parameter for SCAD and MCP penalties. must be $\geq 1$
<code>groups</code>	A vector of describing the grouping of the coefficients. See the example below. All unpenalized variables should be put in group 0
<code>penalty.factor</code>	Separate penalty factors can be applied to each coefficient. This is a number that multiplies lambda to allow differential shrinkage. Can be 0 for some variables, which implies no shrinkage, and that variable is always included in the model. Default is 1 for all variables.
<code>group.weights</code>	penalty factors applied to each group for the group lasso. Similar to <code>penalty.factor</code> , this is a number that multiplies lambda to allow differential shrinkage. Can be 0 for some groups, which implies no shrinkage, and that group is always included in the model. Default is $\sqrt{\text{group size}}$ for all groups.
<code>standardize</code>	Logical flag for x variable standardization, prior to fitting the models. The coefficients are always returned on the original scale. Default is <code>standardize = TRUE</code> . If variables are in the same units already, you might not wish to standardize. Keep in mind that standardization is done differently for sparse matrices, so results (when standardized) may be slightly different for a sparse matrix object and a dense matrix object
<code>intercept</code>	Should intercept(s) be fitted (default = TRUE) or set to zero (FALSE)
<code>maxit</code>	integer. Maximum number of OEM iterations
<code>tol</code>	convergence tolerance for OEM iterations
<code>irls.maxit</code>	integer. Maximum number of IRLS iterations
<code>irls.tol</code>	convergence tolerance for IRLS iterations. Only used if <code>family != "gaussian"</code>
<code>compute.loss</code>	should the loss be computed for each estimated tuning parameter? Defaults to FALSE. Setting to TRUE will dramatically increase computational time
<code>hessian.type</code>	only for logistic regression. if <code>hessian.type = "full"</code> , then the full hessian is used. If <code>hessian.type = "upper.bound"</code> , then an upper bound of the hessian is used. The upper bound can be dramatically faster in certain situations, ie when $n \gg p$

**Value**

An object with S3 class "oem"

**Examples**

```

set.seed(123)
nrows <- 50000
ncols <- 100
bkFile <- "bigmat.bk"
descFile <- "bigmatk.desc"
bigmat <- filebacked.big.matrix(nrow=nrows, ncol=ncols, type="double",
                               backingfile=bkFile, backingpath=".",
                               descriptorfile=descFile,
                               dimnames=c(NULL, NULL))

# Each column value will be the column number multiplied by
# samples from a standard normal distribution.
set.seed(123)
for (i in 1:ncols) bigmat[,i] = rnorm(nrows)*i

y <- rnorm(nrows) + bigmat[,1] - bigmat[,2]

fit <- big.oem(x = bigmat, y = y,
              penalty = c("lasso", "grp.lasso"),
              groups = rep(1:20, each = 5))

fit2 <- oem(x = bigmat[,], y = y,
            penalty = c("lasso", "grp.lasso"),
            groups = rep(1:20, each = 5))

max(abs(fit$beta[[1]] - fit2$beta[[1]]))

layout(matrix(1:2, ncol = 2))
plot(fit)
plot(fit, which.model = 2)

```

---

cv.oem

*Orthogonalizing EM*


---

**Description**

Orthogonalizing EM

**Usage**

```

cv.oem(x, y, penalty = c("elastic.net", "lasso", "ols", "mcp", "scad",
                        "grp.lasso"), weights = numeric(0), lambda = NULL,
       type.measure = c("mse", "deviance", "class", "auc", "mae"), nfolds = 10,
       foldid = NULL, grouped = TRUE, keep = FALSE, parallel = FALSE,
       ncores = -1, ...)

```

**Arguments**

x	input matrix or CsparseMatrix objects of the <b>Matrix</b> (sparse not yet implemented). Each row is an observation, each column corresponds to a covariate
y	numeric response vector of length nobs.
penalty	Specification of penalty type in lowercase letters. Choices include "lasso", "ols" (Ordinary least squares, no penalty), "elastic.net", "scad", "mcp", "grp.lasso"
weights	observation weights. defaults to 1 for each observation (setting weight vector to length 0 will default all weights to 1)
lambda	A user supplied lambda sequence. By default, the program computes its own lambda sequence based on nlambda and lambda.min.ratio. Supplying a value of lambda overrides this.
type.measure	measure to evaluate for cross-validation. The default is type.measure = "deviance", which uses squared-error for gaussian models (a.k.a type.measure = "mse" there), deviance for logistic regression. type.measure = "class" applies to binomial only. type.measure = "auc" is for two-class logistic regression only. type.measure = "mse" or type.measure = "mae" (mean absolute error) can be used by all models; they measure the deviation from the fitted mean to the response.
nfolds	number of folds for cross-validation. default is 10. 3 is smallest value allowed.
foldid	an optional vector of values between 1 and nfold specifying which fold each observation belongs to.
grouped	Like in <b>glmnet</b> , this is an experimental argument, with default TRUE, and can be ignored by most users. For all models, this refers to computing nfolds separate statistics, and then using their mean and estimated standard error to describe the CV curve. If grouped = FALSE, an error matrix is built up at the observation level from the predictions from the nfold fits, and then summarized (does not apply to type.measure = "auc").
keep	If keep = TRUE, a prevalidated list of array is returned containing fitted values for each observation and each value of lambda for each model. This means these fits are computed with this observation and the rest of its fold omitted. The foldid vector is also returned. Default is keep = FALSE
parallel	If TRUE, use parallel foreach to fit each fold. Must register parallel before hand, such as <b>doMC</b> .
ncores	Number of cores to use. If parallel = TRUE, then ncores will be automatically set to 1 to prevent conflicts
...	other parameters to be passed to "oem" function

**Value**

An object with S3 class "cv.oem"

**Examples**

```

set.seed(123)
n.obs <- 1e4
n.vars <- 100

true.beta <- c(runif(15, -0.25, 0.25), rep(0, n.vars - 15))

x <- matrix(rnorm(n.obs * n.vars), n.obs, n.vars)
y <- rnorm(n.obs, sd = 3) + x %*% true.beta

fit <- cv.oem(x = x, y = y,
             penalty = c("lasso", "grp.lasso"),
             groups = rep(1:20, each = 5))

layout(matrix(1:2, ncol = 2))
plot(fit)
plot(fit, which.model = 2)

```

---

logLik.oem

*log likelihood function for fitted oem objects*


---

**Description**

log likelihood function for fitted oem objects  
log likelihood function for fitted cross validation oem objects  
log likelihood function for fitted cross validation oem objects

**Usage**

```

## S3 method for class 'oem'
logLik(object, which.model = 1, ...)

## S3 method for class 'cv.oem'
logLik(object, which.model = 1, ...)

## S3 method for class 'xval.oem'
logLik(object, which.model = 1, ...)

```

**Arguments**

object	fitted "oem" model object.
which.model	If multiple penalties are fit and returned in the same oem object, the which.model argument is used to specify which model to plot. For example, if the oem object "oemobj" was fit with argument penalty = c("lasso", "grp.lasso"), then which.model = 2 provides a plot for the group lasso model.
...	not used

**Examples**

```

set.seed(123)
n.obs <- 2000
n.vars <- 50

true.beta <- c(runif(15, -0.25, 0.25), rep(0, n.vars - 15))
x <- matrix(rnorm(n.obs * n.vars), n.obs, n.vars)
y <- rnorm(n.obs, sd = 3) + x %*% true.beta

fit <- oem(x = x, y = y, penalty = "lasso", compute.loss = TRUE)

logLik(fit)

fit <- cv.oem(x = x, y = y, penalty = "lasso", compute.loss = TRUE)

logLik(fit)

fit <- xval.oem(x = x, y = y, penalty = "lasso", compute.loss = TRUE)

logLik(fit)

```

oem

*Orthogonalizing EM***Description**

Orthogonalizing EM

**Usage**

```

oem(x, y, family = c("gaussian", "binomial"), penalty = c("elastic.net",
  "lasso", "ols", "mcp", "scad", "grp.lasso"), weights = numeric(0),
  lambda = numeric(0), nlambda = 100L, lambda.min.ratio = NULL,
  alpha = 1, gamma = 3, groups = numeric(0), penalty.factor = NULL,
  group.weights = NULL, standardize = TRUE, intercept = TRUE,
  maxit = 500L, tol = 1e-07, irls.maxit = 100L, irls.tol = 0.001,
  accelerate = FALSE, ncores = -1, compute.loss = FALSE,
  hessian.type = c("full", "upper.bound"))

```

**Arguments**

x	input matrix or CsparseMatrix object of the <b>Matrix</b> package. Each row is an observation, each column corresponds to a covariate
y	numeric response vector of length nobs.
family	"gaussian" for least squares problems, "binomial" for binary response.

penalty	Specification of penalty type. Choices include "lasso", "ols" (Ordinary least squares, no penalty), "elastic.net", "scad", "mcp", "grp.lasso"
weights	observation weights. Not implemented yet. Defaults to 1 for each observation (setting weight vector to length 0 will default all weights to 1)
lambda	A user supplied lambda sequence. By default, the program computes its own lambda sequence based on nlambda and lambda.min.ratio. Supplying a value of lambda overrides this.
nlambda	The number of lambda values. The default is 100.
lambda.min.ratio	Smallest value for lambda, as a fraction of lambda.max, the (data derived) entry value (i.e. the smallest value for which all coefficients are zero). The default depends on the sample size nobs relative to the number of variables nvars. If $nobs > nvars$ , the default is 0.0001, close to zero. If $nobs < nvars$ , the default is 0.01. A very small value of lambda.min.ratio will lead to a saturated fit when $nobs < nvars$ .
alpha	mixing value for elastic.net. penalty applied is $(1 - \alpha) * (\text{ridge penalty}) + \alpha * (\text{lasso penalty})$
gamma	tuning parameter for SCAD and MCP penalties. must be $\geq 1$
groups	A vector of describing the grouping of the coefficients. See the example below. All unpenalized variables should be put in group 0
penalty.factor	Separate penalty factors can be applied to each coefficient. This is a number that multiplies lambda to allow differential shrinkage. Can be 0 for some variables, which implies no shrinkage, and that variable is always included in the model. Default is 1 for all variables.
group.weights	penalty factors applied to each group for the group lasso. Similar to penalty.factor, this is a number that multiplies lambda to allow differential shrinkage. Can be 0 for some groups, which implies no shrinkage, and that group is always included in the model. Default is $\sqrt{\text{group size}}$ for all groups.
standardize	Logical flag for x variable standardization, prior to fitting the models. The coefficients are always returned on the original scale. Default is standardize = TRUE. If variables are in the same units already, you might not wish to standardize. Keep in mind that standardization is done differently for sparse matrices, so results (when standardized) may be slightly different for a sparse matrix object and a dense matrix object
intercept	Should intercept(s) be fitted (default = TRUE) or set to zero (FALSE)
maxit	integer. Maximum number of OEM iterations
tol	convergence tolerance for OEM iterations
irls.maxit	integer. Maximum number of IRLS iterations
irls.tol	convergence tolerance for IRLS iterations. Only used if family != "gaussian"
accelerate	boolean argument. Whether or not to use Nesterov acceleration with adaptive restarting
ncores	Integer scalar that specifies the number of threads to be used
compute.loss	should the loss be computed for each estimated tuning parameter? Defaults to FALSE. Setting to TRUE will dramatically increase computational time



`hessian.type` only for logistic regression. if `hessian.type = "full"`, then the full hessian is used. If `hessian.type = "upper.bound"`, then an upper bound of the hessian is used. The upper bound can be dramatically faster in certain situations, ie when  $n \gg p$

## Value

An object with S3 class "oem"

## References

Shifeng Xiong, Bin Dai, Jared Huling, and Peter Z. G. Qian. Orthogonalizing EM: A design-based least squares algorithm. *Technometrics*, 58(3):285-293, 2016. <http://amstat.tandfonline.com/doi/abs/10.1080/00401706.2015.1054436>

## Examples

```
set.seed(123)
n.obs <- 1e4
n.vars <- 50

true.beta <- c(runif(15, -0.25, 0.25), rep(0, n.vars - 15))

x <- matrix(rnorm(n.obs * n.vars), n.obs, n.vars)
y <- rnorm(n.obs, sd = 3) + x %*% true.beta

fit <- oem(x = x, y = y,
          penalty = c("lasso", "grp.lasso"),
          groups = rep(1:10, each = 5))

layout(matrix(1:2, ncol = 2))
plot(fit)
plot(fit, which.model = 2)

# the oem package has support for
# sparse design matrices

library(Matrix)

xs <- rsparsematrix(n.obs * 25, n.vars * 2, density = 0.01)
ys <- rnorm(n.obs * 25, sd = 3) + as.vector(xs %*% c(true.beta, rep(0, n.vars)) )
x.dense <- as.matrix(xs)

system.time(fit <- oem(x = x.dense, y = ys,
                    penalty = c("lasso", "grp.lasso"),
                    groups = rep(1:20, each = 5), intercept = TRUE))

system.time(fits <- oem(x = xs, y = ys,
                      penalty = c("lasso", "grp.lasso"),
                      groups = rep(1:20, each = 5), intercept = TRUE, lambda = fit$lambda))

max(abs(fit$beta[[1]] - fits$beta[[1]]))
```

```

max(abs(fit$beta[[2]] - fits$beta[[2]]))

# logistic
y <- rbinom(n.obs, 1, prob = 1 / (1 + exp(-x %*% true.beta)))

system.time(res <- oem(x, y, intercept = FALSE,
                      penalty = "lasso",
                      family = "binomial",
                      nlambda = 10,
                      irls.tol = 1e-3, tol = 1e-8))

# sparse design matrix
xs <- rsparsematrix(n.obs * 2, n.vars, density = 0.01)
x.dense <- as.matrix(xs)
ys <- rbinom(n.obs * 2, 1, prob = 1 / (1 + exp(-x %*% true.beta)))

system.time(res.gr <- oem(x.dense, ys, intercept = FALSE,
                          penalty = "grp.lasso",
                          family = "binomial",
                          nlambda = 10,
                          groups = rep(1:5, each = 10),
                          irls.tol = 1e-3, tol = 1e-8))

system.time(res.gr.s <- oem(xs, ys, intercept = FALSE,
                            penalty = "grp.lasso",
                            family = "binomial",
                            nlambda = 10,
                            groups = rep(1:5, each = 10),
                            irls.tol = 1e-3, tol = 1e-8))

max(abs(res.gr$beta[[1]] - res.gr.s$beta[[1]]))

```

---

oem.xtx

*Orthogonalizing EM*


---

## Description

Orthogonalizing EM

## Usage

```

oem.xtx(xtx, xty, family = c("gaussian", "binomial"),
        penalty = c("elastic.net", "lasso", "ols", "mcp", "scad", "grp.lasso"),
        lambda = numeric(0), nlambda = 100L, lambda.min.ratio = NULL,
        alpha = 1, gamma = 3, groups = numeric(0), scale.factor = numeric(0),
        penalty.factor = NULL, group.weights = NULL, maxit = 500L,
        tol = 1e-07, irls.maxit = 100L, irls.tol = 0.001)

```

**Arguments**

<code>xtx</code>	input matrix equal to $\text{crossprod}(x) / \text{nrow}(x)$ . where $x$ is the design matrix. It is highly recommended to scale by the number of rows in $x$ . If $xtx$ is scaled, $xty$ must also be scaled or else results may be meaningless!
<code>xty</code>	numeric vector of length <code>nvars</code> . Equal to $\text{crossprod}(x, y) / \text{nobs}$ . It is highly recommended to scale by the number of rows in $x$ .
<code>family</code>	"gaussian" for least squares problems, "binomial" for binary response. (only gaussian implemented currently)
<code>penalty</code>	Specification of penalty type in lowercase letters. Choices include "lasso", "ols" (Ordinary least squares, no penalty), "elastic.net", "scad", "mcp", "grp.lasso"
<code>lambda</code>	A user supplied lambda sequence. By default, the program computes its own lambda sequence based on <code>nlambda</code> and <code>lambda.min.ratio</code> . Supplying a value of lambda overrides this.
<code>nlambda</code>	The number of lambda values - default is 100.
<code>lambda.min.ratio</code>	Smallest value for lambda, as a fraction of <code>lambda.max</code> , the (data derived) entry value (i.e. the smallest value for which all coefficients are zero). The default depends on the sample size <code>nobs</code> relative to the number of variables <code>nvars</code> . The default is 0.0001
<code>alpha</code>	mixing value for elastic.net. penalty applied is $(1 - \alpha) * (\text{ridge penalty}) + \alpha * (\text{lasso penalty})$
<code>gamma</code>	tuning parameter for SCAD and MCP penalties. must be $\geq 1$
<code>groups</code>	A vector of describing the grouping of the coefficients. See the example below. All unpenalized variables should be put in group 0
<code>scale.factor</code>	of length <code>nvars</code> $=== \text{ncol}(xtx) == \text{length}(xty)$ for scaling columns of $x$ . The standard deviation for each column of $x$ is a common choice for <code>scale.factor</code> . Coefficients will be returned on original scale. Default is no scaling.
<code>penalty.factor</code>	Separate penalty factors can be applied to each coefficient. This is a number that multiplies lambda to allow differential shrinkage. Can be 0 for some variables, which implies no shrinkage, and that variable is always included in the model. Default is 1 for all variables.
<code>group.weights</code>	penalty factors applied to each group for the group lasso. Similar to <code>penalty.factor</code> , this is a number that multiplies lambda to allow differential shrinkage. Can be 0 for some groups, which implies no shrinkage, and that group is always included in the model. Default is $\sqrt{\text{group size}}$ for all groups.
<code>maxit</code>	integer. Maximum number of OEM iterations
<code>tol</code>	convergence tolerance for OEM iterations
<code>irls.maxit</code>	integer. Maximum number of IRLS iterations
<code>irls.tol</code>	convergence tolerance for IRLS iterations. Only used if <code>family != "gaussian"</code>

**Value**

An object with S3 class "oem"

**Examples**

```

set.seed(123)
n.obs <- 1e4
n.vars <- 100

true.beta <- c(runif(15, -0.25, 0.25), rep(0, n.vars - 15))

x <- matrix(rnorm(n.obs * n.vars), n.obs, n.vars)
y <- rnorm(n.obs, sd = 3) + x %*% true.beta

fit <- oem(x = x, y = y,
          penalty = c("lasso", "grp.lasso"),
          standardize = FALSE, intercept = FALSE,
          groups = rep(1:20, each = 5))

xtx <- crossprod(x) / n.obs
xty <- crossprod(x, y) / n.obs

fit.xtx <- oem.xtx(xtx = xtx, xty = xty,
                 penalty = c("lasso", "grp.lasso"),
                 groups = rep(1:20, each = 5))

max(abs(fit$beta[[1]][-1,] - fit.xtx$beta[[1]]))
max(abs(fit$beta[[2]][-1,] - fit.xtx$beta[[2]]))

layout(matrix(1:2, ncol = 2))
plot(fit.xtx)
plot(fit.xtx, which.model = 2)

```

---

oemfit

*Deprecated functions*


---

**Description**

These functions have been renamed and deprecated in **oem**: `oemfit()` (use `oem()`), `cv.oemfit()` (use `cv.oem()`), `print.oemfit()`, `plot.oemfit()`, `predict.oemfit()`, and `coef.oemfit()`.

**Usage**

```

oemfit(formula, data = list(), lambda = NULL, nlambda = 100,
       lambda.min.ratio = NULL, tolerance = 0.001, maxIter = 1000,
       standardized = TRUE, numGroup = 1, penalty = c("lasso", "scad", "ols",
       "elastic.net", "ngarrote", "mcp"), alpha = 3, evaluate = 0,
       condition = -1)

cv.oemfit(formula, data = list(), lambda = NULL, type.measure = c("mse",
       "mae"), ..., nfolds = 10, foldid, penalty = c("lasso", "scad",
       "elastic.net", "ngarrote", "mcp"))

```

```
## S3 method for class 'oemfit'
plot(x, xvar = c("norm", "lambda", "loglambda", "dev"),
     xlab = iname, ylab = "Coefficients", ...)

## S3 method for class 'oemfit'
predict(object, newx, s = NULL, type = c("response",
    "coefficients", "nonzero"), ...)

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

## Arguments

formula	an object of 'formula' (or one that can be coerced to that class): a symbolic description of the model to be fitted. The details of model specification are given under 'Details'
data	an optional data frame, list or environment (or object coercible by 'as.data.frame' to a data frame) containing the variables in the model. If not found in 'data', the variables are taken from 'environment(formula)', typically the environment from which 'oemfit' is called.
lambda	A user supplied lambda sequence. Typical usage is to have the program compute its own lambda sequence based on nlambda and lambda.min.ratio. Supplying a value of lambda overrides this. <b>WARNING:</b> use with care. Do not supply a single value for lambda (for predictions after CV use predict() instead). Supply instead a decreasing sequence of lambda values. oemfit relies on its warms starts for speed, and its often faster to fit a whole path than compute a single fit.
nlambda	The number of lambda values - default is 100.
lambda.min.ratio	Smallest value for lambda, as a fraction of lambda.max, the (data derived) entry value (i.e. the smallest value for which all coefficients are zero). The default depends on the sample size nobs relative to the number of variables nvars. If nobs > nvars, the default is 0.0001, close to zero. If nobs < nvars, the default is 0.01. A very small value of lambda.min.ratio will lead to a saturated fit in the nobs < nvars case.
tolerance	Convergence tolerance for OEM. Each inner OEM loop continues until the maximum change in the objective after any coefficient update is less than tolerance. Defaults value is 1E-3.
maxIter	Maximum number of passes over the data for all lambda values; default is 1000.
standardized	Logical flag for x variable standardization, prior to fitting the model sequence. The coefficients are always returned on the original scale. Default is standardize=TRUE. If variables are in the same units already, you might not wish to standardize.
numGroup	Integer value for the number of groups to use for OEM fitting. Default is 1.
penalty	type in lower letters. Different types include 'lasso', 'scad', 'ols' (ordinary least square), 'elastic-net', 'ngarrote' (non-negative garrote) and 'mcp'.

alpha	alpha value for scad and mcp.
evaluate	debugging argument
condition	Debugging for different ways of calculating OEM.
type.measure	type.measure measure to evaluate for cross-validation. type.measure = "mse" (mean squared error) or type.measure = "mae" (mean absolute error)
...	arguments to be passed to oemfit()
nfolds	number of folds for cross-validation. default is 10.
foldid	an optional vector of values between 1 and nfold specifying which fold each observation belongs to.
x	fitted oemfit object
xvar	what is on the X-axis. "norm" plots against the L1-norm of the coefficients, "lambda" against the log-lambda sequence, and "dev" against the percent deviance explained.
xlab	x-axis label
ylab	y-axis label
object	fitted oemfit object
newx	matrix of new values for x at which predictions are to be made. Must be a matrix.
s	Value(s) of the penalty parameter lambda at which predictions are required. Default is the entire sequence used to create the model.
type	not used.
digits	significant digits in print out.

### Details

The sequence of models implied by 'lambda' is fit by OEM algorithm.

### Author(s)

Bin Dai

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plot.oem

*Plot method for Orthogonalizing EM fitted objects*

---

### Description

Plot method for Orthogonalizing EM fitted objects

Plot method for Orthogonalizing EM fitted objects

**Usage**

```
## S3 method for class 'oem'
plot(x, which.model = 1, xvar = c("norm", "lambda",
  "loglambda", "dev"), labsize = 0.6, xlab = iname, ylab = "Coefficients",
  ...)

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

## S3 method for class 'xval.oem'
plot(x, which.model = 1, type = c("cv", "coefficients"),
  xvar = c("norm", "lambda", "loglambda", "dev"), labsize = 0.6,
  xlab = iname, ylab = "Coefficients", sign.lambda = 1, ...)
```

**Arguments**

x	fitted "oem" model object
which.model	If multiple penalties are fit and returned in the same oem object, the which.model argument is used to specify which model to plot. For example, if the oem object "oemobj" was fit with argument penalty = c("lasso", "grp.lasso"), then which.model = 2 provides a plot for the group lasso model.
xvar	What is on the X-axis. "norm" plots against the L1-norm of the coefficients, "lambda" against the log-lambda sequence, and "dev" against the percent deviance explained.
labsize	size of labels for variable names. If labsize = 0, then no variable names will be plotted
xlab	label for x-axis
ylab	label for y-axis
...	other graphical parameters for the plot
sign.lambda	Either plot against log(lambda) (default) or its negative if sign.lambda = -1.
type	one of "cv" or "coefficients". type = "cv" will produce a plot of cross validation results like plot.cv.oem. type = "coefficients" will produce a coefficient path plot like plot.oem()

**Examples**

```
set.seed(123)
n.obs <- 1e4
n.vars <- 100
n.obs.test <- 1e3

true.beta <- c(runif(15, -0.5, 0.5), rep(0, n.vars - 15))

x <- matrix(rnorm(n.obs * n.vars), n.obs, n.vars)
y <- rnorm(n.obs, sd = 3) + x %*% true.beta

fit <- oem(x = x, y = y, penalty = c("lasso", "grp.lasso"), groups = rep(1:10, each = 10))
```

```

layout(matrix(1:2, ncol = 2))
plot(fit, which.model = 1)
plot(fit, which.model = 2)

set.seed(123)
n.obs <- 1e4
n.vars <- 100
n.obs.test <- 1e3

true.beta <- c(runif(15, -0.5, 0.5), rep(0, n.vars - 15))

x <- matrix(rnorm(n.obs * n.vars), n.obs, n.vars)
y <- rnorm(n.obs, sd = 3) + x %*% true.beta

fit <- cv.oem(x = x, y = y, penalty = c("lasso", "grp.lasso"), groups = rep(1:10, each = 10))

layout(matrix(1:2, ncol = 2))
plot(fit, which.model = 1)
plot(fit, which.model = 2)

set.seed(123)
n.obs <- 1e4
n.vars <- 100
n.obs.test <- 1e3

true.beta <- c(runif(15, -0.5, 0.5), rep(0, n.vars - 15))

x <- matrix(rnorm(n.obs * n.vars), n.obs, n.vars)
y <- rnorm(n.obs, sd = 3) + x %*% true.beta

fit <- xval.oem(x = x, y = y, penalty = c("lasso", "grp.lasso"), groups = rep(1:10, each = 10))

layout(matrix(1:4, ncol = 2))
plot(fit, which.model = 1)
plot(fit, which.model = 2)

plot(fit, which.model = 1, type = "coef")
plot(fit, which.model = 2, type = "coef")

```

---

predict.cv.oem

*Prediction function for fitted cross validation oem objects*

---

## Description

Prediction function for fitted cross validation oem objects



**Usage**

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

**Arguments**

object	fitted "cv.oem" model object
newx	Matrix of new values for x at which predictions are to be made. Must be a matrix; can be sparse as in the CsparseMatrix objects of the <b>Matrix</b> package This argument is not used for type = c("coefficients", "nonzero")
which.model	If multiple penalties are fit and returned in the same oem object, the which.model argument is used to specify which model to make predictions for. For example, if the oem object "oemobj" was fit with argument penalty = c("lasso", "grp.lasso"), then which.model = 2 provides predictions for the group lasso model. For predict.cv.oem(), can specify "best.model" to use the best model as estimated by cross-validation
s	Value(s) of the penalty parameter lambda at which predictions are required. Default is the entire sequence used to create the model. For predict.cv.oem(), can also specify "lambda.1se" or "lambda.min" for best lambdas estimated by cross validation
...	used to pass the other arguments for predict.oem

**Value**

An object depending on the type argument

**Examples**

```
set.seed(123)
n.obs <- 1e4
n.vars <- 100
n.obs.test <- 1e3

true.beta <- c(runif(15, -0.5, 0.5), rep(0, n.vars - 15))

x <- matrix(rnorm(n.obs * n.vars), n.obs, n.vars)
y <- rnorm(n.obs, sd = 3) + x %*% true.beta
x.test <- matrix(rnorm(n.obs.test * n.vars), n.obs.test, n.vars)
y.test <- rnorm(n.obs.test, sd = 3) + x.test %*% true.beta

fit <- cv.oem(x = x, y = y,
             penalty = c("lasso", "grp.lasso"),
             groups = rep(1:10, each = 10),
             nlambdas = 10)

preds.best <- predict(fit, newx = x.test, type = "response", which.model = "best.model")

apply(preds.best, 2, function(x) mean((y.test - x) ^ 2))
```

predict.oem

*Prediction method for Orthogonalizing EM fitted objects***Description**

Prediction method for Orthogonalizing EM fitted objects

**Usage**

```
## S3 method for class 'oem'
predict(object, newx, s = NULL, which.model = 1,
        type = c("link", "response", "coefficients", "nonzero", "class"), ...)
```

**Arguments**

object	fitted "oem" model object
newx	Matrix of new values for x at which predictions are to be made. Must be a matrix; can be sparse as in the CsparseMatrix objects of the <b>Matrix</b> package. This argument is not used for type=c("coefficients", "nonzero")
s	Value(s) of the penalty parameter lambda at which predictions are required. Default is the entire sequence used to create the model.
which.model	If multiple penalties are fit and returned in the same oem object, the which.model argument is used to specify which model to make predictions for. For example, if the oem object oemobj was fit with argument penalty = c("lasso", "grp.lasso"), then which.model = 2 provides predictions for the group lasso model.
type	Type of prediction required. type = "link" gives the linear predictors for the "binomial" model; for "gaussian" models it gives the fitted values. type = "response" gives the fitted probabilities for "binomial". type = "coefficients" computes the coefficients at the requested values for s. type = "class" applies only to "binomial" and produces the class label corresponding to the maximum probability.
...	not used

**Value**

An object depending on the type argument

**Examples**

```
set.seed(123)
n.obs <- 1e4
n.vars <- 100
n.obs.test <- 1e3

true.beta <- c(runif(15, -0.5, 0.5), rep(0, n.vars - 15))
```

```

x <- matrix(rnorm(n.obs * n.vars), n.obs, n.vars)
y <- rnorm(n.obs, sd = 3) + x %*% true.beta
x.test <- matrix(rnorm(n.obs.test * n.vars), n.obs.test, n.vars)
y.test <- rnorm(n.obs.test, sd = 3) + x.test %*% true.beta

fit <- oem(x = x, y = y,
          penalty = c("lasso", "grp.lasso"),
          groups = rep(1:10, each = 10),
          nlambda = 10)

preds.lasso <- predict(fit, newx = x.test, type = "response", which.model = 1)
preds.grp.lasso <- predict(fit, newx = x.test, type = "response", which.model = 2)

apply(preds.lasso, 2, function(x) mean((y.test - x) ^ 2))
apply(preds.grp.lasso, 2, function(x) mean((y.test - x) ^ 2))

```

---

predict.xval.oem	<i>Prediction function for fitted cross validation oem objects</i>
------------------	--

---

## Description

Prediction function for fitted cross validation oem objects

## Usage

```

## S3 method for class 'xval.oem'
predict(object, newx, which.model = "best.model",
        s = c("lambda.min", "lambda.1se"), ...)

```

## Arguments

object	fitted "cv.oem" model object
newx	Matrix of new values for x at which predictions are to be made. Must be a matrix; can be sparse as in the <code>CsparseMatrix</code> objects of the <b>Matrix</b> package This argument is not used for <code>type=c("coefficients","nonzero")</code>
which.model	If multiple penalties are fit and returned in the same oem object, the <code>which.model</code> argument is used to specify which model to make predictions for. For example, if the oem object "oemobj" was fit with argument <code>penalty = c("lasso", "grp.lasso")</code> , then <code>which.model = 2</code> provides predictions for the group lasso model. For <code>predict.cv.oem()</code> , can specify "best.model" to use the best model as estimated by cross-validation
s	Value(s) of the penalty parameter lambda at which predictions are required. Default is the entire sequence used to create the model. For <code>predict.cv.oem</code> , can also specify "lambda.1se" or "lambda.min" for best lambdas estimated by cross validation
...	used to pass the other arguments for <code>predict.oem()</code>

**Value**

An object depending on the type argument

**Examples**

```

set.seed(123)
n.obs <- 1e4
n.vars <- 100
n.obs.test <- 1e3

true.beta <- c(runif(15, -0.5, 0.5), rep(0, n.vars - 15))

x <- matrix(rnorm(n.obs * n.vars), n.obs, n.vars)
y <- rnorm(n.obs, sd = 3) + x %>% true.beta
x.test <- matrix(rnorm(n.obs.test * n.vars), n.obs.test, n.vars)
y.test <- rnorm(n.obs.test, sd = 3) + x.test %>% true.beta

fit <- xval.oem(x = x, y = y,
               penalty = c("lasso", "grp.lasso"),
               groups = rep(1:10, each = 10),
               nlambdas = 10)

preds.best <- predict(fit, newx = x.test, type = "response", which.model = "best.model")

apply(preds.best, 2, function(x) mean((y.test - x) ^ 2))

```

---

print.summary.cv.oem *print method for summary.cv.oem objects*

---

**Description**

print method for summary.cv.oem objects

**Usage**

```

## S3 method for class 'summary.cv.oem'
print(x, digits, ...)

```

**Arguments**

x	a "summary.cv.oem" object
digits	digits to display
...	not used

---

summary.cv.oem	<i>summary method for cross validation Orthogonalizing EM fitted objects</i>
----------------	--

---

### Description

summary method for cross validation Orthogonalizing EM fitted objects

summary method for cross validation Orthogonalizing EM fitted objects

### Usage

```
## S3 method for class 'cv.oem'
summary(object, ...)
```

```
## S3 method for class 'xval.oem'
summary(object, ...)
```

### Arguments

object	fitted "cv.oem" object
...	not used

---

xval.oem	<i>Orthogonalizing EM</i>
----------	---------------------------

---

### Description

Orthogonalizing EM

### Usage

```
xval.oem(x, y, nfolds = 10L, foldid = NULL, type.measure = c("mse",
  "deviance", "class", "auc", "mae"), ncores = -1, family = c("gaussian",
  "binomial"), penalty = c("elastic.net", "lasso", "ols", "mcp", "scad",
  "grp.lasso"), weights = numeric(0), lambda = numeric(0), nlambda = 100L,
  lambda.min.ratio = NULL, alpha = 1, gamma = 3, groups = numeric(0),
  penalty.factor = NULL, group.weights = NULL, standardize = TRUE,
  intercept = TRUE, maxit = 500L, tol = 1e-07, irls.maxit = 100L,
  irls.tol = 0.001, compute.loss = FALSE)
```

**Arguments**

x	input matrix (sparse matrices not yet implemented). Each row is an observation, each column corresponds to a covariate
y	numeric response vector of length nobs = nrow(x).
nfolds	integer number of cross validation folds. 3 is the minimum number allowed. defaults to 10
foldid	an optional vector of values between 1 and nfold specifying which fold each observation belongs to.
type.measure	measure to evaluate for cross-validation. The default is type.measure = "deviance", which uses squared-error for gaussian models (a.k.a type.measure = "mse" there), deviance for logistic regression. type.measure = "class" applies to binomial only. type.measure = "auc" is for two-class logistic regression only. type.measure="mse" or type.measure="mae" (mean absolute error) can be used by all models; they measure the deviation from the fitted mean to the response.
ncores	Integer scalar that specifies the number of threads to be used
family	"gaussian" for least squares problems, "binomial" for binary response (not implemented yet).
penalty	Specification of penalty type in lowercase letters. Choices include "lasso", "ols" (Ordinary least squares, no penalty), "elastic.net", "scad", "mcp", "grp.lasso"
weights	observation weights. defaults to 1 for each observation (setting weight vector to length 0 will default all weights to 1)
lambda	A user supplied lambda sequence. By default, the program computes its own lambda sequence based on nlambda and lambda.min.ratio. Supplying a value of lambda overrides this.
nlambda	The number of lambda values - default is 100.
lambda.min.ratio	Smallest value for lambda, as a fraction of lambda.max, the (data derived) entry value (i.e. the smallest value for which all coefficients are zero). The default depends on the sample size nobs relative to the number of variables nvars. If nobs > nvars, the default is 0.0001, close to zero.
alpha	mixing value for elastic.net. penalty applied is (1 - alpha) * (ridge penalty) + alpha * (lasso penalty)
gamma	tuning parameter for SCAD and MCP penalties. must be >= 1
groups	A vector of describing the grouping of the coefficients. See the example below. All unpenalized variables should be put in group 0
penalty.factor	Separate penalty factors can be applied to each coefficient. This is a number that multiplies lambda to allow differential shrinkage. Can be 0 for some variables, which implies no shrinkage, and that variable is always included in the model. Default is 1 for all variables.
group.weights	penalty factors applied to each group for the group lasso. Similar to penalty.factor, this is a number that multiplies lambda to allow differential shrinkage. Can be 0 for some groups, which implies no shrinkage, and that group is always included in the model. Default is sqrt(group size) for all groups.

<code>standardize</code>	Logical flag for x variable standardization, prior to fitting the models. The coefficients are always returned on the original scale. Default is <code>standardize = TRUE</code> . If variables are in the same units already, you might not wish to standardize.
<code>intercept</code>	Should intercept(s) be fitted (default = <code>TRUE</code> ) or set to zero ( <code>FALSE</code> )
<code>maxit</code>	integer. Maximum number of OEM iterations
<code>tol</code>	convergence tolerance for OEM iterations
<code>irls.maxit</code>	integer. Maximum number of IRLS iterations
<code>irls.tol</code>	convergence tolerance for IRLS iterations. Only used if <code>family != "gaussian"</code>
<code>compute.loss</code>	should the loss be computed for each estimated tuning parameter? Defaults to <code>FALSE</code> . Setting to <code>TRUE</code> will dramatically increase computational time

**Value**

An object with S3 class "xval.oem"

**Examples**

```
set.seed(123)
n.obs <- 1e4
n.vars <- 100

true.beta <- c(runif(15, -0.25, 0.25), rep(0, n.vars - 15))

x <- matrix(rnorm(n.obs * n.vars), n.obs, n.vars)
y <- rnorm(n.obs, sd = 3) + x %*% true.beta

system.time(fit <- oem(x = x, y = y,
  penalty = c("lasso", "grp.lasso"),
  groups = rep(1:20, each = 5)))

system.time(xfit <- xval.oem(x = x, y = y,
  penalty = c("lasso", "grp.lasso"),
  groups = rep(1:20, each = 5)))
```

# Index

`big.oem`, [2](#)

`cv.oem`, [4](#), [12](#)

`cv.oemfit (oemfit)`, [12](#)

`logLik.cv.oem (logLik.oem)`, [6](#)

`logLik.oem`, [6](#)

`logLik.xval.oem (logLik.oem)`, [6](#)

`oem`, [7](#), [12](#)

`oem-deprecated (oemfit)`, [12](#)

`oem.txt`, [10](#)

`oemfit`, [12](#)

`plot.cv.oem (plot.oem)`, [14](#)

`plot.oem`, [14](#)

`plot.oemfit (oemfit)`, [12](#)

`plot.xval.oem (plot.oem)`, [14](#)

`predict.cv.oem`, [16](#)

`predict.oem`, [18](#)

`predict.oemfit (oemfit)`, [12](#)

`predict.xval.oem`, [19](#)

`print.oemfit (oemfit)`, [12](#)

`print.summary.cv.oem`, [20](#)

`summary.cv.oem`, [21](#)

`summary.xval.oem (summary.cv.oem)`, [21](#)

`xval.oem`, [21](#)