Package 'glmnet'

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Title Lasso and elastic-net regularized generalized linear models							
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Depends Matrix (>= 1.0-6), utils							
Suggests survival, foreach							
Description Extremely efficient procedures for fitting the entire lasso or elastic-net regularization path for linear regression, logistic and multinomial regression models, poisson regression and the Cox model. Two recent additions are the multiresponse gaussian, and the grouped multinomial. The algorithm uses cyclical coordinate descent in a pathwise fashion, as described in the paper listed below.							
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Description

This package fits lasso and elastic-net model paths for regression, logistic and multinomial regression using coordinate descent. The algorithm is extremely fast, and exploits sparsity in the input x matrix where it exists. A variety of predictions can be made from the fitted models.

Details

Package: glmnet Type: Package Version: 1.0

Date: 2008-05-14

License: What license is it under?

Very simple to use. Accepts x,y data for regression models, and produces the regularization path over a grid of values for the tuning parameter lambda. Only 5 functions: glmnet predict.glmnet plot.glmnet print.glmnet coef.glmnet

Author(s)

Jerome Friedman, Trevor Hastie and Rob Tibshirani Maintainer: Trevor Hastie hastie@stanford.edu>

References

```
Friedman, J., Hastie, T. and Tibshirani, R. (2008) Regularization Paths for Generalized Linear Models via Coordinate Descent, http://www.stanford.edu/~hastie/Papers/glmnet.pdf
Journal of Statistical Software, Vol. 33(1), 1-22 Feb 2010
http://www.jstatsoft.org/v33/i01/
```

Simon, N., Friedman, J., Hastie, T., Tibshirani, R. (2011) Regularization Paths for Cox's Proportional Hazards Model via Coordinate Descent, Journal of Statistical Software, Vol. 39(5) 1-13 http://www.jstatsoft.org/v39/i05/

Examples

```
x=matrix(rnorm(100*20),100,20)
y=rnorm(100)
g2=sample(1:2,100,replace=TRUE)
g4=sample(1:4,100,replace=TRUE)
fit1=glmnet(x,y)
predict(fit1,newx=x[1:5,],s=c(0.01,0.005))
predict(fit1,type="coef")
plot(fit1,xvar="lambda")
fit2=glmnet(x,g2,family="binomial")
predict(fit2,type="response",newx=x[2:5,])
predict(fit2,type="nonzero")
fit3=glmnet(x,g4,family="multinomial")
predict(fit3,newx=x[1:3,],type="response",s=0.01)
```

cv.glmnet

Cross-validation for glmnet

Description

Does k-fold cross-validation for glmnet, produces a plot, and returns a value for lambda

Usage

Arguments

X	x matrix as in glmnet.
у	response y as in glmnet.
weights	Observation weights; defaults to 1 per observation
offset	Offset vector (matrix) as in glmnet
lambda	Optional user-supplied lambda sequence; default is NULL, and glmnet chooses its own sequence
nfolds	number of folds - default is 10. Although nfolds can be as large as the sample size (leave-one-out CV), it is not recommended for large datasets. Smallest value allowable is nfolds=3
foldid	an optional vector of values between 1 and nfold identifying what fold each observation is in. If supplied, nfold can be missing.
type.measure	loss to use for cross-validation. Currently five options, not all available for all models. The default is type.measure="deviance", which uses squared-error for gaussian models (a.k.a type.measure="mse" there), deviance for logistic and poisson regression, and partial-likelihood for the Cox model. type.measure="class" applies to binomial and multinomial logistic regression only, and gives misclas-

sification error. type.measure="auc" is for two-class logistic regression only,

and gives area under the ROC curve. type.measure="mse" or type.measure="mae" (mean absolute error) can be used by all models except the "cox"; they measure the deviation from the fitted mean to the response.

grouped

This is an experimental argument, with default TRUE, and can be ignored by most users. For all models except the "cox", 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"). For the "cox" family, grouped=TRUE obtains the CV partial likelihood for the Kth fold by *subtraction*; by subtracting the log partial likelihood evaluated on the full dataset from that evaluated on the on the (K-1)/K dataset. This makes more efficient use of risk sets. With grouped=FALSE the log

partial likelihood is computed only on the Kth fold

keep If keep=TRUE, a *prevalidated* array is returned containing fitted values for each

observation and each value of lambda. This means these fits are computed with this observation and the rest of its fold omitted. The folid vector is also re-

turned. Default is keep=FALSE

parallel If TRUE, use parallel for each to fit each fold. Must register parallel before hand,

such as doMC or others. See the example below.

... Other arguments that can be passed to glmnet

Details

The function runs glmnet nfolds+1 times; the first to get the lambda sequence, and then the remainder to compute the fit with each of the folds omitted. The error is accumulated, and the average error and standard deviation over the folds is computed. Note that cv.glmnet does NOT search for values for alpha. A specific value should be supplied, else alpha=1 is assumed by default. If users would like to cross-validate alpha as well, they should call cv.glmnet with a precomputed vector foldid, and then use this same fold vector in separate calls to cv.glmnet with different values of alpha.

Value

an object of class "cv.glmnet" 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 length(lambda).

cvsd estimate of standard error of cvm.

cvup upper curve = cvm+cvsd. cvlo lower curve = cvm-cvsd.

nzero number of non-zero coefficients at each lambda.

name a text string indicating type of measure (for plotting purposes).

glmnet.fit a fitted glmnet object for the full data.

lambda.min value of lambda that gives minimum cvm.

lambda.1se largest value of lambda such that error is within 1 standard error of the mini-

mum.

fit.preval if keep=TRUE, this is the array of prevalidated fits. Some entries can be NA, if

that and subsequent values of lambda are not reached for that fold

foldid if keep=TRUE, the fold assignments used

Author(s)

Jerome Friedman, Trevor Hastie and Rob Tibshirani Noah Simon helped develop the 'coxnet' function.

Jeffrey Wong and B. Narasimhan helped with the parallel option

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References

```
Friedman, J., Hastie, T. and Tibshirani, R. (2008) Regularization Paths for Generalized Linear Models via Coordinate Descent, http://www.stanford.edu/~hastie/Papers/glmnet.pdf
Journal of Statistical Software, Vol. 33(1), I-22 Feb 2010
http://www.jstatsoft.org/v33/i01/
Simon, N., Friedman, J., Hastie, T., Tibshirani, R. (2011) Regularization Paths for Cox's Proportional Hazards Model via Coordinate Descent, Journal of Statistical Software, Vol. 39(5) 1-13
http://www.jstatsoft.org/v39/i05/
```

See Also

glmnet and plot, predict, and coef methods for "cv.glmnet" object.

Examples

```
set.seed(1010)
n=1000; p=100
nzc=trunc(p/10)
x=matrix(rnorm(n*p),n,p)
beta=rnorm(nzc)
fx= x[,seq(nzc)] %*% beta
eps=rnorm(n)*5
y=drop(fx+eps)
px=exp(fx)
px=px/(1+px)
ly=rbinom(n=length(px),prob=px,size=1)
set.seed(1011)
cvob1=cv.glmnet(x,y)
plot(cvob1)
coef(cvob1)
predict(cvob1,newx=x[1:5,], s="lambda.min")
title("Gaussian Family",line=2.5)
set.seed(1011)
cvob1a=cv.glmnet(x,y,type.measure="mae")
plot(cvob1a)
title("Gaussian Family", line=2.5)
set.seed(1011)
```

```
par(mfrow=c(2,2), mar=c(4.5,4.5,4,1))
cvob2=cv.glmnet(x,ly,family="binomial")
plot(cvob2)
title("Binomial Family",line=2.5)
frame()
set.seed(1011)
cvob3=cv.glmnet(x,ly,family="binomial",type.measure="class")
plot(cvob3)
title("Binomial Family",line=2.5)
set.seed(1011)
cvob3a=cv.glmnet(x,ly,family="binomial",type.measure="auc")
plot(cvob3a)
title("Binomial Family",line=2.5)
set.seed(1011)
mu=exp(fx/10)
y=rpois(n,mu)
cvob4=cv.glmnet(x,y,family="poisson")
plot(cvob4)
title("Poisson Family",line=2.5)
# Multinomial
n=500; p=30
nzc=trunc(p/10)
x=matrix(rnorm(n*p),n,p)
beta3=matrix(rnorm(30),10,3)
beta3=rbind(beta3,matrix(0,p-10,3))
f3=x%*% beta3
p3=exp(f3)
p3=p3/apply(p3,1,sum)
g3=rmult(p3)
set.seed(10101)
cvfit=cv.glmnet(x,g3,family="multinomial")
plot(cvfit)
title("Multinomial Family",line=2.5)
# Cox
beta=rnorm(nzc)
fx=x[,seq(nzc)]%*%beta/3
hx=exp(fx)
ty=rexp(n,hx)
tcens=rbinom(n=n,prob=.3,size=1)# censoring indicator
y=cbind(time=ty,status=1-tcens) # y=Surv(ty,1-tcens) with library(survival)
foldid=sample(rep(seq(10),length=n))
fit1_cv=cv.glmnet(x,y,family="cox",foldid=foldid)
plot(fit1_cv)
title("Cox Family",line=2.5)
## Not run:
# Parallel
require(doMC)
registerDoMC(cores=4)
x = matrix(rnorm(1e5 * 100), 1e5, 100)
y = rnorm(1e5)
system.time(cv.glmnet(x,y))
system.time(cv.glmnet(x,y,parallel=TRUE))
```

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```
## End(Not run)
```

deviance.glmnet

Extract the deviance from a glmnet object

Description

Compute the deviance sequence from the glmnet object

Usage

```
## S3 method for class 'glmnet'
deviance(object,...)
```

Arguments

object fitted glmnet object
... additional print arguments

Details

A glmnet object has components dev.ratio and nulldev. The former is the fraction of (null) deviance explained. The deviance calculations incorporate weights if present in the model. The deviance is defined to be 2*(loglike_sat - loglike), where loglike_sat is the log-likelihood for the saturated model (a model with a free parameter per observation). Null deviance is defined to be 2*(loglike_sat -loglike(Null)); The NULL model refers to the intercept model, except for the Cox, where it is the 0 model. Hence dev.ratio=1-deviance/nulldev, and this deviance method returns (1-dev.ratio)*nulldev.

Value

```
(1-dev.ratio)*nulldev
```

Author(s)

Jerome Friedman, Trevor Hastie and Rob Tibshirani Maintainer: Trevor Hastie hastie@stanford.edu>

References

Friedman, J., Hastie, T. and Tibshirani, R. (2008) Regularization Paths for Generalized Linear Models via Coordinate Descent

See Also

```
glmnet, predict, print, and coef methods.
```

Examples

```
x=matrix(rnorm(100*20),100,20)
y=rnorm(100)
fit1=glmnet(x,y)
deviance(fit1)
```

glmnet

fit a GLM with lasso or elasticnet regularization

Description

Fit a generalized linear model via penalized maximum likelihood. The regularization path is computed for the lasso or elasticnet penalty at a grid of values for the regularization parameter lambda. Can deal with all shapes of data, including very large sparse data matrices. Fits linear, logistic and multinomial, poisson, and Cox regression models.

Usage

```
glmnet(x, y, family=c("gaussian","binomial","poisson","multinomial","cox","mgaussian"),
    weights, offset=NULL, alpha = 1, nlambda = 100,
    lambda.min.ratio = ifelse(nobs<nvars,0.01,0.0001), lambda=NULL,
    standardize = TRUE, intercept=TRUE, thresh = 1e-07, dfmax = nvars + 1,
    pmax = min(dfmax * 2+20, nvars), exclude, penalty.factor = rep(1, nvars),
    lower.limits=-Inf, upper.limits=Inf, maxit=100000,
    type.gaussian=ifelse(nvars<500,"covariance","naive"),
    type.logistic=c("Newton","modified.Newton"),
    standardize.response=FALSE, type.multinomial=c("ungrouped","grouped"))</pre>
```

Arguments

У

x input matrix, of dimension nobs x nvars; each row is an observation vector. Can be in sparse matrix format (inherit from class "sparseMatrix" as in package Matrix; not yet available for family="cox")

response variable. Quantitative for family="gaussian", or family="poisson" (non-negative counts). For family="binomial" should be either a factor with two levels, or a two-column matrix of counts or proportions. For family="multinomial", can be a nc>=2 level factor, or a matrix with nc columns of counts or proportions. For family="cox", y should be a two-column matrix with columns named 'time' and 'status'. The latter is a binary variable, with '1' indicating

death, and '0' indicating right censored. The function Surv() in package **survival** produces such a matrix. For family="mgaussian", y is a matrix of quantitative responses.

family Response type (see above)

weights observation weights. Can be total counts if responses are proportion matrices.

Default is 1 for each observation

offset A vector of

A vector of length nobs that is included in the linear predictor (a nobs x nc matrix for the "multinomial" family). Useful for the "poisson" family (e.g. log of exposure time), or for refining a model by starting at a current fit. Default is NULL. If supplied, then values must also be supplied to the predict function.

alpha

The elasticnet mixing parameter, with $0 \le \alpha \le 1$. The penalty is defined as

$$(1-\alpha)/2||\beta||_2^2 + \alpha||\beta||_1$$
.

alpha=1 is the lasso penalty, and alpha=0 the ridge penalty.

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. This is undefined for "binomial" and "multinomial" models, and glmnet will exit gracefully when the percentage deviance explained is almost 1.

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. WARNING: 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. glmnet relies on its warms starts for speed, and its often faster to fit a whole path than compute a single fit.

standardize

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. See details below for y standardization with family="gaussian".

intercept

Should intercept(s) be fitted (default=TRUE) or set to zero (FALSE)

thresh

Convergence threshold for coordinate descent. Each inner coordinate-descent loop continues until the maximum change in the objective after any coefficient update is less than thresh times the null deviance. Defaults value is 1E-7.

dfmax

Limit the maximum number of variables in the model. Useful for very large nvars, if a partial path is desired.

pmax

Limit the maximum number of variables ever to be nonzero

exclude

Indices of variables to be excluded from the model. Default is none. Equivalent to an infinite penalty factor (next item).

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 (and implicitly infinity for variables listed in exclude). Note: the penalty factors are internally rescaled to sum to nvars, and the lambda sequence will reflect this change.

lower.limits Vector of lower limits for each coefficient; default -Inf. Each of these must be

non-positive. Can be presented as a single value (which will then be replicated),

else a vector of length nvars

upper.limits Vector of upper limits for each coefficient; default Inf. See lower.limits

maxit Maximum number of passes over the data for all lambda values; default is 10⁵.

type.gaussian Two algorithm types are supported for (only) family="gaussian". The default when nvar<500 is type.gaussian="covariance", and saves all inner-

rault when nvar<500 is type.gaussian="covariance", and saves all inner-products ever computed. This can be much faster than type.gaussian="naive", which loops through nobs every time an inner-product is computed. The latter can be far more efficient for nvar >> nobs situations, or when nvar > 500.

 $type.logistic \quad If "Newton" then the exact hessian is used (default), while "modified.Newton" \\$

uses an upper-bound on the hessian, and can be faster.

standardize.response

This is for the family="mgaussian" family, and allows the user to standardize the response variables

type.multinomial

If "grouped" then a grouped lasso penalty is used on the multinomial coefficients for a variable. This ensures they are all in our out together. The default is "ungrouped"

Details

The sequence of models implied by lambda is fit by coordinate descent. For family="gaussian" this is the lasso sequence if alpha=1, else it is the elasticnet sequence. For the other families, this is a lasso or elasticnet regularization path for fitting the generalized linear regression paths, by maximizing the appropriate penalized log-likelihood (partial likelihood for the "cox" model). Sometimes the sequence is truncated before nlambda values of lambda have been used, because of instabilities in the inverse link functions near a saturated fit. glmnet(...,family="binomial") fits a traditional logistic regression model for the log-odds. glmnet(...,family="multinomial") fits a symmetric multinomial model, where each class is represented by a linear model (on the log-scale). The penalties take care of redundancies. A two-class "multinomial" model will produce the same fit as the corresponding "binomial" model, except the pair of coefficient matrices will be equal in magnitude and opposite in sign, and half the "binomial" values. Note that the objective function for "gaussian" is

$$1/2RSS/nobs + \lambda * penalty$$
,

and for the other models it is

$$-loglik/nobs + \lambda * penalty.$$

Note also that for "gaussian", glmnet standardizes y to have unit variance before computing its lambda sequence (and then unstandardizes the resulting coefficients); if you wish to reproduce/compare results with other software, best to supply a standardized y. The latest two features in glmnet are the family="mgaussian" family and the type.multinomial="grouped" option for multinomial fitting. The former allows a multi-response gaussian model to be fit, using a "group-lasso" penalty on the coefficients for each variable. Tying the responses together like this is called "multi-task" learning in some domains. The grouped multinomial allows the same penalty for the family="multinomial" model, which is also multi-responsed. For both of these the penalty on the coefficient vector for variable j is

$$(1-\alpha)/2||\beta_i||_2^2 + \alpha||\beta_i||_2.$$

When alpha=1 this is a group-lasso penalty, and otherwise it mixes with quadratic just like elasticnet.

Value

An object with S3 class "glmnet", "*", where "*" is "elnet", "lognet", "multnet", "fishnet" (poisson), "coxnet" or "mrelnet" for the various types of models.

call the call that produced this object

a0 Intercept sequence of length length(lambda)

beta For "elnet", "lognet", "fishnet" and "coxnet" models, a nvars x length(lambda)

matrix of coefficients, stored in sparse column format ("CsparseMatrix"). For "multnet" and "mgaussian", a list of nc such matrices, one for each class.

lambda The actual sequence of lambda values used

dev.ratio The fraction of (null) deviance explained (for "elnet", this is the R-square).

The deviance calculations incorporate weights if present in the model. The deviance is defined to be 2*(loglike_sat - loglike), where loglike_sat is the log-likelihood for the saturated model (a model with a free parameter per observa-

tion). Hence dev.ratio=1-dev/nulldev.

nulldev Null deviance (per observation). This is defined to be 2*(loglike_sat -loglike(Null));

The NULL model refers to the intercept model, except for the Cox, where it is

the 0 model.

df The number of nonzero coefficients for each value of lambda. For "multnet",

this is the number of variables with a nonzero coefficient for any class.

dfmat For "multnet" and "mrelnet" only. A matrix consisting of the number of

nonzero coefficients per class

dim dimension of coefficient matrix (ices)

nobs number of observations

npasses total passes over the data summed over all lambda values

offset a logical variable indicating whether an offset was included in the model

jerr error flag, for warnings and errors (largely for internal debugging).

Author(s)

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References

Friedman, J., Hastie, T. and Tibshirani, R. (2008) Regularization Paths for Generalized Linear Models via Coordinate Descent, http://www.stanford.edu/~hastie/Papers/glmnet.pdf
Journal of Statistical Software, Vol. 33(1), 1-22 Feb 2010

http://www.jstatsoft.org/v33/i01/

Simon, N., Friedman, J., Hastie, T., Tibshirani, R. (2011) Regularization Paths for Cox's Proportional Hazards Model via Coordinate Descent, Journal of Statistical Software, Vol. 39(5) 1-13 http://www.jstatsoft.org/v39/i05/

```
Tibshirani, Robert., Bien, J., Friedman, J., Hastie, T., Simon, N., Taylor, J. and Tibshirani, Ryan. (2010) Strong Rules for Discarding Predictors in Lasso-type Problems, 
http://www-stat.stanford.edu/~tibs/ftp/strong.pdf
Stanford Statistics Technical Report
```

See Also

print, predict, coef and plot methods, and the cv.glmnet function.

Examples

```
# Gaussian
x=matrix(rnorm(100*20),100,20)
y=rnorm(100)
fit1=glmnet(x,y)
print(fit1)
coef(fit1,s=0.01) # extract coefficients at a single value of lambda
predict(fit1,newx=x[1:10,],s=c(0.01,0.005)) # make predictions
#multivariate gaussian
y=matrix(rnorm(100*3),100,3)
fit1m=glmnet(x,y,family="mgaussian")
plot(fit1m, type.coef="2norm")
#binomial
g2=sample(1:2,100,replace=TRUE)
fit2=glmnet(x,g2,family="binomial")
#multinomial
g4=sample(1:4,100,replace=TRUE)
fit3=glmnet(x,g4,family="multinomial")
fit3a=glmnet(x,g4,family="multinomial",type.multinomial="grouped")
#poisson
N=500; p=20
nzc=5
x=matrix(rnorm(N*p),N,p)
beta=rnorm(nzc)
f = x[,seq(nzc)]%*%beta
mu=exp(f)
y=rpois(N,mu)
fit=glmnet(x,y,family="poisson")
plot(fit)
pfit = predict(fit,x,s=0.001,type="response")
plot(pfit,y)
#Cox
set.seed(10101)
N=1000; p=30
nzc=p/3
x=matrix(rnorm(N*p),N,p)
beta=rnorm(nzc)
fx=x[,seq(nzc)]%*%beta/3
```

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```
hx=exp(fx)
ty=rexp(N,hx)
tcens=rbinom(n=N,prob=.3,size=1)# censoring indicator
y=cbind(time=ty,status=1-tcens) # y=Surv(ty,1-tcens) with library(survival)
fit=glmnet(x,y,family="cox")
plot(fit)
# Sparse
n=10000; p=200
nzc=trunc(p/10)
x=matrix(rnorm(n*p),n,p)
iz=sample(1:(n*p),size=n*p*.85,replace=FALSE)
x[iz]=0
sx=Matrix(x,sparse=TRUE)
inherits(sx,"sparseMatrix")\#confirm\ that\ it\ is\ sparse
beta=rnorm(nzc)
fx=x[,seq(nzc)]%*%beta
eps=rnorm(n)
y=fx+eps
px=exp(fx)
px=px/(1+px)
ly=rbinom(n=length(px),prob=px,size=1)
system.time(fit1<-glmnet(sx,y))</pre>
system.time(fit2n<-glmnet(x,y))</pre>
```

glmnet.control

internal glmnet parameters

Description

View and/or change the factory default parameters in glmnet

Usage

Arguments

fdev	minimum fractional change in deviance for stopping path; factory default = 1.0e-5
devmax	maximum fraction of explained deviance for stopping path; factory default = 0.999
eps	minimum value of lambda.min.ratio (see glmnet); factory default= 1.0e-6
big	large floating point number; factory default = 9.9e35. Inf in definition of upper.limit is set to big
mnlam	minimum number of path points (lambda values) allowed; factory default = 5
pmin	minimum null probability for any class. factory default = 1.0e-5

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exmx	maximum allowed exponent. factory default = 250.0
prec	convergence threshold for multi response bounds adjustment solution. factory default = $1.0e-10$
mxit	maximum iterations for multiresponse bounds adjustment solution. factory default = 100
factory	If TRUE, reset all the parameters to the factory default; default is FALSE

Details

If called with no arguments, glmnet.control() returns a list with the current settings of these parameters. Any arguments included in the call sets those parameters to the new values, and then silently returns. The values set are persistent for the duration of the R session.

Value

A list with named elements as in the argument list

Author(s)

```
Jerome Friedman, Trevor Hastie

Maintainer: Trevor Hastie <a href="https://doi.org/10.1007/j.jerome-friedman">https://doi.org/10.1007/j.jerome-friedman, Trevor Hastie</a>
```

See Also

glmnet

Examples

```
glmnet.control(fdev=0)#continue along path even though not much changes
glmnet.control() # view current settings
glmnet.control(factory=TRUE) # reset all the parameters to their default
```

```
plot.cv.glmnet plot the cross-validation curve produced by cv.glmnet
```

Description

Plots the cross-validation curve, and upper and lower standard deviation curves, as a function of the lambda values used.

Usage

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

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Arguments

```
x fitted "cv.glmnet" 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, and nothing is returned.

Author(s)

```
Jerome Friedman, Trevor Hastie and Rob Tibshirani Maintainer: Trevor Hastie <a href="mailto:kastie@stanford.edu">hastie@stanford.edu</a>>
```

References

Friedman, J., Hastie, T. and Tibshirani, R. (2008) Regularization Paths for Generalized Linear Models via Coordinate Descent

See Also

```
glmnet and cv.glmnet.
```

Examples

```
set.seed(1010)
n=1000; p=100
nzc=trunc(p/10)
x=matrix(rnorm(n*p),n,p)
beta=rnorm(nzc)
fx= (x[,seq(nzc)] %*% beta)
eps=rnorm(n)*5
y=drop(fx+eps)
px=exp(fx)
px=px/(1+px)
ly=rbinom(n=length(px),prob=px,size=1)
cvob1=cv.glmnet(x,y)
plot(cvob1)
title("Gaussian Family",line=2.5)
frame()
set.seed(1011)
par(mfrow=c(2,2), mar=c(4.5,4.5,4,1))
cvob2=cv.glmnet(x,ly,family="binomial")
plot(cvob2)
title("Binomial Family",line=2.5)
set.seed(1011)
cvob3=cv.glmnet(x,ly,family="binomial",type="class")
plot(cvob3)
title("Binomial Family",line=2.5)
```

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plot.glmnet	plot coefficients from a "glmnet" object
-------------	--

Description

Produces a coefficient profile plot of the coefficient paths for a fitted "glmnet" object.

Usage

```
## S3 method for class 'glmnet'
plot(x, xvar = c("norm", "lambda", "dev"), label = FALSE, ...)
## S3 method for class 'multnet'
plot(x, xvar = c("norm", "lambda", "dev"), label = FALSE,type.coef=c("coef","2norm"), ...)
## S3 method for class 'mrelnet'
plot(x, xvar = c("norm", "lambda", "dev"), label = FALSE,type.coef=c("coef","2norm"), ...)
```

Arguments

х	fitted "glmnet" 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.
label	If TRUE, label the curves with variable sequence numbers.
type.coef	If type.coef="2norm" then a single curve per variable, else if type.coef="coef", a coefficient plot per response
	Other graphical parameters to plot

Details

A coefficient profile plot is produced. If x is a multinomial model, a coefficient plot is produced for each class.

Author(s)

```
Jerome Friedman, Trevor Hastie and Rob Tibshirani Maintainer: Trevor Hastie <a href="mailto:hastie@stanford.edu">hastie@stanford.edu</a>>
```

References

Friedman, J., Hastie, T. and Tibshirani, R. (2008) Regularization Paths for Generalized Linear Models via Coordinate Descent

See Also

glmnet, and print, predict and coef methods.

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Examples

```
x=matrix(rnorm(100*20),100,20)
y=rnorm(100)
g2=sample(1:2,100,replace=TRUE)
g4=sample(1:4,100,replace=TRUE)
fit1=glmnet(x,y)
plot(fit1)
plot(fit1,xvar="lambda",label=TRUE)
fit3=glmnet(x,g4,family="multinomial")
plot(fit3,pch=19)
```

predict.cv.glmnet

make predictions from a "cv.glmnet" object.

Description

This function makes predictions from a cross-validated glmnet model, using the stored "glmnet.fit" object, and the optimal value chosen for lambda.

Usage

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

Arguments

object	Fitted "cv.glmnet" object.
newx	Matrix of new values for x at which predictions are to be made. Must be a matrix; can be sparse as in Matrix package. See documentation for predict.glmnet.
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. Alternatively s="lambda.min" can be used. 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 make a prediction.

Value

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

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Author(s)

Jerome Friedman, Trevor Hastie and Rob Tibshirani Maintainer: Trevor Hastie hastie@stanford.edu>

References

```
Friedman, J., Hastie, T. and Tibshirani, R. (2008) Regularization Paths for Generalized Linear Models via Coordinate Descent, Journal of Statistical Software, Vol. 33, Issue 1, Feb 2010 http://www-stat.stanford.edu/~hastie/Papers/glmnet.pdf http://www.jstatsoft.org/v33/i01/
```

See Also

glmnet, and print, and coef methods, and cv.glmnet.

Examples

```
x=matrix(rnorm(100*20),100,20)
y=rnorm(100)
cv.fit=cv.glmnet(x,y)
predict(cv.fit,newx=x[1:5,])
coef(cv.fit)
coef(cv.fit,s="lambda.min")
predict(cv.fit,newx=x[1:5,],s=c(0.001,0.002))
```

predict.glmnet

make predictions from a "glmnet" object.

Description

Similar to other predict methods, this functions predicts fitted values, logits, coefficients and more from a fitted "glmnet" object.

Usage

```
## S3 method for class 'glmnet'
predict(object, newx, s = NULL,
type=c("link","response","coefficients","nonzero","class"), exact = FALSE, offset, ...)
## S3 method for class 'glmnet'
coef(object,s=NULL, exact=FALSE, ...)
```

Arguments

object Fitted "glmnet" 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 Matrix package. This argument is not used for

type=c("coefficients", "nonzero")

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Value(s) of the penalty parameter lambda at which predictions are required. Default is the entire sequence used to create the model.

type Type of prediction required. Type "link" gives the linear predictors for "binomial",

"multinomial", "poisson" or "cox" models; for "gaussian" models it gives the fitted values. Type "response" gives the fitted probabilities for "binomial" or "multinomial", fitted mean for "poisson" and the fitted relative-risk for "cox"; for "gaussian" type "response" is equivalent to type "link". Type "coefficients" computes the coefficients at the requested values for s. Note that for "binomial" models, results are returned only for the class corresponding to the second level of the factor response. Type "class" applies only to "binomial" or "multinomial" models, and produces the class label corresponding to the maximum probability. Type "nonzero" returns a list of the indices of the popular coefficients for each value of s.

indices of the nonzero coefficients for each value of s.

If exact=TRUE, and predictions are to made at values of s not included in the original fit, these values of s are merged with object\$lambda, and the model is refit before predictions are made. If exact=FALSE (default), then the predict

function uses linear interpolation to make predictions for values of s that do not coincide with those used in the fitting algorithm. Note that exact=TRUE is fragile when used inside a nested sequence of function calls. predict.glmnet() needs

to update the model, and expects the data used to create it to be around.

offset If an offset is used in the fit, then one must be supplied for making predictions

(except for type="coefficients" or type="nonzero")

... Not used. Other arguments to predict.

Details

exact

The shape of the objects returned are different for "multinomial" objects. This function actually calls NextMethod(), and the appropriate predict method is invoked for each of the three model types. coef(...) is equivalent to predict(type="coefficients",...)

Value

The object returned depends on type.

Author(s)

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References

Friedman, J., Hastie, T. and Tibshirani, R. (2008) Regularization Paths for Generalized Linear Models via Coordinate Descent, http://www.stanford.edu/~hastie/Papers/glmnet.pdf Journal of Statistical Software, Vol. 33(1), 1-22 Feb 2010

http://www.jstatsoft.org/v33/i01/

Simon, N., Friedman, J., Hastie, T., Tibshirani, R. (2011) *Regularization Paths for Cox's Proportional Hazards Model via Coordinate Descent, Journal of Statistical Software, Vol. 39*(5) 1-13 http://www.jstatsoft.org/v39/i05/

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See Also

glmnet, and print, and coef methods, and cv.glmnet.

Examples

```
x=matrix(rnorm(100*20),100,20)
y=rnorm(100)
g2=sample(1:2,100,replace=TRUE)
g4=sample(1:4,100,replace=TRUE)
fit1=glmnet(x,y)
predict(fit1,newx=x[1:5,],s=c(0.01,0.005))
predict(fit1,type="coef")
fit2=glmnet(x,g2,family="binomial")
predict(fit2,type="response",newx=x[2:5,])
predict(fit2,type="nonzero")
fit3=glmnet(x,g4,family="multinomial")
predict(fit3,newx=x[1:3,],type="response",s=0.01)
```

print.glmnet

print a glmnet object

Description

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

Usage

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

Arguments

```
x fitted glmnet objectdigits significant digits in printoutadditional print arguments
```

Details

The call that produced the object x is printed, followed by a three-column matrix with columns Df, %dev and Lambda. The Df column is the number of nonzero coefficients (Df is a reasonable name only for lasso fits). %dev is the percent deviance explained (relative to the null deviance).

Value

The matrix above is silently returned

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Author(s)

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References

Friedman, J., Hastie, T. and Tibshirani, R. (2008) Regularization Paths for Generalized Linear Models via Coordinate Descent

See Also

glmnet, predict and coef methods.

Examples

```
x=matrix(rnorm(100*20),100,20)
y=rnorm(100)
fit1=glmnet(x,y)
print(fit1)
```

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