

Package ‘glmnet’

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Type Package

Title Lasso and elastic-net regularized generalized linear models

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Author Jerome Friedman, Trevor Hastie, Rob Tibshirani

Maintainer Trevor Hastie <hastie@stanford.edu>

Depends Matrix, utils

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. The algorithm uses cyclical coordinate descent in a pathwise fashion, as described in the paper listed below.

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URL <http://www.jstatsoft.org/v33/i01/>.

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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 λ . 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

```
cv.glmnet(x, y, weights, offset, lambda, type.measure, ..., nfolds, foldid, grouped)
```

Arguments

x	x matrix as in glmnet.
y	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
...	Other arguments that can be passed to glmnet.
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 misclassification 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 `nfold`s 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 `K`th 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 `K`th fold

Details

The function runs `glmnet` `nfold`s+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.

Value

an object of class "cv.glmnet" 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>cvup</code>	upper curve = <code>cvm+cvsd</code> .
<code>cvlo</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>glmnet.fit</code>	a fitted <code>glmnet</code> object for the full data.
<code>lambda.min</code>	value of <code>lambda</code> that gives minimum <code>cvm</code> .
<code>lambda.1se</code>	largest value of <code>lambda</code> such that error is within 1 standard error of the minimum.

Author(s)

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

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=1000;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)
set.seed(10101)
cvfit=cv.glmnet(x,g3,family="mult",type="mse")
plot(cvfit)
title("Multinomial Family",line=2.5)
set.seed(10101)
cvfit=cv.glmnet(x,g3,family="mult",type="class")
plot(cvfit)
# 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)

```

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 $-2 * (\text{loglike_sat} - \text{loglike})$, 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 $-2 * (\text{loglike_sat} - \text{loglike}(\text{Null}))$; The NULL model refers to the intercept model, except for the Cox, where it is the 0 model. Hence `dev.fraction=1-deviance/nulldev`, and this deviance method returns $(1-\text{dev.ratio}) * \text{nulldev}$.

Value

$(1-\text{dev.ratio}) * \text{nulldev}$

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"), weights, offset=NULL,
  alpha = 1, nlambda = 100, lambda.min.ratio = ifelse(nobs < nvars, 0.01, 0.0001), lambda=NULL,
  standardize = TRUE, thresh = 1e-07, dfmax = nvars + 1, pmax = min(dfmax * 2, nvars),
  exclude, penalty.factor = rep(1, nvars), maxit=100000, type.gaussian=ifelse(nvars < 500, "covariance", "r
```

Arguments

- 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"`)
- y** 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
- 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 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 $0 < \alpha \leq 1$. The penalty is defined as
- $$(1 - \alpha)/2 \|\beta\|_2^2 + \alpha \|\beta\|_1.$$
- `alpha=1` is the lasso 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 warm starts for speed, and its often faster to fit a whole path than compute a single fit.
standardize	Logical flag for 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.
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).
maxit	Maximum number of passes over the data for all lambda values; default is 10^5.
type.gaussian	Two algorithm types are supported for (only) family="gaussian". The default 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.

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

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

Value

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

call	the call that produced this object
a0	Intercept sequence of length length(lambda)
beta	For "elnet" and "lognet" models, a nvars x length(lambda) matrix of coefficients, stored in sparse column format ("CsparseMatrix"). For "multnet", 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 $-2*(\text{loglike_sat} - \text{loglike})$, where loglike_sat is the log-likelihood for the saturated model (a model with a free parameter per observation). Hence dev.fraction=1-dev/nulldev.
nulldev	Null deviance (per observation). This is defined to be $-2*(\text{loglike_sat} - \text{loglike}(\text{Null}))$; 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 <i>any</i> class.
dfmat	For "multnet" 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)

Jerome Friedman, Trevor Hastie and Rob Tibshirani
 Noah Simon helped develop the 'cox' feature.
 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/>
- 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

#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")

#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
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))
system.time(fit2n<-glmnet(x,y))

```

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, ...)

```

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 <hastie@stanford.edu>

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)

```

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, ...)

```

Arguments

x 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.
...	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 <hastie@stanford.edu>

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.

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,type="s",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.

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 <code>type=c("coefficients","nonzero")</code>
s	Value(s) of the penalty parameter λ 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 nonzero coefficients for each value of s .
exact	By default (<code>exact=FALSE</code>) the predict function uses linear interpolation to make predictions for values of s that do not coincide with those used in the fitting algorithm. Currently <code>exact=TRUE</code> is not implemented, but prints an error message telling the user how to achieve the exact predictions. This is done by rerunning the algorithm with the desired values interspersed (in order) with the values used in the original fit. This is easily achieved via the R command <code>lambda=sort(c(object\$lambda, new.lambda))</code>

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

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)

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/>

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 object
digits	significant digits in printout
...	additional 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

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 and coef methods.

Examples

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

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