

# An Overview of glmnet

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## The Package

The `nested.glmnet()` function of the ‘glmnet’ package allows the user to fit multiple machine learning models on a common data set with a single function call allowing an efficient comparison of different modeling approaches. Additionally this function uses cross validation (CV) or a bootstrap method to estimate model performances for these different modeling approaches from the hold out (out of bag) sample partitions. As most of these machine learning models choose hyperparameters informed by a cross validation or some sort or out of bag (OOB) performance measure, the `nested.glmnet()` function provides model performance estimates based upon either a nested cross validation (NCV), bootstrapping or analogous approach. Measures of model performance include concordances for survival time and binomial outcomes and R-squares for quantitative numeric outcomes, as well as deviance ratios, i.e.  $1 - \text{deviance}(\text{model}) / \text{deviance}(\text{nullmodel})$ , and linear calibration coefficients. Too often one sees performance reports including things like sensitivity, specificity or F1 scores in absence of any consideration of calibration. In addition to providing linear calibration coefficient estimates, we also show spline fits on the hold out datasets of the nested cross validation outer folds. The `nested.glmnet()` function also fits the respective models on the whole dataset. Performance measures and fitted models based upon the whole dataset are stored in a single output object.

The `nested.glmnet()` function fits cross validation informed lasso, relaxed lasso, ridge, gradient boosting machine (‘xgboost’), Random Forest (‘RandomForestSRC’), Oblique Random Forest (‘aorsf’), Artificial Neural Network (ANN), Recursive Partitioning and Regression Trees (‘RPART’) and step wise regression models. As run times may be long, the user specifies which of these models to fit. By default only the lasso model suite is fit, including the (standard) lasso, relaxed lasso, fully relaxed lasso ( $\text{gamma}=0$ ) and the ridge regression models. (The program was originally written to simply compare the lasso and stepwise regression models and thus this inclusion of the lasso by default, as well as the program name.) By default model performances are calculated using cross validation but if the goal is to only fit the models this can be done using the option `resample=0`.

As with the ‘glmnet’ package, tabular and graphical summaries can be generated using the `summary` and `plot` functions. Use of the ‘glmnet’ package has many similarities to the ‘glmnet’ package and the user may benefit by a review of the documentation for the ‘glmnet’ package <https://cran.r-project.org/package=glmnet>, with the “An Introduction to ‘glmnet’” and “The Relaxed Lasso” being especially helpful in this regard.

For some data sets, for example when the design matrix is not of full rank, ‘glmnet’ may have very long run times when fitting the relaxed lasso model, from our experience when fitting Cox models on data with many predictors and many patients, making it difficult to get solutions from either `glmnet()` or `cv.glmnet()`. This may be remedied with the ‘`path=TRUE`’ option when calling `cv.glmnet()`. In the ‘glmnet’ package we always take an approach like that of `path=TRUE`.

When fitting not a relaxed lasso model but an elastic-net model, the R-packages ‘`nestedcv`’ <https://cran.r-project.org/package=nestedcv>, ‘`glmnetSE`’ <https://cran.r-project.org/package=glmnetSE> or others may provide greater functionality when performing a nested CV.

## Installing glmnet

Installing glmnet is much like installing other R packages, but with a small wrinkle. As usual one submits

```
install.packages( 'glmnet' )
```

and then loads the package from the library

```
library( glmnet )
```

So far this is like with most packages. However, when loading the package for the first time one may be prompted to load further software needed to run the torch library for the neural network models, for example as in

```
> library( glmnet )  
i Additional software needs to be downloaded and installed for torch to work correctly.  
Do you want to continue? (Yes/no/cancel)
```

where response Yes should allow neural network model fitting. A no answer will not install torch, and if not already installed then attempting to run the neural network models will lead to crashes.

## Data requirements

The basic data elements for input to the *glmnet* analysis programs are similar to those of *glmnet* and include 1) a matrix of predictors and 2) an outcome variable in vector form. For the different machine learning modeling approaches the package is set up to model generalizations of the Cox proportional hazards survival model, the “binomial” outcome logistic model and linear regression with independent identically distributed errors amenable to being treated as if “gaussian”. When fitting the Cox model the outcome model variable is interpreted as the “time” variable, and one must also specify 3) a variable for event, again in vector form, and optionally 4) a variable for start time, also in vector form. Row *i* of the predictor matrix and element *i* of the outcome vector(s) are to include the data for the same sampling unit.

The input vectors may optionally be specified as column matrices (with only one column each) in which case the column name will be kept and expressed in the model summaries.

## An example data set

To demonstrate usage of *glmnet* we first generate a data set for analysis, run an analysis and evaluate using the `plot()`, `summary()` and `predict()` functions.

The code

```
# Simulate data for use in an example for relaxed lasso fit of survival data  
# First, optionally, assign a seed for random number generation to get replicable results  
set.seed(116291949)  
simdata=glmnet.simdata(nrows=1000, ncols=100, beta=NULL)
```

generates simulated data for analysis. We extract data in the format required for input to the *glmnet* programs.

```

# Extract simulated survival data
xs = simdata$xs      # matrix of predictors
y_ = simdata$y_     # vector of Gaussian (normal) outcomes
yb = simdata$yb     # vector of binomial outcomes
yt = simdata$yt     # vector of survival times
event = simdata$event # indicator of event vs. censoring

```

Inspecting the predictor matrix we see

```

# Check the sample size and number of predictors
cat(dim(xs))

```

```
## 1000 100
```

```

# Check the rank of the design matrix, i.e. the degrees of freedom in the predictors
# using function from the Matrix package
Matrix::rankMatrix(xs)[[1]]

```

```
## [1] 94
```

```

# Inspect the first few rows and some select columns
print(xs[1:10,c(1:12,18:20)])

```

```

##      X1 X2 X3 X4 X5 X6 X7 X8 X9 X10 X11 X12      X18      X19      X20
## [1,]  1  1  0  0  0  0  0  0  0  1  0  1  0.1513225 -0.4034383  0.35250844
## [2,]  1  0  0  0  1  0  0  1  0  0  0  0 -1.1610480  0.5533030  0.14578868
## [3,]  1  0  0  1  0  0  1  0  0  0  0  0 -0.3292269  0.3086399 -0.48443836
## [4,]  1  1  0  0  0  0  0  0  0  1  0  0  2.0635214 -0.5500741 -0.02173104
## [5,]  1  0  0  0  1  0  0  1  0  0  0  0  0.3905722 -0.6836452 -0.37643201
## [6,]  1  0  1  0  0  0  0  0  1  0  0  0 -0.2397597  1.6909447  0.49599945
## [7,]  1  0  1  0  0  0  0  1  0  0  0  0 -0.5592424  0.2314638 -0.53198341
## [8,]  1  0  0  1  0  0  0  0  0  0  1  0 -1.0050514  0.5319574  0.54287646
## [9,]  1  0  0  1  0  0  0  0  0  0  1  0  1.2548034  0.8213164  0.17067691
## [10,] 1  0  0  0  1  0  0  0  1  0  0  0 -0.3079151 -0.6105910 -0.88711869

```

## Performance of cross validation (CV) informed relaxed lasso model

Because the values for lambda and gamma informed by CV are specifically chosen to give a best fit, model fit statistics for the CV informed model, when based upon the same train data used to derive the model, will be biased. (Using the common terminology of machine learning, both the “training” and “validation” data inform the model fit.) To address this one can perform a CV on the CV derived estimates, that is a nested cross validation as argued for in SRDM ( Simon R, Radmacher MD, Dobbin K, McShane LM. Pitfalls in the Use of DNA Microarray Data for Diagnostic and Prognostic Classification. J Natl Cancer Inst (2003) 95 (1): 14-18. <https://academic.oup.com/jnci/article/95/1/14/2520188> ). For this second layer of CV, there is no usage of information from the hold out data back to the model fit. (Using the common terminology of machine learning, each of the hold-out subsets of this outer layer of CV is treated as a “test” data set for calculation of model performance, and the results combined across these multiple hold out “test” sets.) We demonstrate the model performance evaluation by nested cross validation first for the lasso models with the evaluation of other machine learning models being similar. For this performance evaluation we use the nested.glmnetr() function which first fits all models based upon all data and then performs the cross validation for calculation of concordances or R-squares, deviance ratios and linear calibration summaries.

```

set.seed(465783345)
nested.cox.fit = nested.glmnet(xs, NULL, yt, event, family="cox",
                              dolasso=1, dostep=1, steps_n=40, folds_n=10, track=1)

```

Note, in the derivation of the relaxed lasso model fits, individual coefficients may be unstable even when the model may be stable which elicits warning messages. We suppress these warnings here. The first term in the call to `nested.glmnet()`, `xs`, is the design matrix for predictors. The second input term, here given the value `NULL`, is for the start time in case the (start, stop) time data setup is used in a Cox survival model fit. The third term, here `yt`, is the outcome variable for the linear regression or logistic regression model and the time of event or censoring in case of the Cox model, and finally the fourth term is the event indicator variable for the Cox model taking the value 1 in case of an event or 0 in case of censoring at time `yt`. The fourth term would be `NULL` for either linear or logistic regression. If one sets `track=1` the program will update progress in the R console, else for `track=0` it will not. We recommend setting `track=1` when running the program interactively. Depending on the size of the data set and the different machine learning models fit, run time can be long and it can be helpful to view the progress in calculations.

As usual with R functions and packages we use the summary function to describe output. Here the summary function displays a brief summary of the input data before proceeding to describe model performances. The data summary includes sample size, number of events, number of candidate model predictors, degrees of freedom in these predictors as well as average deviance and some average minus 2 log likelihoods. Model performances are displayed for the different lasso models, e.g. standard, relaxed, fully relaxed as well as the ridge regression and stepwise regression models. Hyperparameters considered for stepwise regression are degrees of freedom (df) and p, the p-value for entry into the regression equation, as discussed by JWHT (James, Witten, Hastie and Tibshirani, An Introduction to Statistical Learning with applications in R, 2nd ed., Springer, New York, 2021). Performance measures include deviance ratio, linear calibration coefficients and measures of agreement, here for the Cox model framework concordance. Additionally there are the deviance ratio and agreement values naively calculated on the whole data set.

```

# Summarize relaxed lasso model performance informed by cross validation
summary(nested.cox.fit, width=84)

```

```

## Sample information including number of records, events, number of columns in
## design (predictor, X) matrix, and df (rank) of design matrix:
##           family           n           nevent
##           cox             1000           698
##           xs.columns       xs.df       null.dev/nevent
##           100              94           12.43
## null.m2LogLik/nevent  sat.m2LogLik/nevent
##           12.43              0
##
## For LASSO, and Stepwise regression tuned by df and p, average (Ave) model
## performance measures from the 10-fold (NESTED) Cross Validation are given together
## with naive summaries calculated using all data without cross validation
##
##           Ave DevRat Ave Slope Ave Concordance Ave Non Zero
## LASSO min           0.2446   1.0742           0.8705           45.4
## LASSO minR          0.2434   0.9790           0.8701           17.6
## LASSO minR.GO       0.2410   0.9320           0.8691           16.2
## Ridge              0.2244   1.2850           0.8626           99.0
##
##           Naive DevRat Naive Concordance Non Zero
## LASSO min           0.1696           0.8794           42
## LASSO minR          0.1720           0.8793           22
## LASSO minR.GO       0.1728           0.8796           20

```

```

## Ridge                0.1718                0.8822                99
##
##                Ave DevRat Ave Slope Ave Concordance Ave Non Zero
## Stepwise df tuned    0.2501    0.9590                0.8738                14.7
## Stepwise p tuned     0.2527    0.9677                0.8747                14.8
##                Naive DevRat Naive Concordance Non Zero
## Stepwise df tuned    0.1711                0.8785                15
## Stepwise p tuned     0.1711                0.8785                15

```

Here we see, not unexpectedly, that the concordances estimated from the nested CV are smaller than the concordances naively calculated using original data set. Depending on the data the nested CV and naive agreement measures, here concordance, can be very similar or disparate. Possibly surprisingly the deviance ratios are larger for the (nested) cross validation than naively calculated using all data. This flip in direction has not to do with the strength of association being stronger in the hold out data, but from the hold out data set being smaller and how this impacts average risk set size in the partial likelihood of the “cox” model. Such a flip in direction will generally not be the case for “binomial” and “gaussian” data. Despite this flip, the value of the cross validation estimated deviance ratios for comparing the machine learning models remains.

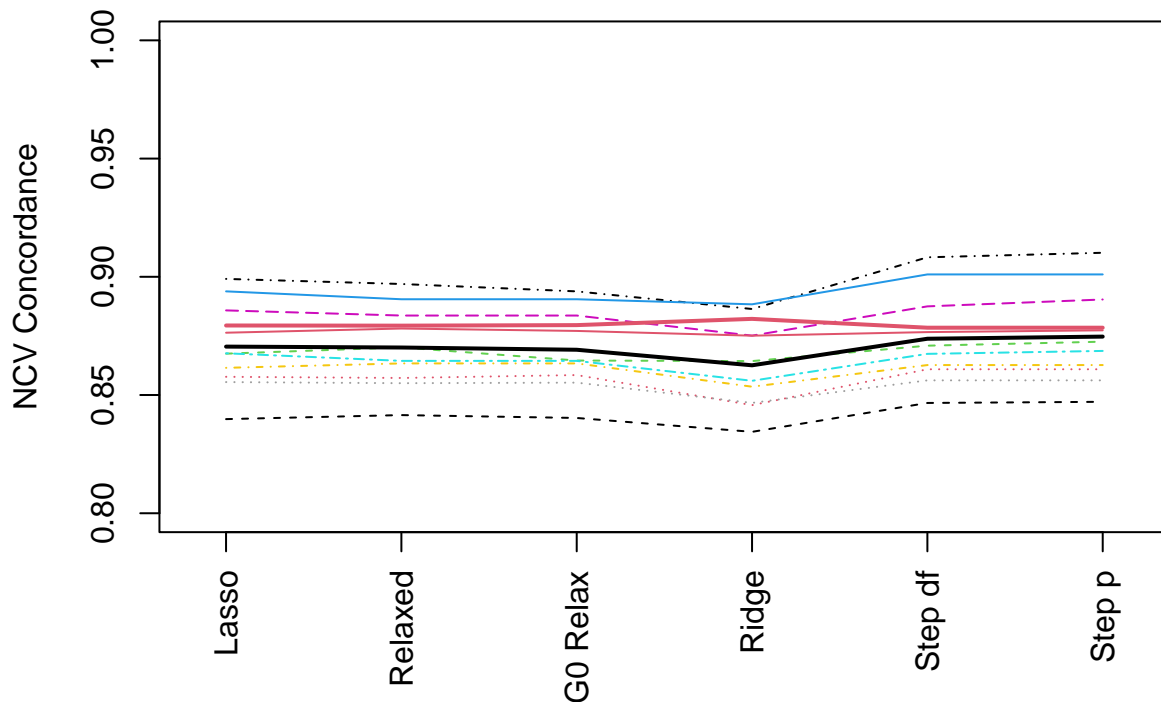
From this output we also see the number of non-zero coefficients in the different models, reflecting model complexity at least for the lasso model, along with the linear calibration coefficients obtained by regressing the outcome on the predicted, i.e. the XBeta or the log(hazard ratio). (Many machine learning fitting routines use as predicted the hazard ratio for “cox” model generalizations or the probability for “binomial” data. We give predicted as the XBeta term which applies to the different data types. For the “binomial” case this is  $\log(P/(1-P))$  where P is the predicted probability.)

Model performance measures from the nested cross validation (NCV) can also be visualized with a plot which shows the calculated performances for the individual folds of the cross validation.

```

# Summarize relaxed lasso model performance informed by cross validation
plot(nested.cox.fit, type="agree", ylim=c(0.8,1.0))

```



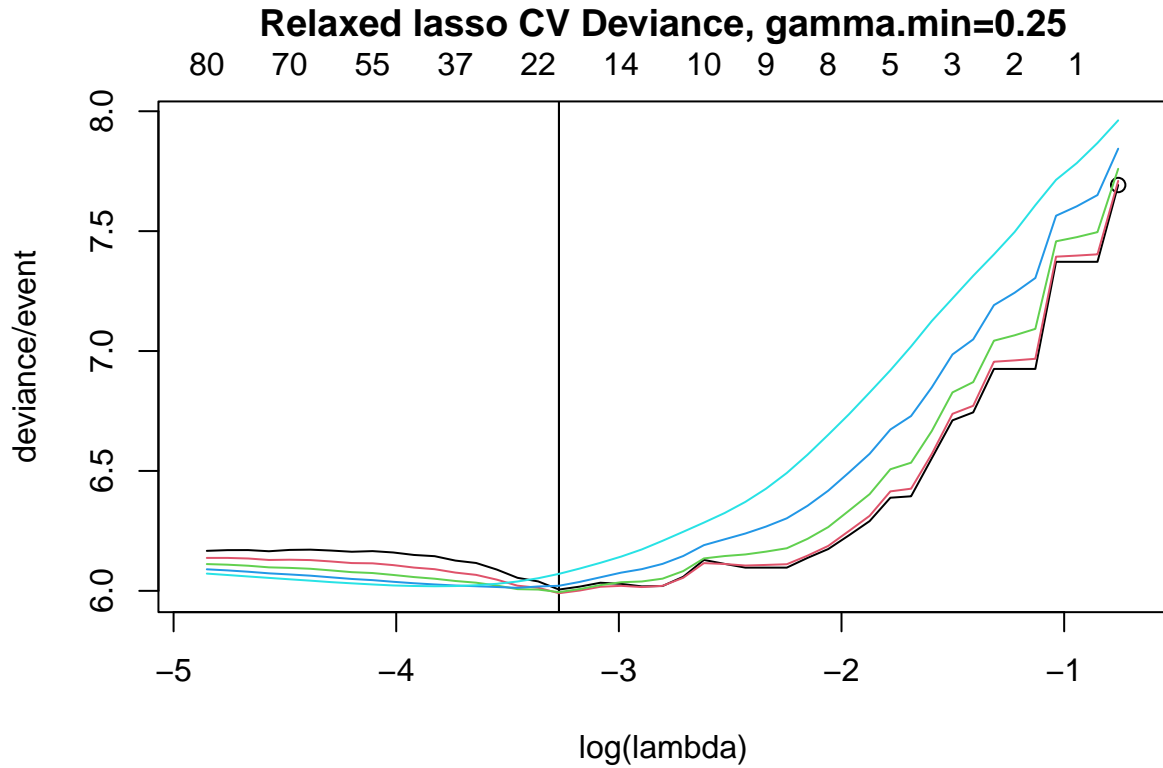
The performance measure estimates, here of concordance, from the individual (outer) cross validation for each fold are depicted by thin lines of different colors and styles, while the composite value from all folds is depicted by a thicker black line, and the performance estimates naively calculated on all data, the same as the data used for model derivation, are depicted in a thicker red line. Here we see that the performance measures for the different models are quite variable across the folds, yet highly correlated with each other. Also, as expected the concordance for the model derived using the the full data set, naively calculated on the same data (i.e. the full data set) as used in model derivation, are larger than the average of the concordances from the different folds. Plots can also be constructed using option “devrat” for deviance ratios, “intcal” for linear calibration intercept coefficients and “lincal” for linear calibration slope coefficients.

## The CV informed relaxed lasso model fit

As mentioned, the `nested.glmnet()` function also derives the models using all data and stores these for further examination and usage. The choice of `lambda` and `gamma` hyperparameters for the relaxed lasso model is based upon a search across two dimensions for the pair that maximizes the likelihood, or similarly minimizes the deviances, as informed by a cross validation. The next plot depicts the deviances across these two dimensions for the full data set.

```
# Plot cross validation average deviances for a relaxed lasso model
plot(nested.cox.fit, type="lasso")
```

```
## min CV average deviance (max log likelihood) for
## relaxed at log(lambda) = -3.269, gamma.min = 0.25, df = 20
## fully relaxed at log(lambda) = -3.269, df = 20
## fully penalized at log(lambda) = -3.827, df = 42
```

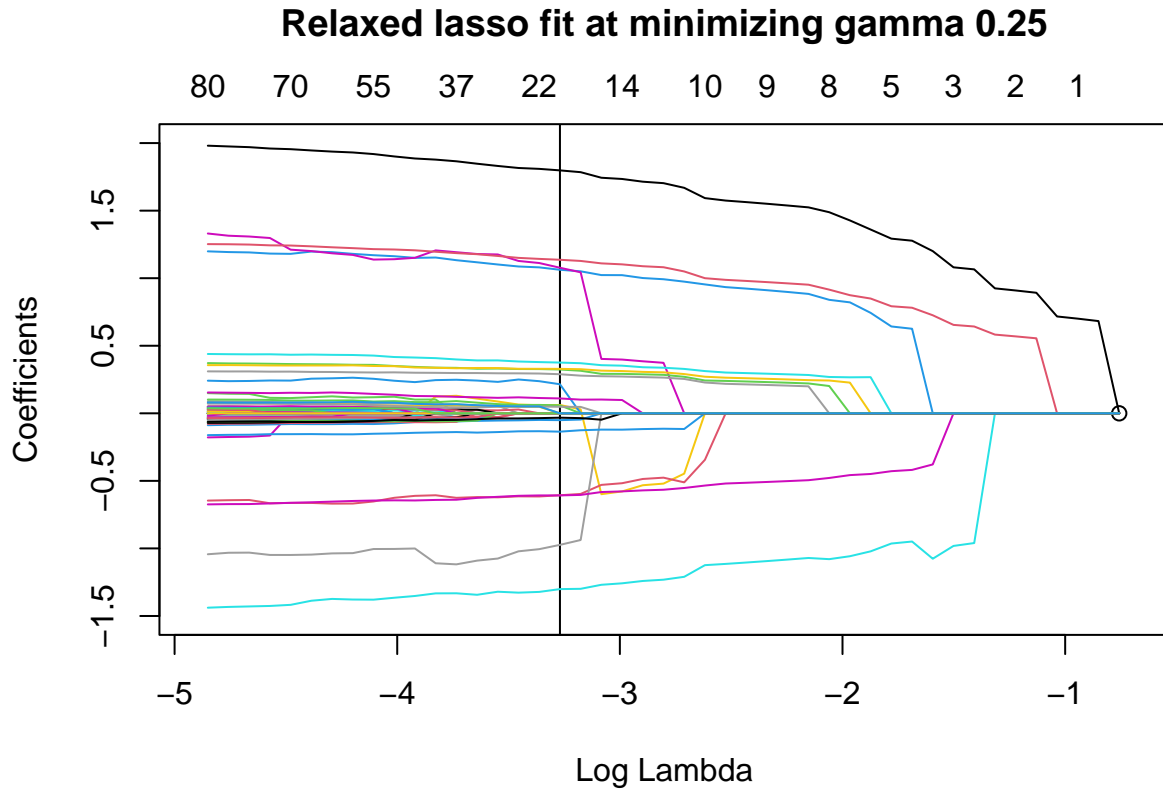


This figure has multiple lines, depicting deviance as a function of lambda for different gamma values. Whereas there is no legend here for gamma, when non-zero coefficients start to enter the model as the penalty is reduced, here shown at the right, deviances tend to be smaller for gamma = 0, greater for gamma = 1 and in between for other gammas values. The minimizing lambda and gamma pair is indicated by the left most vertical line, here about  $\log(\lambda) = -3.27$ . The minimizing lambda can be read from the horizontal axis. Because the different lines depicting deviances for the different values of gamma can be nearly overlapping, the minimizing gamma is described in the title, here 0.25. From this figure we also see that at  $\log(\lambda) = -3.27$  the deviance is hardly distinguishable for gamma ranging from 0 to 0.5. More relevant we see that the fully unpenalized lasso model fits (gamma=0) shown in a black line with a black circle at the largest lambda, achieves a minimal deviance at about  $\log(\lambda) = -3.27$ , and highlighted by the right most vertical line. The minimizing deviance for the fully relaxed lasso model is nearly that of the relaxed lasso model tuning for both lambda and gamma.

A plot depicting model fits as a function of lambda is given in the next figure.

```
# Plot coefficients informed by a cross validation  
plot(nested.cox.fit, type="coef")
```

```
## min CV average deviance (max log likelihood)  
## at log(lambda.min) = -3.269, gamma.min = 0.25, df = 20
```

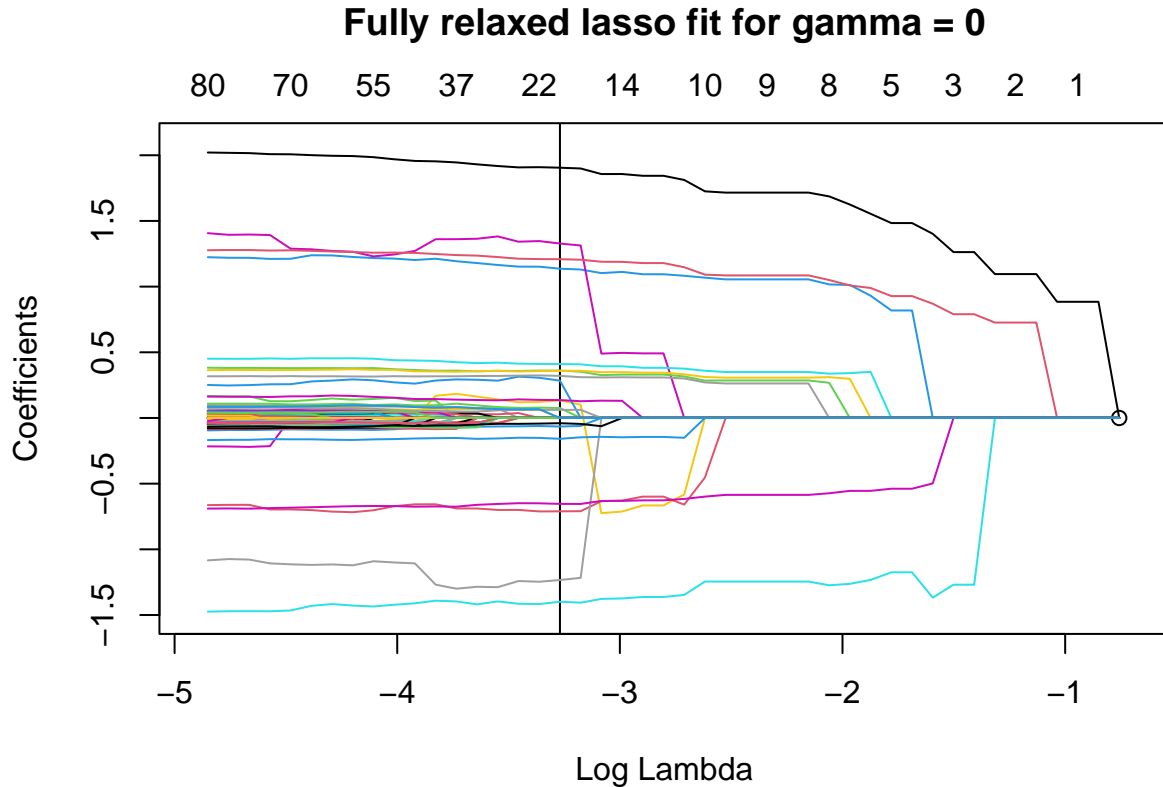


In this plot of coefficients we use the same orientation for lambda as in the plot for deviances with larger values of the lambda penalty to the right and corresponding to fewer non-zero coefficients. The displayed coefficients are for the minimizing gamma=0.25 as noted in the title, and the minimizing lambda indicated by the vertical line. Since the fully relaxed lasso model had a deviance almost that of the relaxed lasso model we also plot the coefficients using the option gam=0.



```
# Plot fully relaxed coefficients informed by a cross validation
plot(nested.cox.fit, type="coef", gam=0)
```

```
## Fully relaxed min CV average deviance (max log likelihood)
## at log(lambda.min) = -3.269, df = 20
```



This plot shows how the coefficients change for the un-penalized (fully relaxed) model with  $\gamma=0$  as lambda decreases. In particular we see the coefficients become slightly larger in magnitude as the lambda penalty decreases and also as additional terms come into the model. This is not unexpected as omitted terms from the Cox model tend to bias coefficients toward 0 more than increase the standard error. We also see, as too indicated in the deviance plot, the number of model non-zero coefficients, 20, to be substantially less than the 20 from the relaxed lasso fit and the 42 from the fully penalized lasso fit.

A summary of the actual lasso model fit can be gotten by using the `cvfit=1` option in the `summary()` call.

```
# Summarize relaxed lasso model fit informed by cross validation
summary(nested.cox.fit, cvfit=1)
```

```
## The relaxed minimum is obtained for lambda = 0.03803792 and gamma = 0.25
## with df (number of non-zero terms) = 20, average deviance = 5.990078 and beta =
## X4 X5 X7 X10 X12
## 1.062428e+00 -1.301185e+00 6.180743e-02 -6.065663e-01 2.150892e-01
## X14 X15 X16 X17 X18
## 1.077432e+00 -3.337106e-16 -9.737055e-01 2.697506e-17 1.136458e+00
## X19 X20 X21 X22 X23
## 3.242268e-01 -1.359473e-01 3.755073e-01 -6.075709e-01 3.290318e-01
```

```

##           X24           X25           X38           X43           X60
## 2.893126e-01 1.797959e+00 1.099952e-01 5.828904e-02 -5.064047e-02
##           X88           X97
## 4.837592e-02 -3.240492e-02
##
## The fully relaxed (gamma=0) minimum is obtained at lambda = 0.03803792
## with df (number of non-zero terms) = 20, average deviance = 6.005437 and beta =
##           X4           X5           X7           X10           X12           X14
## 1.13641535 -1.39885327 0.13367973 -0.71086068 0.28432797 1.32787050
##           X16           X18           X19           X20           X21           X22
## -1.23342009 1.20856642 0.36062709 -0.15870146 0.41060234 -0.65415048
##           X23           X24           X25           X38           X43           X60
## 0.35935105 0.32055505 1.90516232 0.13629323 0.07577501 -0.06497194
##           X88           X97
## 0.06139717 -0.03978756
##
## The UNrelaxed (gamma=1) minimum is obtained at lambda = 0.02176669
## with df (number of non-zero terms) = 42, average deviance = 6.019153
##
## Order coefficients entered into the lasso model (1st to last):
## [1] "X25" "X18" "X5" "X22" "X4" "X21" "X23" "X19" "X24" "X10"
## [11] "X7" "X20" "X14" "X38" "X97" "X16" "X60" "X88" "X12" "X43"
## [21] "X71" "X100" "X34" "X32" "X50" "X58" "X41" "X49" "X64" "X84"
## [31] "X91" "X98" "X39" "X40" "X66" "X73" "X74" "X11" "X61" "X69"
## [41] "X70" "X96"

```

In the summary output we first see the relaxed lasso model fit based upon the (lambda, gamma) pair which minimizes the cross validated average deviance. Next is the model fit based upon the lambda that minimizes the cross validated average deviance along the path where gamma=0, that is among the fully relaxed lasso models. After that is information on the fully penalized lasso fit, but without the actual coefficient estimates. These estimates can be printed using the option `printg1=TRUE`, but are suppressed by default for space. Finally, the order that coefficients enter the lasso model as the penalty is decreased is provided, which gives some indication of relative model importance of the coefficients. Because, though, the differences in successive lambda values used in the numerical algorithms may allow multiple new terms to enter into the model between successive numerical steps, the ordering in this list may not be strict. If the user would want they could read lambda from `output$lambda`, set up a new lambda with finer steps and rerun the model. Our experience though is that this does not generally lead to a meaningfully different model and so is not done by default or as option.

One can use the `predict()` function to get the coefficients for the lasso model, which is done by not specifying a predictor matrix. If one specifies a new design predictor matrix `xs_new`, then the predicted `xs_new*beta` are generated. When used to extract the model coefficients, the `predict()` function provides an output object in vector form (actually a list with two vectors) and so can easily be used for further calculations. By default the `predict()` function will use the (lambda, gamma) pair that minimizes the average CV deviances. One can also specify `lam=NULL` and `gam=1` to use the fully penalized lasso best fit, i.e. select the lambda that minimizes the CV deviance while holding gamma=1, or `gam=0` to use the fully relaxed lasso best fit, that is minimizes while holding gamma=0. One can also numerically specify both lam for lambda and gam for gamma. Within the package lambda and gamma usually denote vectors for the search algorithm and so other names are used in this function.

```

# Get coefficients
beta = predict(nested.cox.fit)

```

```

## (lambda, gamma) pair minimizing CV average deviance is used

```

```
# Print out the non-zero coefficients
beta$beta
```

```
##          X4          X5          X7          X10          X12
## 1.062428e+00 -1.301185e+00  6.180743e-02 -6.065663e-01  2.150892e-01
##          X14          X15          X16          X17          X18
## 1.077432e+00 -3.337106e-16 -9.737055e-01  2.697506e-17  1.136458e+00
##          X19          X20          X21          X22          X23
## 3.242268e-01 -1.359473e-01  3.755073e-01 -6.075709e-01  3.290318e-01
##          X24          X25          X38          X43          X60
## 2.893126e-01  1.797959e+00  1.099952e-01  5.828904e-02 -5.064047e-02
##          X88          X97
## 4.837592e-02 -3.240492e-02
```

```
# Print out all coefficients
beta$beta_
```

```
##          X1          X2          X3          X4          X5
## 0.000000e+00 0.000000e+00 0.000000e+00 1.062428e+00 -1.301185e+00
##          X6          X7          X8          X9          X10
## 0.000000e+00 6.180743e-02 0.000000e+00 0.000000e+00 -6.065663e-01
##          X11          X12          X13          X14          X15
## 0.000000e+00 2.150892e-01 0.000000e+00 1.077432e+00 -3.337106e-16
##          X16          X17          X18          X19          X20
## -9.737055e-01 2.697506e-17 1.136458e+00 3.242268e-01 -1.359473e-01
##          X21          X22          X23          X24          X25
## 3.755073e-01 -6.075709e-01 3.290318e-01 2.893126e-01 1.797959e+00
##          X26          X27          X28          X29          X30
## 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00
##          X31          X32          X33          X34          X35
## 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00
##          X36          X37          X38          X39          X40
## 0.000000e+00 0.000000e+00 1.099952e-01 0.000000e+00 0.000000e+00
##          X41          X42          X43          X44          X45
## 0.000000e+00 0.000000e+00 5.828904e-02 0.000000e+00 0.000000e+00
##          X46          X47          X48          X49          X50
## 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00
##          X51          X52          X53          X54          X55
## 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00
##          X56          X57          X58          X59          X60
## 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00 -5.064047e-02
##          X61          X62          X63          X64          X65
## 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00
##          X66          X67          X68          X69          X70
## 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00
##          X71          X72          X73          X74          X75
## 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00
##          X76          X77          X78          X79          X80
## 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00
##          X81          X82          X83          X84          X85
## 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00
##          X86          X87          X88          X89          X90
## 0.000000e+00 0.000000e+00 4.837592e-02 0.000000e+00 0.000000e+00
```

```
##           X91           X92           X93           X94           X95
## 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00 0.000000e+00
##           X96           X97           X98           X99           X100
## 0.000000e+00 -3.240492e-02 0.000000e+00 0.000000e+00 0.000000e+00
```

```
# Get the predicted (linear predictors) for the original data set
```

```
predicted = predict(nested.cox.fit, xs)
```

```
## (lambda, gamma) pair minimizing CV average deviance is used
```

```
# Print out the first few predicted
```

```
predicted[1:20]
```

```
## [1] -0.4873661 -3.5150833 4.3306965 1.3514012 -0.1708235 1.3846970
## [7] -3.7546765 0.2763778 6.1528498 1.5709802 1.1229711 -1.7437172
## [13] 1.1030609 2.9132104 -0.5753324 0.5351649 0.8883025 3.3120748
## [19] -2.4726732 1.9326001
```

## Nested cross validation (NCV) for multiple models

Here we evaluate multiple machine learning models, in particular the lasso, ridge, XGB, random forest and neural network models. For this example we perform an analysis for the generalizations of linear regression in contrast to the Cox model in the last example. The `glmnet.simdata()` function used above actually creates an output object list containing `xs` for the predictor matrix, `yt` for time to event or censoring and an event indicator, as well as `y_` for “gaussian” and `yb` for “binomial” data.

```
# Nested cross validation evaluating a machine learning model suite with gaussian errors
# Use the same simulated data output object from above, that is from the call
# simdata=glmnet.simdata(nrows=1000, ncols=100, beta=NULL)
#
```

```
# recall linear regression model data generated above with line
# y_ = simdata$y_ # outcome vector with Gaussian (normal) errors
# Get the ML model fits
nested.gau.fit = nested.glmnet(xs=NULL,y_,NULL,family="gaussian",
                             dolasso=1, doxgb=list(nrounds=250), dorf=1, doorf=1, doann=list(bestof=10),
                             folds_n=10, seed=219301029, track=1)
```

```
# Summarize the results
```

```
summary(nested.gau.fit, width=84)
```

```
## Sample information including number of records, number of columns in
## design (predictor, X) matrix, and df (rank) of design matrix:
## family n xs.columns xs.df null.dev/n
## gaussian 1000 100 94 8.09
##
## For LASSO, gradient boosting machine using XGBoost (XGB), Random Forest (RF),
## Oblique Random Forest (ORF), Artificial Neural Networks (ANN), average (Ave) model
## performance measures from the 10-fold (NESTED) Cross Validation are given together
## with naive summaries calculated using all data without cross validation
##
```

```

## Ave DevRat Ave Int Ave Slope Ave R-square Ave Non Zero
## LASSO min 0.8613 -0.0069 1.0211 0.8641 54.7
## LASSO minR 0.8605 -0.0039 1.0150 0.8630 36.4
## LASSO minR.GO 0.8570 0.0171 0.9903 0.8594 24.0
## Ridge 0.8485 -0.0724 1.1046 0.8581 99.0
## Naive DevRat Naive R-square Non Zero
## LASSO min 0.8794 0.9380 65
## LASSO minR 0.8753 0.9357 26
## LASSO minR.GO 0.8695 0.9325 14
## Ridge 0.8806 0.9400 99
##
## Ave DevRat Ave Int Ave Slope Ave R-square Ave Non Zero
## XGB (not tuned) 0.6782 -0.1294 1.1281 0.6891 100
## XGB Tuned 0.8179 -0.0590 1.0717 0.8250 100
## Naive DevRat Naive R-square Non Zero
## XGB (not tuned) 0.9992 0.9993 100
## XGB Tuned 0.9317 0.9349 100
##
## Ave DevRat Ave Int Ave Slope Ave R-square Ave Non Zero
## RF Simple 0.7075 -0.2147 1.214 0.7338 58
## Naive DevRat Naive R-square Non Zero
## RF Simple 0.9267 0.9513 50
##
## Ave DevRat Ave Int Ave Slope Ave R-square Ave Non Zero
## ORF Simple 0.6702 -0.7466 1.6941 0.8116 35
## Naive DevRat Naive R-square Non Zero
## ORF Simple 0.8558 0.9579 40
##
## Ave DevRat Ave Int Ave Slope Ave R-square Ave Non Zero
## ANN Uninformed 0.671 0.0625 0.9557 0.6716 100
## Naive DevRat Naive R-square Non Zero
## ANN Uninformed 0.9803 0.9803 100

```

Here we see a set of machine learning models evaluated together. All evaluations are based upon the same folds for the outer loop of the cross validation. Those models informed by cross validation in identification of hyperparameters, i.e. lasso, XGB, neural network and stepwise, use the same folds in the inner cross validation making the comparisons of model performance between models more stable. For the models based upon other random splittings, i.e. random forest (and sometimes the XGB), the same seed is set using `set.seed()` before each model call facilitating replicability of results. The “Non Zero” column is the number of non zero regression coefficients for the lasso and ridge models, and the number of predictors given to the XGB model. For the random forest model “Non Zero” is the number of predictors randomly selected for possible splitting at each node, a tuned hyperparameter. For the ANN models “non Zero” again is the number of terms given to the ANN for training.

One can save the tabled information from a summary to a data frame using the `table` option as in

```

## put the data into a data frame
dframe = summary( nested.gau.fit , table = 0)
## use the roundperf() function from 'glmnet' to round so long as it doesn't too
## obscure the original values
## print just the first 4 columns to get a more succinct table
roundperf(dframe, digits = 3)[,c(1:4)]

```

```

## Ave DevRat Ave Int Ave Slope Ave R-square

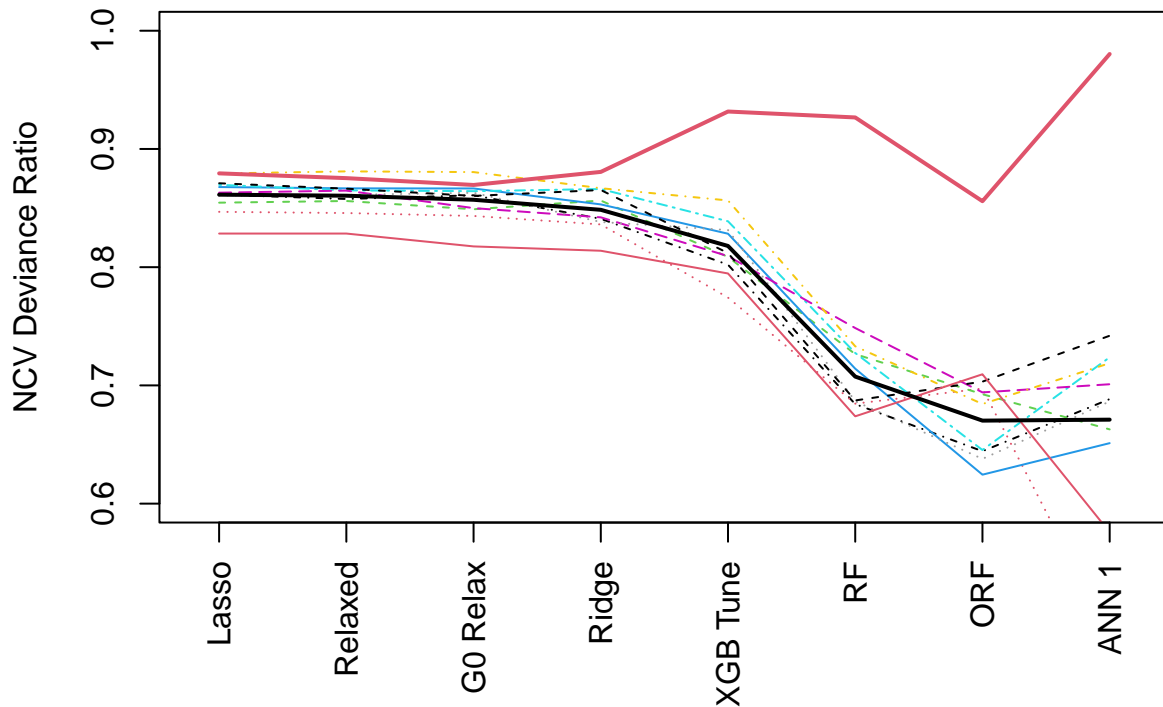
```

## LASSO min	0.861	-0.007	1.021	0.864
## LASSO minR	0.860	-0.004	1.015	0.863
## LASSO minR.GO	0.857	0.017	0.990	0.859
## Ridge	0.849	-0.072	1.105	0.858
## XGB (not tuned)	0.678	-0.129	1.128	0.689
## XGB Tuned	0.818	-0.059	1.072	0.825
## RF Simple	0.707	-0.215	1.214	0.734
## ORF Simple	0.670	-0.747	1.694	0.812
## ANN Uninformed	0.671	0.062	0.956	0.672

This may be helpful when wanting to further process the findings or incorporate into reports.

Model performance measures from the nested cross validation can be plotted individually as shown here for deviance ratio

```
# Summarize relaxed lasso model performance informed by cross validation
plot(nested.gau.fit, type="devrat", ylim=c(0.6,1))
```

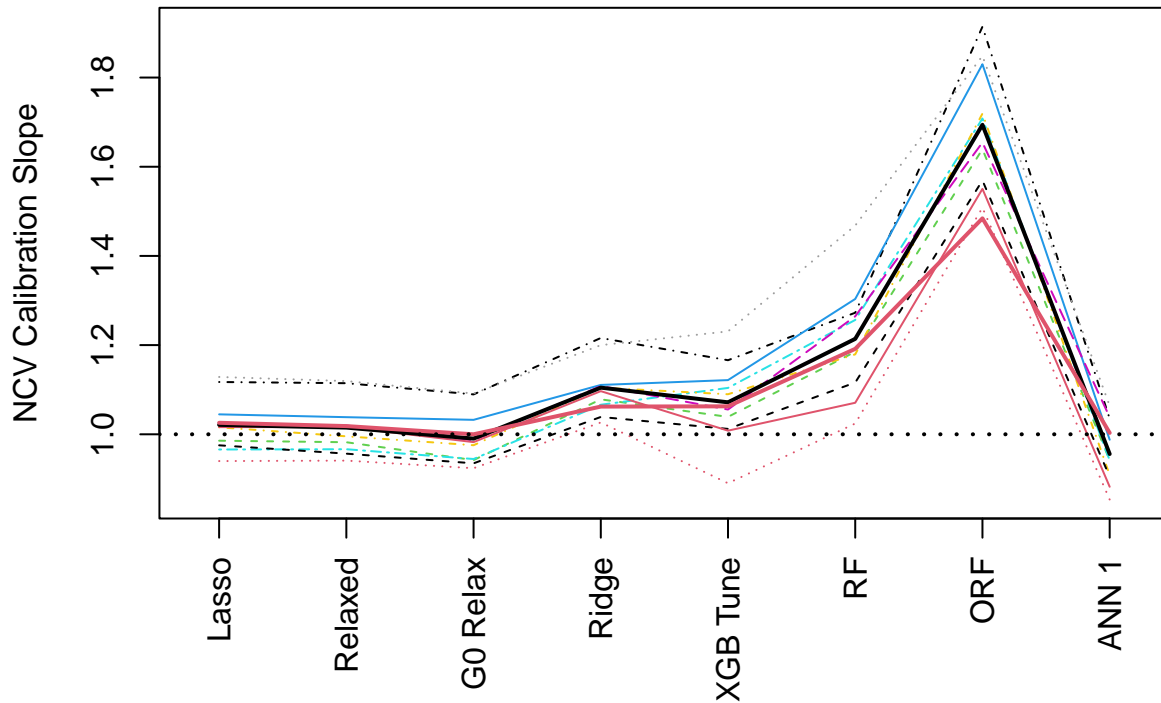


For “gaussian” data, i.e. generalization of the linear regression model, the deviance ratio is the reduction Mean Square Error (MSE) achieved by the machine learning model relative to the MSE of the model based upon only the overall average.

While of less importance than deviance ratio one can (by using the options `type="agree"` and `pow=2`) also describe model performances in terms of R-square for “gaussian” data. Note, from Jensens’ inequality we expect the R-squares for the different folds to be more biased than the R’s (correlation), and thus calculate the CV estimate for R-square as the (average R from the CV folds)-squared instead of average (R-squared from the CVs folds).

To understand how the models might be over or underestimating the strength of the association between the predictors and outcome one can regress the outcome on the model  $X \cdot \beta$  predicted, i.e. to a linear calibration. Slope terms from this linear calibration can be plotted as in

```
# Summarize relaxed lasso model performance informed by cross validation
plot(nested.gau.fit, type="lincal")
```



## Model and Cross Validation storage, and reuse

The individual model fits are all captured in the `nested.glmnet()` output object with names like `cv_glmnet_fit`, `xgb.simple.fit`, `xgb.tuned.fit`, `rf.tuned.fit` and `cv.stepreg.fit`. `cv_glmnet_fit` has a similar yet different format to that of `cv.glmnet()` by including further fit information. The XGB outputs are essentially direct outputs from 'xgboost' `xgb.train()`, but with added information regarding the seed or fold used in the fit as well as the parameters used in the search for the hyperparameters using the 'mlrMBO' package. The `rf.tuned.fit` object contains the output from `rfsrc()` in the object `rf.tuned$rf_tuned` along with information used for tuning. The `ann_fit_X` objects are derived using the R 'torch' package and take on their own format for logistical reasons. See the 'Using ann\_tab\_cv' vignette. Cross validation information from the individual outer folds are contained in data sets like `xx.devian.rep`, `xx.lincal.rep`, `xx.agree.rep` for further processing by the `summary()` function or by the user. For example

```
# Manually calculate CV R_square for lasso models
corr.rep = nested.gau.fit$lasso.agree.rep
corr.rep
```

```
##          1se      min      1seR      minR      1seR.GO      minR.GO      ridge
```

```
## [1,] 0.9275540 0.9333873 0.9300288 0.9312217 0.9310631 0.9310631 0.9320290
## [2,] 0.9168125 0.9221275 0.9179293 0.9215422 0.9188685 0.9216412 0.9147417
## [3,] 0.9229887 0.9277811 0.9229887 0.9287002 0.9209577 0.9259051 0.9312230
## [4,] 0.9335710 0.9324451 0.9335710 0.9316452 0.9340878 0.9314005 0.9283197
## [5,] 0.9266159 0.9331830 0.9257712 0.9306354 0.9273937 0.9313555 0.9324954
## [6,] 0.9294115 0.9292147 0.9289503 0.9300965 0.9220603 0.9220603 0.9220923
## [7,] 0.9341421 0.9382703 0.9316241 0.9392097 0.9334094 0.9393066 0.9357181
## [8,] 0.9350497 0.9337961 0.9350497 0.9332950 0.9286967 0.9322348 0.9285782
## [9,] 0.9354322 0.9348529 0.9329582 0.9330383 0.9316201 0.9311720 0.9320988
## [10,] 0.9131634 0.9103742 0.9131634 0.9103742 0.9110625 0.9043502 0.9059926
```

```
avecorr = colMeans(corr.rep)
R_square = round(avecorr^2, digits = 4)
R_square
```

```
##      1se      min      1seR      minR 1seR.GO minR.GO      ridge
## 0.8602 0.8641 0.8597 0.8630 0.8573 0.8594 0.8581
```

These numbers are consistent with the output from the `summary()` call.

## Further model assessment

### A naive calibration

Further model assessment can be made based upon the predicted values from the `predict()` function. For example, one can model the outcomes based upon a spline for the  $X \cdot \beta$  hats from the predicted values. This may help to understand potential nonlinearities in the model, but may also give inflated hazard ratios.

```
# Get predicted values from CV relaxed lasso model embedded in nested CV outputs & Plot
xb.hat = predict(object=nested.cox.fit, xs_new=xs, lam=NULL, gam=NULL, comment=FALSE)
# describe the distribution of xb.hat
round(1000*quantile(xb.hat, c(0.01, 0.05, 0.1, 0.25, 0.5, 0.75, 0.90, 0.95, 0.99)))/1000
```

```
##      1%      5%      10%      25%      50%      75%      90%      95%      99%
## -5.877 -4.091 -3.245 -1.788 -0.044  1.613  3.236  4.077  5.516
```

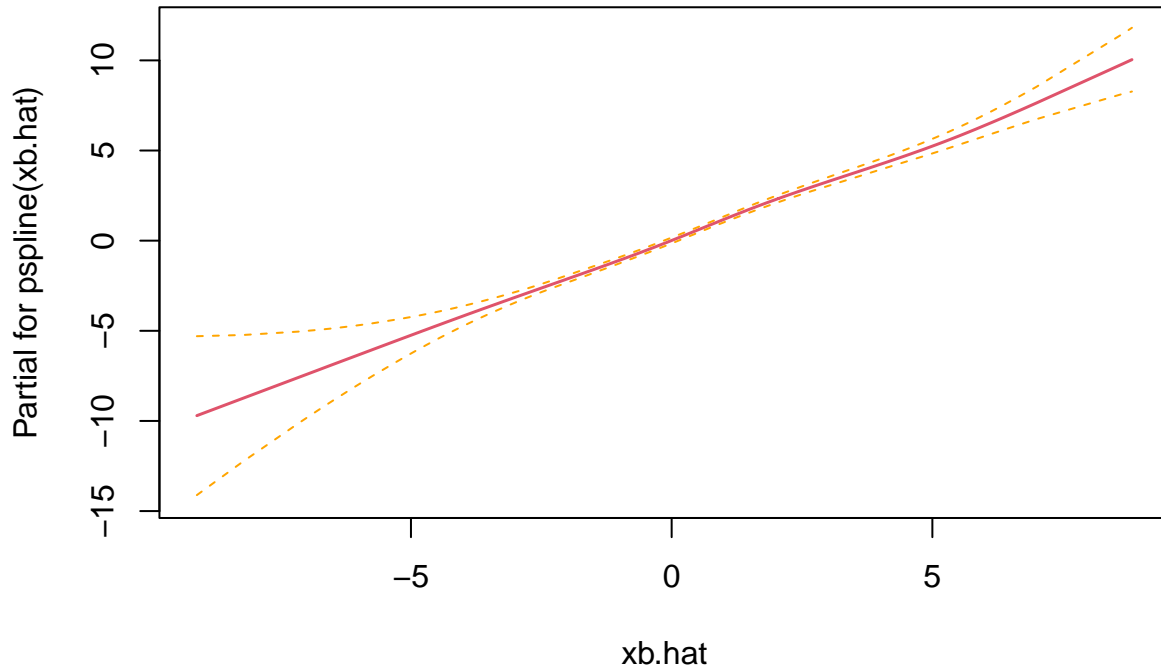
```
# Fit a spline to xb.hat using coxph, and plot
library(survival)
fit1 = coxph(Surv(yt, event) ~ pspline(xb.hat))
print(fit1)
```

```
## Call:
## coxph(formula = Surv(yt, event) ~ pspline(xb.hat))
##
##              coef se(coef)      se2    Chisq  DF      p
## pspline(xb.hat), linear 1.07e+00 3.32e-02 3.32e-02 1.03e+03 1.00 <2e-16
## pspline(xb.hat), nonlin                4.62e+00 3.04  0.21
##
## Iterations: 4 outer, 16 Newton-Raphson
##      Theta= 0.741
```



```
## Degrees of freedom for terms= 4
## Likelihood ratio test=1504 on 4.04 df, p=<2e-16
## n= 1000, number of events= 698
```

```
termplot(fit1,term=1,se=TRUE)
```

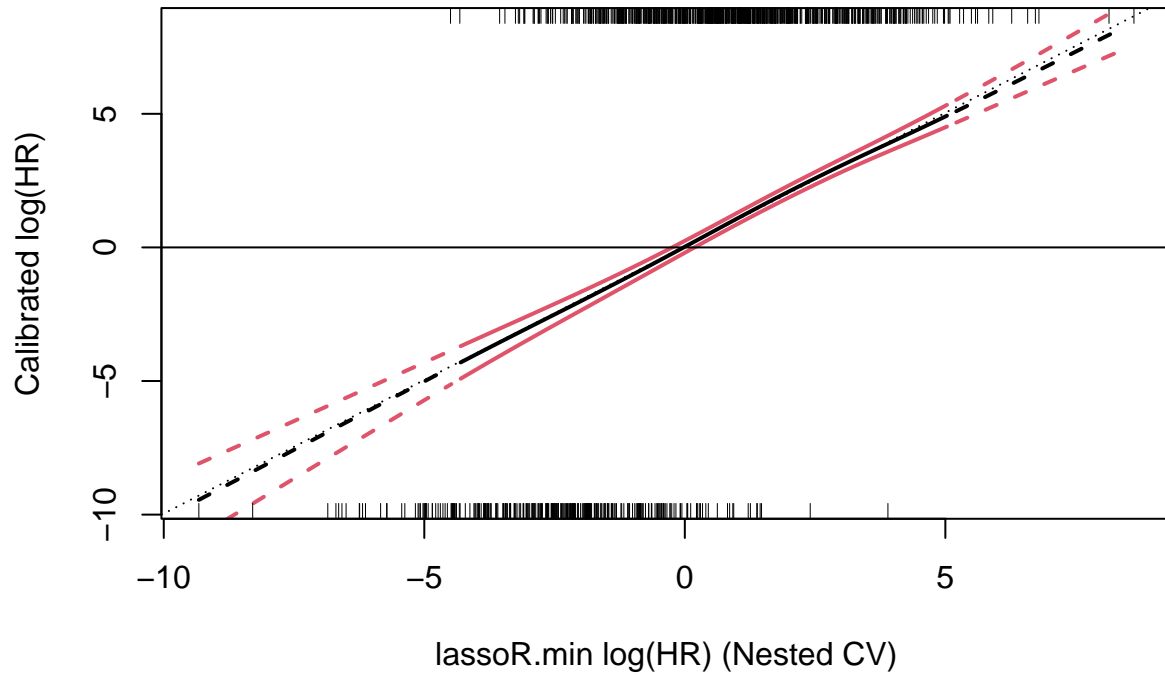


From this spline fit we see the log hazard ratio is roughly linear in the relaxed lasso predicted. Still, a calibration based upon data not used in model derivation would be more meaningful.

## Hold out data calibrations

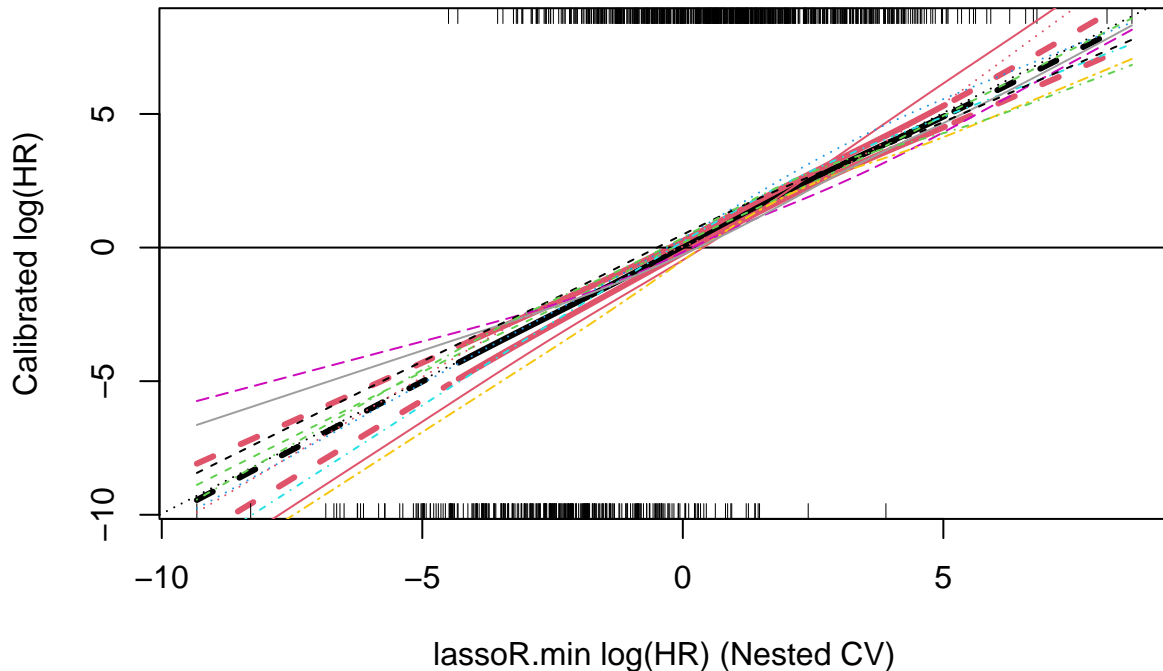
The `calplot()` function generates calibration plots using spline fits for each of the hold out data sets, either from the outer loop of the nested cross validation or the out of bag sample partitions when using a bootstrap approach, for example based upon cross validation

```
calplot(nested.cox.fit,wbeta=5)
```



Here the “overall” calibration and confidence interval is based upon the average and  $\pm 2$  standard error of the spline fits. One can also include the individual calibration curves from the outer cross validation loop with the option `plotfold=1` as in

```
calplot(nested.cox.fit, wbeta=5, plotfold=1 )
```



Calibration is discussed in more detail in the “Calibration of Machine Learning Models” vignette.

## Model comparisons with simulated and observed data

The examples above compare models using simulated data. Simulated data can often be constructed to show one or the other model to do perform the best. For the examples above the lasso models and those informed by the relaxed lasso performed better. When comparing model using observed medical data we have generally seen the lasso models to be consistently among the better performing models, but the differences in performances not as large and consistent as in these examples. Notably, when using observed data, the oblique random forest (ORF) often performed slightly (but not statistically significantly) better than the other models. By its construction we would expect the ORF to fit better when there are stronger correlations or interactions between predictors/features. In our test runs on observed data we often did have meaningful correlations between the predictors/features. Because different models may perform better with different types of data it may be informative to compare model performances for multiple data sets of a particular data type, when available, not only a single data set of immediate interest.

## Model replicability and model comparisons

To facilitate reproducible results the `nested.glmnetr()` function stores the seeds used to generate the pseudo random samples used for assigning observations to folds in the outer loop, as well as the fold ids themselves, for all models. Additionally, the program stores the seeds when generating the folds in the inner loop for

lasso, XGB, neural network and stepwise regressions as well as the seeds used when generating the bootstrap samples for the random forest models. Using the seeds from a prior call to `nested.glmnet()`, one can reproduce the results from this earlier call. Because the seeds are saved for sample partitioning in both the outer cross validation or the bootstrap method along with the folds for inner loop model derivations, results can be reproduced even when rerunning an analysis for a single model. If we did not save and manage the seeds the user might have to run all models included in an earlier run to verify or inspect a single model fit. If the seed option is unspecified in the call to `nested.glmnet()`, one should be cautious when using `set.seed()` in one's own code as this too will effect the pseudo randomness used in the ML calculations, which could unwittingly yield identical results when pseudo independent runs are intended.

Using the same folds, i.e. data splits, for the different models controls for some of the variability in the model performance estimates due to the randomness in the choice of folds, which should reduce variability in the differences in performance estimates between different models. `nested.glmnet()` stores the model performance measures from each iteration of the outer loop allowing calculation of means and standard deviations (SD) for each model performance measure, as well as differences paired on each data split in the outer loop. Because of dependencies between the different model fits from a CV loop, there are concerns about the accuracy of the SDs (Bengio Y & Grandvalet Y, "No Unbiased Estimator of the Variance of K-Fold Cross-Validation", Journal of Machine Learning Research 5 (2004) 1089–1105). Still, people use these SDs routinely as a rough approximation, being cautious of making strong inferences. We expect any dependencies at least between the paired differences should be minimal and reasonable in approximation. This intuition is supported by informal simulations. The reader is invited to perform similar studies using the type of data they encounter.

A simple comparison of model performances based upon deviance ratio can be done as in the example

```
# compare fractional reductions in Mean Square Error between different models
nested.compare(nested.gau.fit, type="devrat")
```

```
## Model performance comparison in terms of ** Deviance Ratio **
##
## Comparison                estimate   (95% CI)      p
##
## lasso.minR - lasso.min      -0.001 (-0.0028, 9e-04)  0.2644
## lasso.minR - lasso.minR0     0.0038 (-4e-04, 0.0079)  0.0692
## lasso.min - lasso.minR0      0.0047 (9e-04, 0.0086)  0.0215
##
## XGBoost (tuned) - XGBoost (simple)  0.1413 (0.1122, 0.1703)  0
##
## step (df) - step (p)        1e-04 (-0.0016, 0.0018)  0.8962
##
## lasso.minR - XGB (tuned)      0.0437 (0.0326, 0.0548)  0
## lasso.minR - Random Forest    0.1527 (0.138, 0.1674)  0
## lasso.minR - Oblique Random Forest  0.186 (0.1581, 0.2139)  0
## lasso.minR - ANN              0.1954 (0.1471, 0.2436)  0
##
## XGBoost (tuned) - RF          0.109 (0.0915, 0.1265)  0
## XGBoost (tuned) - ORF        0.1423 (0.1084, 0.1762)  0
## XGBoost (tuned) - ANN        0.1517 (0.1073, 0.1961)  0
##
## RF - ORF                     0.0333 (0.0033, 0.0633)  0.0331
##
## RF - ANN                     0.0427 (-0.0068, 0.0923)  0.083
##
## ORF - ANN                    0.0094 (-0.056, 0.0748)  0.753
```

As the outer CV loop performance calculations are saved in the `nested.glmntr()` output object (with names like `lasso.agree.rep`, `lasso.lincal.rep`, `xgb.agree.rep`, `xgb.lincal.rep`, etc.) comparisons not provided by `nested.compare()` can be calculated from these stored CV data.

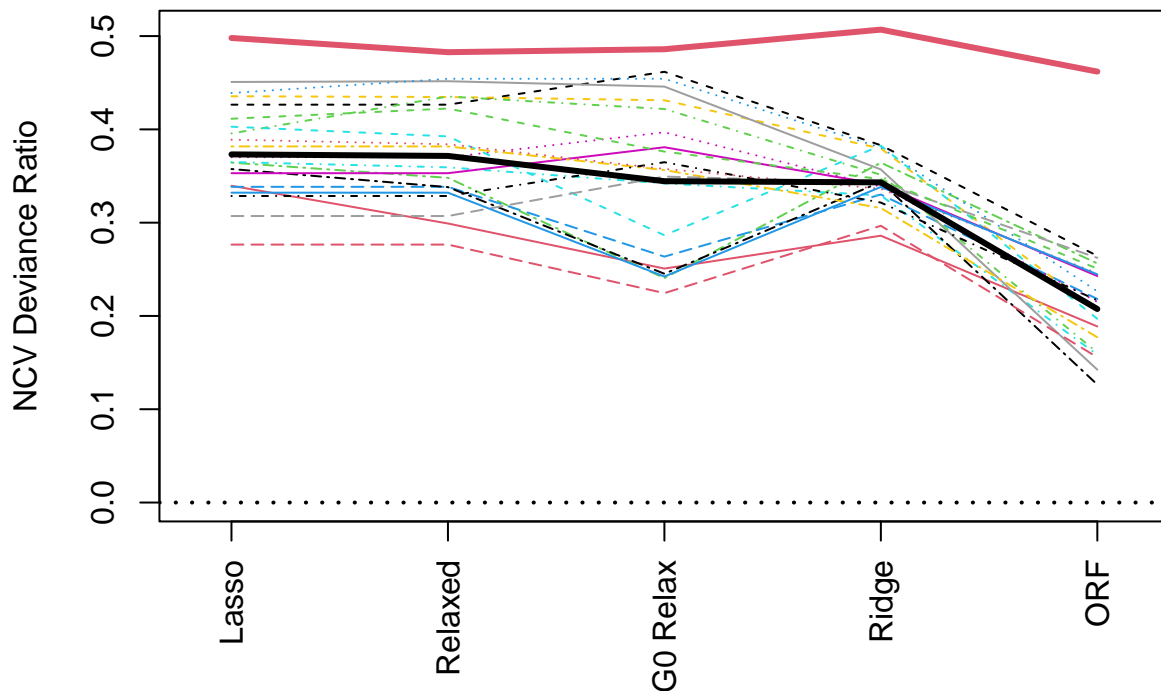
## Bootstrap assessment of model performance

If one is uncertain about using cross validation for assessing model performance, one can implement a bootstrap assessment, where the properties may be better understood. This is done by specifying the bootstrap option for the number of bootstrap samples to be used, e.g. as in

```
nested.bin.boot.fit = nested.glmntr(xs=NULL,yb=NULL,family="binomial",
  dolasso=1, doorf=1,
  folds_n=10, seed=219301029, bootstrap=20, track=1)
```

where models are fit for each of 20 re-samples from the original data. For each re-sample models are evaluated using the respective out of bag (OOB) data, i.e. the original data not included in the re-sample, and these are used to derive overall model performances.

```
plot(nested.bin.boot.fit)
```



With a larger number of resamples than typically used in cross validation the bootstrap approach may require much longer computing times.