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Bootstrap internal validation command for predictive logistic regression models

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Abstract. Overfitting is a common problem in the development of predictive models. It leads to an optimistic estimation of apparent model performance. Internal validation using bootstrapping techniques allows one to quantify the optimism of a predictive model and provide a more realistic estimate of its performance measures. Our objective is to build an easy-to-use command, `bsvalidation`, aimed to perform a bootstrap internal validation of a logistic regression model.

Keywords: `st0644`, `bsvalidation`, bootstrap, internal validation, predictive model, performance, logistic, logit

1 Introduction

A multivariable predictive model is a mathematical equation that relates multiple predictors for a particular individual to the probability of future occurrence of an outcome (Royston et al. 2009). Overfitting is a common problem in the development of these models, and it usually yields an overly optimistic model performance (Steyerberg 2009). In this context, internal validation is essential to provide a more realistic estimate of model ability to predict the risk of the outcome in a new subject. Several solutions have been proposed to correct for this optimism (sample splitting, cross-validation, and its variants leave-one-out cross-validation or leave-pair-out cross-validation). Among these strategies, bootstrapping emerges as a popular strategy to correct for optimistic estimates of the apparent performance.

The transparent reporting of a multivariable prediction model for an individual prognosis or diagnosis (TRIPOD) statement is an evidence-based guide of recommendations to standardize reporting of predictive models. The TRIPOD statement recommends bootstrapping techniques to carry out internal model validation and shrinkage methods to adjust overfitted models (Moons et al. 2015; Collins et al. 2015).

Our objective is to develop a new command, `bsvalidation`, to perform internal model validation using bootstrapping techniques that is executable as a postestimation command after the logistic or logit command. Stata has implemented postestimation commands to assess the apparent performance of the model. First, it has implemented the `lroc` postestimation command to assess model discrimination. It also

has implemented `estat gof` to assess model calibration with a Hosmer–Lemeshow test. To the best of our knowledge, there is no user-defined internal validation command implemented in Stata to date such as the one we are presenting.

2 Methods

`bsvalidation` needs to be executed after `logistic` or `logit`. The command allows one to estimate different performance measures in terms of overall model fit performance (that is, how close our predictions are to the actual outcome, related to the amount of variability that is explained); discrimination (that is, how well the model distinguishes between those with and without the outcome); and calibration (that is, how well predictions and observations agree). These measures can be observed in table 1.

Table 1. Performance measures

Item	Measure	Characteristics
Overall performance (Steyerberg et al. 2010)	Brier _{scaled}	Range: [0, 100] High values indicate predictions are closer to the actual outcome.
Discrimination (Riley et al. 2019)	C-statistic	Range: [0.5, 1] High values indicate better discrimination.
Calibration (Riley et al. 2019)	E:O ratio	Ideal value: 1 E:O < 1 indicates the model underestimates for the total number of events. E:O > 1 indicates the model overestimates for the total number of events.
	Calibration-in-the-large (CITL)	Ideal value: 0 CITL < 0 indicates the predictions are systematically too high. CITL > 0 indicates the predictions are systematically too low.
Calibration slope		Ideal value: 1

Slope < 1 indicates the predictions are too extreme and the model is overfit.
Slope > 1 indicates the predictions are not varied enough and the model is underfit.

NOTE: $\text{Brier}_{\text{scaled}} = 1 - \text{Brier}_{\text{score}} / \text{Brier}_{\text{max}}$

After the user has fit a logistic predictive model in the original sample using either the `logit` or `logistic` command, the validation command goes over the following algorithm:

1. It determines its apparent performance in the original sample (table 1).
2. It draws a bootstrap sample with replacement from the original sample.
3. It builds a new prediction model (bootstrap model) replicating the same modeling strategy used in the model that is being validated, and it determines its apparent performance in the bootstrap sample (bootstrap performance). If the original model is prespecified (that is, fit without variable selection), `bsvalidation` uses original model specification without any strategy for variable selection.
4. It applies the bootstrap model to the original sample to determine its performance (test performance).
5. It calculates the model's optimism as the difference between the bootstrap performance and the test performance.
6. It repeats steps 2–5 a userdefined number of times to obtain a stable averaged estimate of the optimism.
7. Finally, it subtracts the averaged optimism estimate obtained in step 6 from the initial apparent performance estimated in step 1 to obtain the optimismcorrected performance estimate.

Also, uniform shrinkage parameters—heuristic (Van Houwelingen and Le Cessie 1990) and bootstrap (Harrell 2015)—are estimated, and the coefficient of the model can be shrunk.

Our `bsvalidation` command also generates a calibration plot. Calibration is assessed using a lowess smoother function of predicted and observed risks for the overall sample. It also presents pairs of predicted and observed risks for groups defined by the user according to quantiles of predicted risk.

3 The bsvalidation command

3.1 Syntax

The syntax for `bsvalidation` is

102 `bsvalidation` *varlist* , *options*

103 If the final model was prespecified, *varlist* will be empty. If the model was built using
104 selection methods (backward, forward, or stepwise), those predictors previously assessed
105 but excluded from the final model during the selection process should be included in
106 *varlist*.

107 **3.2** **Options**

108 `reps(#)` specifies the number of bootstrap samples. The default is 50 samples. If you
109 are using Stata/IC, up to 800 bootstrap samples are supported. See `help limits`.

110 `rseed(#)` sets the random-number seed. This option can be used to obtain repro-
111 ducible results. `rseed(#)` is equivalent to typing `set seed #` prior to calling
112 `bsvalidation`.

113 `adjust(string)` displays the final model after applying a uniform shrinkage factor to
114 the regression coefficients. *string* is one of the following:

115 heuristic—uniform heuristic shrinkage parameter from Van
116 Houwelingen and Le Cessie (1990).

117 bootstrap—uniform bootstrap shrinkage parameter from Steyerberg (2009).

118 `pr(#)` and `pe(#)` specify the significance level threshold for variables to be removed
119 from or entered into the model, respectively.

120 `pr(#)` is backward elimination. Variables with *p*-value $\geq pr()$ are eligible to be
121 removed.

122 `pe(#)` is forward selection. Variables with *p*-value $< pe()$ are eligible to be entered.

123 `pr(#)` and `pe(#)` indicate backward stepwise.

124 When a predictor-selection approach is considered, a backward elimination strategy is
125 generally preferred (Harrell 2015).

126 Furthermore, `bsvalidation` displays the times each variable is selected in the final
127 model after applying the same selection strategy for each bootstrap sample. Other variable-
128 selection strategies such as lasso (least absolute shrinkage and selection operator) are not
129 included in `bsvalidation`. See `help lasso`.

130 `models` displays the final model for each bootstrap sample. If the final model is pre-
131 specified, this option does not apply.

132 `eform` causes the coefficient table to be displayed in exponentiated form: for each coef-
133 ficient, `exp(b)` rather than `b` is displayed. Standard errors and confidence intervals are
134 also transformed.

135 `graph` produces a calibration plot of observed against expected probabilities. Cali-
136 bration is plotted in groups across the risk spectrum. Confidence intervals for the
137 groupings are displayed as well as a lowess smoother.

138 This allows one to assess the calibration at the individual level. If `adjust()` is considered,
139 then the calibration plot will be adjusted.

140 Other user commands to generate calibration plots can be consulted (Ensor, Snell, and
141 Martin 2018).

142 `group(#)` specifies the number of percentiles to divide the predicted risks into. The
143 default is to divide the predicted risks into 10 equally sized groups.

144 `min(#)` allows one to fix a lower bound of observed and expected probabilities to be
145 plotted.

146 If `min()` is higher than the minimum probability predicted by the model, it is auto
147 matically rounded to the nearest first decimal to minimum.

148 `max(#)` allows one to fix an upper bound of observed and expected probabilities to be
149 plotted.

150 If `max()` is lower than the maximum probability predicted by the model, it is auto
151 matically rounded to the nearest first decimal to maximum.

152

153 **3.3** **Stored results**

154 `bsvalidation` stores the following in `e()`:

155 Scalars	
156 <code>e(N)</code>	number of observations
157 <code>e(k)</code>	number of parameters in the final model
158 <code>e(df_m)</code>	degrees of freedom
159 <code>e(k_max)</code>	number of parameters in the maximum model
160 <code>e(boot)</code>	number of bootstrap samples
161 <code>e(brier)</code>	Brier score for model overall performance
162 <code>e(opt_brier)</code>	optimism of the Brier score
163 <code>e(cstat)</code>	C-statistic for model discrimination
164 <code>e(opt_cstat)</code>	optimism of the C-statistic
165 <code>e(eo_ratio)</code>	ratio between expected and observed events
166 for model calibration	
167 <code>e(citl)</code>	calibration-in-the-large for model calibration
168 <code>e(slope)</code>	calibration slope for model calibration
169 <code>e(heur_shrink)</code>	uniform heuristic shrinkage
170 <code>e(boot_shrink)</code>	uniform

171 bootstrap shrinkage Macros

172 <code>e(cmd)</code>	<code>bsvalidation</code>
173 <code>e(depvar)</code>	dependent variable
174 <code>e(all_vars)</code>	independent variables in the maximum model
175 <code>e(sel_vars)</code>	independent variables in the final model
176 <code>e(model)</code>	regression model
177 <code>e(properties)</code>	<code>b V</code>

178 Matrices

179 <code>e(b)</code>	coefficient vector
180 <code>e(V)</code>	variance–covariance

181 matrix of the estimators Functions

182 <code>e(sample)</code>	marks estimation sample
----------------------------	-------------------------

183

184 **4** **Examples**

185 We illustrate the use of `bsvalidation` with a predictive model developed to estimate

the risk of low birthweight using the dataset lbw.dta from Hosmer, Lemeshow, and Sturdivant (2013).

In the first example, the command bsvalidation runs a bootstrap internal validation of a prespecified model.

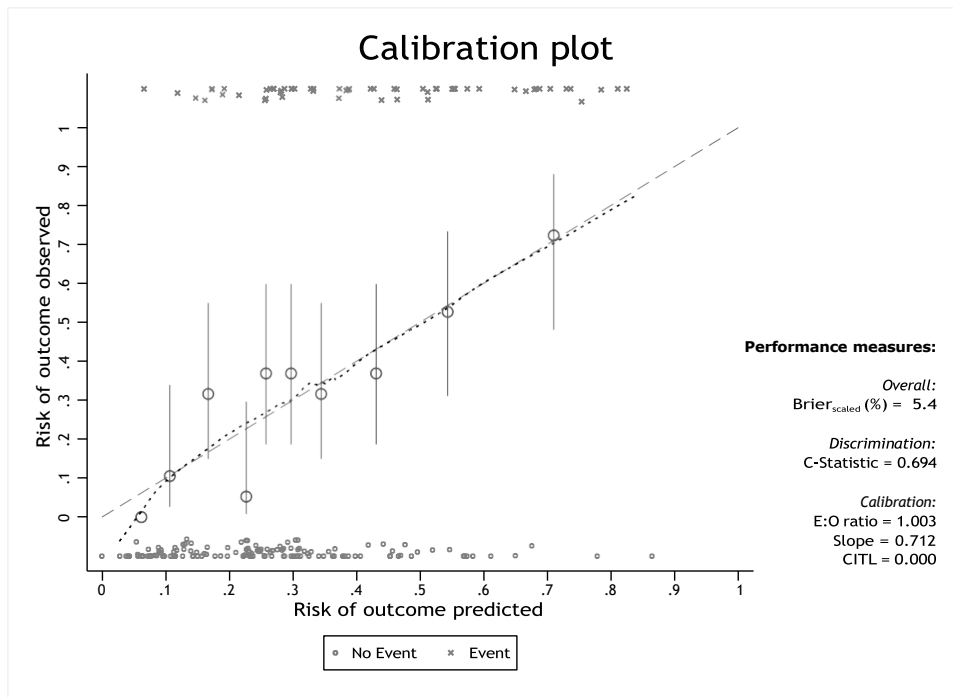


Figure 1. Calibration plot

In this first example, we fit a prespecified logistic model to predict the risk of low birthweight (defined as birthweight lower than 2,500 grams), using the mother's age (age), weight at last menstrual period (lwt), race (race), smoking status during pregnancy (smoke), previous history of premature labor (ptl), hypertension (ht), and uterine irritability (ui) as predictors. The bsvalidation output shows all apparent performance statistics (for example, C-statistic = 0.746). These performance measures are then adjusted for the estimated optimism, which is calculated from 50 (the default number) bootstrap samples (for example, C-statistic = 0.694). Additionally, by using the graph option, we visualize a calibration plot of observed against expected risks of low birthweight in groups defined by deciles of predicted risk, along with a smooth fit line. Further, it shows scatterplots with the distribution of events (x symbol) and nonevents (hollow circle symbol) along the x axis.

In the second example, bsvalidation performs a bootstrap internal validation of a model that was previously built using a backward-selection strategy with significance level ($p = 0.1$). After the backward-selection strategy, the predictors age and ptl were

dropped. The model coefficients are finally adjusted by the bootstrap-estimated uniform shrinkage factor or coefficient.

In the second example, the model is built using a backward-selection strategy in the original data. The predictors selected in the process are lwt, race, smoke, ht, and ui (logistic command). Other candidate predictors (age and ptl) initially assessed, but excluded during the selection process, are added in the *varlist* of the *bsvalidation* command to replicate the same modeling strategy used during the development of the original model. The output shows both apparent and optimism-adjusted performance measures. Additionally, because the backward-selection strategy is replicated in each bootstrap sample, the output also shows the number of times each predictor is selected in the final model (that is, lwt was included in 75 out of 100 bootstrap models). Finally, the coefficients of the final model are adjusted by bootstrap-based uniform shrinkage to correct overfitting. Thus, coefficients are multiplied by 0.712.

5 Conclusion

bsvalidation is a useful command to run bootstrap internal validation of predictive logistic regression models. It makes this internal validation method more accessible to researchers promoting a more complete and better report of predictive models according to TRIPOD guidelines.

6 Limitations

Although *bsvalidation* helps standardize the internal validation process, a disadvantage of bootstrap validation is that it allows validation only of models built following fixed or automated modeling strategies (that is, without dynamic modeling strategies or stepwise modeling strategies). Other important steps during the modeling process, such as collapsing factor variables, assessing nonlinearities, or testing for interaction terms, cannot be handled by *bsvalidation*. The command does not handle other shrinkage methods, such as the least absolute shrinkage and selection operator (Tibshirani 1996), and cannot handle missing values.

7 Future works

In the future, we will work to solve some of the previously mentioned limitations, and we will evolve the command to validate other regression models commonly used in biomedical research, such as Cox regression.

8 Programs and supplemental materials

To install a snapshot of the corresponding software files as they existed at the time of publication of this article, type

```
. net sj 21-2  
. net install st0644 (to install program files, if available)  
. net get st0644 (to install ancillary files, if available)
```

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