Selecting Variables in Multiple Regression

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Introduction

- One problem that can arise in "exploratory" multiple regression studies is which predictors from a set of potential predictor variables should be included in the multiple regression analysis, and in the ultimate prediction formula.
- In this module, we review some traditional and newer approaches to variable selection, pointing out some of the pitfalls involved in selecting a subset of variables to analyze.

The Problem with Redundancy

- A fundamental problem when one has several potential predictors is that some may be largely redundant with others.
- One result of such redundancy is called multicollinearity, which occurs when some predictors are linear combinations of others (or nearly so), resulting in a covariance matrix of predictors that is singular, or nearly so.
- One outcome of multicollinearity is that parameter estimates become subject to wild sampling fluctuations, for theoretical reasons that we investigate on the next slide.

The Problem with Redundancy

Collinearity and Variances of Beta Estimates

• Suppose we have just two predictors, and the mean function is

$$E(Y|X_1 = x_1, X_2 = x_2) = \beta_0 + \beta_1 X_1 + \beta_2 X_2$$
(1)

It can be shown that

$$Var(\hat{\beta}_{j}) = \frac{\sigma^{2}}{1 - r_{12}^{2}} \frac{1}{SX_{j}X_{j}}$$
(2)

where r_{12} is the correlation between X_1 and X_2 , and $SX_jX_j = \sum_i (x_{ij} - \overline{x}_j)^2$.

• From the above formula, we can see that, as r_{12}^2 approaches 1, these variances are greatly inflated.

The Problem with Redundancy

Collinearity and Variances of Beta Estimates

- When the number of predictors exceeds 2, the previous result generalizes.
- Specifically, we have

$$\mathsf{Var}(\hat{\beta}_j) = \frac{\sigma^2}{1 - R_j^2} \frac{1}{SX_j X_j}$$
(3)

where R_i^2 is the squared multiple correlation between X_j and the other predictors.

• It is easy to see why the quantity $1/(1 - R_j^2)$ is called the *j*th *variance inflation factor*, or VIF_j .

Detecting and Dealing with Redundancy

- Simple multicollinearity may be detected in several ways. For example, one might examine the correlation matrix to see if any predictors are highly correlated, and delete some.
- Examining the *varimax-rotated principal component structure* of a set of predictors will reveal more complex forms of multicollinearity, so long as the redundancy is linear.
- Principal component analysis will reveal uncorrelated variables that are linear combinations of the original predictors, and which account for maximum possible variance.
- If there is a lot of redundancy, just a few principal components might be as effective.

Detecting and Dealing with Redundancy

- In some cases, predictors may be redundant with each other, but the redundancy is nonlinear.
- Frank Harrell's Hmisc package includes a function redun to detect such nonlinear redundancy and suggest variables that might be candidates for elimination.

- In this section, we review the classic variable selection procedures that have dominated the social sciences literature.
- These procedures are usually referred to as
 - Forward Selection
 - ② Backward Elimination
 - Stepwise Regression

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- The goal of variable selection is to divide a set of predictors in the columns of a matrix **X** into active and inactive terms.
- The number of partitions is 2^k, which becomes quite large very quickly when k is even moderate.
- There are two fundamental issues:
 - Given a particular candidate for the active terms, what criterion should be used to compare this candidate to other possible choices?
 - e How do we deal computationally with the potentially huge number of comparisons that need to be made?

- Originally, the criteria for model evaluation were purely statistical. In order to be added to a model, a variable had to be significant according to the classic partial *F* test, either with a *p*-value below a certain "*p* to enter" value, or with an *F* statistic specified as the "*F* to enter" value (as in SPSS).
- More recently, attention has shifted to so-called "informational criteria," which appear, at least at first glance, to combine model fit with model complexity in assessing whether a variable should be added to a prediction formula.

The Akaike Information Criterion (AIC)

- Criteria for comparing various candidate subsets are based on the lack of fit of a model and its complexity.
- Ignoring constants that are the same for every candidate subset, the AIC, or *Akaike Information Criterion* for a candidate *C*, is

$$AIC_{\mathcal{C}} = n \log(RSS_{\mathcal{C}}/n) + 2p_{\mathcal{C}}$$
(4)

• According to the Akaike criterion, the model with the smallest AIC is to be preferred.

The Bayesian Information Criterion(BIC)

• The Schwarz Bayesian Informatin Criterion (BIC) is

$$BIC_{\mathcal{C}} = n \log(RSS_{\mathcal{C}}/n) + p_{\mathcal{C}} \log(n)$$
(5)

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Cross-Validation Based Criteria

- The major reason for employing fit indices that correct for complexity is because, for sample data, increasing the complexity of the model can never yield a higher *RSS*, and almost always will yield a lower *RSS*, even when the increase in complexity yields no gain in prediction in the population.
- In genuine cross-validation, the sample is divided into two parts at random, a *construction* (or calibration) set and a validation set.
- The model is fit to the construction set and parameter estimates are obtained.
- That model with those parameter estimates is then used to predict the response variable in the validation set data.
- The RSS is used as a measure of fit, and is not corrected for complexity.

Cross-Validation Based Criteria

- The *PRESS* measure is an attempt to assess the cross-validation capability of a model based on a single sample.
- For a particular model, for each observation,
 - compute fitted values from $\hat{\beta}$ based on all the data other than that observation.
 - compute the squared difference between the response and the predicted values
- These squared errors are summed up across the entire sample.

Cross-Validation Based Criteria

 \bullet The resulting statistic, for model subset candidate $X_{\mathcal{C}}$ is

$$PRESS = \sum_{i=1}^{n} \left(y_i - \mathbf{x}'_{\mathcal{C}i} \hat{\beta}_{\mathcal{C}(i)} \right)^2$$
(6)

• PRESS can be computed as

$$PRESS = \sum_{i=1}^{n} \left(\frac{\hat{\mathbf{e}}_{Ci}}{1 - h_{Cii}} \right)^2 \tag{7}$$

where \hat{e}_{Ci} and h_{Cii} are, respectively, the residual and the leverage for the *i*th case in the subset model.

• This index is relatively straightforward to compute in simple linear regression because of the above computational simplification, but this simplicity does not generalize to more complex models.

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An Example — The Highway Data

- This example employs the highway accident data from ALR Section 8.2.
- The variables (including the response, log(*Rate*)), are described in ALR3 Table 10.5 reproduced below.

Variable	Description			
log(Rate)	Base-two logarithm of 1973 accident rate per million vehicle miles, the response			
log(Len)	Base-two logarithm of the length of the segment in miles			
log(ADT)	Base-two logarithm of average daily traffic count in thousands			
log(Trks)	Base-two logarithm of truck volume as a percent of the total volume			
Slim	1973 speed limit			
Lwid	Lane width in feet			
Shld	Shoulder width in feet of outer shoulder on the roadway			
Itg	Number of freeway-type interchanges per mile in the segment			
log(Sigs1)	Base-two logarithm of (number of signalized interchanges per mile in the segment + 1)/(length of segment)			
Acpt	Number of access points per mile in the segment			
Hwy	A factor coded 0 if a federal interstate highway, 1 if a principal arterial highway, 2 if a major arterial, and 3 otherwise			

TABLE 10.5 Definition of Terms for the Highway Ac	cident Data
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An Example — The Highway Data

- We begin by fitting a model with all terms
 - > data(highway)
 - > a <- highway
 - > a\$logADT <- logb(a\$ADT,2)</pre>
 - > a\$logTrks <- logb(a\$Trks,2)</pre>
 - > a\$logLen <- logb(a\$Len,2)</pre>
 - > a\$logSigs1 <- logb((a\$Sigs*a\$Len+1)/a\$Len,2)</pre>
 - > a\$logRate <- logb(a\$Rate,2)</pre>
 - > # set the contrasts to the R default
 - > options(contrasts=c(factor="contr.treatment",ordered="contr.poly"))
 - > a\$Hwy <- if(is.null(version\$language) == FALSE) factor(a\$Hwy,ordered=FALSE) else factor(a\$Hwy)</pre>
 - > attach(a)
 - > names(a)

+

[1]	"ADT"	"Trks"	"Lane"	"Acpt"	"Sigs"	"Itg"
[7]	"Slim"	"Len"	"Lwid"	"Shld"	"Hwy"	"Rate"
[13]	"logADT"	"logTrks"	"logLen"	"logSigs1"	"logRate"	

```
> cols <- c(17,15,13,14,16,7,10,3,4,6,9,11)</pre>
```

```
> m1 <- lm(logRate ~ logLen+logADT+logTrks+logSigs1+Slim+Shld+</pre>
```

Lane+Acpt+Itg+Lwid+Hwy)

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An Example — The Highway Data

• Here is the table

> xtable(m1)

	Estimate	Std. Error	t value	$\Pr(> t)$
Intercept)	5.7046	2.5471	2.24	0.0342
logLen	-0.2145	0.1000	-2.15	0.0419
logADT	-0.1546	0.1119	-1.38	0.1792
logTrks	-0.1976	0.2398	-0.82	0.4178
logSigs1	0.1923	0.0754	2.55	0.0172
Slim	-0.0393	0.0242	-1.62	0.1172
Shld	0.0043	0.0493	0.09	0.9313
Lane	-0.0161	0.0823	-0.20	0.8468
Acpt	0.0087	0.0117	0.75	0.4622
ltg	0.0515	0.3503	0.15	0.8842
Lwid	0.0608	0.1974	0.31	0.7607
Hwy1	0.3427	0.5768	0.59	0.5578
Hwy2	-0.4123	0.3940	-1.05	0.3053
Hwy3	-0.2074	0.3368	-0.62	0.5437

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Selecting Variables in Multiple Regression

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An Example — The Highway Data

- The R^2 value for the full model is .791, and there are p = 14 terms. Using the anova command, we can determine that RSS = 3.5370 for this model. (There is a typo in ALR, and the value is given as 3.5377.) The estimated error variance is $\partial^2 = 0.1415$.
- Next consider a more compact model, consisting of only 6 terms, an intercept and log(*Len*), *Slim*, *Acpt*, log(*Trks*), *Shld*. This model has an RSS of 5.0159.

```
> m2 <- lm(logRate ~ logLen+logTrks+Slim+Shld+Acpt)</pre>
```

```
> anova(m2)
```

Analysis of Variance Table

```
Response: logRate
         Df Sum Sg Mean Sg F value
                                  Pr(>F)
         1 5.5373 5.5373 36.4297 8.685e-07 ***
logLen
        1 1.5155 1.5155 9.9704 0.003391 **
logTrks
Slim
         1 4.3339 4.3339 28.5128 6.769e-06 ***
Shld
         1 0.1464 0.1464 0.9631
                                  0.333559
Acpt
          1 0.4021 0.4021 2.6452
                                  0.113375
Residuals 33 5.0159 0.1520
____
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

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```
An Example — The Highway Data
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 Calculations are straightforward, and show that the values for the reduced mean function are smaller than those for the full mean function for both AIC and BIC.
 > anova(m2,m1)

```
Analysis of Variance Table
```

```
Model 1: logRate ~ logLen + logTrks + Slim + Shld + Acpt
Model 2: logRate ~ logLen + logADT + logTrks + logSigs1 + Slim + Shld +
    Lane + Acpt + Itg + Lwid + Hwy
  Res.Df RSS Df Sum of Sq F Pr(>F)
1
      33 5.0159
2
      25 3 5370 8 1 479 1 3067 0 2852
> AIC2<-extractAIC(m2)</pre>
> BIC2<-extractAIC(m2,k=log(39))</pre>
> PRESS2 <-sum( (residuals(m2)/(1-hatvalues(m2)))^2 )</pre>
> AIC1 <- extractAIC(m1)</pre>
> BIC1 <- extractAIC(m1.k=log(39))</pre>
> PRESS1 <-sum( (residuals(m1)/(1-hatvalues(m1)))^2 )</pre>
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An Example — The Highway Data

> AIC1

[1] 14.00000 -65.61145

> AIC2

[1] 6.00000 -67.98662

> BIC1

[1] 14.00000 -42.32159

> BIC2

[1] 6.00000 -58.00525

> PRESS1

[1] 11.27222

> PRESS2

[1] 7.688042

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An Example — The Highway Data

- The single most important tool in selecting a subset of variables is the analyst's knowledge of the area under study and of each of the variables.
- In the highway accident data, *Hwy* is a factor, so all of its levels should probably either be in the candidate subset or excluded.
- Weisberg also makes the case that the variable log(*Len*) should be treated differently from the others, since its inclusion in the active predictors may be required by the way highway segments are defined.

Forward Selection

- In the preceding section, we simply compared two models.
- However, these two models represented only two of hundreds of possible models.
- Automated subset selection procedures can sort through many possible models and choose one.
- Our first example of such an procedure is Forward Selection.
- In forward selection, we start with a base model and consider a set of additional possible regressors. In what follows, assume that the base model is empty, i.e., has no regressors.
 - Consider all candidate subsets consisting of one term beyond the intercept, and find the subset that minimizes the criterion of interest. If an information criterion is used, then this amounts to finding the term that is most highly correlated with the response because its inclusion in the subset gives the smallest residual sum of squares. Regardless of the criterion, this step requires examining *k* candidate subsets.
 - For all remaining steps, consider adding one term to the subset selected at the previous step. Using an information criterion, this will amount to adding the term with the largest partial correlation with the response given the terms already in the subset, and so this is a very easy calculation. Using cross-validation, this will require fitting all subsets consisting of the subset selected at the previous step plus one additional term. At step j, k j + 1 subsets need to be considered.
 - Stop when all the terms are included in the subset, or when addition of another term increases the value of the selection criterion.

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Forward Selection

- If the number of terms beyond the intercept is k, this algorithm will consider at most $k + (k 1) + \ldots + 1 = k(k 1)/2$ of the 2k possible subsets.
- For k = 10, the number of subsets actually considered is only 45 of the 1024 possible subsets. The subset among these 45 that has the best value of the criterion selected is tentatively selected as the candidate.
- The algorithm requires modification if a group of terms is to be treated as all included or all not included, as would be the case with a factor.
- In such a case, we would have to consider adding the term *or the group of terms* that produces the best value on the criterion of interest.
- Each of the information criteria can now give different best choices because at each step, as we are no longer necessarily examining mean functions with p_{C} fixed.

Backward Elimination

- Backward selection works in the reverse order. You start with a candidate subset and then decide which terms can be eliminated.
 - If the full candidate subset.
 - At the next step, consider all possible subsets obtained by removing one term other than those to be forced to be in all mean functions from the candidate subset selected at the last step. Using an information criterion, this amounts to removing the term with the smallest *t*-value in the regression summary because this will give the smallest increase in residual sum of squares. Using cross-validation, all subsets formed by deleting one term from the current subset must be considered.
 - Ontinue until all terms but those forced into all mean functions are deleted, or until the next deletion increases the value of the criterion.

Once again, we need consider only k(k-1)/2 subsets. The subsets considered by forward selection and backward elimination can be different.

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Stepwise Regression

- The forward and backward algorithms can be combined into a stepwise method, where at each step, a term is either deleted *or* added so that the resulting candidate mean function minimizes the criterion function of interest. This will have the advantage of allowing consideration of more subsets, without the need for examining all 2^k subsets.
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- At each stage, the possibility is that a variable entered at a previous stage has now become superfluous because of additional variables now in the model that were not in the model when this variable was selected.
- To check on this, in the classic approach implemented in programs like SPSS, at each step a partial *F* test for each variable in the model is made as if it were the variable entered last. We look at the lowest of these *F*s and if the lowest one is sufficiently low (i.e., below the "*F*-to-remove" value, we remove the variable from the model, recompute all the partial *F*s, and keep going until we can remove no more variables.

Computational Examples

- We illustrate forward selection with the Highway data.
- We start by defining some special variables and then defining, for convenience, our maximal model.
 - > Highway\$sigs1 <- with(Highway, (sigs * len + 1)/len)</pre>
 - > f <- \sim log(len) + shld + log(adt) + log(trks) + lane + slim + lwid +
 - + itg + log(sigs1) + acpt + htype
- Next are the commands that set up a base model and start the Forward Selection procedure. They produce extensive output, so we'll execute them in class and examine the results there.
 - > m0 <- lm(log(rate) ~ log(len), Highway) # the base model
 - > m.forward <- step(m0, scope=f, direction="forward")</pre>
- Here are the commands that produce Backwards Elimination.

```
> m1 <- update(m0, f)</pre>
```

```
> m.backward <- step(m1, scope = c(lower = ~ log(len)), direction="backward")
```

• Here is the command for Stepwise Regression.

```
> m.stepup <- step(m0, scope=f)</pre>
```

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Caution about Selection Methods

- It is easy to demonstrate that subset selection methods can overstate significance, and that, after subset selection, the *p*-values printed in output are no longer valid.
- In some of my courses, we generate completely random data and then subject it to forward selection, finding a highly significant R^2 when in fact the population squared multiple correlation is zero.
- Not only are *p*-values wrong, but of course the $\hat{\beta}$ values are badly biased too.
- Our online code file provides a demonstration, in which 99 completely random predictors are assessed via Forward Selection.
- A number of "significant" regressors are found, and the R^2 value of 0.3591 has a listed *p*-value of 0.0000174.
- What has gone wrong? (C.P.)