

# Model selection II

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# The Swiss fertility data set

- Today we will explore how to calculate and apply model selection criteria in R and SAS
- We will look at a data set describing fertility in Switzerland in 1888
- Some background: fertility (number of children a woman gives birth to over the course of her life) is high in developing nations and low in developed countries
- In 1888, Switzerland was at the critical point in its development where it was undergoing the “demographic transition” in which its fertility was falling to levels seen in developed nations

## The Swiss fertility data set (cont'd)

The following variables were collected (primarily from military records) for each of the 47 French-speaking provinces in Switzerland:

- Fertility (standardized)
- Agriculture: Percent of males involved in agriculture as an occupation
- Examination: Percent of draftees who received the highest mark on their army examination
- Education: Percent of draftees educated beyond primary school
- Catholic: Percent Catholic (as opposed to Protestant)
- InfantMortality: Percent of live births who live less than one year

# Model selection criteria in R

- In R,  $R^2$  and  $R_{\text{adj}}^2$  are given by summary
- AIC, BIC, and  $C_p$  can all be obtained from the `extractAIC` function:

```
fit <- lm(Fertility~.,data=swiss)
extractAIC(fit)
extractAIC(fit,k=log(n)) ## BIC
extractAIC(fit,scale=sig2) ## Cp
```

Note: there is also a function `AIC`, though be aware that the two functions do not return exactly the same number (AIC drops constant terms)

- There is no default function to calculate PRESS, but you can calculate it via:

```
sum((fit$resid/(1-hatvalues(fit)))^2)
```

# Model selection criteria in SAS

- In SAS, unfortunately, none of these options are available in PROC GLM – you have to use PROC REG
- In PROC REG, you can get all the criteria with

```
PROC REG DATA=swiss OUTEST=fits;  
    MODEL Fertility = Agriculture Examination Education  
    Catholic InfantMortality / ADJRSQ CP AIC BIC PRESS;  
RUN;
```

although be aware that this estimate of CP is meaningless without an external estimate of  $\hat{\sigma}^2$
- Note that PROC REG allows multiple model statements; all these models show up in fits, so you can select from among a list of models by, say, running PROC SORT on fits

# Automatic model selection: Overview

- Given that we have an objective way of choosing between models, it is possible to automate the model selection process by fitting a number of models and ranking them according to some criterion
- There are two common strategies to searching through the set of all possible models:
  - Best subsets selection, an exhaustive search of all possible models
  - Stepwise selection, in which the search is simplified by taking it one step at a time

# Best subsets selection in SAS

- We will start with best subsets regression and illustrate its use on the Swiss fertility data set
- PROC REG in SAS allows for best subsets regression based on  $C_p$  through the SELECTION option in a MODEL statement:

```
PROC REG DATA=swiss;  
    MODEL Fertility = Agriculture Examination Education  
          Catholic InfantMortality / SELECTION = CP;  
RUN;
```

# Best subsets selection in R

- In R, best subsets selection is available through the `leaps` package, which you will have to install:

```
install.packages("leaps")  
require(leaps)
```

- Once installed, you can perform best subsets selection via:

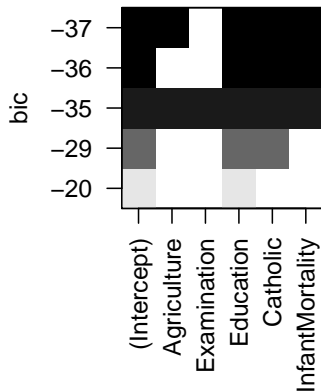
```
regs <- regsubsets(Fertility~., data=swiss)
```

where the `.` in a model formula means “all the variables not already in the formula”



# Best subsets selection in R

These models can be plotted via `plot(regs)`:



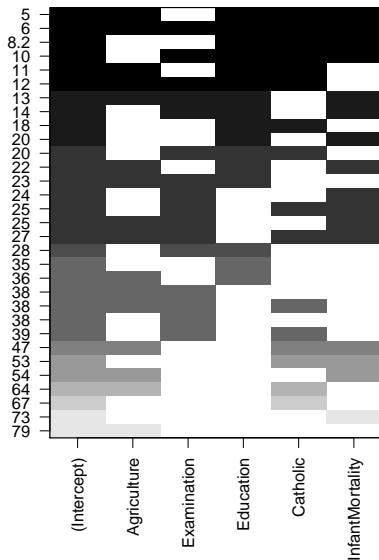
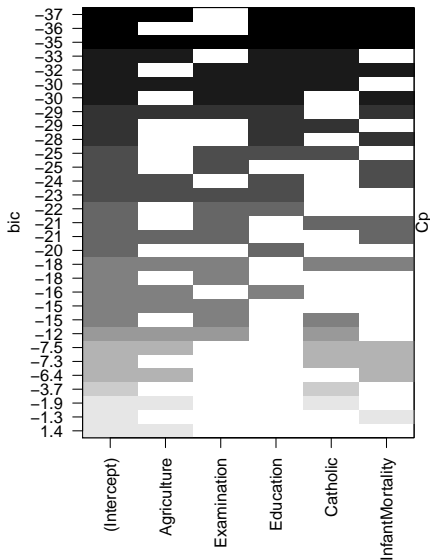
# Options in leaps

- By default, the package only returns the best model of each size (the best one-variable model, two-variable model, etc)
- This can be modified with the `nbest` option:  

```
regs <- regsubsets(Fertility~., data=swiss, nbest=10)
```
- The `leaps` package calculates BIC and  $C_p$  by default; to rank by  $C_p$ , we can submit  

```
plot(regs, scale="Cp")
```
- `regsubsets` returns RSS and  $p$ , so in principle, one could calculate and sort by AIC as well

# Best subsets



# Interpreting the results

- Note that the best subsets approach selects all of the variables except Examination
- This is somewhat interesting, as Examination is highly correlated with Fertility ( $-0.65$ ,  $p = 7 \times 10^{-7}$ )
- However, Examination is also highly correlated with Education and Agriculture, and seems to add little to the model if those two variables are already present

# Stepwise selection

- Note that there are  $2^p$  total subsets in a model with  $p$  candidate explanatory variables
- This number gets extremely large very quickly as  $p$  increases, to the point where, if  $p$  is above 40 or so, it is computationally infeasible to fit all these models
- Thus, *stepwise* approaches have been proposed, in which you find the best one-variable model, then find the best two-variable model that can be constructed by adding a variable to the best one-variable model, and so on
- In other words, we don't look at all two-variable models, only those that contain the best one-variable model

## Stepwise selection (cont'd)

- Specifically, this approach is known as *forward* stepwise selection
- An alternative approach is to start with the full model and successively eliminate variables, which is known as *backward* stepwise selection
- There are other variants as well, capable of moving both forward and backward, allowing for a variable to be added, but then removed later if it no longer seems necessary

# Stepwise selection in R

- In R, stepwise selection can be requested via the `method` option, as in

```
regs <- regsubsets(Fertility~., data=swiss, method="forward")
```
- Other options for `method` include ‘‘backward’’ and ‘‘seqrep’’ for sequential replacement (an approach which considers both forward and backward steps)
- In this case, of course, stepwise selection is unnecessary, as we can perform the exhaustive search
- However, it is worth noting that the stepwise approach in this case selects the same model (this is not always the case)

# Stepwise selection in SAS

- In SAS, stepwise selection can be requested via the SELECTION option in the model statement (e.g., SELECTION=FORWARD, SELECTION=BACKWARD, SELECTION=STEPWISE)
- A subtle distinction between SAS and R is that SAS considers “stepwise” to be synonymous with choosing variables based on  $p$ -values
- In other words, it moves forward by adding the variable with the most significant  $p$ -value, and backwards by dropping the variable with the least significant  $p$ -value
- This approach is popular because, by construction, it leads to models with significant terms, but is on a weak foundation, as the construction of the model is not guided by any meaningful model selection criterion



# Greedy algorithms

- In the computer science literature, stepwise approaches are known as *greedy algorithms*, in the sense that they operate according to grabbing the variable that will help most in the short term
- However, just as this is not always the best strategy in other areas of life, there is no guarantee that the stepwise approach will find the best model

# Example

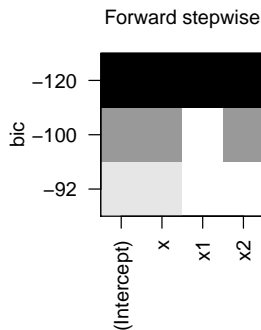
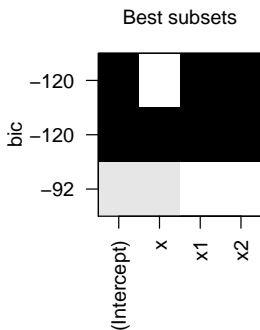
For example, consider:

$$X_1 \sim N(0, 1)$$

$$X_2 \sim N(0, 1)$$

$$X|X_1, X_2 \sim N(X_1 + X_2, 0.5)$$

$$Y|X_1, X_2, X \sim N(X_1 + X_2, 1)$$



# Best subsets vs. stepwise approaches

- In low dimensions, forward stepwise algorithms and best subset approaches often agree
- Even if they do not (as in the previous example), simple tweaks such as sequential replacement usually fix the problem
- In higher dimensions, however, stepwise approaches only investigate an exceptionally small fraction of the possible models, and rarely find the optimal model