

Model Building 1

Lecture 14

STA 371G





Topics

35 Texas Counties Have Zero Physicians

John Commins, May 6, 2015















Even if you don't live in Texas, these numbers should scare anyone who cares about rural healthcare, because this crisis is not unique to Texas.

How bad is the provider shortage in Texas?

What might explain this?

- Small counties
- Poverty
- Health insurance

- Unemployment
- Large rural areas
- Something else?



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- This type of analysis is an exploratory study.

An exploratory study of the Texas physician shortage

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An exploratory study of the Texas physician shortage

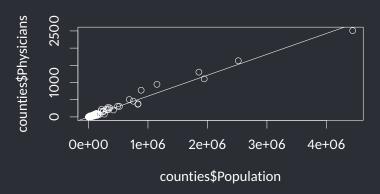
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- Exploratory studies require the most in terms of model selection. Automated tools are helpful, but judgement is still needed!

Population as a predictor of number of physicians

plot(counties\$Population, counties\$Physicians)
popmodel <- lm(counties\$Physicians ~ counties\$Population)
abline(popmodel)</pre>



Transform and Subset the data

```
# Create a variable for physicians per 10,000 people
counties$PhysiciansPer10000 <-
  counties$Physicians / counties$Population * 10000
# Remove the very small and very large counties
mcounties <- counties[counties$Population < 500000 &
                      counties$Population > 10000,]
# Which medium counties have no physicians?
mcounties[mcounties$Physicians == 0, c(1,5,12)]
      County Population Physicians
157 Live Oak
                  12091
                                 0
159
       Duval
                  11533
                                 0
```

Potential predictor variables

- LandArea: Area in square miles
- PctRural: Percentage rural land
- MedianIncome: Median household income
- Population: Population
- PctUnder18: Percent children
- PctOver65: Percent seniors
- PctPoverty: Percent below the poverty line
- PctUninsured: Percent without health insurance
- PctSomeCollege: Percent with some higher education
- PctUnemployed: Percent unemployed

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- This is where R's automated model building tools help.

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- But R² is not good for comparing models with different numbers of variables because it tends to increase a little bit with each additional variable, just due to randomness.
- Adjusted-R² is better because it multiplies R² by a penalty that depends on the number of variables, but the penalty is somewhat arbitrary and increases as the number of variables increases.

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- No method is ideal in all situations, so it is generally best to use multiple methods and look at the results.
- AIC (Akaike's Information Criterion) and the very similar BIC (your reading calls it SBC) are other widely used criterion that often gives different results than Adjusted-R².



Stepping forwards

The step function uses the AIC criterion to compare models. First we'll build a "null model" with no variables, and a "full model" with all variables:

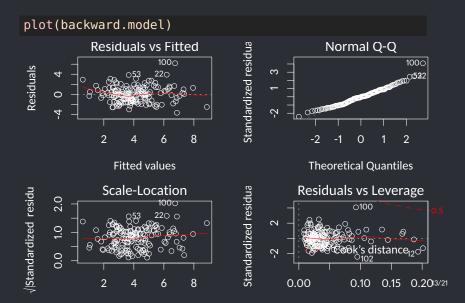


Stepping backwards and both ways

You can also step backwards (similar to what we have been doing manually), or in both directions:



Check assumptions



Check for multicollinearity

vif(backward.mod	lel)			
PctRural 1.911623 PctUnemployed 1.125032	Population 1.843085	Pct0ver65 1.776352	PctUninsured 1.029993	PctSomeCollege 1.541539

We can't be sure this is the best possible model.

Sometimes, stepwise regression leads you down a suboptimal path and you end up discarding a valuable variable (or keeping a variable that is only marginally useful), because of the order in which the variables are considered.

Best-subsets regression

• **Best-subsets regression** compares every possible model containing some subset of the predictor variables!

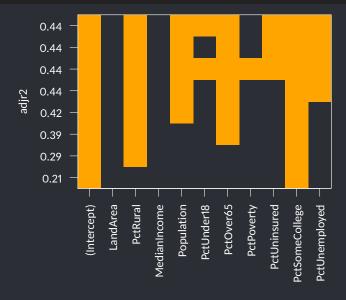
Best-subsets regression

- Best-subsets regression compares every possible model containing some subset of the predictor variables!
- Then we can compare the models using different model selection criteria and select the most parsimonious one

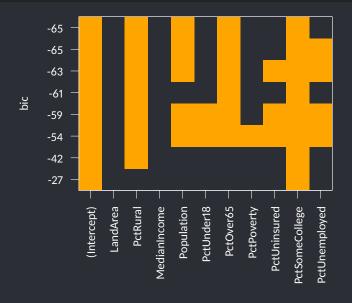
Best-subsets regression

Let's compare models using Adjusted R^2 . Each row is a candidate model; filled-in squares indicate the variable is included in that model:

plot(regsubsets.output, scale="adjr2")



plot(regsubsets.output, scale="bic")



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- Find the parsimonious middle ground between an underspecified model and extraneous variables.
- Fine-tune the model to ensure the model meets assumptions and captures key relationships: you may need to transform predictors and/or add interactions.
- Think about logical reasons why certain predictors might be useful, don't just focus on p-values.

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- Otherwise, you can select the ones that happen to fit the data the best and essentially create a spurious correlation!
- Remember to check for multicolliearity and the LINE assumptions!