

Sta 111 - Summer II 2017
Probability and Statistical Inference
22. Model selection and conditions for MLR

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Outline

1. Model selection

1. Model selection criterion depends on goal: significance vs. prediction
2. Backward-elimination
3. Forward-selection

2. Conditions for MLR

1. Checking model conditions using graphs

Professor rating

Data: Student evaluations of instructors' teaching quality for 463 courses at the University of Texas.

| | Estimate | Std. Error | t value | Pr(> t) |
|----------------------------------|----------|------------|---------|----------|
| (Intercept) | 4.6282 | 0.1720 | 26.90 | 0.00 |
| beauty ¹ | 0.1080 | 0.0329 | 3.28 | 0.00 |
| gender.male | 0.2040 | 0.0528 | 3.87 | 0.00 |
| age | -0.0089 | 0.0032 | -2.75 | 0.01 |
| formal.yes ² | 0.1511 | 0.0749 | 2.02 | 0.04 |
| lower.yes ³ | 0.0582 | 0.0553 | 1.05 | 0.29 |
| native.non english | -0.2158 | 0.1147 | -1.88 | 0.06 |
| minority.yes | -0.0707 | 0.0763 | -0.93 | 0.35 |
| students ⁴ | -0.0004 | 0.0004 | -1.03 | 0.30 |
| tenure.tenure track ⁵ | -0.1933 | 0.0847 | -2.28 | 0.02 |
| tenure.tenured | -0.1574 | 0.0656 | -2.40 | 0.02 |

¹beauty: the beauty judgements were made by six students who had not attended the classes and were not aware of the course evaluations.

²formal: picture wearing tie&jacket/blouse, levels: yes, no

³lower: lower division course, levels: yes, no

⁴students: number of students

⁵tenure: tenure status, levels: non-tenure track, tenure track, tenured

Hypotheses

Just as the interpretation of the slope parameters take into account all other variables in the model, the hypotheses for testing for significance of a predictor also takes into account all other variables.

$H_0 : \beta_i = 0$ when other explanatory variables are included in the model.

$H_A : \beta_i \neq 0$ when other explanatory variables are included in the model.

Assessing significance: numerical variables

The p-value for age is 0.01. What does this indicate?

| | Estimate | Std. Error | t value | Pr(> t) |
|-----|----------|------------|---------|----------|
| ... | | | | |
| age | -0.0089 | 0.0032 | -2.75 | 0.01 |
| ... | | | | |

- (a) Since p-value is positive, higher the professor's age, the higher we would expect them to be rated.
- (b) If we keep all other variables in the model, there is strong evidence that professor's age is associated with their rating.
- (c) Probability that the true slope parameter for age is 0 is 0.01.
- (d) There is about 1% chance that the true slope parameter for age is -0.0089.

Assessing significance: categorical variables

Tenure is a categorical variable with 3 levels: non tenure track, tenure track, tenured. Based on the model output given, which of the below is false?

| | Estimate | Std. Error | t value | Pr(> t) |
|---------------------|----------|------------|---------|----------|
| ... | | | | |
| tenure.tenure track | -0.1933 | 0.0847 | -2.28 | 0.02 |
| tenure.tenured | -0.1574 | 0.0656 | -2.40 | 0.02 |

- (a) Reference level is non tenure track.
- (b) All else being equal, tenure track professors are rated, on average, 0.19 points lower than non-tenure track professors.
- (c) All else being equal, tenured professors are rated, on average, 0.16 points lower than non-tenure track professors.
- (d) All else being equal, there is a significant difference between the average ratings of tenure track and tenured professors.

Model selection strategies

Based on what we've learned so far, what are some ways you can think of that can be used to determine which variables to keep in the model and which to leave out?

- ▶ If the goal is to find the set of statistically significant predictors of $y \rightarrow$ use p-value selection.
- ▶ If the goal is to do better prediction of $y \rightarrow$ use adjusted R^2 selection.
- ▶ Either way, can use *backward elimination* or *forward selection*.
- ▶ Expert opinion and focus of research might also demand that a particular variable be included in the model.

Backward-elimination

1. R_{adj}^2 approach:

- Start with the full model
- Drop one variable at a time and record R_{adj}^2 of each smaller model
- Pick the model with the highest increase in R_{adj}^2
- Repeat until none of the models yield an increase in R_{adj}^2

2. p-value approach:

- Start with the full model
- Drop the variable with the highest p-value and refit a smaller model
- Repeat until all variables left in the model are significant

Backward-elimination: R^2_{adj} approach

| Step | Variables included | R^2_{adj} |
|--|--|---------------|
| Full | beauty + gender + age + formal + lower + native + minority + students + tenure | 0.0839 |
| Step 1 | gender + age + formal + lower + native + minority + students + tenure | 0.0642 |
| | beauty + age + formal + lower + native + minority + students + tenure | 0.0557 |
| | beauty + gender + formal + lower + native + minority + students + tenure | 0.0706 |
| | beauty + gender + age + lower + native + minority + students + tenure | 0.0777 |
| | beauty + gender + age + formal + native + minority + students + tenure | 0.0837 |
| | beauty + gender + age + formal + lower + minority + students + tenure | 0.0788 |
| | <i>beauty + gender + age + formal + lower + native + students + tenure</i> | <i>0.0842</i> |
| | beauty + gender + age + formal + lower + native + minority + tenure | 0.0838 |
| | beauty + gender + age + formal + lower + native + minority + students | 0.0733 |
| Step 2 | gender + age + formal + lower + native + students + tenure | 0.0647 |
| | beauty + age + formal + lower + native + students + tenure | 0.0543 |
| | beauty + gender + formal + lower + native + students + tenure | 0.0708 |
| | beauty + gender + age + lower + native + students + tenure | 0.0776 |
| | <i>beauty + gender + age + formal + native + students + tenure</i> | <i>0.0846</i> |
| | beauty + gender + age + formal + lower + native + tenure | 0.0844 |
| Step 3 | beauty + gender + age + formal + lower + native + students | 0.0725 |
| | gender + age + formal + native + students + tenure | 0.0653 |
| | beauty + age + formal + native + students + tenure | 0.0534 |
| | beauty + gender + formal + native + students + tenure | 0.0707 |
| | beauty + gender + age + native + students + tenure | 0.0786 |
| | beauty + gender + age + formal + students + tenure | 0.0756 |
| | <i>beauty + gender + age + formal + native + tenure</i> | <i>0.0855</i> |
| beauty + gender + age + formal + native + students | 0.0713 | |
| Step 4 | gender + age + formal + native + tenure | 0.0667 |
| | beauty + age + formal + native + tenure | 0.0553 |
| | beauty + gender + formal + native + tenure | 0.0723 |
| | beauty + gender + age + native + tenure | 0.0806 |
| | beauty + gender + age + formal + tenure | 0.0773 |
| | beauty + gender + age + formal + native | 0.0713 |

best model →

Selected model

| | Estimate | Std. Error | t value | Pr(> t) |
|---------------------|----------|------------|---------|----------|
| (Intercept) | 4.6284 | 0.1673 | 27.66 | 0.00 |
| beauty | 0.1055 | 0.0328 | 3.21 | 0.00 |
| gender.male | 0.2081 | 0.0519 | 4.01 | 0.00 |
| age | -0.0088 | 0.0032 | -2.75 | 0.01 |
| formal.yes | 0.1324 | 0.0714 | 1.85 | 0.06 |
| native:non english | -0.2430 | 0.1080 | -2.25 | 0.02 |
| tenure:tenure track | -0.2068 | 0.0839 | -2.46 | 0.01 |
| tenure:tenured | -0.1760 | 0.0641 | -2.74 | 0.01 |

Backward-elimination: p – value approach

| Step | Variables included & p-value | | | | | | | | | |
|--------|------------------------------|----------------|------|---------------|--------------|-----------------------|-----------------|----------|------------------------|------------------------|
| Full | beauty | gender male | age | formal yes | lower yes | native non english | minority yes | students | tenure tenure track | tenure tenure track |
| | 0.00 | 0.00 | 0.01 | 0.04 | 0.29 | 0.06 | 0.35 | 0.30 | 0.02 | 0.01 |
| Step 1 | beauty | gender male | age | formal yes | lower yes | native non english | | students | tenure tenure track | tenure tenure track |
| | 0.00 | 0.00 | 0.01 | 0.04 | 0.38 | 0.03 | | 0.34 | 0.02 | 0.01 |
| Step 2 | beauty | gender male | age | formal yes | | native non english | | students | tenure tenure track | tenure tenure track |
| | 0.00 | 0.00 | 0.01 | 0.05 | | 0.02 | | 0.44 | 0.01 | 0.01 |
| Step 3 | beauty | gender male | age | formal yes | | native non english | | | tenure tenure track | tenure tenure track |
| | 0.00 | 0.00 | 0.01 | 0.06 | | 0.02 | | | 0.01 | 0.01 |
| Step 4 | beauty | gender male | age | | | native non english | | | tenure tenure track | tenure tenure track |
| | 0.00 | 0.00 | 0.01 | | | 0.06 | | | 0.01 | 0.01 |
| Step 5 | beauty | gender male | age | | | | | | tenure tenure track | tenure tenure track |
| | 0.00 | 0.00 | 0.01 | | | | | | 0.01 | 0.01 |

Best model: beauty + gender + age + tenure

Forward-selection

1. R_{adj}^2 approach:

- Start with regressions of response vs. each explanatory variable
- Pick the model with the highest R_{adj}^2
- Add the remaining variables one at a time to the existing model, and once again pick the model with the highest R_{adj}^2
- Repeat until the addition of any of the remaining variables does not result in a higher R_{adj}^2

2. p – value approach:

- Start with regressions of response vs. each explanatory variable
- Pick the variable with the lowest significant p-value
- Add the remaining variables one at a time to the existing model, and pick the variable with the lowest significant p-value
- Repeat until any of the remaining variables does not have a significant p-value

In forward-selection the p-value approach isn't any simpler (you still need to fit a bunch of models), so there's almost no incentive to use it.

Using the p-value approach, which variable would you remove from the model next?

| | Estimate | Std. Error | t value | Pr(> t) |
|---------------|-----------|------------|---------|----------|
| (Intercept) | -14022.48 | 11137.08 | -1.26 | 0.21 |
| hrs_work | 1045.85 | 149.05 | 7.02 | 0.00 |
| raceblack | -7636.32 | 6177.50 | -1.24 | 0.22 |
| raceasian | 29944.35 | 9137.13 | 3.28 | 0.00 |
| raceother | -7212.57 | 7212.25 | -1.00 | 0.32 |
| age | 559.51 | 133.27 | 4.20 | 0.00 |
| genderfemale | -17010.85 | 3699.19 | -4.60 | 0.00 |
| citizenyes | -13059.46 | 8219.99 | -1.59 | 0.11 |
| time_to_work | 88.77 | 79.73 | 1.11 | 0.27 |
| langother | -10150.41 | 5431.15 | -1.87 | 0.06 |
| marriedyes | 5400.41 | 3896.12 | 1.39 | 0.17 |
| educollege | 16214.46 | 4089.17 | 3.97 | 0.00 |
| edugrad | 59572.20 | 5631.33 | 10.58 | 0.00 |
| disabilityyes | -14201.11 | 6628.26 | -2.14 | 0.03 |

(a) married

(b) race

(c) race:other

(d) race:black

(e) time_to_work

Conditions for MLR are (almost) the same as conditions for SLR

Important regardless of doing inference

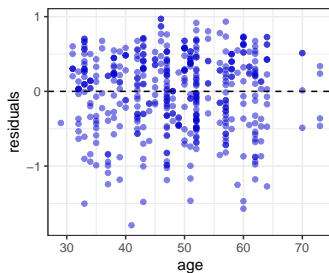
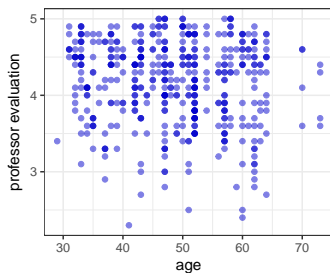
- ▶ *Linearity* → each variable is linearly related to the outcome

Important for doing inference

- ▶ *Nearly normally distributed residuals* → primary concern relates to residuals that are outliers
- ▶ *Constant variability of residuals (homoscedasticity)*
- ▶ *Independence of observations (and hence residuals)*
- ▶ Also important to make sure that your explanatory variables are *not collinear*

(1) linear relationships

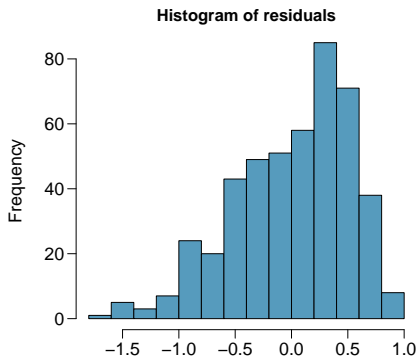
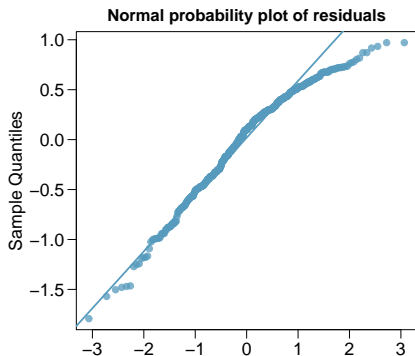
- ▶ For categorical variable, using boxplot of the residuals against each level to check whether variability fluctuates across levels.
- ▶ Using scatterplot of residuals vs. each numerical predictor to check if there is some possible structure such as curvature in the residuals.



Does this condition appear to be satisfied?

(2) nearly normal residuals

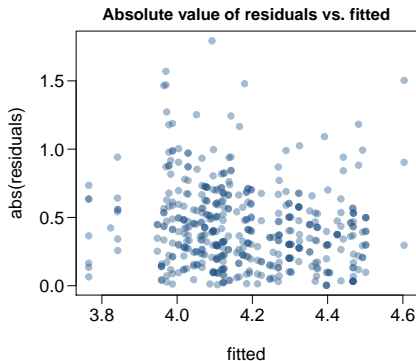
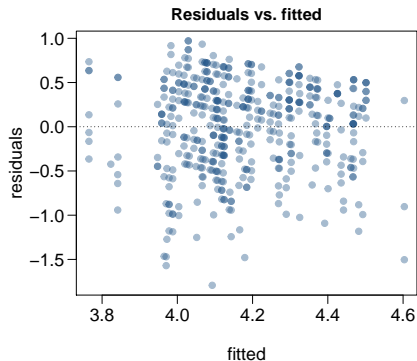
Q-Q plot and/or histogram of residuals:



Does this condition appear to be satisfied?

(3) constant variability in residuals

scatterplot of residuals and/or absolute value of residuals vs. fitted (predicted):



Does this condition appear to be satisfied?

Checking constant variance - recap

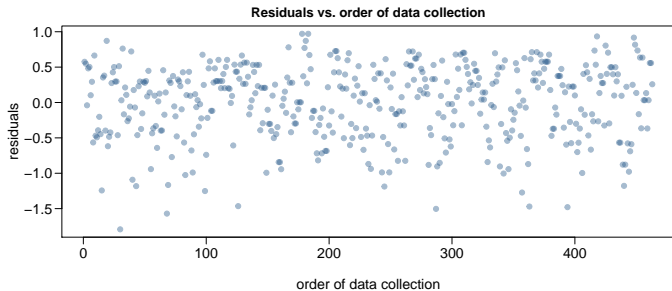
- ▶ When we did simple linear regression (one predictor) we checked the constant variance condition using a plot of *residuals vs. x*.
- ▶ With multiple linear regression (2+ predictors) we checked the constant variance condition using a plot of *residuals vs. fitted*.

Why are we using different plots?

In multiple linear regression there are many explanatory variables, so a plot of residuals vs. one of them wouldn't give us the complete picture.

(4) independent residuals

scatterplot of residuals vs. order of data collection:



Does this condition appear to be satisfied?

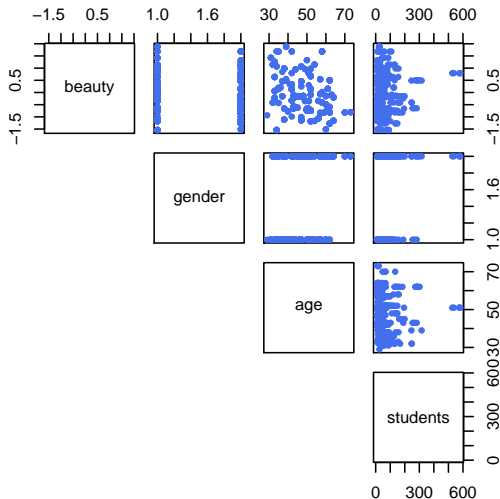
More on the condition of independent residuals

- ▶ Checking for independent residuals allows us to indirectly check for independent observations.
- ▶ If observations and residuals are independent, we would not expect to see an increasing or decreasing trend in the scatterplot of residuals vs. order of data collection.
- ▶ This condition is often violated when we have time series data. Such data require more advanced time series regression techniques for proper analysis.

(5) Checking collinearity among predictors

Use pairwise correlations to check collinearity.

```
pairs(~beauty+gender+age+students, pch = 19,  
      col = "royalblue", lower.panel = NULL)
```



Which of the following is the appropriate plot for checking the homoscedasticity condition in MLR?

- (a) scatterplot of residuals vs. \hat{y}
- (b) scatterplot of residuals vs. x
- (c) histogram of residuals
- (d) Q-Q plot of residuals
- (e) scatterplot of residuals vs. order of data collection

Plotting residuals against \hat{y} (predicted, or fitted, values of y) allows us to evaluate the whole model as a whole as opposed to homoscedasticity with regards to just one of the explanatory variables in the model.