# Stat 306: Finding Relationships in Data.

Lecture 17 Sections 4.4 and 4.5

## **BONUS FOR LEAVE-ONE-OUT Cross Validation**

The cross-validated root mean square (prediction) error is:

(4.26) 
$$CVRMSE_{leaveoneout}(x_1, \dots, x_p) = \sqrt{n^{-1} \sum_{i=1}^{n} (y_i - \hat{y}_{i|-i})^2}.$$

 $CVRMSE_{leaveoneout}(\mathbf{x}_J)$  can be also defined for the subset of the explanatory variables indexed by  $J \subset \{1,\ldots,p\}$ .

It turns out there is a simple formula for cross-validation leave-one-out residuals (without having to compute n different regressions). An identity is

(4.27) 
$$y_i - \hat{y}_{i|-i} = (y_i - \hat{y}_i)/(1 - P_{ii}), \quad P_{ii} = \mathbf{x}_i^T (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{x}_i,$$

where  $\mathbf{x}_i^T = (1, x_{i1}, \dots, x_{ip})$ . Alternatively,  $P_{ii}$  is the *i*th diagonal element of the projection matrix  $\mathbf{P} = \mathbf{X}(\mathbf{X}^T\mathbf{X})^{-1}\mathbf{X}^T$ . See Section 4.3 for why this quantity is called the projection matrix.

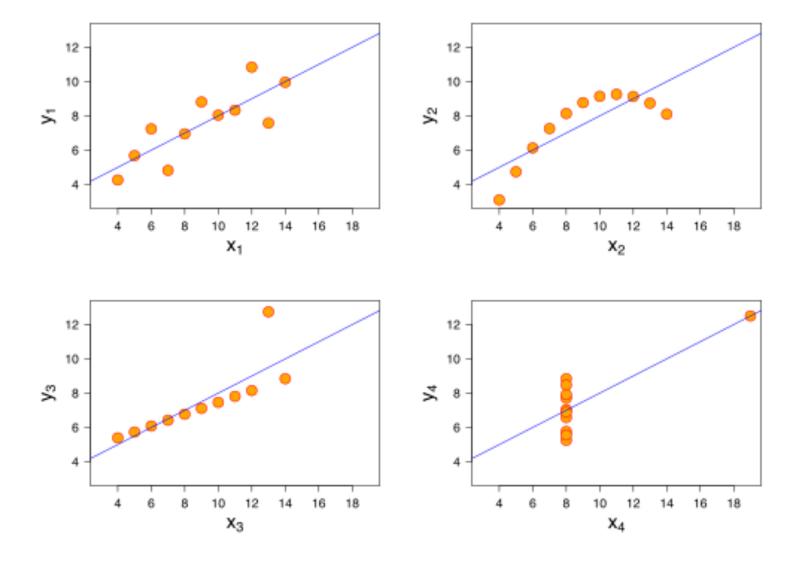
# Chapter 4 – Variable selection and additional diagnostics

- 4.1 Variable Selection algorithms
- 4.2 Cross-validation and out-of sample assessment

## 4.3 Additional diagnostics

- 4.4 Transforms and nonlinearity
- 4.5 Diagnostics for data collected sequentially in time

## Classic example: Anscombe's quartet



 https://www.refsmmat.com/regression/ regression.html

# Chapter 4 – Variable selection and additional diagnostics

- 4.1 Variable Selection algorithms
- 4.2 Cross-validation and out-of sample assessment
- 4.3 Additional diagnostics
- 4.4 Transforms and nonlinearity
- 4.5 Diagnostics for data collected sequentially in time

In this section, we discuss when transforms of variables might be needed and how to handle diagnostics that suggest that g in (3.33) is nonlinear in the explanatory variables. The general prediction model is:

$$(4.32) Y_i = g(x_{i1}, \dots, x_{ip}) + \epsilon_i, \quad i = 1, \dots, n, \quad \epsilon_i \sim N(0, \sigma^2(\mathbf{x}_i)) \text{ independently},$$

where now we indicate heteroscedasticity with  $\sigma^2(\mathbf{x})$  being a function of the explanatory variables.

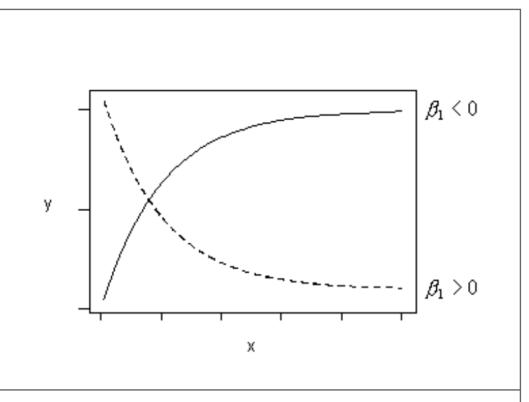
Sometimes after a transform of y or one or more of the explanatory variables might lead to the homoscedastic model with g linear being a better approximation to the data.

In particular, if diagnostics suggest heteroscedasticity with more variability and y increases, then a log or square root or power transform of y can be useful (assuming y is positive-valued). Examples of this are shown in Exercise 2.7 and in the case study in Chapter 5.

If the trend in your data follows either of these patterns, you could try fitting this regression function:

$$\mu_Y = \beta_0 + \beta_1 e^{-x}$$

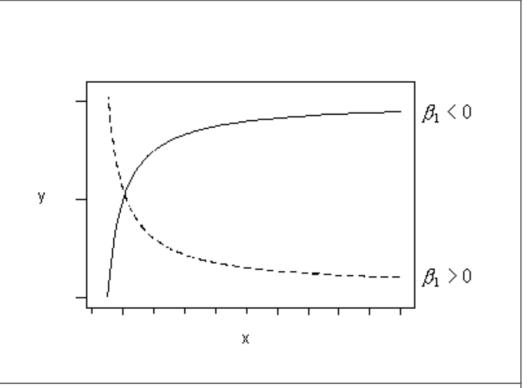
to your data.



Or, if the trend in your data follows either of these patterns, you could try fitting this regression function:

$$\mu_Y = \beta_0 + \beta_1 \left(\frac{1}{x}\right)$$

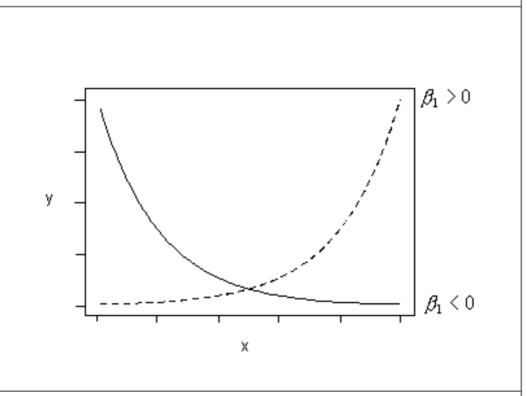
to your data. (This is sometimes called a "reciprocal" transformation.)



Or, if the trend in your data follows either of these patterns, try fitting this regression function:

$$\mu_{lnY} = \beta_0 + \beta_1 x$$

to your data. That is, fit the model with ln(y) as the response and x as the predictor.

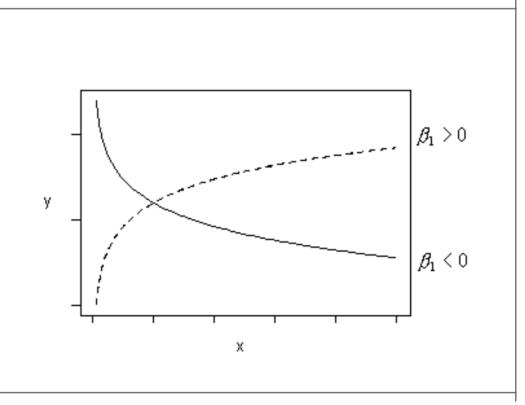


source: https://onlinecourses.science.psu.edu/stat501/node/322

Or, try fitting this regression function:

$$\mu_Y = \beta_0 + \beta_1 \ln(x)$$

if the trend in your data follows either of these patterns. That is, fit the model with y as the response and ln(x) as the predictor.

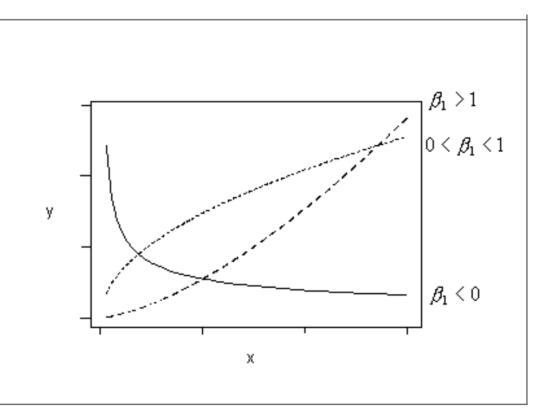


source: https://onlinecourses.science.psu.edu/stat501/node/322

And, finally, try fitting this regression function:

$$\mu_{lnY} = \beta_0 + \beta_1 ln(x)$$

if the trend in your data follows any of these patterns. That is, fit the model with ln(y) as the response and ln(x) as the predictor.



source: https://onlinecourses.science.psu.edu/stat501/node/322

## It's easy to make transformations.

If you see that the variance of the residuals is not constant with the mean of the fitted values... you just try a transform. For example:  $y \rightarrow \log(y)$  Do the residuals now look any better?

## Interpretation is where things get tough.

Good explanations at:

https://stats.idre.ucla.edu/other/mult-pkg/faq/general/faqhow-do-i-interpret-a-regression-model-when-somevariables-are-log-transformed/

The linear model: 
$$y_i = \beta_0 + \beta_1 x_i + \epsilon_i$$
 ,  $i = 1, ..., n$ 

## Interpretation:

For every one unit increase in x, the mean of y increases by  $\beta_1$  units.

The linear model: 
$$y_i = \beta_0 + \beta_1 x_i + \epsilon_i$$
 ,  $i = 1, ..., n$ 

## Interpretation:

 $> lm(y\sim x1+x2)$ 

For every one unit increase in x, the mean of y increases by  $\beta_1$  units.

**Conclusion:** For every additional year in age, an individual's amount of money increases on average by an estimated of 0.54 (95% C.I. = 0.01, 1.06), when adjusted for income (x2).

The linear-log model: 
$$y_i = \beta_0 + \beta_1 log(x_i) + \epsilon_i$$
 ,  $i = 1, ..., n$ 

## Interpretation:

For every one unit increase in log(x), the mean of y increases by  $\beta_1$  units.

or:

A p% increase in X is associated with an increase in the mean of y of  $\beta_1 \log([100+p]/100)$  units.

The linear-log model: 
$$y_i = \beta_0 + \beta_1 \log(x_i) + \epsilon_i$$
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## Interpretation:

A p% increase in X is associated with an increase in the mean of y of  $\beta_1 \log([100+p]/100)$  units.

```
> lm(y~I(log(x1))+x2)
```

Call:  

$$lm(formula = y \sim I(log(x1)) + x2)$$

### Consider a 22 year old: $exp(y_{22}) = exp(\beta_0) + exp(\beta_1 \log(22))$

$$\exp(y_{20}) = \exp(\beta_0) + \exp(\beta_1 \log(20))$$

#### We have:

```
\exp(y_{22})/\exp(y_{20}) = (22/20)^{\beta 1}
```

## or equivalently:

$$\exp(y_{22}-y_{20}) = (22/20)^{\beta 1}$$

## or equivalently:

$$y_{22}$$
- $y_{20}$  = log[(22/20) <sup>$\beta$ 1</sup>] = **1.7**

**Conclusion:** For any 10% increase in age, an individual's amount of money increases on average by an estimated of \$1.7 (95% C.I. = [\$-2, \$37]), when adjusted for income (x2).

The log-linear model: 
$$log(y_i) = \beta_0 + \beta_1 x_i + \epsilon_i$$
,  $i = 1, ..., n$ 

## Interpretation:

For every one unit increase in x, the mean of log(y) increases by  $\beta_1$  units. or:

The effect of a p-unit increase in X is to multiply the mean of y by  $exp(p\beta_1)$ .

The log-linear model: 
$$log(y_i) = \beta_0 + \beta_1 x_i + \epsilon_i$$
 ,  $i = 1, ..., n$ 

## Interpretation:

The effect of a p-unit increase in X is to multiply the mean of y by  $exp(p\beta_1)$ .

> lm(log(y)~x1+x2)

i(tog(y)~xi+xz)

Call:  $lm(formula = log(y) \sim x1 + x2)$ 

ffi ai anta.

Coefficients:
(Intercept)

tercept) x1 2.71759 0.01580 x2 0.00484 Consider a 21 year old:  $y_{22} = \exp(\beta_0 + \beta_1 21)$ 

and a 20 year old:

 $y_{20} = \exp(\beta_0 + \beta_1 20)$ 

We have:

 $y_{21}/y_{20} = \exp(\beta_1 (21-20))$ 

or equivalently:

 $y_{21} = y_{20} (\exp(\beta_1(1)))$ 

or equivalently:  $y_{21} = y_{20} (1.02)$ 

**Conclusion:** For every additional year in age, an individual's amount of money increases on average by a factor of 1.02 (95% C.I. = [1.00, 1.04]), when adjusted for income (x2).

The log-linear model: 
$$log(y_i) = \beta_0 + \beta_1 x_i + \epsilon_i$$
 ,  $i = 1, ..., n$ 

## Interpretation:

The effect of a p-unit increase in X is to multiply the mean of y by  $exp(p\beta_1)$ .

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**Conclusion:** For every additional year in age, an individual's amount of money increases on average by a factor of 1.02 (95% C.I. = [1.00, 1.04]), when adjusted for income (x2).

The log-log model: 
$$log(y_i) = \beta_0 + \beta_1 log(x_i) + \epsilon_i$$
 ,  $i = 1, ..., n$ 

## Interpretation:

For every one unit increase in log(x), the mean of log(y) increases by  $\beta_1$  units.

## or:

For every p-unit increase in X, the mean of y is multiplied by  $\exp(a\beta_1)$ , where  $a=\log([100+p]/100)$ .

The log-log model: 
$$log(y_i) = \beta_0 + \beta_1 log(x_i) + \epsilon_i$$
 ,  $i = 1, ..., n$ 

## Interpretation:

For every p-unit increase in X, the mean of y is multiplied by  $\exp(a\beta_1)$ , where  $a=\log([100+p]/100)$ .

## **Exercise:**

How would you interpret this?

How would you obtain a the proper confidence interval?

Good explanations and examples from economics:

http://kenbenoit.net/assets/courses/ME104/logmodels2.pdf

## 4.5 Diagnostics for data collected sequentially in time

If data are collected sequentially in time, the assumption of independent  $\epsilon_i$  in model (3.36) might not be appropriate. In this situation, sometimes  $\epsilon_i$  are serially correlated. If the  $\epsilon_i$  are positive serially correlated, then SEs of  $\hat{\beta}$ 's derived under the assumption of independent  $\epsilon_i$  are generally too small.

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As before, suppose least square has been applied to get  $\hat{\beta}$  and residuals  $e_1, \ldots, e_n$ . One additional residual plot is  $e_t$  versus t for  $t = 1, \ldots, n$ . If there are trends or patterns in this plot, then the assumption of independent  $\epsilon_t$  is violated. There can be several patterns, and two of them are as follows; see also Figure 4.5.

- (a) A positive  $e_t$  is often followed by a negative residual and a negative  $e_t$  is often followed by a positive residual. This is negative serial correlation, and the plot shows with many crossings of the horizontal line if consecutive residuals are joined by line segments.
- (b) A positive  $e_t$  is often followed by a positive residual and a negative  $e_t$  is often followed by a negative residual. This is positive serial correlation, and the plot shows alternating runs of several positive residuals and runs of several negative residuals, with few crossings of the horizontal line if consecutive residuals are joined by line segments.

As before, suppose least square has been applied to get  $\hat{\beta}$  and residuals  $e_1, \ldots, e_n$ . One additional residual plot is  $e_t$  versus t for  $t = 1, \ldots, n$ . If there are trends or patterns in this plot, then the assumption of independent  $\epsilon_t$  is violated.

Important Question: Is there any "serial correlation"?

To find out, first plot the residuals vs. time.

Below is the table of the lagged residual  $e_{t-1}$  versus  $e_t$ 

$e_1$	$e_2$
$e_2$	$e_3$
÷	:
$e_{n-1}$	$e_n$

The lag 1 serial correlation (with  $\bar{e} = 0$ ) is

$$\hat{\rho}_1 = \frac{\sum_{t=2}^n e_{t-1} e_t}{\sqrt{\sum_{t=2}^n e_{t-1}^2 \sum_{t=2}^n e_t^2}} \in (-1, 1).$$

One could also look at scatterplot of  $e_{t-1}$  versus  $e_t$  to check if there appears to be significant serial correlation

The Durbin-Watson statistic is

(4.33) 
$$DW = \frac{\sum_{t=2}^{n} (e_t - e_{t-1})^2}{\sum_{t=1}^{n} e_t^2}.$$

If n is large, then

$$DW = \frac{\sum_{t=2}^{n} e_t^2 + \sum_{t=2}^{n} e_{t-1}^2 - 2\sum_{t=2}^{n} e_{t-1}e_t}{\sum_{t=1}^{n} e_t^2} \approx 2 - 2\hat{\rho}_1.$$

Hence  $DW \approx 2$  for serially uncorrelated residuals,  $DW \approx 0$  for strong positive serial correlation ( $\hat{\rho}_1$  near 1) and  $DW \approx 4$  for strong negative serial correlation ( $\hat{\rho}_1$  near -1), since

$$(4.34) -1 \le \hat{\rho}_1 \le 1 \Rightarrow 0 \le 2 - 2\hat{\rho}_1 \le 4.$$

https://www.youtube.com/watch?v=QGN073tpDXE&ab\_channel=BenLambert

### 4.7 Summary of variable selection

Variable selection methods are algorithms for finding a good subset of explanatory variables. Cross-validation, either leave-one-out or with training/holdout subsets, provide an out-of-sample comparison of different prediction equations; the assessment of predictability is applied to data that are not used in the estimation of regression coefficients. Cross-validated and out-of-sample root mean squared errors combined with residual plots and other diagnostics could provide information that transforms of variables are needed or that the response variables in nonlinear in the explanatory variables.