

# Econometrics Part 2

## The linear model in practice

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# Outline

- 1 Multicollinearity
- 2 Handling categories
- 3 Omitted variables
- 4 Predictions in logs
- 5 Influential observations

- This problem arises if some variable is equal to an exact linear combination of some other variables (e.g. if we have variables like *income*, *income after tax* and *tax*)
- This variable is unnecessary because it contains repetitive information
- Worse, it prevents the computation of the OLS estimator, since  $X$  is no longer full rank and we cannot compute  $\hat{b} = (X'X)^{-1}X'y$
- If the software detects multicollinearity, it arbitrarily removes one repetitive variable

- This problem arises if some variable is *almost* equal to an exact linear combination of some other variables
- In other words, it means that they are very strongly correlated
- We *can* compute  $\hat{b} = (X'X)^{-1}X'y$
- But :  $\hat{b}$  is very unstable and the estimator is not very reliable (great variance)

# How to detect near multicollinearity

- The VIF (Variance Inflation Factor) indicates by how much the variance of a parameter  $b_j$  is inflated due to the correlation of variable  $X_j$  with all the other explanatory variables
- Python command : `from statsmodels.stats.outliers_influencers import variance_inflation_factor`
- Say our model has  $k$  variables : constant,  $X_1, X_2, \dots, X_{k-1}$
- Python will run the regression of each  $X_i$  on all the other  $X_j$ 's, save its  $R^2$  called  $R_i^2$
- Example :  $X_1 = a_1 + a_2 X_2 + \dots + a_{k-1} X_{k-1} + \varepsilon$
- Next, VIF for each variable  $i$  is computed as :  $VIF_i = \frac{1}{1-R_i^2}$
- One can prove that in the original regression  
 $y = b_0 + b_1 X_1 + \dots + b_{k-1} X_{k-1} :$
- $V(\hat{b}_j) = \frac{\sigma^2}{(N-1)Var(X_j)} VIF_j$
- $VIF_i > 5$  indicates a high risk of multicollinearity between variable  $i$  and the other variables
- If you don't see any big issue, do not remove variables : risk of omitted variable bias (see below)

# Dummy variables : 2 categories

- Let's say we have  $N$  individual observations providing income, gender and the number of years of education.
- We want to explain individual income  $y$  by education  $x$  and gender  $z$  :  
$$y_i = a + b.x_i + c.z_i + u_i.$$
- $z$  is a *categorical variable*, ideally coded with 0/1
- Say  $z = 1$  codes for males
- $c$  represents the extra money that provides the fact of being a male with respect to being a female
- It would be irrelevant to have a male *and* a female variable because it would cause *exact multicollinearity* (besides the fact that it is unnecessary)
- Remark : setting a categorical variable as the dependent variable needs other type of modeling, such as Probit and Logit (see later chapters)

# Dummy variables : more than 2 categories

- Let's say we have  $N$  individual observations providing an index of life satisfaction, income and type of environment (*big city*, *small city*, *rural*), respectively coded with values 1, 2 and 3.
- We want to explain life satisfaction  $y$  by income  $x$  and environment  $z$
- $z$  is still a *categorical variable*, but we have 3 categories : we have to choose a category with respect to which parameters will be computed, say *rural*
- The model should be written :  
$$y_i = a + b.x_i + c_1.z_{1i} + c_2.z_{2i} + u_i$$
, with  $z_1$  and  $z_2$  are dummy variables that code for *big city* and *small city* respectively
- These dummy variables should be created by hand or through pandas
- Python command : `pd.get_dummies(df['environment'], drop_first=True)` or using statsmodels formula with `C(environment)`
- Do not keep original coding (1, 2, 3), which would be meaningless in a regression

# Dummy variables : outliers

- Let's say we have  $N$  observations for a country that experienced a shock.
- We want to explain life expectancy  $y$  by number of doctors per capita  $x$  :  
$$y_t = a + b.x_t + u_t.$$
- Let's say that the shock took place on year  $i$  : this unusual observation is likely to bias the whole regression, but we don't want to delete it
- We will take this event into account with a dummy variable :
- $D_t = 1$  if  $t = i$ ,  $D_t = 0$  otherwise.
- The model is now written :  $y_t = a + b.x_t + c.D_t + u_t.$
- $c$  represents the impact of the shock



- It can be easily understood that estimating the previous model with and without dummy  $D$  can lead to changes even in parameters  $a$  and  $b$
- Omitting a relevant variable (here  $D$ ) leads to *omitted variable bias* that affects all estimated parameters
- Proof : in the finite sample case with the Frish-Waugh theorem (see Dormont), and proof involving consistency in later chapters (on endogeneity)
- This is a very important issue in applied work, so try not to forget important variables in models

# The Frish-Waugh theorem

- Let  $Z$  be a vector of variables.
- $M_Z$  is the orthogonal projection matrix on  $L^\perp(Z)$ .
- Let's define 2 models :
  - ①  $y = Xb_{(i)} + Zc + u$
  - ②  $M_Z y = M_Z X b_{(ii)} + M_Z u$ .
- The Frish-Waugh theorem states that  $\hat{b}_{(i)} = \hat{b}_{(ii)}$
- Important consequence : estimating the model with both sets of variables  $X$  and  $Z$  **or** with  $X$  alone will not provide the same results.
- This is called the "omitted variable bias"

*On which condition will results be the same ?*

- Estimating the model with both sets of variables  $X$  and  $Z$  **or** with  $X$  alone will yield the same results only if vectors  $X$  and  $Z$  are perpendicular, i.e. totally uncorrelated
- i.e.  $X'Z = 0$
- Which is quite unlikely





# About the log transformation

- The average predicted  $y$  should be equal to the average observed  $y$
- Here, we can compute a prediction of  $\log(y)$
- Can we get back to raw predicted wage simply by taking the exponential of the predicted  $y$  in  $\log$  ?

# Log and predictions (1)

- Let's use model  $\forall i, \log(y_i) = a + bx_i + u_i$  (what matters is that we take the *log* of  $y$ )
- Taking the exponential :  
$$y_i = \exp(a + bx_i + u_i) = \exp(a + bx_i) \cdot \exp(u_i)$$
- So if the  $u$ 's are independent from the  $x$ 's, we get :  
$$E[y_i] = E[\exp(a + bx_i) \cdot \exp(u_i)] = E[\exp(a + bx_i)] \cdot E[\exp(u_i)]$$
- We usually consider the  $x$ 's as fixed, so :  
$$E[y_i] = \exp(a + bx_i) \cdot E[\exp(u_i)]$$
- A prediction for  $y$  is thus :  $\hat{y}_i = \exp(\hat{a} + \hat{b}x_i) \cdot \hat{E}[\exp(u_i)]$ ,
- But how to estimate  $\hat{E}[\exp(u_i)]$  (the retransformation parameter) ?

## Log and predictions (2)

It can be proven that if  $u \hookrightarrow N(0, \sigma^2)$ , then  $E[\exp(u_i)] = \exp(\frac{\sigma^2}{2})$

This can be estimated using  $\exp(\frac{\hat{\sigma}^2}{2}) = \hat{\psi}$

We thus have :  $\hat{y}_i = \exp(\hat{a} + \hat{b}x_i) \cdot \hat{\psi}$

To get back to levels when using a prediction in *logs*, we cannot simply take the exponential of the prediction : there is an additional retransformation parameter to take into account.

The preceding formula holds only if the  $u$ 's are normal : if their distribution is unknown but homoscedastic (same variance), we can use the non-parametric<sup>1</sup> *smearing* estimator :

$$\hat{\psi} = \frac{\sum \exp(\hat{u}_i)}{N}$$

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<sup>1</sup>Because no assumptions on the type of distribution



# Are all observations given the same weight ?

We have :

$$\hat{b}_1 = \frac{\sum (x_i - \bar{x})(y_i - \bar{y})}{\sum (x_i - \bar{x})^2}$$

So :

$$\hat{b}_1 = \frac{\sum (x_i - \bar{x})^2 \frac{(y_i - \bar{y})}{(x_i - \bar{x})}}{\sum (x_i - \bar{x})^2} = \sum p_i \frac{(y_i - \bar{y})}{(x_i - \bar{x})}$$

Calling  $p_i = \frac{(x_i - \bar{x})^2}{\sum (x_i - \bar{x})^2}$ .

$\frac{(y_i - \bar{y})}{(x_i - \bar{x})}$  is the slope of the line drawn from the point corresponding to individual  $i$  to the sample average, and  $p_i$  is an increasing function of  $(x_i - \bar{x})$ .

Estimate  $\hat{b}_1$  is thus highly influenced by extreme points (see Anscombe's quartet of identical regressions).

# Outliers (1)

- Outlier : observation with a large residual compared to other observations
- They indicate poor goodness of fit
- How to detect them : using a residuals plot
- Goal : estimate how far the observed value is from the predicted value
- Or more rigorously : using the *studentized* residuals, that adjust for the *leverage* of that particular point

## Outliers (2)

- Outliers are observations that have a large residual
- However, if the observation is highly influential, it has pulled the predicted value towards itself
- In other words, the variances of the residuals differ, even though the variances of the true errors are all equal to each other
- Consequence : need for studentization
- The *studentized* residuals adjust for the leverage of that particular point (see what  $h$  is below)

$$r_{stud_i} = \frac{\hat{u}_i}{\hat{\sigma}\sqrt{1-h_i}}$$

- High leverage observation : has an explanatory variable that takes a value very different from the sample mean
- In a simple regression model, could be (for example) at the far right and would influence the slope of the regression line
- How to detect them : the "hat statistic"

## Leverage (2)

- We know that  $\hat{y} = P_X y$
- $P_X$  can be called the "hat" matrix (puts a hat on  $y$ )
- Let's call  $h_i$  the  $i^{th}$  element of the diagonal of the hat matrix
- $h_i$  is the *leverage* of observation  $i$  : if it is large, then observation  $i$  has a high leverage
- $0 < h_i < 1$

- We know that  $\hat{u} = (I - P_X)u$ , so  $V(\hat{u}) = \sigma^2(I - P_X)$
- So :  $V(\hat{u}_i) = \sigma^2(1 - h_i)$
- The greater the leverage  $h_i$ , the smaller the variance of the residual for observation  $i$
- Meaning : high leverage observations tend to bring the regression line close to them
- High leverage observations are not necessarily influential, but low leverage observations won't be influential
- For the simple linear model :  $h_i = \frac{1}{N} + \frac{(x_i - \bar{x})^2}{\sum (x_i - \bar{x})^2}$

# Influence measures (1)

- Influential observation : if removed from the regression, estimated parameters vary a lot
- How to detect them : the *Dfbeta* statistic
- For every variable  $k$ , its *Dfbeta* measures how much its coefficient would change if we removed one observation
- Every *Dfbeta* possible is computed, and the software provides its maximum and minimum
- The change in  $\hat{b}_k$  if observation  $i$  is dropped is :

$$Dfbeta_{i,k} = \frac{\hat{b}_k - \hat{b}_{k,(-i)}}{\hat{\sigma}_{k,(-i)}}$$

with  $\hat{b}_{k,(-i)}$  and  $\hat{\sigma}_{k,(-i)}$  computed without observation  $i$ . It represents the number of se's by which the  $\hat{b}_k$  changes when observation  $i$  is removed.

## Influence measures (2)

- *Dffit* : measures how much the prediction for observation  $i$  changes when the estimation is made *without* observation  $i$
- $Dffit = \frac{\hat{y}_i - \hat{y}_{i,(-i)}}{\hat{\sigma}_{(-i)}\sqrt{h_i}}$
- *Covratio* statistic : measures the change in the determinant of the covariance matrix of the estimates by deleting the  $i$ th observation (indicates loss in precision)
- *Cook's distance* : measures the influence of observation  $i$  on the model as a whole, not on a single coefficient (it combines all the *Dfbeta*'s)
- $D_i = \frac{(\hat{b} - \hat{b}_{(-i)})'(X'X)(\hat{b} - \hat{b}_{(-i)})}{k\hat{\sigma}^2}$



- Leverage  $h_i$  of observation  $i$  :  $\bar{h} = k/N$ ,  $0 \leq h_i \leq 1$ . If for observation  $i$ ,  $h_i > \bar{h}$  then it might be influential.
- Change in predicted value  $Dffit$  : observation  $i$  is influential if  $|Dffit_i| > 2$  (general cutoff) or  $> 2\sqrt{k/N}$
- Change in a parameter  $Dfbeta$  : observation  $i$  is influential if  $|Dfbeta_{k,i}| > 2$  (general cutoff) or  $> 2/\sqrt{N}$  (adj. for large sample)
- Impact on the whole model (Cook's distance  $D$ ) : observation  $i$  is influential if  $D_i > 1$
- "Covratio" statistic : observation  $i$  is influential if  $|covratio - 1| > 3k/N$