Regression 1: Framework

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Introduction

Observational Study (観察研究)

- Researchers in social science cannot always conduct a randomized control trial.
- Instead, we need to use **observational data** in which treatment assignment may not be random.
- An approach in this case is **controlling observable characteristics** that causes a selection bias.
- This approach is essentially estimation of linear regression model (線形回帰モデル) by ordinally least squares (OLS, 最小二乗法).

Overview

- Introduce an idea of matching (マッチング) estimator.
 - Identification of treatment effect under **selection on observable** assumption.
 - Linear regression is a special case of matching estimator.
- Linear regression: framework, practical topics, inference

Selection on Observables, or Matching

Matching to eliminate a selection bias

- Idea: Compare individuals with the same observed characteristics X across treatment and control groups
- If treatment choice is driven by observed characteristics (such as age, income, gender, etc), controlling for such factor would eliminate the selection.
- Two key assumptions in matching

Assumption 1: Selection on observables

- Let X_i denote the observed characteristics (sometimes called **covariates (共変量)**) \circ age, income, education, race, etc..
- Assumption 1:

 $D_i \perp (Y_{0i},Y_{1i}) \ket{X_i}$

- **Conditional on** X_i , treatment assignment is random.
- This assumption is often referred in a different name:
 - $\circ~$ Selection on observables
 - Ignorability
 - \circ Unconfoundedness

Assumption 2: Overlapping assumption

• Assumption 2:

$$P(D_i=1|X_i=x)\in (0,1) \ orall x$$

- Given *x*, we should be able to observe people from both control and treatment group.
- The probability $P(D_i = 1 | X_i = x)$ is called **propensity score (傾向スコア)**.

Identification of Treatment Effect Parameters

• Assumption 1 (unconfoundedness) implies that

$$egin{aligned} &E[Y_{1i}|D_i=1,X_i]=E[Y_{1i}|D_i=0,X_i]=E[Y_{1i}|X_i]\ &E[Y_{0i}|D_i=1,X_i]=E[Y_{0i}|D_i=0,X_i]=E[Y_{0i}|X_i] \end{aligned}$$

• Once you conditioning on X_i , the argument is essentially the same as the one in RCT.

• The ATT conditional on $X_i = x$ is given by

$$egin{aligned} E[Y_{1i}-Y_{0i}|D_i=1,X_i] &= E[Y_{1i}|D_i=1,X_i] - E[Y_{0i}|D_i=1,X_i] \ &= E[Y_{1i}|D_i=1,X_i] - E[Y_{0i}|D_i=0,X_i] \end{aligned}$$

• Assumption 2 (overlapping) is needed to use the following

$$E[Y_{di}|D_i = d, X_i] = E[Y_i|D_i = d, X_i] ext{ for } d = 0, 1$$

- Why? Overlapping assumption $P(D_i = 1 | X_i = x) \in (0, 1)$ means that for each x, we should have people in both treatment and control group.
- If not, we cannot observe both $E[Y_i|D_i=d,X_i]$ for d=0,1.
- With two assumptions,

$$E[Y_{1i} - Y_{0i} | D_i = 1, X_i] = \underbrace{E[Y_i | D_i = 1, X_i]}_{avg \ with \ X_i \ in \ treatment} - \underbrace{E[Y_i | D_i = 0, X_i]}_{avg \ with \ X_i \ in \ control}$$

ATT $E[Y_{1i} - Y_{0i}|D_i = 1]$

• ATT is given by

$$egin{aligned} ATT &= E[Y_{1i} - Y_{0i} | D_i = 1] \ &= \int E[Y_{1i} - Y_{0i} | D_i = 1, X_i = x] f_{X_i}(x | D_i = 1) dx \ &= E[Y_i | D_i = 1] - \int \left(E[Y_i | D_i = 0, X_i = x]
ight) f_{X_i}(x | D_i = 1) \end{aligned}$$

ATT $E[Y_{1i}-Y_{0i}]$

• ATE is

$$egin{aligned} ATE =& E[Y_{1i}-Y_{0i}] \ &= \int E[Y_{1i}-Y_{0i}|X_i=x]f_{X_i}(x)dx \ &= \int E[Y_{1i}|D_i=1, X_i=x]f_{X_i}(x)dx + \int E[Y_{0i}|D_i=0, X_i=x]f_{X_i}(x)dx \ &= \int E[Y_i|D_i=1, X_i=x]f_{X_i}(x)dx + \int E[Y_i|D_i=0, X_i=x]f_{X_i}(x)dx \end{aligned}$$

From Identification to Estimation

- We need to estimate two conditional expectations $E[Y_i | D_i = 1, X_i = x]$ and $E[Y_i | D_i = 0, X_i = x]$
- Several ways to implement this.

1. Regression: Nonparametric and Parametric 2. Nearest neighborhood matching (最近傍マッチング)

- 3. Propensity Score Matching (傾向スコアマッチング)
- Here, I only explain a parametric regression as a way to implement the matching method.
- See Appendix and textbooks for the details of matching estimators.

From Matching to Linear Regression Model

• Assume that

$$E[Y_i|D_i=0,X_i=x]=eta'x_i
onumber \ E[Y_i|D_i=1,X_i=x]=eta'x_i+ au$$

- Here, treament effect is given by τ .
- You will have a linear regression model

$$y_i = eta' x_i + au D_i + \epsilon_i, E[\epsilon_i | D_i, x_i] = 0$$

• Running a linear regression to obtain the treatment effect parameter τ .

Linear Regression: Framework

Regression (回帰) framework

• Linear regression model (線形回帰モデル) is defined as

$$Y_i = eta_0 + eta_1 X_{1i} + \dots + eta_K X_{Ki} + \epsilon_i$$

- $\circ~i$: index for observations. $i=1,\cdots,N$.
- $\circ Y_i$: dependent variable (被説明変数)
- X_{ki} : explanatory variable (説明変数)
- *ϵ_i*: error term (誤差項)
- *β*: coefficients (係数)
- Data (sample): $\{Y_i, X_{i1}, \ldots, X_{iK}\}_{i=1}^N$
- We want to estimate coefficients β .

Ordinaly Least Squares (最小二乗法、OLS)

• OLS estimators are the minimizers of the sum of squared residuals:

$$\min_{eta_0,\cdots,eta_K}rac{1}{N}\sum_{i=1}^N(Y_i-(eta_0+eta_1X_{i1}+\cdots+eta_KX_{iK}))^2$$

• First order conditions characterize the OLS estimator. Denote it by $\hat{\beta}$.

Residual Regression (残差回帰)

• Consider the model

$$Y_i = \beta_0 + \alpha D_i + \beta_1 X_{1i} + \dots + \beta_K X_{Ki} + \epsilon_i$$

- Suppose that you are interested in α (say treatment effect parameter).
- Residual regression characterizes the OLS estimator of $\hat{\alpha}$ in the following way.

Frisch–Waugh–Lovell Theorem

- 1. Run OLS regression of D_i on all other explanatory variables $1, X_{1i}, \dots, X_{Ki}$. Obtain the residual $\hat{u_i^D}$.
- 2. Run OLS regression of Y_i on all other explanatory variables $1, X_{1i}, \cdots, X_{Ki}$. Obtain the residual $\hat{u_i^Y}$.
- 3. Run OLS regression of \hat{u}_i^Y on \hat{u}_i^D without constant term. The OLS estimator \hat{lpha} is

$$\hat{lpha} = rac{\sum \hat{u}_i^Y \hat{u}_i^D}{\sum (\hat{u}_i^D)^2}$$

How to use FWL theorem

- 1. Computational advantage if you are interested in a particular coefficient. Use this idea in estimation of panel data model.
- 1. Useful to see how the coefficient of interest is estimated. We will see this later in relation to multicolinearity (多重共線性).
- 1. Double machine learning (Chernozhukov et al 2018): Estimation of treatment effect parameters when so many covariates are available.

Assumptions for OLS

- 1. Random sample (ランダムサンプル): $\{Y_i, X_{i1}, \ldots, X_{iK}\}$ is i.i.d. (identically and independently distributed) drawn sample
- 2. **mean independence**: ϵ_i has zero conditional mean

 $E[\epsilon_i|X_{i1},\ldots,X_{iK}]=0$

- 3. Large outliers are unlikely: The random variable Y_i and X_{ik} have finite fourth moments.
- 4. No perfect multicollinearity (多重共線性): No linear relationship between explanatory variables.

Theoretical Properties of OLS estimator

1. **Unbiasedness**: Conditional on the explantory variables X, the expectation of the OLS estimator $\hat{\beta}$ is equal to the true value β .

$$E[\hat{\beta}|X]=\beta$$

2. Consistency: As the sample size N goes to infinity, the OLS estimator $\hat{\beta}$ converges to β in probability

$$\hat{\beta} \stackrel{p}{\longrightarrow} \beta$$

3. Asymptotic normality (漸近正規性): discuss later.

Linear Regression: Practical Topics

Interpretation of Regression Coefficients

• Remember that

$$Y_i = eta_0 + eta_1 X_{1i} + \dots + eta_K X_{Ki} + \epsilon_i$$

- The coefficient β_k : the effect of X_k on Y ceteris paribus (all things being equal)
- Equivalently, if X_k is continuous random variable,

$$\frac{\partial Y}{\partial X_k} = \beta_k$$

• If we can estimate β_k without bias, can obtain **causal effect** of X_k on Y.

Common Specifications in Linear Regression Model

- Several specifications frequently used in empirical analysis.
 - 1. Nonlinear term
 - 2. log specification
 - 3. dummy (categorical) variables
 - 4. interaction terms (交差項)

Nonlinear term (非線形項)

• Non-linear relationship between \boldsymbol{Y} and \boldsymbol{X} in a linearly additive form

$$Y_i = \beta_0 + \beta_1 X_i + \beta_2 X_i^2 + \beta_3 X_i^3 + \epsilon_i$$

- As long as the error term ϵ_i appreas in a additively linear way, we can estimate the coefficients by OLS.
 - Multicollinarity could be an issue if we have many polynomials (多項式).
 - $\circ~$ You can use other non-linear variables such as $\log(x)$ and \sqrt{x} .

log specification

• Using log changes the interpretation of the coefficient β in terms of scales.

Dependent	Explanatory	interpretation
Y	X	1 unit increase in X causes eta units change in Y
$\log Y$	X	1 unit increase in X causes $100 eta\%$ change in Y
Y	$\log X$	1% increase in X causes $eta/100$ unit change in Y
$\log Y$	$\log X$	1% increase in X causes $\beta\%$ change in Y

Dummy variable (ダミー変数)

- A **dummy variable** takes only 1 or 0. This is used to express qualititative information
- Example: Dummy variable for race

$$white_i = egin{cases} 1 & if white \ 0 & otherwise \end{cases}$$

• The coefficient on a dummy variable captures the difference of the outcome ${\boldsymbol Y}$ between categories

$$Y_i = eta_0 + eta_1 white_i + \epsilon_i$$

The coefficient β_1 captures the difference of *Y* between white and non-white people.

Interaction term (交差項)

- You can add the interaction of two explanatory variables in the regression model.
- For example:

$$wage_i = eta_0 + eta_1 educ_i + eta_2 white_i + eta_3 educ_i imes white_i + \epsilon_i$$

where $wage_i$ is the earnings of person i and $educ_i$ is the years of schooling for person i.

• The effect of $educ_i$ is

$$rac{\partial wage_i}{\partial educ_i}=eta_1+eta_3 white_i,$$

• This allows for heterogeneous effects of education across races.

Measures of Fit

- We often use R^2 (決定係数) as a measure of the model fit.
- Denote **the fitted value** as \hat{y}_i

$$\hat{y}_i = \hat{eta}_0 + \hat{eta}_1 X_{i1} + \dots + \hat{eta}_K X_{iK}$$

 $\,\circ\,$ Also called prediction from the OLS regression.

•
$$\,R^2$$
 is defined as

$$R^2 = rac{SSE}{TSS},$$

where

$$SSE = \sum_{i} ({\hat y}_i - {ar y})^2, \ TSS = \sum_{i} (y_i - {ar y})^2$$

- R^2 captures the fraction of the variation of Y explained by the regression model.
- Adding variables always (weakly) increases R^2 .

• In a regression model with multiple explanatory variables, we often use **adjusted** R^2 that adjusts the number of explanatory variables

$${ar R}^2 = 1 - rac{N-1}{N-(K+1)} rac{SSR}{TSS}$$

where

$$SSR = \sum_i ({\hat y}_i - y_i)^2 (= \sum_i {\hat u}_i^2)$$

Linear Regression: Inference

Statistical Inference of OLS Estimator

- The OLS estimator is **random variables** as it depends on a drawn sample.
- We need to conduct **statistical inference** to evaluate statistical uncertainty of the OLS estimates.
- Plan
 - Asymptotic distribution (漸近分布) of OLS estimator
 - Statistical inference:
 - Homoskedasticity (均一分散) vs Heteroskedasticity (不均一分散)

Asymptotic Normality (漸近正規性) of OLS Estimator

• Under the OLS assumption, the OLS estimator has **asymptotic normality**

$$\sqrt{N}(\hat{eta}-eta)\stackrel{d}{
ightarrow}N\left(0,V
ight)$$

• *V* is called **asymptotic variance (matrix)** given by

$$\underbrace{V}_{(K+1) imes (K+1)} = E[\mathbf{x}_i'\mathbf{x}_i]^{-1}E[\mathbf{x}_i'\mathbf{x}_i\epsilon_i^2]E[\mathbf{x}_i'\mathbf{x}_i]^{-1}$$

• $\mathbf{x}_i = (1, X_{i1}, \cdots, X_{iK})'$ is (K+1) imes 1 vector.

• We can **approximate** the distribution of $\hat{\beta}$ by

$$\hat{eta} \sim N(eta,V/N)$$

• The individual coefficient β_k follows

$${\hat eta}_k \sim N(eta_k,V_{kk}/N)$$

Estimation of Asymptotic Variance (漸近分散)

- V is an unknown object. Need to be estimated.
- Consider the estimator \hat{V} for V using sample analogues

$$\hat{V} = \left(\frac{1}{N}\sum_{i=1}^{N}\mathbf{x}_{i}'\mathbf{x}_{i}\right)^{-1} \left(\frac{1}{N}\sum_{i=1}^{N}\mathbf{x}_{i}'\mathbf{x}_{i}\hat{\epsilon}_{i}^{2}\right) \left(\frac{1}{N}\sum_{i=1}^{N}\mathbf{x}_{i}'\mathbf{x}_{i}\right)^{-1}$$

where $\hat{\epsilon}_i = y_i - (\hat{\beta}_0 + \dots + \hat{\beta}_K X_{iK})$ is the residual.

- Technically speaking, \hat{V} converges to V in probability.
- We often use the (asymptotic) standard error $SE(\hat{eta}_k) = \sqrt{\hat{V}_{kk}/N}$.
- The standard error is an estimator for the standard deviation of the OLS estimator $\hat{eta_k}.$

Hypothesis testing

- You might want to test a particular hypothesis regarding those coefficients.
 - Does x really affects y?
 - Is the production technology the constant returns to scale?

3 Steps in Hypothesis Testing

• Step 1: Consider the null hypothesis H_0 and the alternative hypothesis H_1

$$H_0:eta_1=k, H_1:eta_1
eq k$$

where k is the known number you set by yourself.

• Step 2: Define **t-statistic** by

$$t_n = rac{\hat{eta_1} - k}{SE(\hat{eta_1})}$$

• Step 3: We reject H_0 is at lpha-percent significance level if

$$|t_n| > C_{lpha/2}$$

where $C_{\alpha/2}$ is the $\alpha/2$ percentile of the standard normal distribution. We say we fail to reject H_0 if the above does not hold.

Caveats on Hypothesis Testing

- We often say $\hat{\beta}$ is **statistically significant (統計的有意)** at 5% level if $|t_n| > 1.96$ when we set k = 0.
- You should also discuss economic significance (経済的有意) of the coefficient in analysis.
- Case 1: Small but statistically significant coefficient. $\circ\,$ As the sample size N gets large, the SE decreases.
- Case 2: Large but statistically insignificant coefficient.
 - The variable might have an important (economically meaningful) effect.
 - But you may not be able to estimate the effect precisely with the sample at your hand.

F test

• We often test a composite hypothesis that involves multiple parameters such as

$$H_0:eta_1+eta_2=0,\ H_1:eta_1+eta_2
eq 0$$

• We use **F test** in such a case.

Confidence interval (信頼区間)

• 95% confidence interval

$$egin{aligned} CI_n &= \left\{k: |rac{\hat{eta}_1-k}{SE(\hat{eta}_1)}| \leq 1.96
ight\} \ &= \left[\hat{eta}_1 - 1.96 imes SE(\hat{eta}_1), \hat{eta}_1 + 1.96 imes SE(\hat{eta}_1)
ight] \end{aligned}$$

• Interpretation: If you draw many samples (dataset) and construct the 95% CI for each sample, 95% of those CIs will include the true parameter.

Homoskedasticity vs Heteroskedasticity

• The error term ϵ_i has **heteroskedasticity** (不均一分散) if $Var(u_i|X_i)$ depends on X_i . The asymptotic variance is

$$V = E[\mathbf{x}_i'\mathbf{x}_i]^{-1}E[\mathbf{x}_i'\mathbf{x}_i\epsilon_i^2]E[\mathbf{x}_i'\mathbf{x}_i]^{-1}$$

• If not, we call ϵ_i has **homoskedasticity** (均一分散). In this case,

$$V = E[\mathbf{x}_i'\mathbf{x}_i]^{-1}\sigma^2$$

where $\sigma^2 = V(\epsilon_i).$

Standard Errors in Practice

- Standard errors under heteroskedasticity assumption is called heteroskedasticity robust standard errors (不均一分散に頑健な標準誤差)
- In many statistical packages (including R and Stata), the standard errors for the OLS estimators are calculated under homoskedasticity assumption as a default.
- However, if the error has heteroskedasticity, the standard error under homoskedasticity assumption will be **underestimated**.
- In OLS, we should always use heteroskedasticity robust standard error.

Appendix: Matching Estimator

Estimation Methods

- We need to estimate $E[Y_i | D_i = 1, X_i = x]$ and $E[Y_i | D_i = 0, X_i = x]$
- Several ways to implement the above idea
 - 1. Regression: Nonparametric and Parametric
 - 2. Nearest neighborhood matching
 - 3. Propensity Score Matching

Approach 1: Regression, or Analogue Approach

- Let $\hat{\mu}_k(x)$ be an estimator of $\mu_k(x) = E[Y_i | D_i = k, X_i = x]$ for $k \in \{0,1\}$
- The analog estimators are

$$egin{aligned} &A\hat{T}E = rac{1}{N}\sum_{i=1}^{N} \left(\hat{\mu}_1(X_i) - \hat{\mu}_0(X_i)
ight) \ &A\hat{T}T = rac{N^{-1}\sum_{i=1}^{N}D_i(Y_i - \hat{\mu}_0(X_i))}{N^{-1}\sum_{i=1}^{N}D_i} \end{aligned}$$

• How to estimate $\mu_k(x) = E[Y_i | D_i = k, X_i = x]$?

Nonparametric Estimation

- Suppose that $X_i \in \{x_1, \cdots, x_K\}$ is discrete with small K
 - $\circ~$ Ex: two demographic characteristics (male/female, white/non-white). K=4 \bigskip
- Then, a nonparametric binning estimator is

$$\hat{\mu}_k(x) = rac{\sum_{i=1}^N \mathbf{1}\{D_i = k, X_i = x\}Y_i}{\sum_{i=1}^N \mathbf{1}\{D_i = k, X_i = x\}}$$

\bigskip

• Here, I do not put any parametric assumption on $\mu_k(x) = E[Y_i | D_i = k, X_i = x].$

Curse of dimensionality

- Issue: Poor performance if K is large due to many covariates.
 - $\circ~$ So many potential groups, too few observations for each group.
 - \circ With K variables, each of which takes L values, L^K possible groups (bins) in total.
- This is known as **curse of dimensionality**.
- Relatedly, if X is a continuous random variable, can use kernel regression.

Parametric Estimation, or going back to linear regression

• If you put parametric assumption such as

$$E[Y_i | D_i = 0, X_i = x] = eta' x_i \ E[Y_i | D_i = 1, X_i = x] = eta' x_i + au_0$$

then, you will have a model

$$y_i = eta' x_i + au D_i + \epsilon_i$$

• You can think the matching estimator as controlling for omitted variable bias by adding (many) covariates (control variables) x_i .

Approach 2: M-Nearest Neighborhood Matching

- Idea: Find the counterpart in other group that is close to me.
- Define $\hat{y}_i(0)$ and $\hat{y}_i(1)$ be the estimator for (hypothetical) outcomes when treated and not treated.

$${\hat y}_i(0) = egin{cases} y_i & if \ D_i = 0 \ rac{1}{M} \sum_{j \in L_M(i)} y_j & if \ D_i = 1 \end{cases}$$

- $L_M(i)$ is the set of M individuals in the opposite group who are "close" to individual i
- Several ways to define the distance between X_i and X_j , such as

$$dist(X_i,X_j) = \left|\left|X_i - X_j
ight|\right|^2$$

• Need to choose (1) M and (2) the measure of distance

Approach 3: Propensity Score Matching

- Use propensity score $P(D_i = 1 | X_i = x)$ as a distance to define who is the closest to me.
- Step 1: Estimate propensity score function by logit or probit using a flexible function of X_i .
- Step 2: Calculate the propensity score for each observation. Use it to define the pair.