

Lecture 9/10

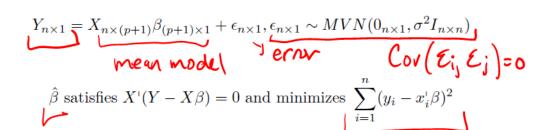
Model Checking and Key Extensions

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Review of where we left off

1. We have established the multiple linear regressio model:



2. We know that:

3. We have defined:

•
$$\hat{R} = Y - \hat{Y} = Y - X\hat{\beta} = (I - H)Y$$

4. Then we showed that:

•
$$\hat{\beta} \sim MVN(\beta, \sigma^2(X^{\scriptscriptstyle \dagger}X)^{-1})$$

• $\hat{Y} \sim MVN(X\beta, \sigma^2H)$
• $\hat{R} \sim MVN(0, \sigma^2(I-H))$

•
$$\hat{Y} \sim MVN(X\beta, \sigma^2 H)$$

•
$$\hat{R} \sim MVN(0, \sigma^2(I-H))$$

Review of where we left off

	Target	Estimate \sim Sampling Distn	95% CI for target	Test statistic for H0: Target $= 0$
	β_j	$\hat{\beta}_{j} \sim N(\beta_{j}, [\sigma^{2}(X^{T}X)^{-1})]_{jj})$	$\hat{\beta}_j \pm t \times \hat{se}(\hat{\beta}_j)$	$\frac{\tilde{\beta}_{j}}{\hat{s}c(\tilde{\beta}_{j})}$
	Αβ	$A\hat{\beta} \sim N(A\beta, \sigma^2 A(X^{\scriptscriptstyle \dagger} X)^{-1} A^{\scriptscriptstyle \dagger})$	$A\hat{eta} \pm t \times \hat{se}(A\hat{eta})$	$\frac{A\hat{\beta}_{1}}{\hat{s}c(A\hat{\beta}_{1})}$
	$g(\beta_J)$	$g(\hat{\beta}_{\mathcal{I}}) \sim N(g(\beta_{\mathcal{I}}), [g^{\shortmid}(\beta_{\mathcal{I}})]^2 [\sigma^2(X^{\shortmid}X)^{-1}]_{\mathcal{I}\mathcal{I}})$	$g(\hat{\beta}_{f}) \pm t \times \hat{se}(g(\hat{\beta}_{f}))$	$\frac{g(\hat{\beta}_{j})}{\hat{se}(g(\hat{\beta}_{j}))}$
	g(eta)	$g(\hat{\beta}) \sim N(g(\beta), g^{\shortmid}(\beta)^{\shortmid} [\sigma^2(X^{\shortmid}X)^{-1}] g^{\shortmid}(\beta))$	$g(\hat{\beta}) \pm t \times \hat{se}(g(\hat{\beta}))$	$\frac{g(\hat{\beta})}{\text{Ac}(g(\hat{\beta}))}$
	$\mu_i = E(Y_i X_i)$	$\hat{Y}_i \sim N(\mu_i, \sigma^2[H]_{ii})$	$\hat{Y}_i \pm t \times \hat{se}(\hat{Y}_i)$	$\frac{\hat{Y}_i}{\hat{se}(\hat{Y}_i)}$
	$\mu(x_0) = E(Y x_0)$	$x_0^{\scriptscriptstyle ext{i}}\hat{eta}\sim N(x_0^{\scriptscriptstyle ext{i}}eta,\hat{\sigma}^2x_0^{\scriptscriptstyle ext{i}}(X^{\scriptscriptstyle ext{i}}X)^{-1}x_0)$	$x_0^{\scriptscriptstyle \rm i} \hat{\beta} \pm t \times \hat{se}(x_0^{\scriptscriptstyle \rm i} \hat{\beta})$	$\frac{x_0^*\hat{\beta}}{se(x_0^*\hat{\beta})}$
l		Xo		

Key Assumptions by Order of Importance

Estimation of and interpretation of β

E(Y|X) = Xβ => we have "Correctly" specified the mean model

- onitted a key confinder / covariate

Confident Covariate

Confident Covariate - incorrectly specified functual form for a continuous X - missed key interactions - error measurement in X G Design of the study: how is the data generated Estiman Longitudinal study / clustered design rotal 3. Variance of residuals is constant Var (E;) = 6 Pt(XI) -Residuals are normally distributed > var (\$) => bootstrap procedur ZEstmatu/nterpetatuot Var (B) There are not a small number of highly influencial observations

Omitted Variable Bias

Exposure of interest: XI, Confinder X2 We will fit: Yi = do + di Xii + Ui The touth population: $Y_i = \beta_0 + \beta_1 X_{ii} + \beta_2 X_{2i} + \epsilon_i$ $\hat{A}_i = \frac{\text{Cov}(X_i, Y)}{\text{Var}(X_i)} = \frac{\text{Cov}(X_i, \beta_0 + \beta_1 X_i + \beta_2 X_2 + \epsilon)}{\text{Var}(X_i)}$ Var(X1) Cov(X1, po) + Cov(X1, piX1) + Cov(X1, paX1) + Cov(X1, E) Var(X) (or (X1/80) = 0 Cov (X1, E) = 0

Omitted Variable Bias

$$Z_{1} = \frac{\text{Cor}(X_{1}, \beta_{1}X_{1}) + (\text{or}(X_{1}, \beta_{3}X_{2})}{\text{Var}(X_{1})} + \beta_{2} \text{Cov}(X_{1}, X_{3})$$

$$= \beta_{1} \text{Var}(X_{1}) + \beta_{2} \text{Cov}(X_{1}, X_{3}) + \beta_{2} \text{Cov}(X_{1}, X_{3$$

Simulation exercise

- Within your breakout group, design a simulation study that would numerically demonstrate the result we just derived.
 - ▶ You go work for 15 minutes and then we will review together

Correct Functional Form for Continuous X

- To explore the assumption that $E(Y|X) = X\beta$, you can make the following plots:
 - Plot \hat{R} vs. X_j , j = 1, ..., p. • Recall that the residuals are independent of X if the model is correctly specified Plot \hat{R} vs. \hat{Y} .

 - The residuals and predicted values are independent if the model is correctly specified
- Never plot \hat{R} vs. Y because these are correlated!
- Based on the figures from 1. and 2., you could modify the model to increase/decrease the complexity of the functional form of the variables.

Example: Nepali Anthropometry Data

```
reg0<-lm(data=d.cc, arm~age)
d.cc$residuals = residuals(reg0)
ggplot(d.cc,aes(x=age, y=residuals)) +
    geom_jitter(alpha = 0.7) +
    geom_smooth() +
                               geom_hline(yintercept=0,color="red") +
    labs(y="Residuals: linear age")
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
         2-
     Residuals: linear age
        -3-
                                     20
                                                              40
                                                                                       60
                                                  age
```

Example: Nepali Anthropometry Data

Update the model to include a smooth function of age via linear splines or natural splines. reg1 = lm(arm~age + agesp6 + agesp12,data=d.cc) reg2 = lm(arm~ns(age,3),data=d.cc) d.cc\$residuals1 = residuals(reg1) d.cc\$residuals2 = residuals(reg2) Residuals: linear spline age 20 40 Residuals: natural spline age -3-

20

40

60

Independence Assumption

MLR => E; are independent Driven by the design of the study ► Longitudinal design : enoll units/people/cell/etc measure the outcome and covariates on the unit over time at various assessments

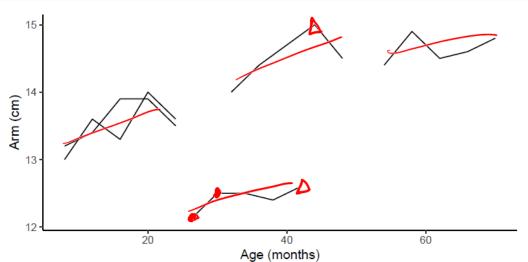
Yij j= follow-up, 1,..., m

i at assessment j. units/clusters are sampled - individuals within assessed male of villages - individuals within Clustered design sample of village - interview auch nousehold in village Sample of clinics - interview patients Why do we care? Yis = outrone for individual; from Var (B) = are ways

Example: Nepali Anthropometry Data

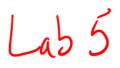
Design: i = 1, ..., m = 200 children each measured at baseline (j = 1) and then every 4 months for 4 follow-up visits (j = 2, 3, 4, 5).

```
ggplot(d5,aes(x=age,y=arm,group = factor(id))) +
  geom_line() +
  labs(x='Age (months)', y ='Arm (cm)') +
  theme_classic()
```



Checking the Independence Assumption

- ▶ Don't need to, we know the independence assumption is violated based on knowledge of the design
- ▶ We can explore covariance/correlation in the observed data
- Example: Consider the Nepali Anthropometry data where we have data for i = 1, ..., m = 200 children each measured at baseline (j = 1) and then every 4 months for 4 follow-up visits (j = 2, 3, 4, 5).
 - ▶ Step 1: Regress Y on X assuming independence and estimate β and R
 - ► Step 2: Plot \hat{R}_{ij} vs. \hat{R}_{ik} for all j,k
 - Compute $Cov(\hat{R}_{ij}, \hat{R}_{ik}) = \sqrt{Var(\hat{R}_{ij})} \times \sqrt{Var(\hat{R}_{ik})} \times Corr(\hat{R}_{ij}, \hat{R}_{ik})$
 - Or standardize the residuals and plot $Corr(\hat{R}_{sij},\hat{R}_{sik})$



How do we rethink the model?

What if we apply least squares to correlated data?

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Solution: Weighted least squares

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Next time....

- ► More model checking....
 - Robust variance estimation
 - Non-constant variance
 - Non-normal residuals
 - Influence and leverage statistics