

Lab 6

Lecture 11

Lab 7 and 8

Finish Model Checking

Implementation of WLS and robust variance estimation in R

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## Key Assumptions by Order of Importance

- E(Y|X) = Xβ, i.e. the mean model is "correctly" specified
  - $\triangleright$  Misspecification of Xβ can lead to biased β / misinterpretations
  - Omitted variable Bias
  - Correct functional form for continuous X
- Residuals are independent
  - ▶ This assumption is violated due to the design of the study
  - Longitudinal study
  - Clustered design
  - Show today: ignoring the correlation will impact  $Var(\hat{\beta})$  and derive weighted least squares
- Variance of residuals is constant
  - Often the variance is a function of some X
  - ▶ Show today: same impact and solution as violation of independence
- Residuals are normally distributed
  - CLT, bootstrap procedure

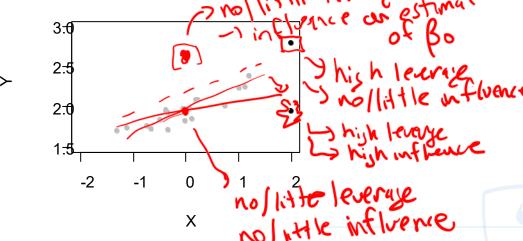
There are not a small number of highly influencial observations
Sensitivity analyses

observations -> (example People

## Leverage and Influence

- Leverage: A measure of how far an individual's predictors (X\_i) are from the mean X\_i
  - Hat matrix:  $h_{ii} = \frac{(X_i \bar{X})^2}{\sum (X_i \bar{X})^2}$

Influence: An observation  $(Y_i, X_i)$  such that including this value would greatly change the fitted values:  $\hat{\beta}$  and  $\hat{Y}$ .



#### Influence statistics

There are several influence statistics that are used in practice: i= observation

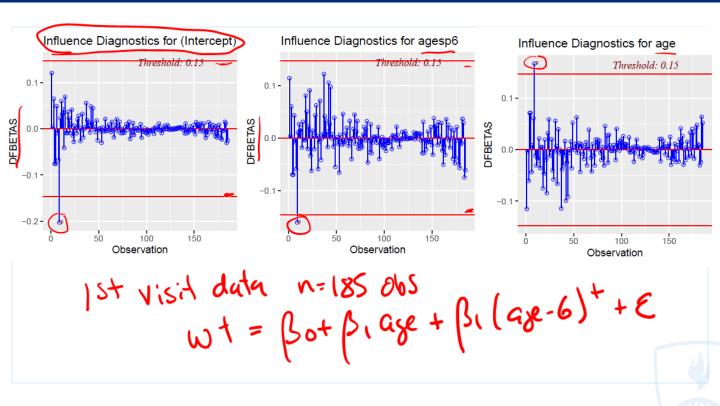
$$DBETA_{ij} = \hat{\beta}_j - \hat{\beta}_{j(-i)}$$

$$DFIT_i = \hat{Y}_i - \hat{Y}_{i(-i)}$$

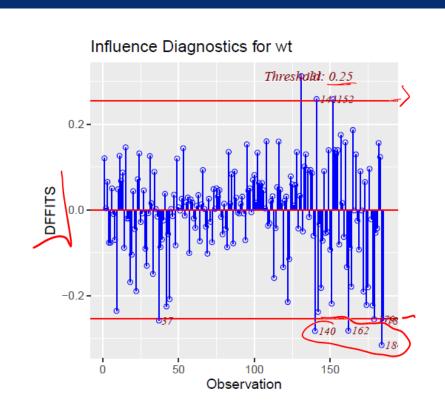
$$TS_i = \frac{DFIT_i}{\hat{se}(\hat{Y}_{i(-i)})}$$

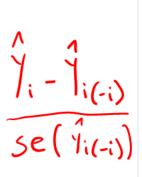
j = coefficient, 0,1,2,...p

## Example: Nepali Anthropometry Data

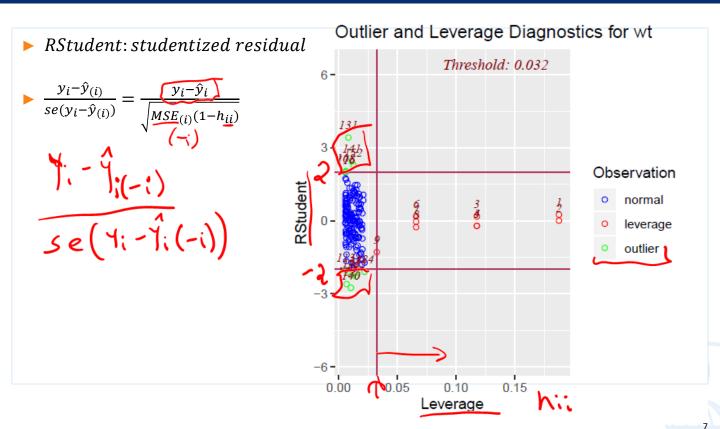


# Example: Nepali Anthropometry Data





## Example: Nepali Anthropometry Data



## Implementation of WLS in R

- For the remainder of the lecture, we will work through some analyses to demonstrate how to fit WLS in R
- In addition, I will show one approach for obtained robust variance estimates for different working correlation models. Here we will use the *gee* package in R

