# **Biostatistics I: Linear Regression**

#### **Model Diagnostics V: Independence**

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### **Linear Regression & Assumptions**

#### **Linear Regression Model:**

$$y_i = \mathbf{x}_i^ op oldsymbol{eta} + arepsilon_i, \quad \mathrm{E}(arepsilon_i) = 0, \quad \mathrm{var}(arepsilon_i) = \sigma^2$$

We need to **evaluate assumptions** about

#### the error terms:

- homoscedastic
- uncorrelated
- (normally distributed)

#### covariates and effects:

- linear effects (i.e., linear in the parameters)
- no (multi)collinearity between covariates

and check for **outliers and influential observations**.

### **Example: Child Growth**



#### **Child Growth: Model for Height**

$$\mathrm{height}_i = eta_0 + eta_1 \mathrm{age}_i + eta_2 \mathrm{kcal\_sd}_i + arepsilon_i$$



Mis-specified covariate effects and left-out predictors can cause correlated error terms.

#### **Child Growth: Model for Weight**

 $\text{weight}_i = \beta_0 + \beta_1 \text{age}_i + \beta_2 \text{height}_i + \beta_3 \text{kcal\_sd}_i + \varepsilon_i$ 



#### **Heteroscedasticity?**



#### Heteroscedasticity?



### Weighted Least Squares?

We fit a model for the residual variance,

$$\log(\hat{arepsilon}_i^2) = lpha_0 + lpha_1 \mathrm{age}_i + lpha_2 \mathrm{height}_i + lpha_3 \mathrm{kcal\_sd}_i + v_i,$$

to obtain weights

$$w_i = rac{1}{ \exp( \widehat{\log(\widehat{arepsilon}_i)})}.$$

### Weighted Least Squares?



### **Normality Assumption?**



### **Linearity Assumption?**



### **Child Growth: Model for Weight**

What is going wrong?

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#### What is going wrong?



#### **Child Growth: Clustered Data**



### **Linear Model Assumption**

#### Assumption of linear regression:

The  $y_i$  are

- all from the same distribution,
- except for a **shift** in the expected value, given by  $\mathbf{X}oldsymbol{eta}$ ,
- and are **independent** of each other.

#### Here:

Children from the same group are more similar to each other than to children from other groups.

The data has a **clustered** structure ⇒ **correlated residuals**.

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Can we "fix" the problem by including "group" in the model?

### **Example: Weight Loss Study**

Longitudinal study in adults to measure weight over time:



### **Example: Weight Loss Study**

Apparently, participants **gained weight** over time:



#### **Example: Weight Loss Study**

But: each participant **lost weight** over time:



### **Consequences of Correlated Error Terms**

The repeated observations from the same participant are **correlated**!

⇒The data has a **clustered** structure.

⇒ **Violation** of the assumption of independent error terms.

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⇒ Violation of the assumption of independent error terms.

Ignoring correlation of error terms results in

- potentially **biased estimates** and
- wrong standard errors.

### **Example: Impact of Correlated Error Terms**

Results from 1000 simulated datasets:



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Results from 1000 simulated datasets:



## **Settings with Correlated Error Terms**

**Common settings** with (likely) correlated observations are, for example,

- multi-center studies,
- studies with multiple family members,
- repeated measurements of the same subjects (longitudinal studies), and
- studies on matched data.

A linear regression model fitted with OLS is **not appropriate** in these settings.

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**Common settings** with (likely) correlated observations are, for example,

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Instead, use models that take into account the correlation, e.g.,

- mixed models (via random effects), or
- marginal models (via a correlation structure for  $\boldsymbol{\varepsilon}$ ).

### Modelling Approaches after Diagnosis

Model diagnosis

- is necessary to **identify violations** of (model) assumptions and
- can indicate how the model can be improved, e.g.,
  - changes in the assumed shape of association (⇒ transformation)
  - re-estimation without **suspicious** (outlying / influential) **observations**
  - use of weighted least squares or robust methods.

Usually, there is no perfect/correct model for real data.

Sensitivity analysis can help to evaluate robustness of the results/conclusions.