

# Pre/post measurements

Nora Wickelmaier

Last modified: October 16, 2024

# Analysis of longitudinal data

## Advantages

- More power than cross sectional studies
- Each subject is its own control person
- Information about individual changes

## Challenges

- Missing values
- Predictors can change over time
- In cross over designs: carry over effects

## Data schema

Person	Time	Observation	Covariates		
1	1	$y_{11}$	$x_{111}$	...	$x_{11p}$
1	2	$y_{12}$	$x_{121}$	...	$x_{12p}$
.	.	.	.	...	.
1	$n_1$	$y_{1n_1}$	$x_{1n_11}$	...	$x_{1n_1p}$
.	.	.	.	...	.
.	.	.	.	...	.
$N$	1	$y_{N1}$	$x_{N11}$	...	$x_{N1p}$
$N$	2	$y_{N2}$	$x_{N21}$	...	$x_{N2p}$
.	.	.	.	...	.
$N$	$n_N$	$y_{Nn_N}$	$x_{Nn_N1}$	...	$x_{Nn_Np}$

## Data schema

Person	Time	Observation	Covariates		
1	1	$y_{11}$	$x_{111}$	...	$x_{11p}$
1	2	$y_{12}$	$x_{121}$	...	$x_{12p}$
.	.	.	.	...	.
1	$n_1$	$y_{1n_1}$	$x_{1n_11}$	...	$x_{1n_1p}$
.	.	.	.	...	.
.	.	.	.	...	.
$N$	1	$y_{N1}$	$x_{N11}$	...	$x_{N1p}$
$N$	2	$y_{N2}$	$x_{N21}$	...	$x_{N2p}$
.	.	.	.	...	.
$N$	$n_N$	$y_{Nn_N}$	$x_{Nn_N1}$	...	$x_{Nn_Np}$

Today we will consider the simplest case with only two time points: pre and post score

# Outline

- ①  $t$  test
- ② Change score analysis
- ③ Analysis of Covariance
- ④ Example: Acupuncture for shoulder pain

1  $t$  test

## t test for dependent samples (comparing two time points)

- Straightforward analysis for two time points  $n = 2$
- Consider for each person the change score  $d_i = y_{i2} - y_{i1}$  and the linear model

$$d_i = \beta_0 + \varepsilon_i$$

$$y_{i2} = \beta_0 + y_{i1} + \varepsilon_i$$

with  $\varepsilon_i \sim N(0, \sigma^2)$  i.i.d.

- Interpretation:

$\beta_0$     average change score

- Hypothesis:  
Is there change between the first and second time point ( $H_0: \beta_0 = 0$ )?
- Which effects are not considered by this?

## t test for dependent samples (comparing two time points)

- Straightforward analysis for two time points  $n = 2$
- Consider for each person the change score  $d_i = y_{i2} - y_{i1}$  and the linear model

$$d_i = \beta_0 + \varepsilon_i$$

$$y_{i2} = \beta_0 + y_{i1} + \varepsilon_i$$

with  $\varepsilon_i \sim N(0, \sigma^2)$  i.i.d.

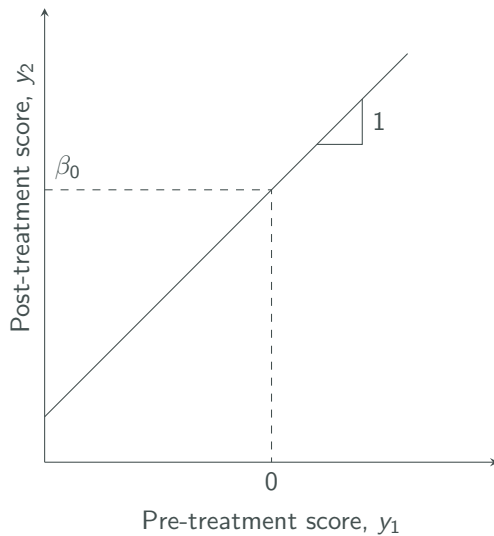
- Interpretation:

$\beta_0$  average change score

- Hypothesis:  
Is there change between the first and second time point ( $H_0: \beta_0 = 0$ )?
- Which effects are not considered by this?
- It is unclear if change is due to the treatment or just appeared over time



## t test for comparing two time points



## t test for comparing two time points

```
# simulate two dependent time points
cg <- MASS::mvrnorm(n = 50, mu = c(10,10),
                   Sigma = matrix(c(1,.9,.9,1), 2))
tg <- MASS::mvrnorm(n = 50, mu = c(10,15),
                   Sigma = matrix(c(1,.9,.9,1), 2))

sim <- as.data.frame(rbind(cg, tg))
names(sim) <- c("t1", "t2")

# add group variable
sim$group <- factor(rep(c("CG", "TG"), each = 50))

# group means
aggregate(cbind(t1, t2) ~ group, data = sim, FUN = mean)
```

## t test for comparing two time points

```
# t test
t.test(sim$t2, sim$t1, paired = TRUE)

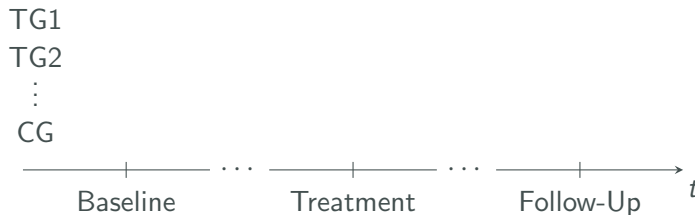
# add change score
sim$d <- sim$t2 - sim$t1

# linear model
lm1 <- lm(d ~ 1, data = sim)
lm2 <- lm(t2 ~ offset(t1), data = sim)

# visualization
plot(t2 ~ t1, data = sim)
abline(coef(lm2), 1)
```

## Better design

- At least two groups are observed before ( $y_{i1}$ ) and after ( $y_{i2}$ ) a treatment



- Research question:  
Do groups differ in the strength of their change?

## t test of follow-up score of two groups

- In regression notation with  $x_i = 1$ , if  $i$ th person is part of the treatment group, and else 0:

$$y_{i2} = \beta_0 + \beta_1 x_i + \varepsilon_i$$

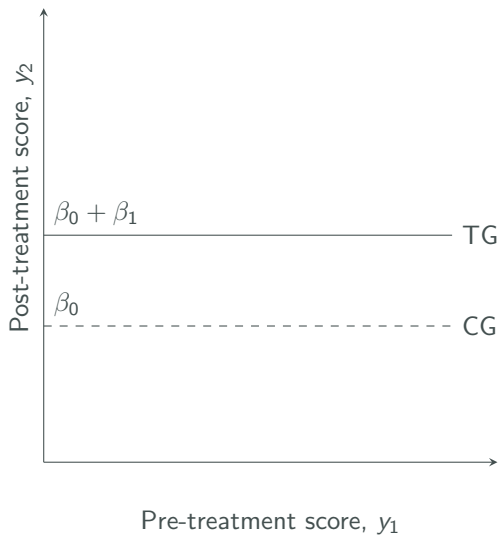
with  $\varepsilon_i \sim N(0, \sigma^2)$  i.i.d.

- Interpretation:

$\beta_0$     average follow-up score of reference group  
 $\beta_1$     effect of treatment group

- Hypothesis:  
Do groups differ at the second time point ( $H_0: \beta_1 = 0$ )?
- Since the baseline score is not considered, estimate of the change is biased

## t test of follow-up score of two groups



## t test of follow-up score of two groups

```
# t test
t.test(t2 ~ group, data = sim, var.equal = TRUE)

# linear model
lm3 <- lm(t2 ~ group, data = sim)

# visualization
plot(t2 ~ t1, sim)
abline(h = cumsum(coef(lm3)))
```

## ② Change score analysis



## Change score analysis

- Regression model

$$d_i = \beta_0 + \beta_1 x_i + \varepsilon_i$$

$$y_{i2} - y_{i1} = \beta_0 + \beta_1 x_i + \varepsilon_i$$

$$y_{i2} = \beta_0 + y_{i1} + \beta_1 x_i + \varepsilon_i$$

with  $\varepsilon_i \sim N(0, \sigma^2)$  i.i.d.

- Interpretation:

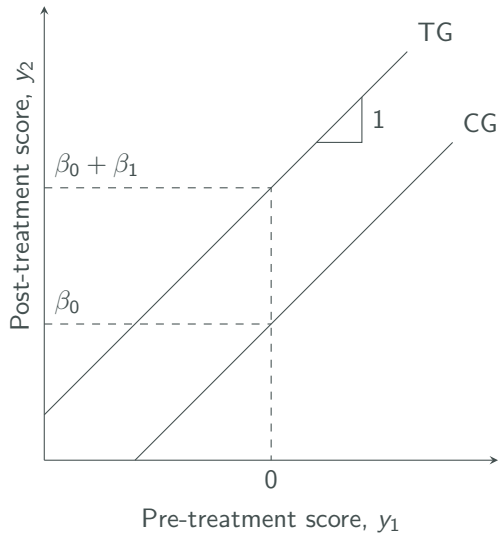
$\beta_0$  average change score for reference group

$\beta_1$  difference to  $\beta_0$  in treatment group

- Hypotheses:

- Is there change in the reference group ( $H_0: \beta_0 = 0$ )?
- Does the change differ between groups ( $H_0: \beta_1 = 0$ )?

## Change score analysis



## Change score analysis

```
# regression model
lm4 <- lm(t2 ~ offset(t1) + group, data = sim)

# visualization
plot(t2 ~ t1, data = sim)
abline(coef(lm4)[1], 1)
abline(coef(lm4)[1] + coef(lm4)[2], 1)
```

## Regression to the mean

- Change score analysis is based on the often too restrictive assumption that the follow-up score depends on the baseline score with a slope of 1
- Often baseline scores are negatively correlated with the change scores: Persons with low (bad) scores improve more than persons with high scores
- This regression to the mean lets us expect a slope of  $< 1$  which has to be estimated from the data

### ③ Analysis of Covariance

# Analysis of Covariance (ANCOVA)

- Regression model

$$y_{i2} = \beta_0 + \beta_1 y_{i1} + \beta_2 x_i + \varepsilon_i$$

with  $\varepsilon_i \sim N(0, \sigma^2)$  i.i.d.

- Interpretation:

$\beta_0$  average follow-up score in the reference group for  $y_{i1} = 0$

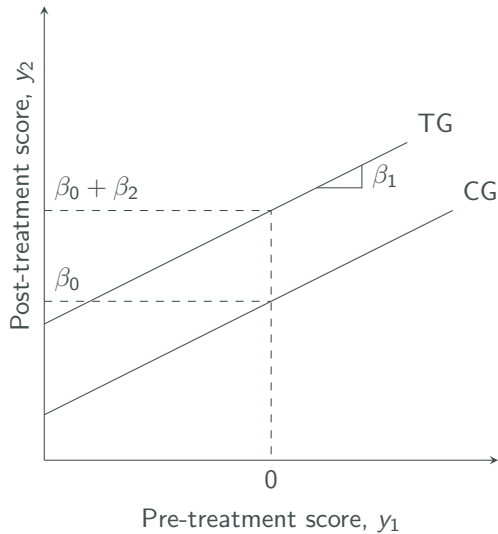
$\beta_1$  effect of baseline score

$\beta_2$  difference to follow-up score in the treatment group

- Hypotheses:

- Is there a relationship between baseline and follow-up score ( $H_0: \beta_1 = 0$ )?
- Do the follow-up scores in the groups differ for persons with identical baseline scores ( $H_0: \beta_2 = 0$ )?

# Analysis of Covariance (ANCOVA)



## Adjusted means

- With an ANCOVA model we can predict average follow-up scores for persons with the same baseline score
- For example, we get

$$\hat{y}_{i2} = \hat{\beta}_0 + \hat{\beta}_1 \bar{y}_1 + \hat{\beta}_2 x_i$$

for an average baseline score  $\bar{y}_1$

- These conditional means are sometimes called (baseline) adjusted means



## Analysis of Covariance (ANCOVA)

```
# ancova
lm5 <- lm(t2 ~ t1 + group, data = sim)

# adjusted means
predict(lm5, newdata = data.frame(t1 = mean(sim$t1),
                                   group = c("CG", "TG")))

# ancova with change score
lm5a <- lm(d ~ t1 + group, data = sim)

# visualization
plot(t2 ~ t1, data = sim)
abline(coef(lm5)[1], coef(lm5)[2])
abline(coef(lm5)[1] + coef(lm5)[3], coef(lm5)[2])
```

## Analysis of Covariance with change score

- Regression model

$$d_i = \beta_0 + \beta_1 y_{i1} + \beta_2 x_i + \varepsilon_i$$

$$y_{i2} - y_{i1} = \beta_0 + \beta_1 y_{i1} + \beta_2 x_i + \varepsilon_i$$

$$y_{i2} = \beta_0 + (1 + \beta_1) y_{i1} + \beta_2 x_i + \varepsilon_i$$

- For testing the difference of change in the groups ( $\beta_2$ ) it is irrelevant if the dependent variable is the follow-up score or the change score

## Different slopes for groups

- Regression model

$$y_{i2} = \beta_0 + \beta_1 y_{i1} + \beta_2 x_i + \beta_3 (y_{i1} \cdot x_i) + \varepsilon_i$$

with  $\varepsilon_i \sim N(0, \sigma^2)$  i.i.d.

- Interpretation:

$\beta_0$  average follow-up score in reference group for  $y_{i1} = 0$

$\beta_1$  effect of baseline score

$\beta_2$  difference to follow-up score in treatment group

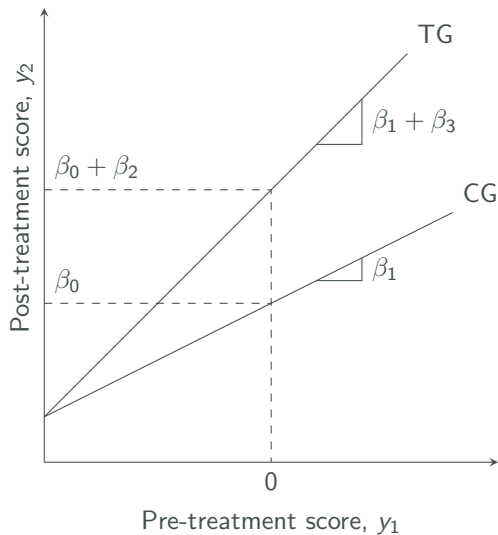
$\beta_3$  difference between slopes in reference and treatment groups

- Hypothesis:

Does effect of baseline score depend on group ( $H_0: \beta_3 = 0$ )?

- Interpretation of adjusted means independently of baseline score implies  $\beta_3 = 0$

## Interaction between baseline score and group



## Interaction between baseline score and group

```
# regression model with interaction
lm6 <- lm(t2 ~ t1 * group, data = sim)

# visualization
plot(t2 ~ t1, data = sim)
abline(coef(lm6)[1], coef(lm6)[2])
abline(coef(lm6)[1] + coef(lm6)[3], coef(lm6)[2] + coef(lm6)[4])

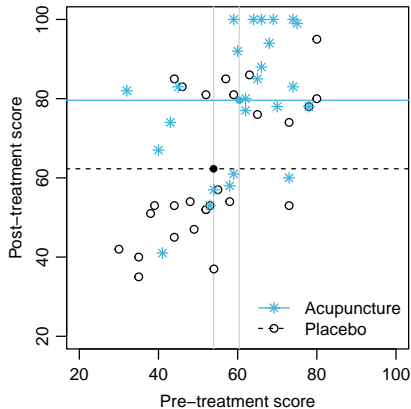
# models are nested
anova(lm4, lm5, lm6)
```

#### ④ Example: Acupuncture for shoulder pain

## Example: Acupuncture for shoulder pain

- Kleinhenz et al. (1999) investigate the effect of acupuncture on the improvement of mobility for 52 patients with shoulder pain
- Patients are randomly assigned to two groups (placebo vs. acupuncture)
- Before and after the treatment a mobility score is measured
- Vickers and Altman (2001) show advantages of an analysis of covariance compared to other methods based on these data

# Acupuncture: Follow-up analysis



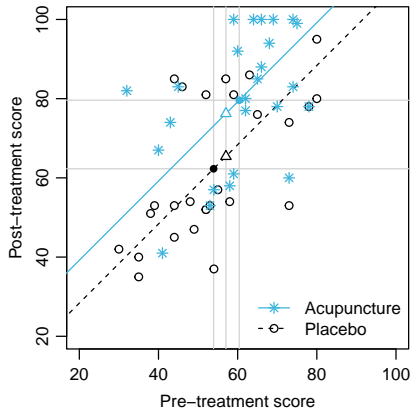
	Pla	Acu	Diff
Baseline	53.9	60.4	6.5
Follow-up	62.3	79.6	17.3
Change sc.	8.4	19.2	10.8
ANCOVA			12.7

$$y_{i2} = \beta_0 + \beta_1 x_i + \varepsilon_i$$

$$\hat{\beta}_1 = 17.3, 0.95\text{-CI: } (7.5, 27.1)$$



# Acupuncture: Change score analysis

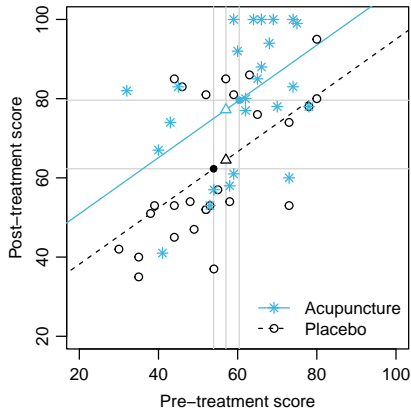


	Pla	Acu	Diff
Baseline	53.9	60.4	6.5
Follow-up	62.3	79.6	17.3
Change sc.	8.4	19.2	10.8
ANCOVA			12.7

$$y_{i2} = \beta_0 + y_{i1} + \beta_1 x_i + \varepsilon_i$$

$$\hat{\beta}_1 = 10.8 (2.3, 19.4)$$

# Acupuncture: ANCOVA



	Pla	Acu	Diff
Baseline	53.9	60.4	6.5
Follow-up	62.3	79.6	17.3
Change sc.	8.4	19.2	10.8
ANCOVA			12.7

$$y_{i2} = \beta_0 + \beta_1 y_{i1} + \beta_2 x_i + \varepsilon_i$$

$$\hat{\beta}_2 = 12.7 (4.1, 21.3)$$

## Example: Acupuncture for shoulder pain

```
# read data
dat <- read.table("kleinhenz.txt", header = TRUE)
dat$grp <- factor(dat$grp, levels = c("plac", "acu"))

# follow-up analysis
m1 <- lm(post ~ grp, data = dat)
summary(m1)
confint(m1)

# change score analysis
m2 <- lm(post ~ offset(pre) + grp, data = dat)

# anova
m3 <- lm(post ~ pre + grp, data = dat)
```

## Example: Acupuncture for shoulder pain

```
# testing if slopes differ
m4 <- lm(post ~ pre * grp, data = dat)
anova(m3, m4)

# adjusted means
predict(m3, newdata = data.frame(pre = mean(dat$pre),
                                   grp = c("plac", "acu")))

# visualization
plot(post ~ pre, data = dat, xlim = c(20,100), ylim = c(20,100))
abline(coef(m3)[1], coef(m3)[2])
abline(coef(m3)[1] + coef(m3)[3], coef(m3)[2])
```

## Summary

- We considered the basic case when several groups are observed at two time points (baseline and follow-up)
- Analysis of these kind of data considers the difference

$$d_i = y_{i2} - y_{i1}$$

or the adjusted follow-up scores which are considered to be independent

- Analysis of Covariance
  - has the highest power to detect differences for average change compared to the other methods
  - must be cautiously interpreted when groups are not randomly assigned

## ⑤ Linear mixed-effects model

## Linear mixed-effects model

- Regression notation with indicator variable for time ( $t_{i1} = 0$ ,  $t_{i2} = 1$ ) and group ( $x_i$ )

$$y_{ij} = \beta_0 + \beta_1 t_{ij} + \beta_2 x_i + \beta_3 (t_{ij} \cdot x_i) + v_i + \varepsilon_{ij}$$

with  $v_i \sim N(0, \sigma_v^2)$  and  $\varepsilon_{ij} \sim N(0, \sigma^2)$

- Interpretation:

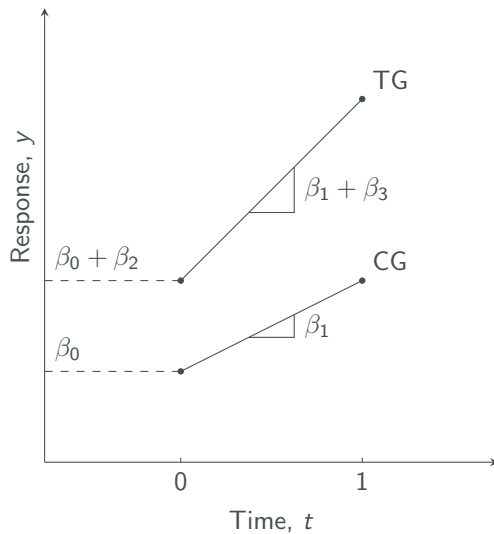
$\beta_0$  mean baseline value in reference group

$\beta_1$  time effect (slope) in reference group

$\beta_2$  effect of treatment group

$\beta_3$  effect on the slope of treatment group

# Linear mixed-effects model





## Linear mixed-effects model

- For the two groups with  $j = 1, 2$ , we get

$$y_{i1} = \beta_0 + \beta_2 x_i + v_i + \varepsilon_{i1}$$

$$y_{i2} = \beta_0 + \beta_1 + \beta_2 x_i + \beta_3 x_i + v_i + \varepsilon_{i2}$$

- For the change score, we then get

$$y_{i2} - y_{i1} = \beta_1 + \beta_3 x_i + (\varepsilon_{i2} - \varepsilon_{i1})$$

- Since  $\varepsilon_{ij}$  are independent, this results in the equation for the change score analysis

$$y_{i2} - y_{i1} = \beta_1 + \beta_3 x_i + \varepsilon_i$$

with

$$\varepsilon_i = \varepsilon_{i2} - \varepsilon_{i1} \sim N(0, \sigma_d^2 = 2\sigma^2)$$

→ LMM for two time points is equivalent to change score analysis!

## Example: Acupuncture for shoulder pain

```
# read data
dat <- read.table("kleinhenz.txt", header = TRUE)
dat$grp <- factor(dat$grp, levels = c("plac", "acu"))

# change score analysis
m1 <- lm(post ~ offset(pre) + grp, data = dat)
# LMM
dat1 <- reshape(dat, direction = "long",
                varying = list(1:2), v.names = "score")
dat1$time <- factor(dat1$time, levels = 1:2, labels = c("pre", "post"))
m2 <- lmer(score ~ grp * time + (1 | id), data = dat1)
# compare residual variances
sigma(m1)^2
2 * sigma(m2)^2
```

## References

- Kleinhenz, J., Streitberger, K., Windeler, J., Güßbacher, A., Mavridis, G., & Martin, E. (1999). Randomised clinical trial comparing the effects of acupuncture and a newly designed placebo needle in rotator cuff tendinitis. *Pain*, 83(2), 235–241.
- Vickers, A. J., & Altman, D. G. (2001). Analysing controlled trials with baseline and follow up measurements. *BMJ*, 323(7321), 1123–1124.