

Featback analyses: PHQ

Data preparation

Primaire uitkomstmaat

- Eetstoornispathologie (EDE-Q_TOT)

Secundaire uitkomstmaten

- Self-efficacy (GSES_TOT)
- Angst en depressie (PHQ_TOT)
- Sociale steun (SSL_TOT)

Conditie indicatoren

- 1 = Featback
- 2 = Featback + ondersteuning van een ervaringsdeskundige via chat of email
- 3 = Ondersteuning van een ervaringsdeskundige via chat of email
- 4 = Wachttijd controle conditie (hen werd conditie 2 aangeboden na 12 maanden + 8 weken wachttijd)

Tijdsindicatoren

- 1 = baseline
- 2 = post-interventie (8 weken)
- 3 = 3 maanden follow-up (i.e., 3 maanden + 8 weken)
- 4 = 6 maanden FU
- 5 = 9 maanden FU
- 6 = 12 maanden FU

Mogelijke moderatoren

- Leeftijd (Age)
- Educatie (moet nog omgezet worden naar 3 levels; laag, middel, hoog) (T0_edu)
- Behandelgeschiedenis (T0_treatment)
- BMI bij Baseline (BMI)
- Eetstoornispathologie bij baseline (EDEQ_TOT)
- (Evt. duur eetstoornis, maar is zeer hoog gecorreleerd met leeftijd; T0_yrsED)
- Motivatie om te veranderen bij baseline (T0_Motiv_TOT)
- Zelfwaardering bij baseline (T0_RSES_TOT)
- Self-efficacy bij baseline (GSES_TOT)
- Angst en depressie bij baseline (PHQ_TOT)
- Type eetstoornis bij baseline (staat nog niet tussen de variabelen; nog even kijken hoe we dit moeten aanpakken ...)

Toevoeging 06-08-2021:

- T0_EDEQ_ObjEet_Keer (= aantal (objectieve) eetbuien in de afgelopen 28 dagen)
- T0_SSL_TOT

```
## Read in data
library("foreign")
data <- read.spss("20210316_Featback_AllMerged_LongFormat_imputed.sav",
                  to.data.frame = TRUE)
```

```

#names(data)
## Set appropriate variable classes
data$ID <- factor(data$ID)
data$T0_edu <- ordered(data$T0_edu)
## Abbreviate condition levels
levels(data$Condition)[levels(data$Condition)=="Waiting list"] <- "WL"
levels(data$Condition)[levels(data$Condition)=="Featback"] <- "Fb"
levels(data$Condition)[levels(data$Condition)=="Featback + expert-patient support"] <- "Fb+eps"
levels(data$Condition)[levels(data$Condition)=="expert-patient support"] <- "eps"
data$Condition <- factor(data$Condition, levels = c("WL", "eps", "Fb", "Fb+eps"))
## Check if every subject has a time 1
all(table(data$ID, data$Time == 1)[,2] == 1L)

## [1] FALSE

## Select only pre- and post assessments
#data <- data[data$Time %in% 1:2, ]
#data$Time <- factor(data$Time)
## Construct T0 variables
for (i in unique(data$ID)) {
  data$T0_EDEQ_TOT[data$ID == i] <- data$EDEQ_TOT[data$ID == i & data$Time == 1]
  data$T0_BMI[data$ID == i] <- data$BMI[data$ID == i & data$Time == 1]
  data$T0_GSES_TOT[data$ID == i] <- data$GSES_TOT[data$ID == i & data$Time == 1]
  data$T0_PHQ_TOT[data$ID == i] <- data$PHQ_TOT[data$ID == i & data$Time == 1]
  data$T0_SSL_TOT[data$ID == i] <- data$SSL_TOT[data$ID == i & data$Time == 1]
  data$T0_eetbuien[data$ID == i] <- data$EDEQ_ObjEet_Keer[data$ID == i & data$Time == 1]
}
## Check for missings and remove
unlist(sapply(data[, c("Age", "T0_edu", "T0_BMI", "T0_EDEQ_TOT", "T0_Motiv_TOT",
                      "T0_GSES_TOT", "T0_PHQ_TOT")], function(x) table(is.na(x))))

##           Age.FALSE      T0_edu.FALSE      T0_edu.TRUE      T0_BMI.FALSE
##           215130           215124           6           215124
##           T0_BMI.TRUE  T0_EDEQ_TOT.FALSE  T0_EDEQ_TOT.TRUE  T0_Motiv_TOT.FALSE
##           6           215112           18           215112
##           T0_Motiv_TOT.TRUE  T0_GSES_TOT.FALSE  T0_GSES_TOT.TRUE  T0_PHQ_TOT.FALSE
##           18           215112           18           215130

data <- data[!is.na(data$T0_BMI), ]
data <- data[!is.na(data$T0_EDEQ_TOT), ]
data <- data[!is.na(data$T0_edu), ]
data <- data[!is.na(data$T0_yrsED), ]
unlist(sapply(data[, c("Age", "T0_edu", "T0_BMI", "T0_EDEQ_TOT", "T0_Motiv_TOT",
                      "T0_GSES_TOT", "T0_PHQ_TOT")], function(x) table(is.na(x))))

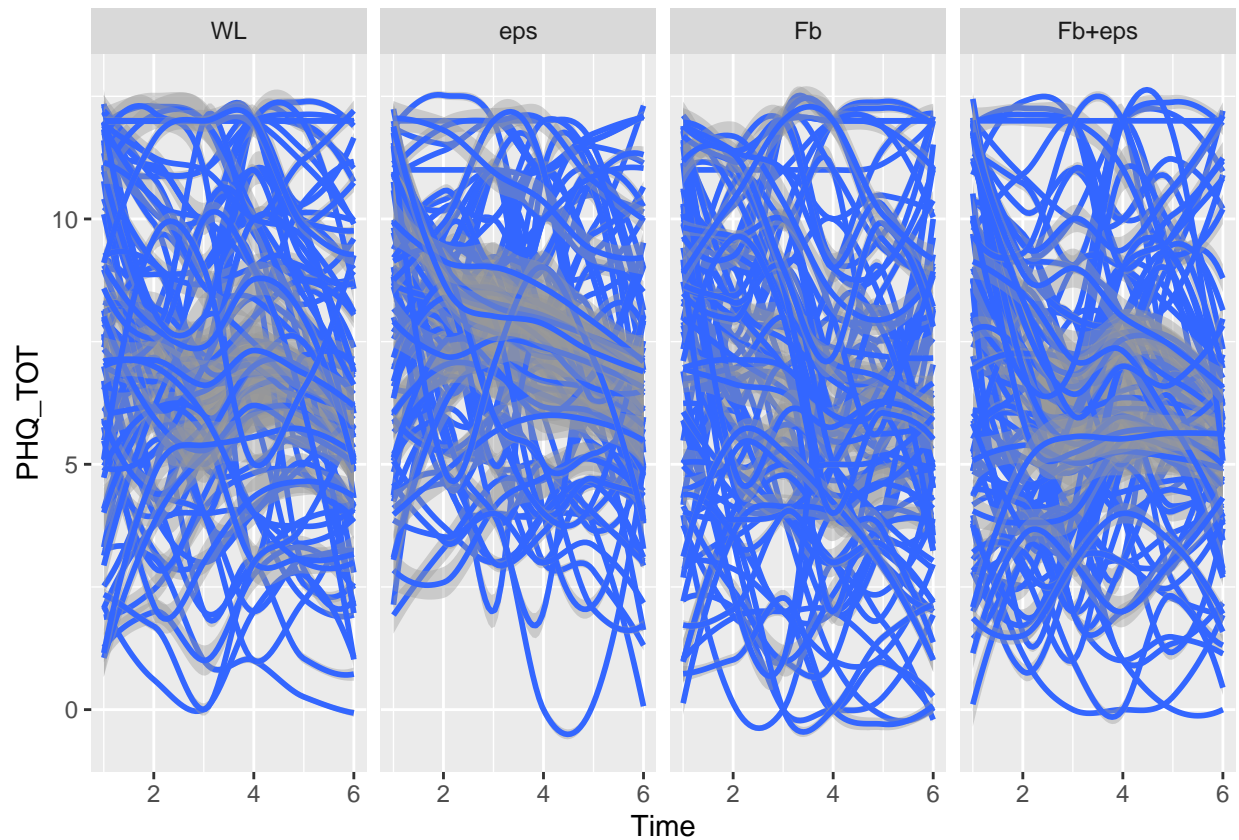
##           Age.FALSE      T0_edu.FALSE      T0_BMI.FALSE  T0_EDEQ_TOT.FALSE
##           215058           215058           215058           215058
##           T0_Motiv_TOT.FALSE  T0_Motiv_TOT.TRUE  T0_GSES_TOT.FALSE  T0_PHQ_TOT.FALSE
##           215043           15           215058           215058

## Exploratory plot
library("ggplot2")
ggplot(data = data, aes(x = Time, y = PHQ_TOT, group = ID)) +
  facet_grid(. ~ Condition) + geom_smooth()

## `geom_smooth()` using method = 'loess' and formula 'y ~ x'

```

```
## Warning: Removed 477 rows containing non-finite values (stat_smooth).
```



De trajecten lijken niet erg lineair.

Splits based only on treatment-time-subgroup interactions

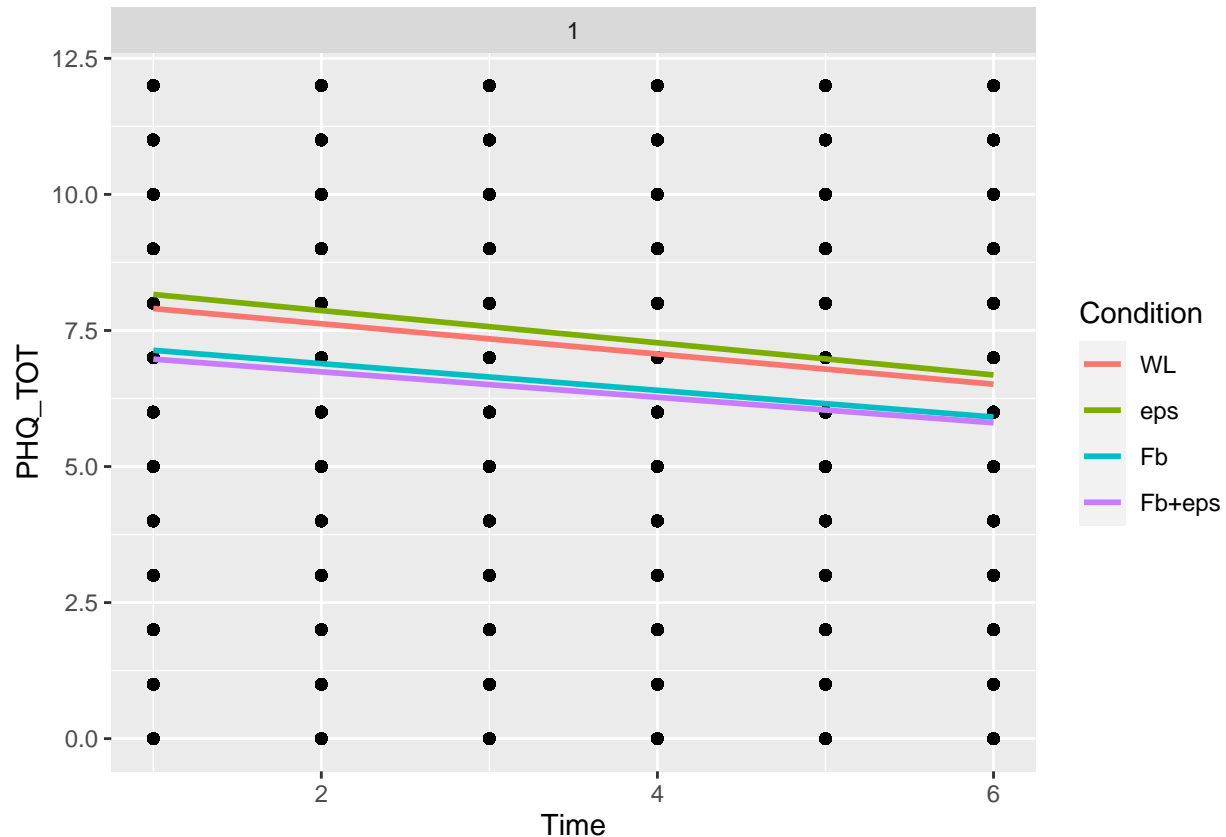
```
library("glmertree")
library("lmerTest")
library("strucchange")
dat1 <- data[!is.na(data$PHQ_TOT),]
lt1 <- lmertree(PHQ_TOT ~ Time*Condition | (1|ID) | Age + TO_edu + TO_BMI + TO_EDEQ_TOT +
               TO_Motiv_TOT + TO_GSES_TOT + TO_PHQ_TOT + TO_treatment + TO_yrsED +
               TO_SSL_TOT + TO_eetbuien,
               data = dat1, cluster = ID, parm = 6:8)
if (length(lt1$tree) > 1L) {
  plot(lt1, which = "tree", fitted = "marginal", gp = gpar(cex = .5))
  #fixef(lt1)
  apply(fixef(lt1), 2, sd)
  VarCorr(lt1)
  tmp <- summary(lmer(attr(lt1$lmer, "call")$formula, data = lt1$data))
  round(tmp$coefficients[tmp$coefficients[, "Pr(>|t|)"] < 0.05, -3], digits = 4)
} else {
  sctest(lt1$tree)
}
```

```
##           Age      TO_edu    TO_BMI TO_EDEQ_TOT TO_Motiv_TOT TO_GSES_TOT
```

```
## statistic 5.501296 22.3922270 5.9747359 3.704816 5.124871 3.881059
## p.value 1.000000 0.9998674 0.9999996 1.000000 1.000000 1.000000
##          TO_PHQ_TOT TO_treatment TO_yrsED TO_SSL_TOT TO_eetbuien
## statistic 6.8747576 7.8146840 9.0920960 12.9439134 4.187099
## p.value 0.9999682 0.4312064 0.9889012 0.6453678 1.000000
```

```
ggplot(data = lt1$data, aes(x = Time, y = PHQ_TOT, group = ID)) + geom_point() +
  facet_grid(. ~ .tree) + geom_smooth(method = "lm", se = FALSE,
  aes(x = Time, y = PHQ_TOT, group = Condition, color = Condition))
```

```
## `geom_smooth()` using formula 'y ~ x'
```



```
dat2 <- dat1[dat1$Time %in% 1:2, ]
lt3 <- lmertree(PHQ_TOT ~ Time*Condition | (1|ID) | Age + TO_edu + TO_BMI + TO_EDEQ_TOT +
  TO_Motiv_TOT + TO_GSES_TOT + TO_PHQ_TOT + TO_treatment + TO_yrsED +
  TO_SSL_TOT + TO_eetbuien,
  data = dat2, cluster = ID, parm = 6:8)
```

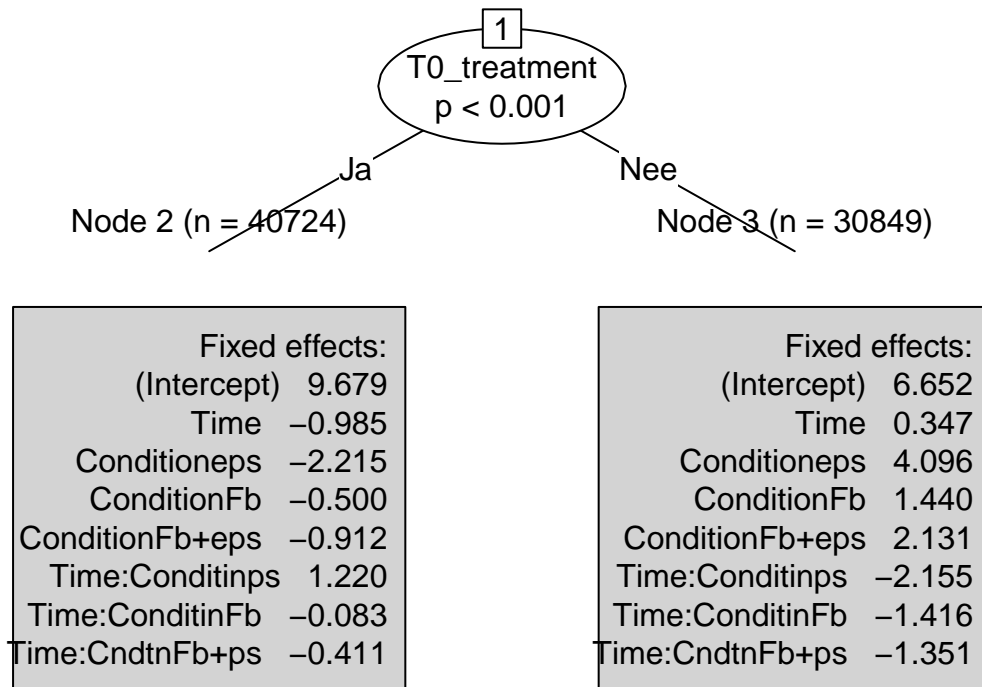
```
## Warning in lmertree(PHQ_TOT ~ Time * Condition | (1 | ID) | Age + TO_edu + :
## 'data' contains missing values, note that listwise deletion will be employed.
```

```
if (length(lt3$tree) > 1L) {
  plot(lt3, which = "tree", type = "simple", fitted = "marginal", gp = gpar(cex = .7))
  #fixef(lt3)
  apply(fixef(lt3), 2, sd)
  VarCorr(lt3)
  tmp <- summary(lmer(attr(lt3$lmer, "call")$formula, data = lt3$data))
  round(tmp$coefficients[tmp$coefficients[, "Pr(>|t|)"] < 0.05, -3], digits = 4)
```

```

} else {
  sctest(lt3$tree)
}

```



```

##               Estimate Std. Error  t value Pr(>|t|)
## (Intercept)      9.6790    0.3904  24.7957  0.0000
## .tree3           -3.0268    0.5783  -5.2336  0.0000
## .tree2:Time       -0.9856    0.0321 -30.6770  0.0000
## .tree3:Time        0.3470    0.0351   9.8823  0.0000
## .tree2:Conditioneps -2.2159    0.5415  -4.0920  0.0001
## .tree3:Conditioneps  4.0987    0.6338   6.4669  0.0000
## .tree3:ConditionFb   1.4405    0.5999   2.4012  0.0168
## .tree3:ConditionFb+eps  2.1251    0.6241   3.4052  0.0007
## .tree2:Time:Conditioneps  1.2209    0.0446  27.3963  0.0000
## .tree3:Time:Conditioneps -2.1567    0.0521 -41.3577  0.0000
## .tree3:Time:ConditionFb  -1.4159    0.0494 -28.6866  0.0000
## .tree2:Time:ConditionFb+eps -0.4123    0.0444  -9.2951  0.0000
## .tree3:Time:ConditionFb+eps -1.3466    0.0514 -26.2193  0.0000

```

```

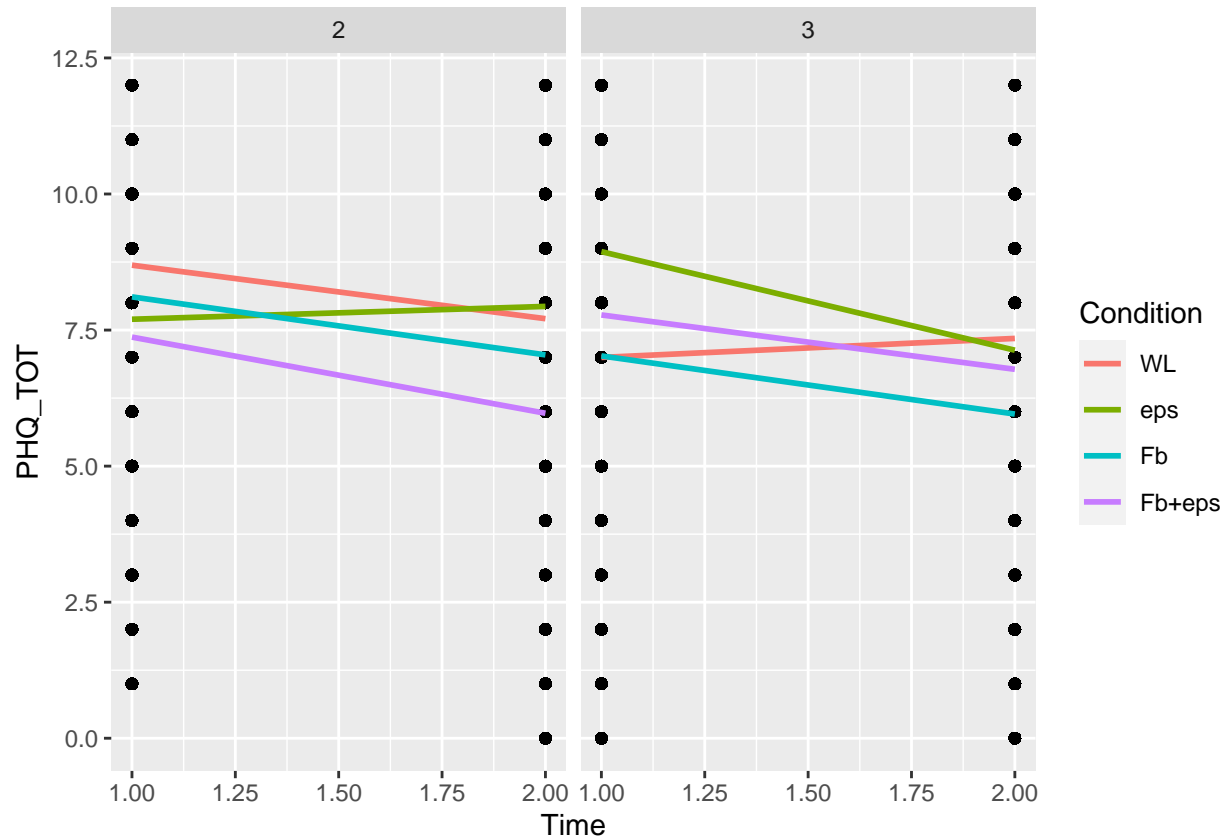
ggplot(data = lt3$data, aes(x = Time, y = PHQ_TOT, group = ID)) + geom_point() +
  facet_grid(. ~ .tree) +
  geom_smooth(method = "lm", se = FALSE,
    aes(x = Time, y = PHQ_TOT, group = Condition, color = Condition))

```

```

## `geom_smooth()` using formula 'y ~ x'

```



Waitlist versus all treatment groups

```
dat1$Condition2 <- factor(dat1$Condition == "WL")
lt2 <- lmertree(PHQ_TOT ~ Time*Condition2 | (1|ID) | Age + TO_edu + TO_BMI + TO_EDEQ_TOT +
  TO_Motiv_TOT + TO_GSES_TOT + TO_PHQ_TOT + TO_treatment + TO_yrsED +
  TO_SSL_TOT + TO_eetbuien,
  data = dat1, cluster = ID, parm = 4, verbose = TRUE, maxdepth = 4)
```

```
## 'log Lik.' -472211.1 (df=6)
```

```
## 'log Lik.' -472211.1 (df=6)
```

```
if (length(lt2$tree) > 1L) {
  plot(lt2, which = "tree", fitted = "marginal", gp = gpar(cex = .5))
  #fixef(lt2)
  apply(fixef(lt2), 2, sd)
  VarCorr(lt2)
  tmp <- summary(lmer(attr(lt2$lmer, "call")$formula, data = lt1$data))
  round(tmp$coefficients[tmp$coefficients[, "Pr(>|t|)"] < 0.05, -3], digits = 4)
} else {
  sctest(lt2$tree)
}
```

```
##           Age      TO_edu    TO_BMI TO_EDEQ_TOT TO_Motiv_TOT TO_GSES_TOT
## statistic 2.8231826 8.5429341 1.647902    1.54926    3.9387850    3.422081
## p.value   0.9999939 0.9950053 1.000000    1.00000    0.9984789    0.999804
##           TO_PHQ_TOT TO_treatment TO_yrsED TO_SSL_TOT TO_eetbuien
```

```
## statistic 4.4193402 6.2313699 1.28882 3.7812250 1.852299
## p.value 0.9937755 0.1297119 1.00000 0.9991278 1.000000

dat2 <- dat1[dat1$Time %in% 1:2, ]
lt4 <- lmertree(PHQ_TOT ~ Time*Condition2 | (1|ID) | Age + TO_edu + TO_BMI + TO_EDEQ_TOT +
  TO_Motiv_TOT + TO_GSES_TOT + TO_PHQ_TOT + TO_treatment + TO_yrsED +
  TO_SSL_TOT + TO_eetbuien,
  data = dat2, cluster = ID, parm = 4, maxdepth = 4)
if (length(lt4$tree) > 1L) {
  plot(lt4, which = "tree", fitted = "marginal", gp = gpar(cex = .7))
  #fixef(lt4)
  apply(fixef(lt4), 2, sd)
  VarCorr(lt4)
  tmp <- summary(lmer(attr(lt4$lmer, "call")$formula, data = lt3$data))
  round(tmp$coefficients[tmp$coefficients[, "Pr(>|t|)"] < 0.05, -3], digits = 4)
} else {
  sctest(lt4$tree)
}

##           Age    TO_edu    TO_BMI TO_EDEQ_TOT TO_Motiv_TOT TO_GSES_TOT
## statistic 4.3853843 4.212643 3.5113754 2.240749 1.893448 4.1582003
## p.value 0.9942949 1.000000 0.9997063 1.000000 1.000000 0.9969604
##           TO_PHQ_TOT TO_treatment TO_yrsED TO_SSL_TOT TO_eetbuien
## statistic 5.2884318 7.63052874 3.0496848 6.2082979 3.7344621
## p.value 0.9636033 0.06134676 0.9999729 0.8816751 0.9992681
```