STOPPD cortical thickness analysis (left hemisphere only)

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This script analyses cortical thickness in the left hemisphere.

```
#load libraries
library(tidyverse)
library(lme4)
library(lmerTest)
library(growthmodels)
#bring in data
df <- read_csv('../generated_csvs/STOPPD_participantsCT_20181111.csv') #generated by 05_STOPPD_error in
#make sure that STUDYID is an interger not a number
 df$STUDYID <- as.character(df$STUDYID)</pre>
#make sure that dateDiff is a number, not an interger
  df$dateDiff <- as.numeric(df$dateDiff)</pre>
# label the randomization variable
df$RandomArm <- factor(df$randomization,</pre>
                       levels = c("0", "P"),
                       labels = c("Olanzapine", "Placebo"))
#restructure data for RCT completers' only (N=40)
 RCT_CT <- df %>%
    filter(category == "RCT")
#write out clean dataframe
# write.csv(RCT_CT, '../generated_data/df_leftCT.csv', row.names=FALSE)
```

RCT only

```
RCT_CT %>% count(randomization) %>% knitr::kable()
```

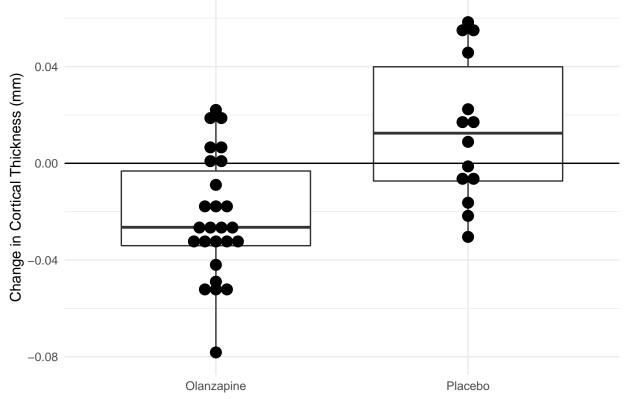
randomization	n
O	26

```
randomization n
P 14
```

```
#boxplot of difference in thickness (y axis) by randomization group (x axis)
ggplot(RCT_CT, aes(x= RandomArm, y = LThickness_change)) +
    geom_boxplot(outlier.shape = NA) +
    geom_dotplot(binaxis = 'y', stackdir = 'center') +
    geom_hline(yintercept = 0) +
    ggtitle("Cortical thickness (left hemisphere)") +
    xlab(NULL) +
    ylab("Change in Cortical Thickness (mm)") +
    theme_minimal()
```

`stat_bindot()` using `bins = 30`. Pick better value with `binwidth`.

Cortical thickness (left hemisphere)



```
#run linear model without covariates
fit_rct <- lm(LThickness_change ~ RandomArm, data= RCT_CT)
print(fit_rct)</pre>
```

```
##
## Call:
## lm(formula = LThickness_change ~ RandomArm, data = RCT_CT)
##
## Coefficients:
## (Intercept) RandomArmPlacebo
## -0.02253 0.03659
```

```
summary(fit_rct)
##
## Call:
## lm(formula = LThickness_change ~ RandomArm, data = RCT_CT)
## Residuals:
##
        Min
                   1Q
                         Median
## -0.055651 -0.018822 -0.003541 0.022344 0.044589
## Coefficients:
                    Estimate Std. Error t value Pr(>|t|)
##
                               0.005294 -4.255 0.000131 ***
                   -0.022529
## (Intercept)
## RandomArmPlacebo 0.036591
                               0.008949
                                         4.089 0.000217 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.027 on 38 degrees of freedom
## Multiple R-squared: 0.3055, Adjusted R-squared: 0.2873
## F-statistic: 16.72 on 1 and 38 DF, p-value: 0.0002168
#run linear model with covariates of sex and age
 fit_rct <- lm(LThickness_change ~ RandomArm + sex + age, data= RCT_CT)</pre>
 print(fit_rct)
##
## lm(formula = LThickness_change ~ RandomArm + sex + age, data = RCT_CT)
##
## Coefficients:
##
        (Intercept) RandomArmPlacebo
                                                  sexM
                                                                     age
##
         -0.0053535
                           0.0389342
                                             0.0087658
                                                              -0.0004063
  summary(fit rct)
##
## Call:
## lm(formula = LThickness_change ~ RandomArm + sex + age, data = RCT_CT)
## Residuals:
                         Median
        Min
                   1Q
                                       30
## -0.054543 -0.019130 -0.001897 0.019599 0.050858
##
## Coefficients:
##
                     Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                   -0.0053535 0.0168885 -0.317 0.753081
## RandomArmPlacebo 0.0389342 0.0090734
                                          4.291 0.000128 ***
## sexM
                    0.0087658 0.0087344
                                          1.004 0.322272
                   -0.0004063 0.0003132 -1.297 0.202857
## age
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.02691 on 36 degrees of freedom
## Multiple R-squared: 0.3465, Adjusted R-squared: 0.292
## F-statistic: 6.363 on 3 and 36 DF, p-value: 0.001424
```

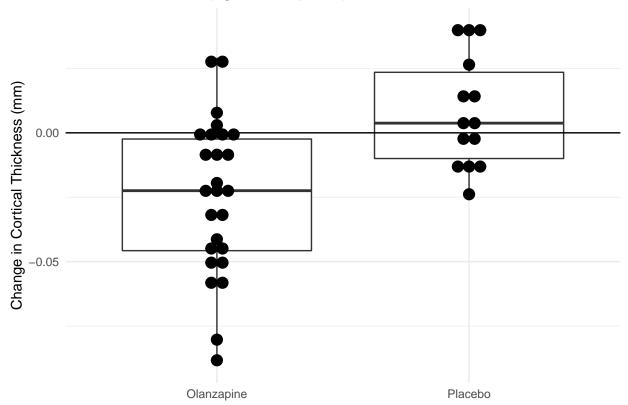
```
#run linear model with covariates of sex and age
 fit_rct <- lm(LThickness_change ~ RandomArm + sex + age + site, data= RCT_CT)
 print(fit_rct)
##
## Call:
## lm(formula = LThickness_change ~ RandomArm + sex + age + site,
##
      data = RCT_CT)
##
## Coefficients:
       (Intercept) RandomArmPlacebo
                                                  sexM
                                                                     age
         0.0003982
##
                    0.0404923
                                             0.0127657
                                                              -0.0003164
##
           siteMAS
                             {	t site}{	t NKI}
                                               sitePMC
                          -0.0212567
##
        -0.0193116
                                            -0.0287115
 summary(fit_rct)
##
## Call:
## lm(formula = LThickness_change ~ RandomArm + sex + age + site,
      data = RCT_CT)
##
## Residuals:
##
        Min
                   1Q
                         Median
                                       3Q
                                                Max
## -0.045029 -0.013484 -0.001428 0.017054 0.052285
##
## Coefficients:
                     Estimate Std. Error t value Pr(>|t|)
##
                    0.0003982 0.0177658 0.022
## (Intercept)
                                                   0.9823
## RandomArmPlacebo 0.0404923 0.0086868
                                         4.661 4.99e-05 ***
## sexM
                  0.0127657 0.0083335 1.532 0.1351
                   -0.0003164 0.0003244 -0.975
## age
                                                 0.3365
## siteMAS
                   -0.0193116 0.0110783 -1.743
                                                  0.0906 .
## siteNKI
                   -0.0212567 0.0105872 -2.008
                                                   0.0529 .
## sitePMC
                   -0.0287115  0.0130066  -2.207
                                                   0.0343 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.02527 on 33 degrees of freedom
## Multiple R-squared: 0.4717, Adjusted R-squared: 0.3757
## F-statistic: 4.911 on 6 and 33 DF, p-value: 0.001094
```

looking at the same thing for Right CT

```
#boxplot of difference in thickness (y axis) by randomization group (x axis)
ggplot(RCT_CT, aes(x= RandomArm, y = RThickness_change)) +
    geom_boxplot(outlier.shape = NA) +
    geom_dotplot(binaxis = 'y', stackdir = 'center') +
    geom_hline(yintercept = 0) +
    ggtitle("Cortical thickness (right hemisphere)") +
    xlab(NULL) +
    ylab("Change in Cortical Thickness (mm)") +
    theme_minimal()
```

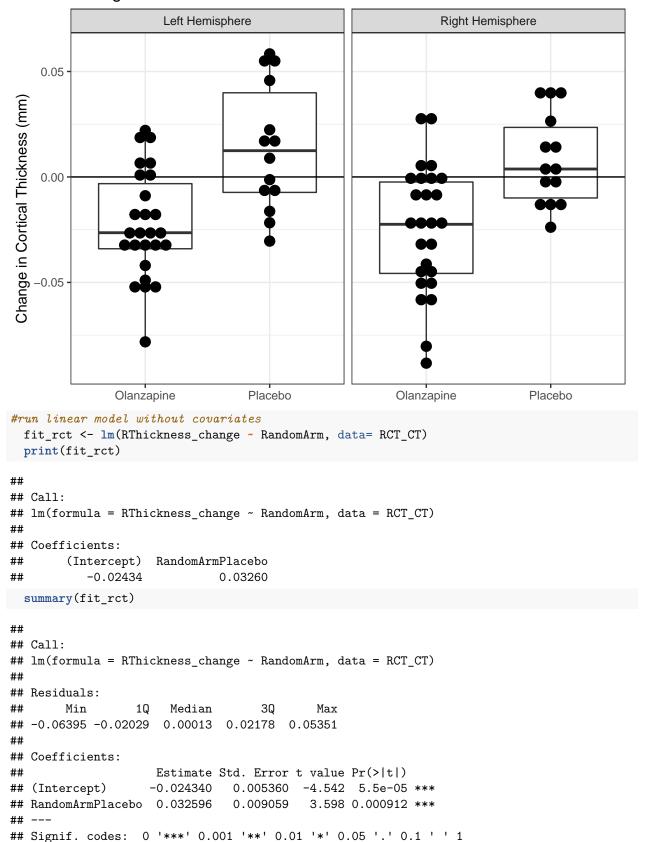
`stat_bindot()` using `bins = 30`. Pick better value with `binwidth`.

Cortical thickness (right hemisphere)



`stat_bindot()` using `bins = 30`. Pick better value with `binwidth`.

Change in Cortical Thickness



```
##
## Residual standard error: 0.02733 on 38 degrees of freedom
## Multiple R-squared: 0.2541, Adjusted R-squared: 0.2345
## F-statistic: 12.95 on 1 and 38 DF, p-value: 0.0009116
#run linear model with covariates of sex and age
  fit_rct <- lm(RThickness_change ~ RandomArm + sex + age, data= RCT_CT)
 print(fit_rct)
##
## Call:
## lm(formula = RThickness_change ~ RandomArm + sex + age, data = RCT_CT)
##
## Coefficients:
##
        (Intercept) RandomArmPlacebo
                                                   sexM
                                                                       age
         -0.0097657
                            0.0336641
                                             -0.0044117
                                                                -0.0002401
##
  summary(fit_rct)
##
## Call:
## lm(formula = RThickness_change ~ RandomArm + sex + age, data = RCT_CT)
## Residuals:
##
       Min
                  1Q
                     Median
                                    30
                                            Max
## -0.06700 -0.01799 -0.00128 0.01847 0.04950
## Coefficients:
                      Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                    -0.0097657 0.0173882 -0.562 0.577851
## RandomArmPlacebo 0.0336641 0.0093419
                                           3.604 0.000941 ***
                    -0.0044117 0.0089928 -0.491 0.626698
                    -0.0002401 0.0003225 -0.744 0.461500
## age
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.0277 on 36 degrees of freedom
## Multiple R-squared: 0.2739, Adjusted R-squared: 0.2134
## F-statistic: 4.527 on 3 and 36 DF, p-value: 0.008575
#run linear model with covariates of sex and age
 fit_rct <- lm(RThickness_change ~ RandomArm + sex + age + site, data= RCT_CT)
 print(fit_rct)
##
## lm(formula = RThickness_change ~ RandomArm + sex + age + site,
       data = RCT CT)
##
##
## Coefficients:
##
        (Intercept) RandomArmPlacebo
                                                   sexM
                                                                       age
##
         -1.111e-02
                            3.389e-02
                                             -1.632e-03
                                                                -9.043e-05
##
                                                sitePMC
            siteMAS
                              {	t site}{	t NKI}
                                             -2.463e-02
         -1.010e-02
                           -1.044e-02
 summary(fit_rct)
```

```
##
## Call:
## lm(formula = RThickness_change ~ RandomArm + sex + age + site,
      data = RCT_CT)
##
##
## Residuals:
                   10
                         Median
                                      30
                                               Max
## -0.072836 -0.015554 0.000948 0.019346 0.044262
##
## Coefficients:
                     Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                   -1.111e-02 1.939e-02 -0.573 0.57042
## RandomArmPlacebo 3.389e-02 9.480e-03
                                         3.575 0.00111 **
                  -1.632e-03 9.095e-03 -0.179 0.85869
## sexM
                   -9.043e-05 3.540e-04 -0.255 0.79998
## age
## siteMAS
                   -1.010e-02 1.209e-02 -0.835 0.40968
## siteNKI
                   -1.044e-02 1.155e-02 -0.903 0.37287
## sitePMC
                   -2.463e-02 1.419e-02 -1.735 0.09208 .
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.02758 on 33 degrees of freedom
## Multiple R-squared: 0.3405, Adjusted R-squared: 0.2206
## F-statistic: 2.84 on 6 and 33 DF, p-value: 0.02429
```

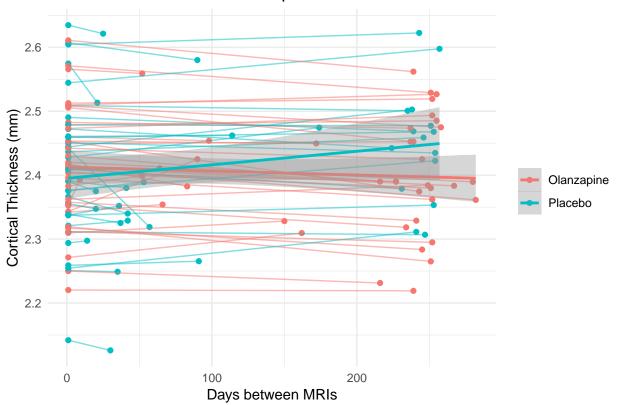
RCT & Relapse (with time as factor)

```
#restructure data for RCT & Relapse participants (N=72)
RCTRelapse_LCT <- df %>%
    gather(thick_oldcolname, thickness, LThickness_01, LThickness_02) %>%
    mutate(model_days = if_else(thick_oldcolname == "LThickness_01", 1, dateDiff))
RCTRelapse_LCT %>% filter(model_days == 1) %>% count(randomization) %>% knitr::kable()
```

${\it randomization}$	n
O	38
P	34

```
#plot all data, including outlier (participant 210030)
RCTRelapse_LCT %>%
    ggplot(aes(x=model_days, y=thickness, colour=RandomArm)) +
    geom_point() +
    geom_line(aes(group=STUDYID), alpha = 0.5) +
    geom_smooth(method="lm", formula=y~poly(x,1)) +
    ggtitle("Cortical thickness in left hemisphere over time") +
    labs(x = "Days between MRIs", y = "Cortical Thickness (mm)", colour = NULL) +
    theme_minimal()
```

Cortical thickness in left hemisphere over time



```
#run mixed linear model, with covariates
  fit_all <- lmer(thickness ~ RandomArm*model_days + sex + age + (1|STUDYID), data= RCTRelapse_LCT)
  summary(fit_all)
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: thickness ~ RandomArm * model_days + sex + age + (1 | STUDYID)
##
      Data: RCTRelapse_LCT
##
## REML criterion at convergence: -396
## Scaled residuals:
                      Median
                                    3Q
                 1Q
## -2.89073 -0.39603 -0.02082 0.40944
##
## Random effects:
  Groups
           Name
                         Variance Std.Dev.
  STUDYID (Intercept) 0.0054535 0.07385
   Residual
                         0.0004953 0.02225
## Number of obs: 144, groups: STUDYID, 72
##
```

Estimate Std. Error

2.639e+00 3.484e-02 6.864e+01 75.756

-6.099e-03 1.792e-02 6.784e+01 -0.340

-2.035e-03 1.816e-02 7.233e+01

-8.012e-05 2.340e-05 7.056e+01

df t value

-0.112

-3.424

Fixed effects:

RandomArmPlacebo

(Intercept)

model_days

sexM

##

```
-4.053e-03 5.853e-04 6.785e+01 -6.924
## RandomArmPlacebo:model_days 1.297e-04 3.942e-05 7.148e+01 3.291
                              Pr(>|t|)
                               < 2e-16 ***
## (Intercept)
## RandomArmPlacebo
                               0.91106
## model days
                               0.00103 **
## sexM
                               0.73470
## age
                              1.96e-09 ***
## RandomArmPlacebo:model_days 0.00155 **
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
              (Intr) RndmAP mdl_dy sexM
##
## RndmArmPlcb -0.206
## model_days -0.077 0.133
## sexM
              -0.171 0.036 0.001
## age
              -0.901 -0.053 0.008 -0.079
## RndmArmPl:_ 0.043 -0.176 -0.594 0.004 -0.003
#run mixed linear model, with covariates
 fit_all <- lmer(thickness ~ RandomArm*model_days + sex + age + site + (1|STUDYID), data= RCTRelapse_L
 summary(fit all)
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: thickness ~ RandomArm * model_days + sex + age + site + (1 |
      STUDYID)
##
##
     Data: RCTRelapse_LCT
##
## REML criterion at convergence: -380.4
##
## Scaled residuals:
       Min
                1Q
                     Median
                                   3Q
## -2.93368 -0.37860 -0.00129 0.40530 2.72488
##
## Random effects:
## Groups Name
                        Variance Std.Dev.
## STUDYID (Intercept) 0.0055987 0.07482
## Residual
                        0.0004952 0.02225
## Number of obs: 144, groups: STUDYID, 72
## Fixed effects:
                               Estimate Std. Error
                                                           df t value
                               2.640e+00 3.703e-02 6.556e+01 71.309
## (Intercept)
## RandomArmPlacebo
                              -1.963e-03 1.863e-02 6.894e+01 -0.105
## model days
                              -8.021e-05 2.341e-05 7.051e+01 -3.427
## sexM
                              -8.192e-03 1.840e-02 6.485e+01 -0.445
                              -4.101e-03 5.970e-04 6.486e+01 -6.869
## age
## siteMAS
                              -6.325e-03 2.335e-02 6.486e+01 -0.271
                              1.359e-04 2.587e-02 6.486e+01 0.005
## siteNKI
## sitePMC
                               2.516e-02 2.662e-02 6.485e+01
                                                              0.945
## RandomArmPlacebo:model_days 1.296e-04 3.942e-05 7.143e+01
                                                                3.286
##
                              Pr(>|t|)
```

< 2e-16 ***

(Intercept)

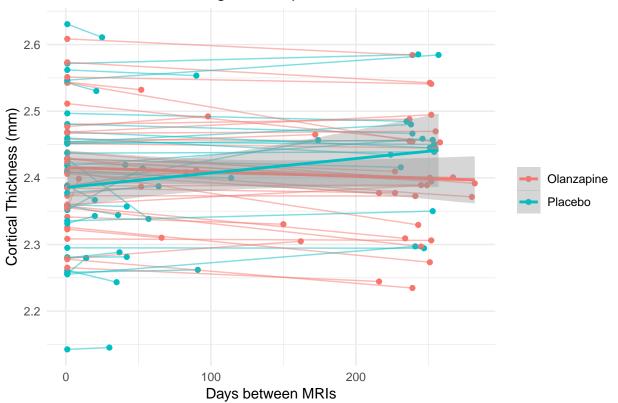
```
0.91640
## RandomArmPlacebo
## model_days
                              0.00102 **
                              0.65759
## sexM
                             2.98e-09 ***
## age
## siteMAS
                              0.78731
## siteNKI
                              0.99582
## sitePMC
                              0.34814
## RandomArmPlacebo:model_days 0.00158 **
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Correlation of Fixed Effects:
              (Intr) RndmAP mdl_dy sexM
                                         age sitMAS sitNKI sitPMC
## RndmArmPlcb -0.145
## model_days -0.074 0.129
## sexM
              -0.130 0.054 0.002
             -0.880 -0.064 0.009 -0.076
## age
## siteMAS -0.291 -0.145 0.009 -0.066 0.108
## siteNKI
            -0.174 -0.119 -0.010 -0.119 0.010 0.357
             -0.153 -0.088  0.000 -0.144 -0.009  0.343  0.319
## sitePMC
## RndmArmPl:_ 0.040 -0.172 -0.594 0.003 -0.003 0.000 0.009 0.002
```

Running the right hemisphere RCTRelapse

```
#restructure data for RCT & Relapse participants (N=72)
RCTRelapse_RCT <- df %>%
    gather(thick_oldcolname, thickness, RThickness_01, RThickness_02) %>%
    mutate(model_days = if_else(thick_oldcolname == "RThickness_01", 1, dateDiff))

#plot all data, including outlier (participant 210030)
RCTRelapse_RCT %>%
    ggplot(aes(x=model_days, y=thickness, colour=RandomArm)) +
    geom_point() +
    geom_line(aes(group=STUDYID), alpha = 0.5) +
    geom_smooth(method="lm", formula=y~poly(x,1)) +
    ggtitle("Cortical thickness in right hemisphere over time") +
    labs(x = "Days between MRIs", y = "Cortical Thickness (mm)", colour = NULL) +
    theme_minimal()
```

Cortical thickness in right hemisphere over time



```
#run mixed linear model, with covariates
  fit_all <- lmer(thickness ~ RandomArm*model_days + sex + age + (1|STUDYID), data= RCTRelapse_RCT)
  summary(fit_all)
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]</pre>
```

```
## Formula: thickness ~ RandomArm * model_days + sex + age + (1 | STUDYID)
## Data: RCTRelapse_RCT
##
## REML criterion at convergence: -409
##
## Scaled residuals:
## Min 1Q Median 3Q Max
## -2.34720 -0.42608 -0.01215 0.43733 2.27881
##
## Random effects:
```

Groups Name Variance Std.Dev.
STUDYID (Intercept) 0.0057442 0.07579
Residual 0.0003947 0.01987
Number of obs: 144, groups: STUDYID, 72
##

Fixed effects:

(Intercept) 2.618e+00 3.554e-02 6.847e+01 73.658 ## RandomArmPlacebo -1.455e-02 1.847e-02 7.131e+01 -0.788 ## model_days -8.813e-05 2.090e-05 7.041e+01 -4.216 ## sexM -7.789e-03 1.830e-02 6.786e+01 -0.426

```
-3.588e-03 5.975e-04 6.786e+01 -6.004
## age
## RandomArmPlacebo:model_days 1.281e-04 3.524e-05 7.112e+01 3.635
                             Pr(>|t|)
                               < 2e-16 ***
## (Intercept)
## RandomArmPlacebo
                              0.433361
## model days
                              7.28e-05 ***
## sexM
                              0.671706
## age
                              8.40e-08 ***
## RandomArmPlacebo:model_days 0.000522 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
              (Intr) RndmAP mdl_dy sexM
##
## RndmArmPlcb -0.205
## model_days -0.067 0.117
              -0.172 0.036 0.001
## sexM
## age
              -0.902 -0.053 0.007 -0.079
## RndmArmPl:_ 0.037 -0.155 -0.593 0.003 -0.002
#run mixed linear model, with covariates
 fit_all <- lmer(thickness ~ RandomArm*model_days + sex + age + site + (1|STUDYID), data= RCTRelapse_R
 summary(fit all)
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: thickness ~ RandomArm * model_days + sex + age + site + (1 |
      STUDYID)
##
##
     Data: RCTRelapse_RCT
##
## REML criterion at convergence: -393.2
##
## Scaled residuals:
       Min
                1Q
                     Median
                                   3Q
## -2.37660 -0.44552 -0.00537 0.43115 2.24882
##
## Random effects:
## Groups Name
                        Variance Std.Dev.
## STUDYID (Intercept) 0.0059298 0.07701
## Residual
                        0.0003947 0.01987
## Number of obs: 144, groups: STUDYID, 72
## Fixed effects:
                               Estimate Std. Error
                                                           df t value
                               2.619e+00 3.788e-02 6.541e+01 69.150
## (Intercept)
## RandomArmPlacebo
                              -1.446e-02 1.900e-02 6.800e+01 -0.761
## model days
                              -8.825e-05 2.091e-05 7.037e+01 -4.221
## sexM
                              -9.731e-03 1.883e-02 6.487e+01 -0.517
                              -3.638e-03 6.112e-04 6.487e+01 -5.952
## age
## siteMAS
                              -7.805e-03 2.390e-02 6.488e+01 -0.327
## siteNKI
                               4.352e-03 2.649e-02 6.488e+01 0.164
## sitePMC
                               2.024e-02 2.725e-02 6.487e+01
                                                                0.743
## RandomArmPlacebo:model_days 1.280e-04 3.524e-05 7.108e+01
                                                                3.631
##
                              Pr(>|t|)
```

< 2e-16 ***

(Intercept)

```
## RandomArmPlacebo
                   0.449288
                      7.14e-05 ***
## model_days
## sexM
                      0.607158
## age
                       1.18e-07 ***
## siteMAS
                       0.745036
## siteNKI
                       0.870002
## sitePMC
                       0.460300
## RandomArmPlacebo:model_days 0.000529 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Correlation of Fixed Effects:
           (Intr) RndmAP mdl_dy sexM age sitMAS sitNKI sitPMC
## RndmArmPlcb -0.144
## model_days -0.064 0.113
## sexM
          -0.130 0.055 0.002
## RndmArmPl:_ 0.035 -0.151 -0.593 0.003 -0.003 0.000 0.008 0.002
```