# Martel Paper - Daily OA Pain

### 2024-03-18

## **Data Wrangling**

```
## Load packages
library(tidyverse)
library(ggplot2)
library(tidyr)
library(haven) ## This library provides functions to read sav file into R
library(lme4)
library(lmerTest)
## Read data
data_paper <- read_sav("Dataset; 2024.1.sav")</pre>
checkdf3 <- data_paper |>
  subset(ID == 2072) |>
  select(c(ID,
           Level1 Even DateIn,
           Level1_Even_TimeIn,
           Wave_Day))
## Delete Weird ID 2072 those weird reporting days
data_paper <- data_paper |>
  filter(!(ID == 2072 & Wave Day >= 7))
```

#### Adjusting Wave Day

- $DT_i$  combines  $D_i$  and  $T_i$ : 'DateTime' variable
- $DT_0$  is the first response 'DateTime' for each patient
- $W_i$  is the adjusted 'Wave\_Day' variable
- Add a grace period G for calculating the adjusted  $W_i$ , in our case G=6 hours
- Calculate the datetime difference  $H_i$  in \*\*hours\*\* from the first response, incorporating the grace period:

$$H_i = DT_i - DT_{i-1}$$

ungroup()

• Then apply the grace period indicator  $I_i$ :

$$I_i = \begin{cases} 1 & \text{if } H_i \le 24 + G \\ \left\lceil \frac{H_i - G}{24} \right\rceil & \text{otherwise} \end{cases}$$

• The initial response for 'Wave\_Day' is 1, i.e.,  $W_0 = 1$ , then the adjusted 'Wave\_Day'  $W_i$  is

$$W_i = \sum_{i=0}^{i-1} I_i$$

```
## Consecutive Days - Grace Period 6 hours
data_paper <- data_paper |>
 mutate(Lev1_DateTimeIn = as.POSIXct(strptime(paste(Level1_Even_DateIn,
                                                     Level1 Even TimeIn),
                                          format="%Y-%m-%d %H:%M:"))) |>
 arrange(ID, Lev1_DateTimeIn) |>
 group_by(ID) |>
 mutate(
    TimeDiffHours = as.numeric(difftime(Lev1_DateTimeIn,
                                        lag(Lev1_DateTimeIn,
                                            default = first(Lev1_DateTimeIn)),
                                        units = "hours")), # T diff
    WithinGracePeriod = if_else(TimeDiffHours <= 30,
                                1,
                                ceiling((TimeDiffHours - 6) / 24)), # Check grace perio
   Wave_Day_Adjusted = cumsum(WithinGracePeriod) # Adjusted Wave_Day
 ) |>
 ungroup()
###### Check if the above approach is correct ######
checkdf <- data_paper |> select(c(ID,
                                  Lev1_DateTimeIn,
                                  TimeDiffHours,
                                  WithinGracePeriod,
                                  Wave_Day_Adjusted,
                                  Baseline_Demog_BMI))
checkdf2 <- checkdf |> subset(ID == 2072) # Weird ID 2072
max(checkdf$Wave_Day_Adjusted, na.rm = TRUE)
## [1] 21
## Fill in the gap of Wave_Day
data paper2 <- data paper |>
 group_by(ID) |>
 complete(Wave_Day = 1:14) |>
```

## Check the aberrant values

summary(data\_paper2\$IndexLev1\_NegativeAffect\_Total) # Lev 1 Negative Affect

```
##
      Min. 1st Qu.
                    Median
                              Mean 3rd Qu.
                                              Max.
                                                      NA's
                            25.189 37.000 97.333
##
     0.000
            8.333 20.667
                                                      1228
summary(data paper2$IndexLev1 Catastrophizing Total) # Lev 1 Catas
##
     Min. 1st Qu.
                    Median
                              Mean 3rd Qu.
                                              Max.
                                                      NA's
##
     0.000
            2.667 12.667 23.085 38.000 100.000
                                                      1228
summary(data paper2$IndexLev2 QST BaselinePPTh) # Lev 2 PPThs?
      Min. 1st Qu. Median
                              Mean 3rd Qu.
##
                                              Max.
                                                      NA's
##
      67.0
             235.5
                     369.0
                             395.8
                                     482.5 1200.0
                                                      1321
summary(data_paper2$IndexLev2_QST_TSPAve) # Lev 2 TSP?
                    Median
                                                      NA's
##
      Min. 1st Qu.
                              Mean 3rd Qu.
                                              {\tt Max.}
##
     -2.50
              1.50
                      7.50
                             13.97
                                     15.00
                                             94.00
                                                      1284
summary(data_paper2$IndexLev2_QST_CpmTrialAve) # Lev 2 CPM?
##
      Min. 1st Qu. Median
                              Mean 3rd Qu.
                                                      NA's
                                              Max.
##
     37.56 104.00 119.74 123.10 135.13 251.76
                                                      1329
summary(data_paper2$IndexLev1_PainAverage) # Lev 1 Pain?
##
      Min. 1st Qu. Median
                              Mean 3rd Qu.
                                              Max.
                                                      NA's
##
      0.00
             24.00
                     38.00
                             41.73
                                     59.75 100.00
                                                      1234
###### Check if lagged value is correct #####
data paper2 check <- data paper2 |>
 select(c(ID, Wave Day Adjusted, IndexLev1 PainAverage, IndexLev1 PainAverage Lagged))
                  APE(t_i) = I(PAIN(t_i) - PAIN(t_i - 1) \ge 20)
## APE index
data_paper2 <- data_paper2 |>
 group_by(ID) |>
 mutate(APE = ifelse(IndexLev1 PainAverage - IndexLev1 PainAverage Lagged >= 20, 1, 0))
###### Check if it is correctly coded ######
data paper2 check <- data paper2 |>
 select(c(ID, Wave_Day_Adjusted,
           IndexLev1 PainAverage,
           IndexLev1 PainAverage Lagged,
           APE))
```

For calculating the RPE based on the within person mean, define the indicator that the pain is above the average pain for person i on day  $t_i$ . Note that  $n_{t_i} = \max t_i$  for patient i.

$$A(t_i) = I\left(PAIN(t_i) > \frac{1}{n_{t_i}} \sum_{t_i=1}^{n_{t_i}} PAIN(t_i)\right)$$

Then define the RPE given  $A(t_i) = 1$  for person i on day  $t_i$ .

$$RPE(t_i) = I\left(PAIN(t_i) \le \frac{1}{n_{t_i}} \sum_{t_i=1}^{n_{t_i}} PAIN(t_i)\right) \times A(t_i - 1)$$

For calculating the RPE based on the APE,

$$RPE(t_i) = I\left(PAIN(t_i) \le \frac{1}{n_{t_i}} \sum_{t_i=1}^{n_{t_i}} PAIN(t_i)\right) \times APE(t_i - 1)$$

## Analysis

```
## Unadjusted APE to Negative Affect
model_APE.NA <- glmer(APE ~ IndexLev1_NegativeAffect_Total + (1|ID), family = binomial()</pre>
summary(model_APE.NA)
## Generalized linear mixed model fit by maximum likelihood (Laplace
    Approximation) [glmerMod]
## Family: binomial (logit)
## Formula: APE ~ IndexLev1_NegativeAffect_Total + (1 | ID)
##
      Data: data paper2
##
##
        AIC
                 BIC
                       logLik deviance df.resid
                       -222.8
##
      451.6
               465.7
                                 445.6
                                            809
##
## Scaled residuals:
      Min
                1Q Median
                                3Q
                                       Max
## -0.5907 -0.2780 -0.2335 -0.2048 3.9737
##
## Random effects:
## Groups Name
                       Variance Std.Dev.
           (Intercept) 0.7268
                                0.8525
## Number of obs: 812, groups:
                                ID, 157
##
## Fixed effects:
##
                                   Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                                  -3.124537
                                              0.329004 -9.497
                                                                  <2e-16 ***
## IndexLev1_NegativeAffect_Total 0.014167
                                              0.006951
                                                         2.038
                                                                  0.0415 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
               (Intr)
## IndxL1_NA_T -0.732
## Unadjusted APE to Catastrophizing
model_APE.Cata <- glmer(APE ~ IndexLev1_Catastrophizing_Total + (1 ID), family = binomia</pre>
summary(model APE.Cata)
## Generalized linear mixed model fit by maximum likelihood (Laplace
##
     Approximation) [glmerMod]
## Family: binomial (logit)
## Formula: APE ~ IndexLev1_Catastrophizing_Total + (1 | ID)
##
      Data: data_paper2
##
##
        AIC
                 BIC
                       logLik deviance df.resid
```

```
##
      444.2
              458.3 -219.1
                                 438.2
                                            809
##
## Scaled residuals:
##
       Min
                1Q Median
## -0.6057 -0.2835 -0.2176 -0.1865 3.9271
##
## Random effects:
## Groups Name
                       Variance Std.Dev.
## ID
           (Intercept) 0.7367
                                0.8583
## Number of obs: 812, groups: ID, 157
## Fixed effects:
                                    Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                                               0.316961 -10.261 < 2e-16 ***
                                   -3.252303
                                                          3.292 0.000994 ***
## IndexLev1_Catastrophizing_Total 0.019020
                                               0.005777
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Correlation of Fixed Effects:
##
               (Intr)
## IndxLv1 C T -0.720
## Unadjusted APE to PPThs
model APE.PPThs <- glmer(APE ~ IndexLev2_QST_BaselinePPTh + (1|ID), family = binomial()</pre>
## Warning in checkConv(attr(opt, "derivs"), opt$par, ctrl = control$checkConv, :
## Model failed to converge with max|grad| = 0.00456593 (tol = 0.002, component 1)
## Warning in checkConv(attr(opt, "derivs"), opt$par, ctrl = control$checkConv, : Model
## - Rescale variables?; Model is nearly unidentifiable: large eigenvalue ratio
## - Rescale variables?
summary(model APE.PPThs)
## Generalized linear mixed model fit by maximum likelihood (Laplace
     Approximation) [glmerMod]
##
## Family: binomial (logit)
## Formula: APE ~ IndexLev2_QST_BaselinePPTh + (1 | ID)
      Data: data paper2
##
##
##
       AIC
                 BIC
                       logLik deviance df.resid
##
      415.4
              429.2
                       -204.7
                                 409.4
                                            726
##
## Scaled residuals:
       Min
                1Q Median
                                3Q
## -0.4954 -0.2973 -0.2563 -0.2168 4.4125
##
```

```
## Random effects:
## Groups Name
                      Variance Std.Dev.
## ID
          (Intercept) 0.49
                              0.7
## Number of obs: 729, groups: ID, 142
##
## Fixed effects:
##
                              Estimate Std. Error z value Pr(>|z|)
                             ## (Intercept)
## IndexLev2 QST BaselinePPTh -0.0021978 0.0009134 -2.406
                                                            0.0161 *
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##
              (Intr)
## IL2 QST BPP -0.807
## optimizer (Nelder Mead) convergence code: 0 (OK)
## Model failed to converge with max|grad| = 0.00456593 (tol = 0.002, component 1)
## Model is nearly unidentifiable: very large eigenvalue
## - Rescale variables?
## Model is nearly unidentifiable: large eigenvalue ratio
## - Rescale variables?
## Unadjusted APE to TSP
model APE.TSP <- glmer(APE ~ IndexLev2 QST TSPAve + (1 ID), family = binomial(), data =
summary(model_APE.TSP)
## Generalized linear mixed model fit by maximum likelihood (Laplace
    Approximation) [glmerMod]
## Family: binomial (logit)
## Formula: APE ~ IndexLev2_QST_TSPAve + (1 | ID)
##
     Data: data paper2
##
##
       AIC
                BIC
                      logLik deviance df.resid
##
     433.9
              447.8 -213.9
                               427.9
                                          758
##
## Scaled residuals:
               1Q Median
                               3Q
                                     Max
## -0.4841 -0.2851 -0.2630 -0.2260 5.6749
##
## Random effects:
## Groups Name
                      Variance Std.Dev.
          (Intercept) 0.512
## ID
                              0.7155
## Number of obs: 761, groups: ID, 146
## Fixed effects:
```

```
##
                       Estimate Std. Error z value Pr(>|z|)
                       -2.40370
                                   0.24326 -9.881 <2e-16 ***
## (Intercept)
## IndexLev2_QST_TSPAve -0.01805
                                   0.01029 - 1.754
                                                     0.0794 .
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
              (Intr)
## IL2_QST_TSP -0.450
## Unadjusted APE to CPM
model_APE.CPM <- glmer(APE ~ IndexLev2_QST_CpmTrialAve + (1|ID), family = binomial(), da</pre>
## Warning in checkConv(attr(opt, "derivs"), opt$par, ctrl = control$checkConv, : Model
## - Rescale variables?
summary(model APE.CPM)
## Generalized linear mixed model fit by maximum likelihood (Laplace
    Approximation) [glmerMod]
## Family: binomial (logit)
## Formula: APE ~ IndexLev2 QST CpmTrialAve + (1 | ID)
##
     Data: data_paper2
##
##
       AIC
                BIC
                      logLik deviance df.resid
     415.4
              429.2 -204.7
##
                                409.4
                                           719
##
## Scaled residuals:
      Min
               1Q Median
                               3Q
                                      Max
## -0.4942 -0.2957 -0.2365 -0.2284 3.3972
##
## Random effects:
## Groups Name
                      Variance Std.Dev.
          (Intercept) 0.7285
                               0.8536
## ID
## Number of obs: 722, groups: ID, 141
##
## Fixed effects:
##
                             Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                            -2.895586
                                        0.682405 -4.243 2.2e-05 ***
## IndexLev2_QST_CpmTrialAve 0.001580
                                        0.005063
                                                  0.312
                                                            0.755
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Correlation of Fixed Effects:
              (Intr)
## IL2_QST_CTA -0.939
```

```
## optimizer (Nelder_Mead) convergence code: 0 (OK)
## Model is nearly unidentifiable: very large eigenvalue
## - Rescale variables?
```