Data analysis pilots

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2024-01-19

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Dataset loading

Dataset preparation

```
d <- d %>% mutate(Subject=rep(seq_len(length(datasets)), each=100))
```

Then we remove total dwell times <=1500ms, scale the Agreement feature and take its square, and log-transform Dwell. Time. We do the same with Agreement. Contacts

Variables plot (violin & boxplot + density)

```
violin_fixation <- ggplot(d_final, aes(x="Dwell time", y=Dwell.Time.Log)) +
  ylab("Dwell Time (log)") + theme_minimal() +
  theme(legend.position = "none") +
  geom_violin(width=1.2) + geom_boxplot(width=0.1, alpha=0.2)</pre>
```

```
violin_agreement <- ggplot(d_final, aes(x="Agreement", y=Agreement.Scaled)) +</pre>
  ylab("Agreement (scaled)") + theme_minimal() +
  theme(legend.position = "none") +
  geom_violin(width=1.2) + geom_boxplot(width=0.1, alpha=0.2)
violin_agreement_contacts <- ggplot(d_final, aes(x="Agreement (contacts)", y=Agreement.Contacts.Scaled)</pre>
  ylab("Agreement (constacts; scaled)") + theme_minimal() +
  theme(legend.position = "none") +
  geom_violin(width=1.2) + geom_boxplot(width=0.1, alpha=0.2)
density_fixation <- ggplot(d_final, aes(x=Dwell.Time.Log)) + geom_density()</pre>
density_agreement <- ggplot(d_final, aes(x=Agreement.Scaled)) + geom_density()</pre>
density_agreement_contacts <- ggplot(d_final, aes(x=Agreement.Contacts.Scaled)) + geom_density()</pre>
grid.arrange(violin_fixation, violin_agreement, violin_agreement_contacts,
              density_fixation, density_agreement, density_agreement_contacts,
              nrow=2)
                                                                    Agreement (constacts; scaled
    10
                                  Agreement (scaled)
Dwell Time (log)
                                      -2
               Dwell time
                                                 Agreement
                                                                              Agreement (contacts)
                   Χ
                                                     Χ
                                                                                       Х
                                                                        0.5
                                      0.4 -
    0.6
                                                                        0.4 -
                                     0.3
                                  density
                                                                     density
0.2 -
density
   0.4
                                      0.2
    0.2
                                     0.1 -
                                                                        0.1
                                     0.0
                                                                       0.0
    0.0
                     9
                             10
                                                       0
            Dwell.Time.Log
                                            Agreement.Scaled
                                                                         Agreement.Contacts.Scale
```

Statistical models

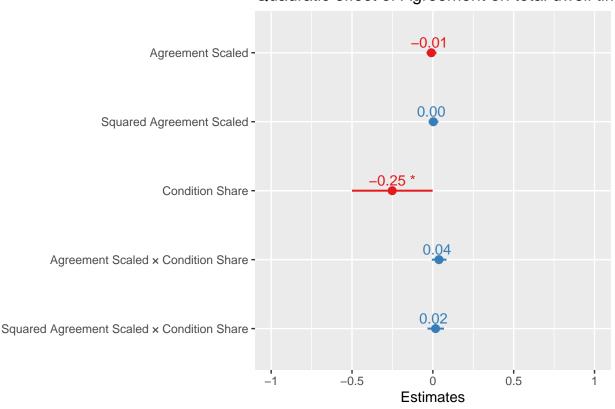
Agreement

```
res_mixed <- lmer(Dwell.Time.Log ~ Agreement.Scaled + Squared.Agreement.Scaled + Condition_Share + Agre (1 | Index) +
```

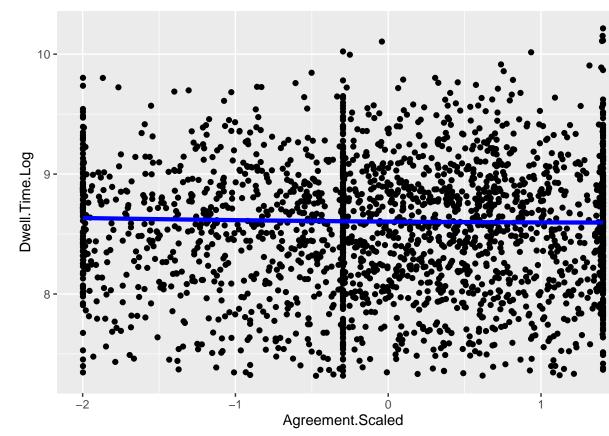
```
(Agreement.Scaled + Squared.Agreement.Scaled + Condition_Share + Agreement.Scaled*C
                  data=d_final, control = lmerControl(optimizer ="bobyga"))
## boundary (singular) fit: see help('isSingular')
## Warning: Model failed to converge with 2 negative eigenvalues: -1.3e-05
## -4.7e-04
summary(res_mixed)
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: Dwell.Time.Log ~ Agreement.Scaled + Squared.Agreement.Scaled +
       Condition_Share + Agreement.Scaled * Condition_Share + Squared.Agreement.Scaled *
       Condition_Share + (1 | Index) + (Agreement.Scaled + Squared.Agreement.Scaled +
##
       Condition_Share + Agreement.Scaled * Condition_Share + Squared.Agreement.Scaled *
##
       Condition_Share | Subject)
##
      Data: d final
##
## Control: lmerControl(optimizer = "bobyqa")
## REML criterion at convergence: 2820.1
##
## Scaled residuals:
##
       Min
                1Q Median
                                3Q
                                       Max
## -3.2546 -0.6495 -0.0344 0.6160 3.7541
##
## Random effects:
## Groups
                                                      Variance Std.Dev. Corr
                                                      0.0200991 0.14177
##
  Index
             (Intercept)
   Subject (Intercept)
                                                      0.0820210 0.28639
                                                      0.0002062 0.01436
##
             Agreement.Scaled
                                                                           0.45
##
             Squared.Agreement.Scaled
                                                      0.0004584 0.02141
                                                                          -0.84
##
             Condition_Share
                                                      0.0990397 0.31471
                                                                          -0.36
##
             Agreement.Scaled:Condition_Share
                                                      0.0010536 0.03246
                                                                           0.26
##
             Squared.Agreement.Scaled:Condition_Share 0.0030229 0.05498
                                                                           0.56
                                                       0.1823998 0.42708
##
   Residual
##
##
##
##
##
   -0.86
  -0.53 0.53
   -0.52 0.17 0.07
##
##
    0.32 -0.51 -0.66 0.59
## Number of obs: 2256, groups: Index, 100; Subject, 26
## Fixed effects:
##
                                             Estimate Std. Error
                                                                         df t value
## (Intercept)
                                             8.603539
                                                        0.082744 12.749158 103.978
## Agreement.Scaled
                                            -0.009477
                                                        0.014720 38.665423
                                                                            -0.644
## Squared.Agreement.Scaled
                                             0.002797
                                                        0.014496 23.461014
                                                                              0.193
## Condition_Share
                                            -0.251466
                                                        0.126338 23.200045
## Agreement.Scaled:Condition_Share
                                             0.037930
                                                        0.022303 38.706171
                                                                              1.701
## Squared.Agreement.Scaled:Condition_Share 0.016859
                                                        0.025027 18.692890
```

```
Pr(>|t|)
##
## (Intercept)
                                              <2e-16 ***
## Agreement.Scaled
                                              0.5235
## Squared.Agreement.Scaled
                                              0.8487
## Condition_Share
                                              0.0585 .
## Agreement.Scaled:Condition_Share
                                              0.0970 .
## Squared.Agreement.Scaled:Condition_Share
                                              0.5088
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##
               (Intr) Agrm.S Sq.A.S Cndt_S A.S:C_
## Agrmnt.Scld 0.058
## Sqrd.Agrm.S -0.479 0.170
## Conditn_Shr -0.635 -0.038 0.309
## Agrmn.S:C_S -0.042 -0.586 -0.098 0.068
## Sqr.A.S:C_S 0.273 -0.095 -0.556 -0.343 0.425
## optimizer (bobyqa) convergence code: 0 (OK)
## boundary (singular) fit: see help('isSingular')
plot_model(res_mixed,
           #axis.labels=c("Agreement (Quadratic)", "Agreement"),
          show.values=TRUE, show.p=TRUE,
           title="Quadratic effect of Agreement on total dwell times")
```

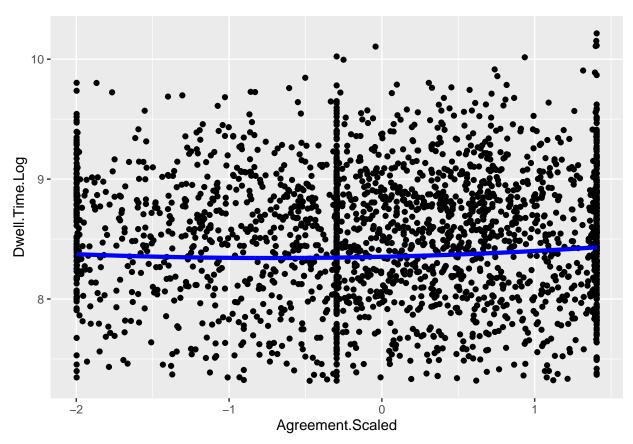
Quadratic effect of Agreement on total dwell tim



```
quad_function <- function(x, alpha, beta1, beta2, beta3, beta4, beta5, condition)
{</pre>
```



Fixed effects plot



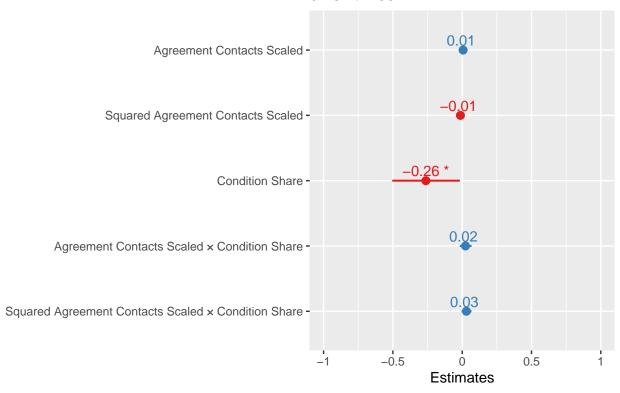
Agreement (contacts)

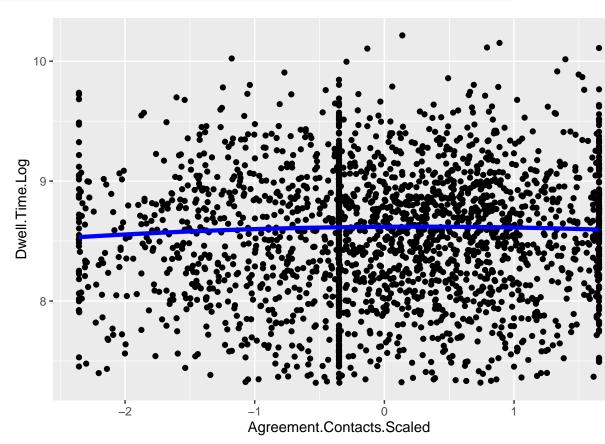
```
res_mixed <- lmer(Dwell.Time.Log ~ Agreement.Contacts.Scaled + Squared.Agreement.Contacts.Scaled + Cond
                    (1 \mid Index) +
                    (Agreement.Contacts.Scaled + Squared.Agreement.Contacts.Scaled + Condition_Share +
                  data=d_final, control = lmerControl(optimizer ="bobyqa"))
## boundary (singular) fit: see help('isSingular')
## Warning: Model failed to converge with 4 negative eigenvalues: -3.6e-05
## -5.0e-05 -5.8e-05 -1.1e+01
summary(res_mixed)
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula:
## Dwell.Time.Log ~ Agreement.Contacts.Scaled + Squared.Agreement.Contacts.Scaled +
       Condition_Share + Agreement.Contacts.Scaled * Condition_Share +
##
##
       Squared.Agreement.Contacts.Scaled * Condition_Share + (1 |
       Index) + (Agreement.Contacts.Scaled + Squared.Agreement.Contacts.Scaled +
##
       Condition_Share + Agreement.Contacts.Scaled * Condition_Share +
##
```

```
##
       Squared.Agreement.Contacts.Scaled * Condition_Share | Subject)
##
      Data: d final
## Control: lmerControl(optimizer = "bobyqa")
## REML criterion at convergence: 2824.4
##
## Scaled residuals:
##
       Min
                1Q Median
                                3Q
                                        Max
## -3.2045 -0.6569 -0.0214 0.6181 3.7307
##
## Random effects:
  Groups
                                                                 Variance Std.Dev.
##
             Name
##
   Index
             (Intercept)
                                                                 2.015e-02 0.141950
             (Intercept)
##
   Subject
                                                                7.082e-02 0.266116
##
                                                                 1.136e-05 0.003370
             Agreement.Contacts.Scaled
##
             Squared.Agreement.Contacts.Scaled
                                                                 3.750e-05 0.006124
##
             Condition_Share
                                                                1.353e-01 0.367824
##
             Agreement.Contacts.Scaled:Condition_Share
                                                                7.142e-04 0.026724
##
             Squared.Agreement.Contacts.Scaled:Condition_Share 2.013e-04 0.014188
##
   Residual
                                                                 1.834e-01 0.428298
##
   Corr
##
##
     1.00
##
##
    0.50 0.50
   -0.45 -0.45 -0.39
   -0.41 -0.41 -0.04 -0.63
##
   -0.49 -0.49 -0.41 -0.49 0.92
##
##
## Number of obs: 2256, groups: Index, 100; Subject, 26
##
## Fixed effects:
##
                                                        Estimate Std. Error
                                                                   0.076945
## (Intercept)
                                                        8.618766
## Agreement.Contacts.Scaled
                                                        0.006745
                                                                   0.013970
## Squared.Agreement.Contacts.Scaled
                                                       -0.012674
                                                                   0.011433
## Condition Share
                                                       -0.261895
                                                                    0.123285
## Agreement.Contacts.Scaled:Condition_Share
                                                        0.024094
                                                                    0.021005
## Squared.Agreement.Contacts.Scaled:Condition_Share
                                                        0.030468
                                                                    0.017316
##
                                                               df t value Pr(>|t|)
## (Intercept)
                                                       12.830448 112.013
                                                                            <2e-16
## Agreement.Contacts.Scaled
                                                      417.875079
                                                                   0.483
                                                                            0.6295
## Squared.Agreement.Contacts.Scaled
                                                        6.253790
                                                                  -1.109
                                                                            0.3084
                                                                  -2.124
## Condition_Share
                                                       22.393086
                                                                            0.0449
## Agreement.Contacts.Scaled:Condition_Share
                                                       94.336630
                                                                   1.147
                                                                            0.2543
## Squared.Agreement.Contacts.Scaled:Condition_Share
                                                       29.905233
                                                                    1.760
                                                                            0.0887
##
## (Intercept)
## Agreement.Contacts.Scaled
## Squared.Agreement.Contacts.Scaled
## Condition_Share
## Agreement.Contacts.Scaled:Condition_Share
## Squared.Agreement.Contacts.Scaled:Condition_Share .
## ---
```

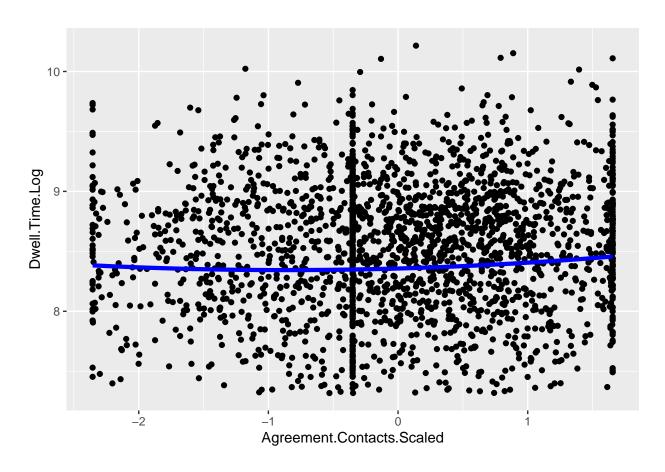
```
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
              (Intr) Ag.C.S Sq.A.C.S Cndt_S A.C.S:
##
## Agrmnt.Cn.S 0.030
## Sqrd.Ag.C.S -0.070 0.238
## Conditn Shr -0.603 -0.019 0.041
## Agr.C.S:C_S -0.021 -0.610 -0.149
                                     -0.273
## S.A.C.S:C_S 0.043 -0.159 -0.641
                                     -0.260 0.354
## optimizer (bobyqa) convergence code: 0 (OK)
## boundary (singular) fit: see help('isSingular')
plot_model(res_mixed,
           #axis.labels=c("Agreement (Quadratic)", "Agreement"),
          show.values=TRUE, show.p=TRUE,
          title="Quadratic effect of contacts Agreement on total dwell times")
```

Quadratic effect of contacts Agreement of dwell times





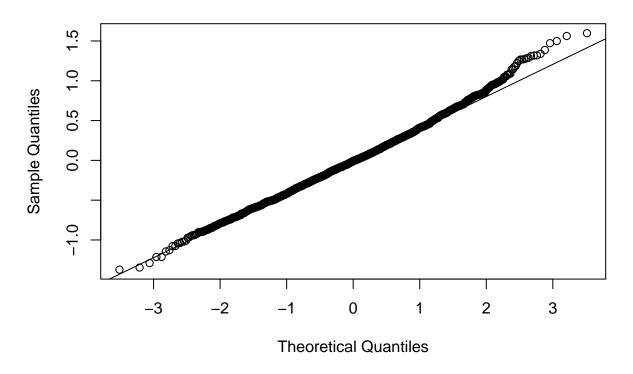
Fixed effects plot



Residuals analysis

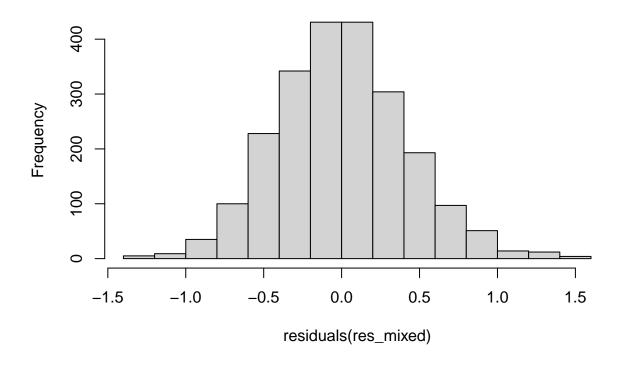
qqnorm(residuals(res_mixed))
qqline(residuals(res_mixed))

Normal Q-Q Plot



hist(residuals(res_mixed))

Histogram of residuals(res_mixed)



```
ks.test(x=residuals(res_mixed), y='pnorm')
```

```
##
## Asymptotic one-sample Kolmogorov-Smirnov test
##
## data: residuals(res_mixed)
## D = 0.20997, p-value < 2.2e-16
## alternative hypothesis: two-sided</pre>
```