LMM considerations

One factor rm ANOVA, categorical fixed effect

Model works fine with random intercept only:

model1 <- lmer(measure ~ situation + (1|id),df1)</pre>

lmer model, random intercept

summary(model1)

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: measure ~ situation + (1 | id)
##
     Data: df1
##
## REML criterion at convergence: 73.5
##
## Scaled residuals:
       \mathtt{Min}
                  1Q
                      Median
## -1.92330 -0.49814 0.05354 0.51792 1.54811
##
## Random effects:
                         Variance Std.Dev.
## Groups Name
## id
             (Intercept) 0.1627
                                  0.4034
## Residual
                         0.1237
                                  0.3517
## Number of obs: 48, groups: id, 32
##
## Fixed effects:
                                         df t value Pr(>|t|)
##
              Estimate Std. Error
## (Intercept) 1.21798 0.10055 41.36437 12.114 3.45e-15 ***
## situation2 0.09838
                           0.11339 18.12464
                                            0.868
                                                       0.397
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Correlation of Fixed Effects:
## situation2 -0.471
Adding a random slope makes it unidentifiable
# lmer model, random intercept + random slope
model2 <- lmer(measure ~ situation + (1+situation|id),df1)</pre>
## Error: number of observations (=48) <= number of random effects (=64) for term (1 + situation | id);
```

Continuous predictor (fixed effect) works

Adding a random slope works when the predictor is continuous

```
model3 <- lmer(delta_distance ~ acceleration_s + (1+acceleration_s|id),df2) #standardized predictor nee
summary(model3)</pre>
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: delta_distance ~ acceleration_s + (1 + acceleration_s | id)
##
     Data: df2
## REML criterion at convergence: 1304.3
## Scaled residuals:
       Min
              10
                     Median
                                   30
## -1.68947 -0.40064 0.02119 0.40142 1.80638
## Random effects:
                           Variance Std.Dev. Corr
## Groups Name
            (Intercept)
                           7637.8
                                    87.39
## id
##
            acceleration_s 685.5
                                    26.18
                                             0.64
## Residual
                           1349.5
                                    36.74
## Number of obs: 118, groups: id, 39
## Fixed effects:
##
                 Estimate Std. Error
                                          df t value Pr(>|t|)
## (Intercept)
                  405.917
                              14.500 38.379
                                               28.00
## acceleration_s 165.635
                               5.729 32.585
                                               28.91
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
              (Intr)
## accelertn_s 0.476
```

two factor mixed anova

model, random intercept

Random intercept only works:

```
model4 <- lmer(dv ~ group*stage+(1|id),df3)</pre>
summary(model4)
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: dv ~ group * stage + (1 | id)
      Data: df3
##
## REML criterion at convergence: 309.1
## Scaled residuals:
      Min
                1Q Median
                                3Q
## -1.8639 -0.4909 -0.1308 0.3467 4.4436
##
## Random effects:
## Groups Name
                         Variance Std.Dev.
## id
            (Intercept) 0.4347 0.6593
```

```
## Residual
                      0.3205
                               0.5661
## Number of obs: 141, groups: id, 36
## Fixed effects:
##
               Estimate Std. Error
                                      df t value Pr(>|t|)
                ## (Intercept)
## group2
                -1.1729 0.3209 80.6371 -3.655 0.000457 ***
                ## stage2
## stage3
                -0.8985
## stage4
                -1.0456 0.2476 96.4297 -4.223 5.46e-05 ***
## stage5
                -0.4726
                           0.2574 96.1694 -1.836 0.069457 .
                -0.9400
                           0.2668 97.8314 -3.523 0.000650 ***
## stage6
## group2:stage2 0.3424
                           0.3722 96.5480 0.920 0.359858
## group2:stage3 0.4927
                                         1.265 0.208743
                           0.3894 97.3911
## group2:stage4 0.7411
                           0.3313 96.5682 2.237 0.027569 *
## group2:stage5
                0.3342
                          0.3218 95.7638 1.039 0.301570
                0.7345
                          0.3314 96.9035
## group2:stage6
                                          2.217 0.028990 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Correlation of Fixed Effects:
             (Intr) group2 stage2 stage3 stage4 stage5 stage6 grp2:2 grp2:3
## group2
             -0.787
## stage2
             -0.455 0.359
## stage3
             -0.490 0.386 0.392
## stage4
             -0.565 0.445 0.483 0.493
## stage5
             -0.538  0.423  0.450  0.483  0.561
## stage6
             -0.528  0.415  0.409  0.497  0.538  0.512
## group2:stg2 0.355 -0.433 -0.779 -0.305 -0.376 -0.351 -0.319
## group2:stg3 0.356 -0.428 -0.285 -0.727 -0.359 -0.351 -0.361 0.354
## group2:stg4 0.422 -0.507 -0.361 -0.369 -0.747 -0.419 -0.402
                                                           0.429 0.415
## group2:stg5 0.430 -0.519 -0.360 -0.387 -0.448 -0.800 -0.409 0.433 0.430
## group2:stg6 0.425 -0.510 -0.329 -0.400 -0.433 -0.412 -0.805 0.403 0.437
             grp2:4 grp2:5
## group2
## stage2
## stage3
## stage4
## stage5
## stage6
## group2:stg2
## group2:stg3
## group2:stg4
## group2:stg5 0.515
## group2:stg6 0.493 0.506
```

model, random intercept + random slope per factor

```
This model is already unidentifiable

model5 <- lmer(dv ~ group*stage+(1+group|id)+(1+stage|id),df3)

## Error: number of observations (=141) <= number of random effects (=216) for term (1 + stage | id); to
```

maximum model?random slope for interaction Also unidentifiable

```
model6 <- lmer(dv ~ group*stage+(group*stage|id),df3)</pre>
```

Error: number of observations (=141) <= number of random effects (=432) for term (group * stage | id