

## Specifying Contrasts for linear model

At the moment my motivation for making this available is so that someone can tell me if I'm doing what I want to do.

Get `tidyverse` for data wrangling and `lme4` for modelling and load some data that I prepared earlier.

```
library(tidyverse)
library(lme4)

load("rti_narrative_data.Rda")
```

This loads a data frame called `Df` that looks like this:

##	Subno	T1234	Condition	score	cond_by_task
## 2	: 4	T1:161	comp:500	Min. : 0.000	T1.comp:125
## 3	: 4	T2:161	int :144	1st Qu.: 4.000	T2.comp:125
## 5	: 4	T3:161		Median : 6.000	T3.comp:125
## 7	: 4	T4:161		Mean : 5.571	T4.comp:125
## 8	: 4			3rd Qu.: 8.000	T1.int : 36
## 9	: 4			Max. :10.000	T2.int : 36
## (Other):620				NA's :5	(Other): 72

T1234 is a factor representing tests at four different time points.

Condition comprises two groups, `comp` and `int`

I am constructing comparisons as follows.

Create a “flat” single factor representing all cells in the design.

```
Df <- Df %>%
  mutate(cond_by_task = interaction(T1234,Condition)) %>%
  filter(complete.cases(.))
summary(Df$cond_by_task)
```

##	T1.comp	T2.comp	T3.comp	T4.comp	T1.int	T2.int	T3.int	T4.int
##	125	125	125	121	36	36	36	35

Then define some contrasts:

Main effect of group

```
cond <- cbind(c(-1,-1,-1,-1,1,1,1,1))
colnames(cond) <- c('group')
```

Main effects of time with separate contrasts giving slope, averaged across groups, between T1 and T2, T2 and T3, and T3 and T4.

```
time <- cbind(rep(c(-1,1,0,0),2),
              rep(c(0,-1,1,0),2),
              rep(c(0,0,-1,1),2))
colnames(time) <- c('T12','T23','T34')
```

Interaction effects. This asks whether there the slopes for the two groups differ, looking separately at slopes between T1 and T2, T2 and T3, and T3 and T4.

```
inter <- cbind(c(1,-1,0,0,-1,1,0,0),
               c(0,1,-1,0,0,-1,1,0),
               c(0,0,1,-1,0,0,-1,1))
colnames(inter) <- c('T12:group','T23:group','T34:group')
```

Then put these all together and assign them to the `cond_by_task` factor.

```
conds <- cbind(cond,time,inter)
contrasts(Df$cond_by_task, how.many = ncol(conds)) <- conds
contrasts(Df$cond_by_task)
```

```
##           group  T12  T23  T34  T12:group  T23:group  T34:group
## T1.comp      -1  -1   0   0           1           0           0
## T2.comp      -1   1  -1   0          -1           1           0
## T3.comp      -1   0   1  -1           0          -1           1
## T4.comp      -1   0   0   1           0           0          -1
## T1.int        1  -1   0   0          -1           0           0
## T2.int        1   1  -1   0           1          -1           0
## T3.int        1   0   1  -1           0           1          -1
## T4.int        1   0   0   1           0           0           1
```

You don't need an intercept because `lmer` gives you it for free (and won't allow you to use your own contrast but specify no intercept in the model).

Then run the model and get the coefficients.

```
m <- lmer(score ~ cond_by_task + (1|Subno),
          data = Df)

summary(m)$coef
```

```
##           Estimate Std. Error  t value
## (Intercept)      5.0411258   0.1375121 36.659499
## cond_by_task group    -0.9775574   0.1375121 -7.108881
## cond_by_task T12      2.8194592   0.1346592 20.937743
## cond_by_task T23      3.3848072   0.1559507 21.704337
## cond_by_task T34      1.7137108   0.1362447 12.578185
## cond_by_task T12:group  0.1607760   0.1346592  1.193947
## cond_by_task T23:group  0.6034408   0.1559507  3.869433
## cond_by_task T34:group  0.3936612   0.1362447  2.889370
```

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So, the question is: Is this set of contrasts giving me what I've claimed it's giving me? If not, what am I doing wrong?

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One way of answering this question - of checking that the contrasts that I've specified really do test the statistical model that I think they are testing, is to separate out the contrasts and get model fits for each, then look to see if the fits for these models make sense. I'm going to do this without the random by-subject effect, because this makes things clearer (because subjects are only in one condition, so if you exclude a condition factor but include by-subject effects, then this effectively reintroduces the condition effect).

So, fit some models and use them to predict scores, and save these into the data frame: `Df$fit = fitted(mymodel)`.

```
# grand mean
m <- lm(score ~ 1, data = Df)
Df$fitGrandMean = fitted(m)

# just main effect of condition (group)
conds <- cond
contrasts(Df$cond_by_task, how.many = ncol(conds)) = conds
```

```

m = lm(score ~ cond_by_task, data = Df)
Df$fitCond = fitted(m)

# just main effect of time
conds <- time
contrasts(Df$cond_by_task, how.many = ncol(conds)) = conds
m = lm(score ~ cond_by_task, data = Df)
Df$fitTime = fitted(m)

# main effects of cond and time
conds = cbind(time,cond)
contrasts(Df$cond_by_task, how.many = ncol(conds)) = conds
m = lm(score ~ cond_by_task, data = Df)
Df$fitTime_Cond = fitted(m)

# just interaction
conds = inter
contrasts(Df$cond_by_task, how.many = ncol(conds)) = conds
m = lm(score ~ cond_by_task, data = Df)
Df$fitTime_X_Cond = fitted(m)

# main effects and interaction
conds = cbind(time,cond,inter)
contrasts(Df$cond_by_task, how.many = ncol(conds)) = conds
m = lm(score ~ cond_by_task, data = Df)
Df$fitTime_Cond_Time_X_Cond = fitted(m)

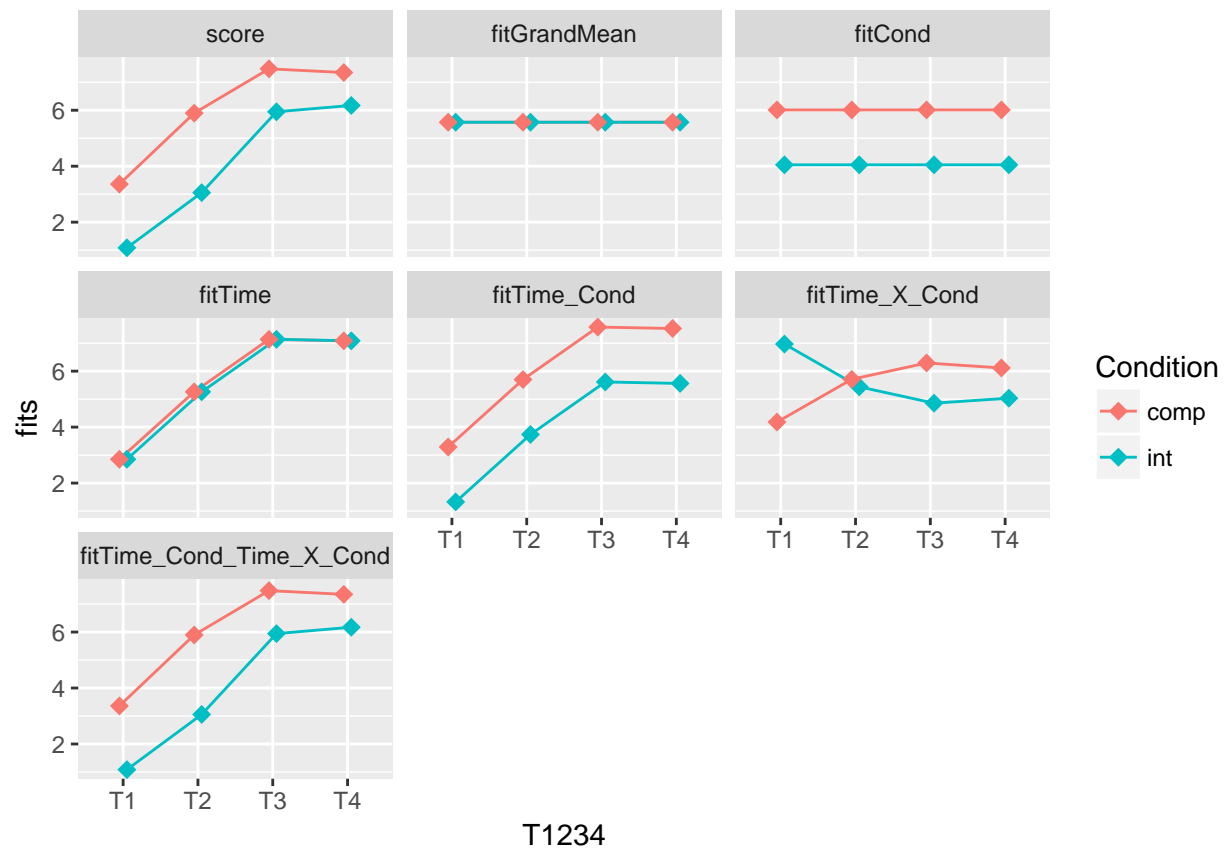
```

And then have a look at them.

```

pd = position_dodge(.2)
Df %>% gather(model, fits, -Subno, -T1234, -Condition, -cond_by_task) %>%
  mutate(model = factor(model,levels = unique(model))) %>%
  ggplot(aes(y = fits, x = T1234, colour = Condition)) +
  geom_line(aes(group = Condition),
            stat = 'summary', fun.y = "mean", na.rm = T,
            position = pd) +
  geom_point(stat = 'summary', fun.y = "mean", na.rm = T,
            size = 3, shape = 18,
            position = pd) +
  facet_wrap(~model)

```



Score is the raw score. So, this suggests that the contrasts are doing what they are meant to be doing.