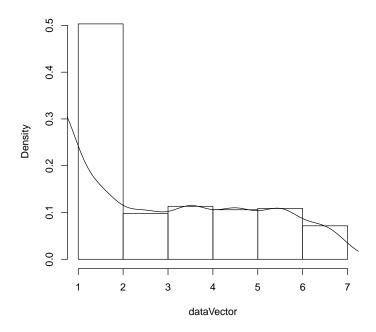
1 Raw data acquisition

```
> data = read.csv("study3-emotions-data.csv")
> dataSelf<-data[,c(1,2:19)]
> dataSelf1<-data[,c(1,20:37)]
> dataSelf2<-data[,c(1,38:55)]

    Density distribution of the scores:
> dataVector <- na.omit(as.vector(t(dataSelf1[,-1])))
> hist(dataVector,probability = T, breaks = "FD")
> lines(density(dataVector - 0.5))
```

Histogram of dataVector



This is used to simulate random scoring with a similar distribution. This simulation has been performed with the Python script random-scores-simulator.py using the following formula for the distribution:

```
random() < 0.5 ? 1 : randint(2,7).

> # 8999 samples gradings, generated by random-scores-simulator.pp
> randomdata = read.csv("random-data.csv")
> randomDataSelf<-randomdata[,c(1,2:19)]
> randomDataSelf1<-randomdata[,c(1,20:37)]
> randomDataSelf2<-randomdata[,c(1,38:55)]
>
```

```
> # check the density distribution of the random data
> #randomdataVector <- na.omit(as.vector(t(randomDataSelf1[,-1])))
> #hist(randomdataVector,probability = T, breaks = "FD")
> #lines(density(randomdataVector - 0.5))
```

2 Computation of the degree of accuracy of partners' models

```
 \mathcal{M}^{\circ}(B,A) \text{ and } \mathcal{M}^{\circ}(C,A). 
 > d1 < -abs(dataSelf-dataSelf1) 
 > d1 * meanDiff < -rowMeans(d1[,c(2:19)],na.rm=T) 
 > d2 < -abs(dataSelf-dataSelf2) 
 > d2 * meanDiff < -rowMeans(d2[,c(2:19)],na.rm=T) 
 > d < -as.data.frame(cbind(d1 * meanDiff,d2 * meanDiff)) 
 > names(d) < -c("scoreB","scoreC") 
 > \# \text{ same thing for the random data} 
 > randomd1 < -abs(randomDataSelf-randomDataSelf1) 
 > randomd1 * meanDiff < -rowMeans(randomd1[,c(2:19)],na.rm=T) 
 > randomd2 < -abs(randomDataSelf-randomDataSelf2) 
 > randomd2 * meanDiff < -rowMeans(randomd2[,c(2:19)],na.rm=T) 
 > randomd < -as.data.frame(cbind(randomd1 * meanDiff,randomd2 * meanDiff))
```

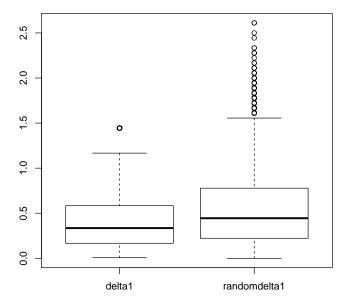
3 Video/Audio conditions

Subjects in the audio-only condition were (marginally) significantly better at modelling than the subjects in the video + audio condition.

4 Computation of Δ_1

```
> delta1<-c(1:60)
> for(i in seq(1, 60, by = 3)) {
    delta1[i] <- abs(d[i,1]-d[i+1,1])
    delta1[i+1] < -abs(d[i+2,1]-d[i,2])
    delta1[i+2] < -abs(d[i+1,2]-d[i+2,2])
+ }
   We test \Delta_1 against randomly attributed scores (following the same distri-
bution as in the actual data) to check if we observe any symmetrical modelling
effects (that would translate into significantly lower \Delta_1).
> randomdelta1 <- c(1:8999)
> for(i in seq(1, 8999, by = 3)) {
    randomdelta1[i] <- abs(randomd[i,1] -randomd[i+1,1])</pre>
    randomdelta1[i+1] <- abs(randomd[i+2,1] -randomd[i,2])</pre>
    randomdelta1[i+2] <- abs(randomd[i+1,2] -randomd[i+2,2])</pre>
+ }
> boxplot(cbind(delta1, randomdelta1))
> # are variances equal?
> var.test(as.vector(delta1), as.vector(randomdelta1))
        F test to compare two variances
data: as.vector(delta1) and as.vector(randomdelta1)
F = 0.5863, num df = 59, denom df = 8997, p-value = 0.009486
alternative hypothesis: true ratio of variances is not equal to 1
95 percent confidence interval:
0.4206348 0.8729522
sample estimates:
ratio of variances
         0.5862648
> t.test(delta1, randomdelta1,var.equal = F)
        Welch Two Sample t-test
data: delta1 and randomdelta1
t = -3.3032, df = 60.35, p-value = 0.00161
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -0.21396938 -0.05257898
sample estimates:
```

mean of x mean of y 0.4032339 0.5365081

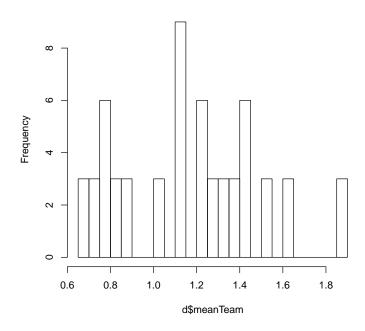


The symmetry hypothesis tells that good modelling should be symmetrical, but does not say much about bad modelling.

hence, the effect should be even stronger if we check the average Δ_1 of the best modellers against random data.

To do so, we first compute the average grading score per triad.

Histogram of d\$meanTeam



A dip test confirms that the distribution is not uni-modal:

> diptest::dip.test(d\$meanTeam)

Hartigans' dip test for unimodality / multimodality

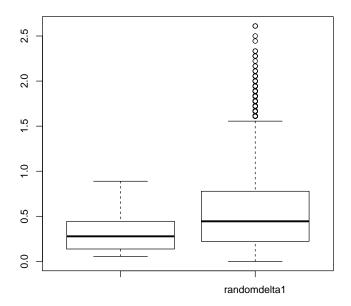
```
data: d$meanTeam
D = 0.0656, p-value = 0.0444
```

alternative hypothesis: non-unimodal, i.e., at least bimodal

We use a K-Mean clustering to know where to cut our group of modellers, provinding an initial estimate of the clusters' centers:

- > centers <- c(1:2)
- > centers[1] = 0.8
- > centers[2] = 1.2
- > clusters<-kmeans(d\$meanTeam,centers)</pre>
- > # alternatively, let kmeans find the centers. But this may change between runs due to the
- > #clusters<-kmeans(d\$meanTeam,2)</pre>
- >
- > clusters\$centers
 - [,1]
- 1 0.8270308
- 2 1.3575446

```
> top <- max(d$meanTeam[clusters$cluster == 1])</pre>
> bottom <- min(d$meanTeam[clusters$cluster == 2])</pre>
> d$teamRating<-ifelse(d$meanTeam<=(top+bottom)/2,"good","bad")</pre>
> # or using a simple median cut...
> #d$teamRating<-ifelse(d$meanTeam<=median(d$meanTeam),"good","bad")
   We finally re-run a t-test on this subset of the modellers:
> boxplot(cbind(delta1[d$teamRating=="good"], randomdelta1))
> t.test(delta1[d$teamRating=="good"], randomdelta1,var.equal = F)
        Welch Two Sample t-test
data: delta1[d$teamRating == "good"] and randomdelta1
t = -4.554, df = 20.296, p-value = 0.0001866
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -0.3322192 -0.1236169
sample estimates:
mean of x mean of y
0.3085901 0.5365081
```

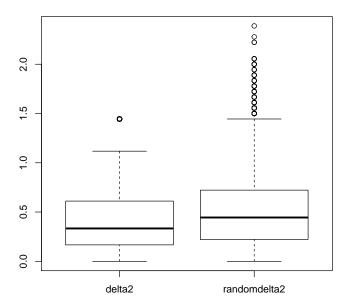


This confirms the symmetry hypothesis: the difference of scores for good modellers with random data is even stronger.

5 Computation of Δ_2

```
> delta2<-c(1:60)
> for(i in seq(1, 60, by = 3)) {
+    delta2[i]<-abs(d[i,1]-d[i,2])
+    delta2[i+1]<-abs(d[i+1,1]-d[i+1,2])
+    delta2[i+2]<-abs(d[i+2,1]-d[i+2,2])
+ }
> sd(delta2)
[1] 0.3407969
```

As previously, we test Δ_2 against randomly attributed scores to check if we observe any similar modelling behaviours between two persons judging the same third one (which would translate into significantly lower Δ_2 compared to chance):



In average, Δ_2 is not significantly smaller than chance: this does not support the hypothesis of $\mathcal{M}^{\circ}(A, C)$ and $\mathcal{M}^{\circ}(B, C)$ being correlated, ie modelling accuracy would either comes for the overall quality of the interaction, or C fosters good models by making his emotional state easily readable.

6 Computation of Δ_3

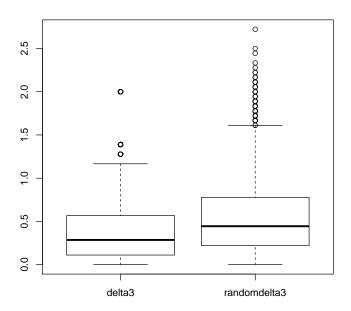
```
> delta3<-c(1:60)
> for(i in seq(1, 60, by = 3)) {
+    delta3[i]<-abs(d[i+1,1]-d[i+2,1])
+    delta3[i+1]<-abs(d[i,1]-d[i+2,2])
+    delta3[i+2]<-abs(d[i,2]-d[i+1,2])
+ }
> sd(delta3)
[1] 0.4074252
```

As previously, we test Δ_3 against randomly attributed scores to check if we observe any similar modelling behaviours between two persons judging the same third one (which would translate into significantly lower Δ_3 compared to chance):

```
> boxplot(cbind(delta3, randomdelta3))
> t.test(delta3, randomdelta3)
```

Welch Two Sample t-test

data: delta3 and randomdelta3 t = -2.625, df = 59.788, p-value = 0.01098 alternative hypothesis: true difference in means is not equal to 0 95 percent confidence interval: -0.24410153 -0.03296222 sample estimates: mean of x mean of y 0.3982911 0.5368230



We obtain a significantly lower Δ_3 , which supports the hypothesis of $\mathcal{M}^{\circ}(C, A)$ and $\mathcal{M}^{\circ}(C, B)$ being correlated: it supports the hypothesis that some persons are better modellers than others.