Synchrony in Psychotherapy, example with F1044 patient data

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Aim and Hypothesis

The aim of this project is to evaluate if it is possible to detect automatic signals that could predict the outcomes of a familial psychotherapy. Here are the analysis of the F1044 subject and his family (2 parents). Is there synchrony signals between him self, his parents and the therapist? His there synchrony signal between his parents and the therapist?

Lists

Functions list

MeanMotionByTime

Function that takes raw motion history data and make mean on a given interval. Intervals don't overlap, so the frequency of the data change (from 25 frames by seconde to 25 frames/interval by second).

Arguments:

- subject: Subject studied (patient, mother, father or therapist)
 indexOfvideos: List of videos studied (element eg 3 or list eg 1:3 or c(1,2,4))
- interval: number of frames in the studied interval
- data : data frame where there is data

```
MeanMotionByTime <- function(subject, indexOfvideos=1:NumberOfvideos, interval, data){
    x <- c()
    for (file in indexlist[indexOfvideos]){
        dataVector <- data[which(data$file==file), subject]
        ## with ceiling : superior limit of the round
        IntervalNumbersVideo <- ceiling(length(dataVector)/interval)
        for (i in 1:IntervalNumbersVideo){
            borneinf<- 1+(i-1)*interval
            bornesup <-i*interval
            dataVector[borneinf:bornesup]
            mean <- mean(dataVectorInterval, na.rm=TRUE)
            x <- c(x, mean)}}
    return (x)}</pre>
```

Slidinginterval

Function that takes raw motion history data and make mean on a given interval. The interval overlap, so the frequency of the data don't change. It stays at 25 frames/s.

Arguments:

- subject : subject studied (patient, mother, father or therapist)
- indexOfvideos : list of videos studied (element eg. 3 or list eg 1:3 or c(1,2,4))
- interval : number of frames in the studied interval
- data: data frame where there is data

```
SlidingInterval <- function(subject, indexOfvideos=1:NumberOfvideos, interval, data)
  {x <- c()
  for (file in indexlist[indexOfvideos]){
     dataVector <- data[which(data$file==file), subject]
     NBofAnalysedFrames <- length(dataVector)-interval+1
        for (i in 1:NBofAnalysedFrames){
          borneinf<- (i)
          bornesup <-(interval-1+i)
          dataVectorInterval <- dataVector[borneinf:bornesup]
          mean <- mean(dataVectorInterval, na.rm=TRUE)
          x <- c(x, mean)}}
return (x)}</pre>
```

Nomenclature

F1044 is the name of the subject studied (called pa for patient) He has a mother (mo) a father (fa) helped by a therapist (th) When a variable is referring to several participants, it is organised in alphabetical order separated by underscores, eg. SSI_fa_mo refers to the synchrony index between the father and the mother. SSI_mo_fa doesn't exist. This familly had several consultations with the psychotherapist. Some of them were video recorded. These videos are names with the name of the subject + an index letter. They can subdivised after that with numbers (eg F1044C).

File lists

Participants list

```
## [1] "/Users/Ofix/Documents/Fac/internat/Recherche/projets/synchro/synchroData"
## [1] "father" "mother" "patient" "therapist"
```

Presentation of the data

This data comes from the INCANT study. The F1044 subject which as dependence with cannabis, had a familial therapy (TAU and MDFT). The goal of this pilot study is to extract meaningfull motion history from 17 videos called F1044C, F1044D1... to F1044R2 (see listsindexlist)

```
str(data)
```

```
'data.frame':
                    477258 obs. of 7 variables:
   $ frame
               : int
                     1 2 3 4 5 6 7 8 9 10 ...
##
               : num
                      0.01996 0.00915 0.01355 0.01787 0.01758 ...
   $ father
   $ mother
               : num 1.82e-05 1.82e-05 3.64e-05 1.82e-05 9.09e-05 ...
   $ patient : num NA ...
##
   $ therapist: num  0.00162 0.00506 0.00349 0.00223 0.00249 ...
               : Factor w/ 17 levels "F1044C.VOB", "F1044D1.VOB",..: 1 1 1 1 1 1 1 1 1 1 ...
##
   $ file
   $ timeMin : num  0.000667 0.001333 0.002 0.002667 0.003333 ...
```

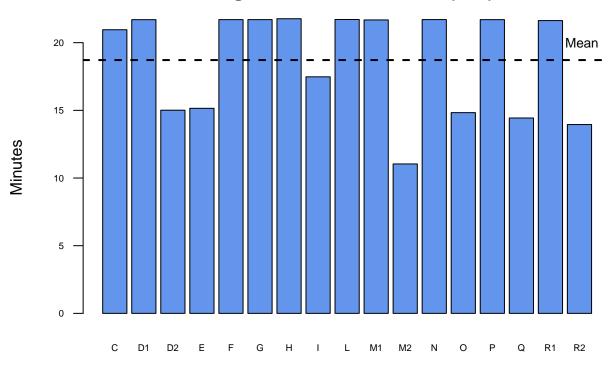
summary(data)

```
##
        frame
                         father
                                            mother
                                                            patient
##
    Min.
           :
                     Min.
                             :0.00
                                       Min.
                                               :0.00
                                                        Min.
                                                                :0.00
                 1
##
    1st Qu.: 7019
                     1st Qu.:0.00
                                       1st Qu.:0.00
                                                        1st Qu.:0.00
    Median :14038
                     Median:0.00
                                       Median:0.00
                                                        Median:0.00
##
                             :0.00
                                               :0.00
                                                                :0.01
##
    Mean
           :14576
                     Mean
                                       Mean
                                                        Mean
##
    3rd Qu.:21364
                     3rd Qu.:0.00
                                       3rd Qu.:0.00
                                                        3rd Qu.:0.01
##
    Max.
           :32656
                     Max.
                             :0.19
                                       Max.
                                               :0.49
                                                        Max.
                                                                :0.54
##
                     NA's
                             :265686
                                       NA's
                                               :91545
                                                        NA's
                                                                :189317
##
      therapist
                              file
                                              timeMin
##
           :0.0
                     F1044H.VOB: 32656
                                                  : 0.000667
    Min.
                                          Min.
##
    1st Qu.:0.0
                     F1044L.VOB: 32570
                                          1st Qu.: 4.679333
##
    Median:0.0
                     F1044N.VOB: 32562
                                          Median : 9.358333
##
    Mean
           :0.0
                     F1044G.VOB: 32556
                                                  : 9.717052
                     F1044F.VOB: 32555
                                          3rd Qu.:14.242667
##
    3rd Qu.:0.0
##
    Max.
            :0.8
                     F1044P.VOB: 32554
                                          Max.
                                                  :21.770667
    NA's
##
            :77972
                     (Other)
                                :281805
```

The timeMin is calculated with a frame rate of 25/sec.

Length of the videos in minutes

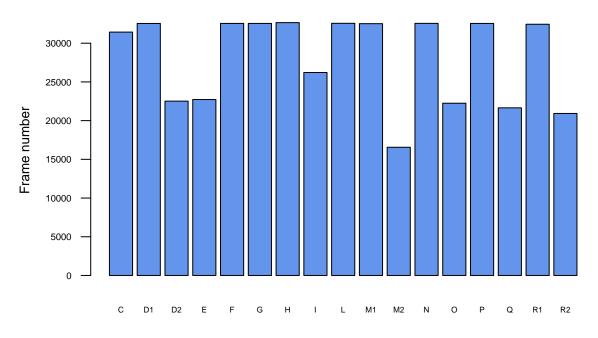
Length in each F1044 video (min)



Video Name

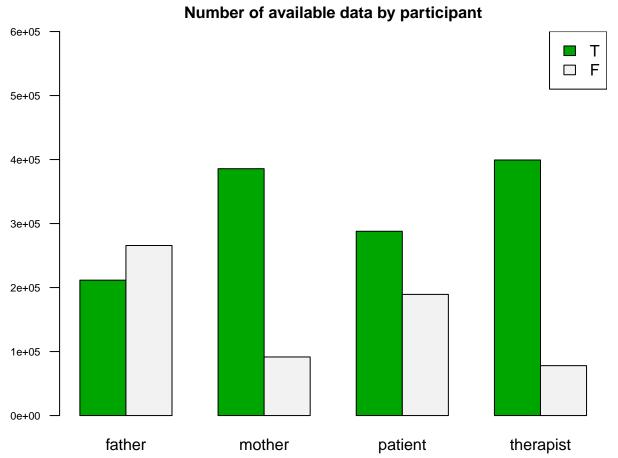
Length of the videos in number of frames

Number of frames in each F1044 video



Video Name

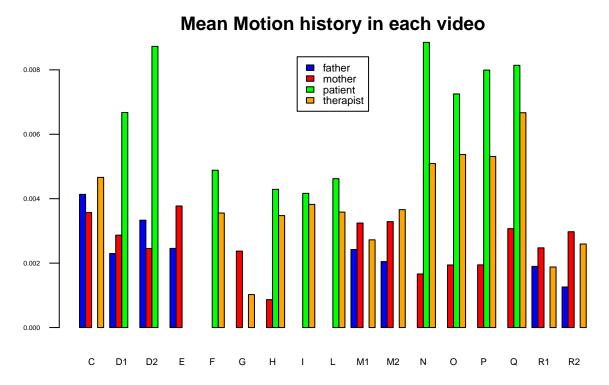
Number of Available (True) and Not Available (False) data for each participant



Some participants are not filmed (eg therapist) or don't come in the sessions. Mother and therapist are the more often present participants.

Global Motion history

Mean Motion history by video by participant



Raw data and mean of Motion History on sliding and non overlapping intervals on F1044C video

F1044C video

It is the first video of F10044C. The father, mother and the rapist are present. The patient is absent. ## Raw data

```
rawdatafather <- data[which(data$file=="F1044C.VOB"),]$father
rawdataMother <- data[which(data$file=="F1044C.VOB"),]$mother
rawdataTherapist <- data[which(data$file=="F1044C.VOB"),]$therapist</pre>
summary(rawdatafather)
##
             1st Qu.
                       Median
                                   Mean 3rd Qu.
                                                               NA's
## 0.000000 0.000000 0.000196 0.004135 0.003488 0.092340
                                                                 10
summary(rawdataMother)
##
       Min. 1st Qu.
                       Median
                                   Mean 3rd Qu.
                                                               NA's
## 0.000000 0.000036 0.000127 0.003570 0.002200 0.159600
                                                                 10
```

```
summary(rawdataTherapist)
##
       Min. 1st Qu.
                       Median
                                   Mean 3rd Qu.
                                                     Max.
                                                              NA's
## 0.001179 0.002750 0.003405 0.004662 0.004234 0.236000
                                                                 10
Sliding interval
## REMINDER:
\# SlidingInterval <- function(subject, indexOfvideos=1:NumberOfvideos, interval, data) with :
# subject : subject studied (patient, mother, father or therapist)
# indexOfvideos : list of videos studied (element eq. 3 or list eq 1:3 or c(1,2,4))
# interval : number of frames in the studied interval
# data : data frame where there is data
slidedfather <- SlidingInterval("father", 1 , 5, data)</pre>
slidedmother <- SlidingInterval("mother", 1 , 5, data)</pre>
slidedtherapist <- SlidingInterval("therapist", 1 , 5, data)</pre>
slidedpatient <- SlidingInterval("patient", 1 , 5, data)</pre>
summary(slidedfather)
                                                              NA's
       Min. 1st Qu.
                       Median
                                  Mean 3rd Qu.
                                                     Max.
## 0.000000 0.000005 0.000335 0.004139 0.003691 0.091450
summary(slidedmother)
       Min. 1st Qu.
                       Median
                                   Mean 3rd Qu.
                                                              NA's
                                                     Max.
## 0.000000 0.000036 0.000156 0.003574 0.002520 0.145900
                                                                  6
summary(slidedpatient)
      Min. 1st Qu. Median
                              Mean 3rd Qu.
##
                                               Max.
                                                       NA's
##
       NA
                NA
                        NA
                               \mathtt{NaN}
                                         NA
                                                      31431
                                                 NA
summary(slidedtherapist)
       Min. 1st Qu.
                       Median
                                  Mean 3rd Qu.
                                                               NA's
## 0.001702 0.003029 0.003352 0.004672 0.003955 0.219600
par(mar=c(3,3,2,2))
boxplot(slidedfather, slidedmother, slidedtherapist,
        col=colOrderList[c(1,2,4)],
        names=ParticipantsList[c(1,2,4)],
       main= "Box plot of motion history sliding interval on F1044C video", las=1)
par(mar=c(1,0.5,0.5,1))
legend("topleft", ParticipantsList[c(1,2,4)], fill=colOrderList, cex=0.7)
```

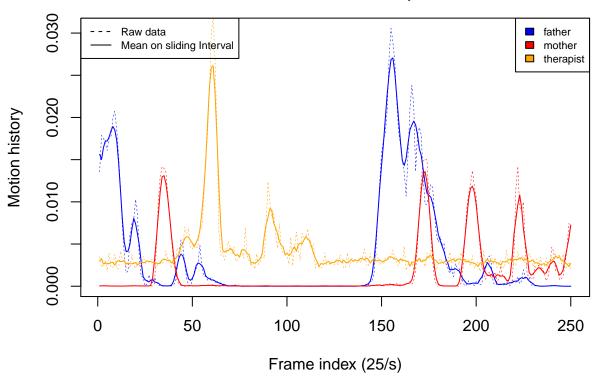
Non overlapping interval

```
fatherFive<- MeanMotionByTime("father", indexOfvideos=1, interval=5, data)</pre>
motherFive <- MeanMotionByTime("mother", indexOfvideos=1, interval=5, data)</pre>
therapistFive <- MeanMotionByTime("therapist", indexOfvideos=1, interval=5, data)
summary(fatherFive)
##
               1st Qu.
                          Median
                                              3rd Qu.
                                                                      NA's
        Min.
                                       Mean
                                                            Max.
## 0.0000000 0.0000049 0.0003305 0.0041350 0.0036840 0.0883000
summary(motherFive)
               1st Qu.
                          Median
                                       Mean
                                              3rd Qu.
                                                                      NA's
## 0.0000000 0.0000364 0.0001564 0.0035700 0.0025310 0.1459000
                                                                         2
summary(therapistFive)
       Min. 1st Qu.
                       Median
                                   Mean 3rd Qu.
                                                               NA's
                                                      Max.
## 0.001807 0.003029 0.003352 0.004662 0.003972 0.219600
```

Focus on the motion history of the first 20 seconds of the first video(C)

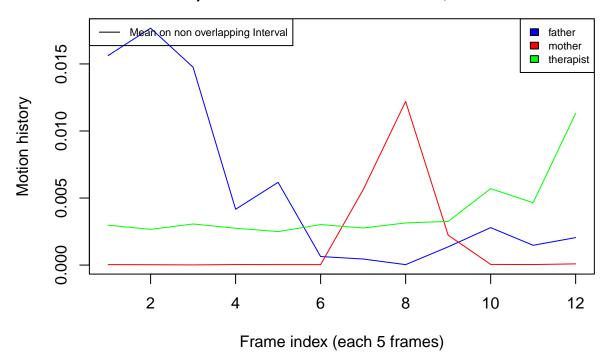
Sliding interval function on a 5 frames interval

Mean motion history (Sliding 5 frames interval) for father on F1044C video, 10 seconds



Non overlapping interval function on a 5 frames interval

Mean Motion history (non overlapping 5 frames intervals) for father on F1044C video, first 10 seconds



Non overlapping interval function on a 5 frames interval with shifting of therapist (substraction of min value of therapist)

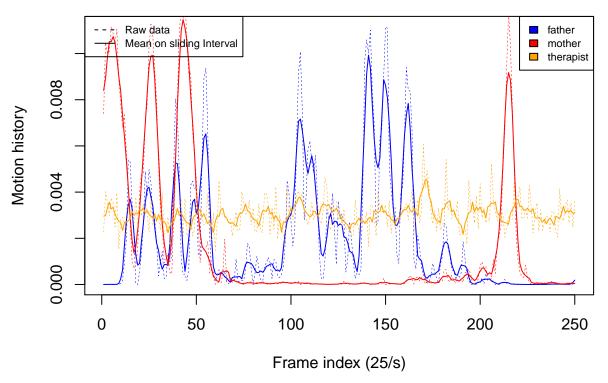
Motion history of the father during 10-20 seconds of the first video(C)

Non overlapping interval function on a 5 frames interval

```
par(mar=c(4,4,4,2))
plot(1:250, data$father[253:502], main="Mean motion history (Sliding 5 frames
    interval) for father on F1044C video, 10-20 seconds", xlab="Frame index (25/s)",
    ylab="Motion history", col="blue", type="l", lty=2, lwd=0.5)
lines(slidedfather[251:500], col="blue", lty=1)
lines(data$mother[253:502], col="red", lty=2, lwd=0.5)
lines(slidedmother[251:500], col="red", lty=1)
lines(data$therapist[253:502], col="orange", lty=2, lwd=0.5)
```

```
lines(slidedtherapist[251:500], col="orange", lty=1)
legend("topleft", c("Raw data", "Mean on sliding Interval") , lty=c(2, 1), cex=0.7)
legend("topright", ParticipantsList[c(1,2,4)], fill=colOrderList[c(1,2,4)], cex=0.7)
```

Mean motion history (Sliding 5 frames interval) for father on F1044C video, 10–20 seconds

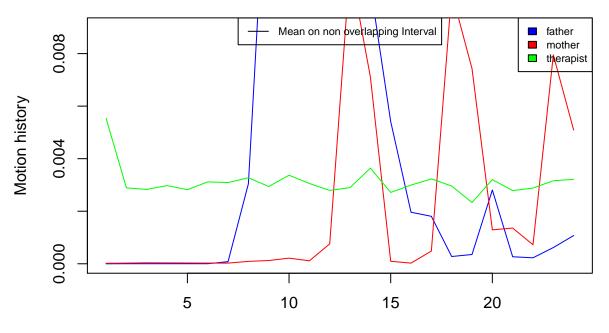


Non overlapping interval function on a 5 frames interval

```
plot (1:24, fatherFive[23:46], type="1", col="blue",
main="Mean motion history (non overlapping 5 frames intervals) for
father on F1044C video, between 10-20 seconds",
```

```
ylab="Motion history", xlab="Frame index (each 5 frames)", ylim=c(0, 0.009))
lines(motherFive[23:46], col="red", lty=1)
lines(therapistFive[23:46], col="green", lty=1)
legend("top", "Mean on non overlapping Interval" , lty=1, cex=0.7)
legend("topright", ParticipantsList[c(1,2,4)], fill=colOrderList, cex=0.7)
```

Mean motion history (non overlapping 5 frames intervals) for father on F1044C video, between 10–20 seconds



Frame index (each 5 frames)

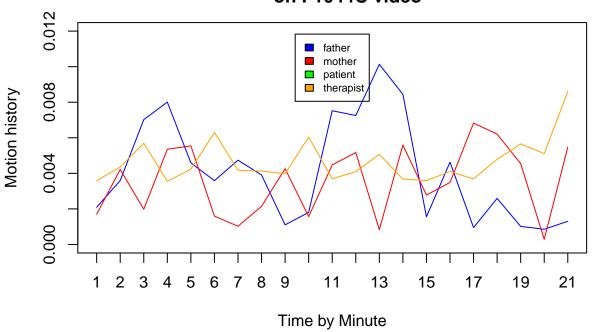
```
### Non overlapping interval function on a 5 frames interval with shifting of therapist (substraction o
plot (1:24, fatherFive[23:46], type="l", col="blue",
main="Mean motion history (non overlapping 5 frames intervals) for
father on F1044C video, between 10-20 seconds,
data therapist shifted",
ylab="Motion history", xlab="Frame index (each 5 frames)", ylim=c(0, 0.009))
lines(motherFive[23:46], col="red", lty=1)
lines(therapistFive[23:46]-min(slidedtherapist), col="green", lty=1)
legend("top", "Mean on non overlapping Interval" , lty=1, cex=0.7)
legend("topright", ParticipantsList[c(1,2,4)], fill=colOrderList, cex=0.7)
```

Mean motion history by minute plots

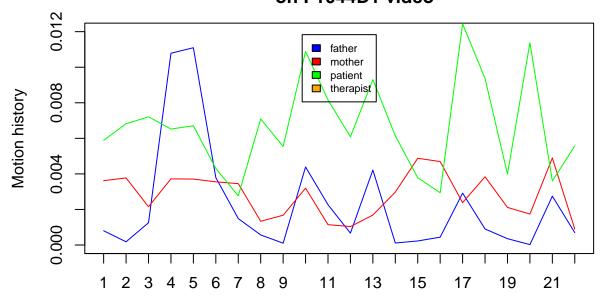
```
for (i in 1:NumberOfvideos){
  fatherMinute<- MeanMotionByTime("father", indexOfvideos=i, interval=1500, data)

MotherMinute<- MeanMotionByTime("mother", indexOfvideos=i, interval=1500, data)</pre>
```

Mean motion history (non overlaping minute intervals) on F1044C video



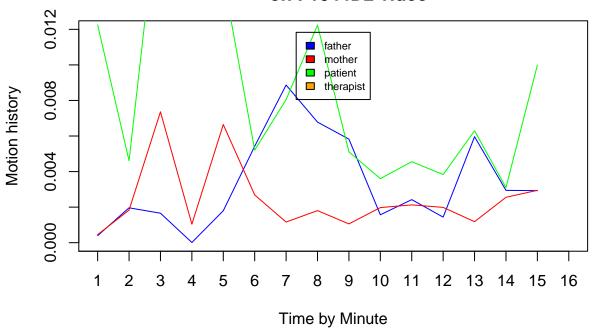
Mean motion history (non overlaping minute intervals) on F1044D1 video



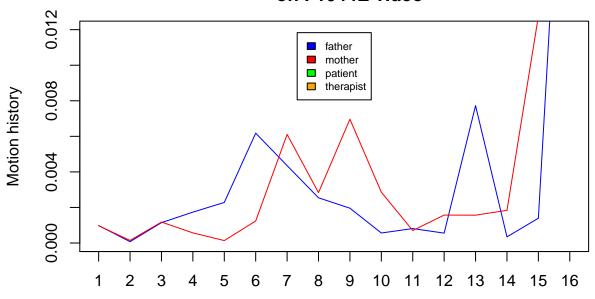
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044D2 video



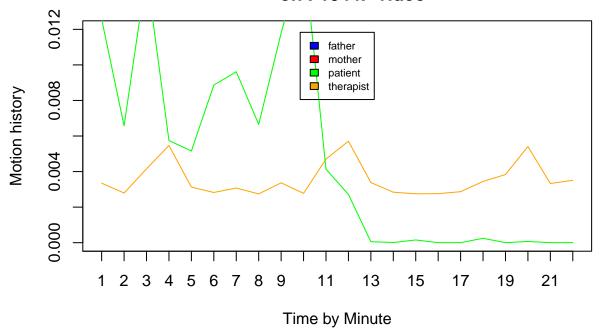
Mean motion history (non overlaping minute intervals) on F1044E video



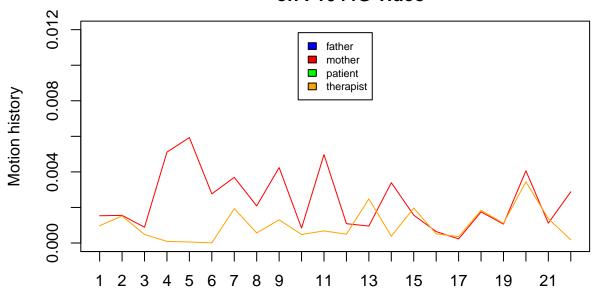
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044F video



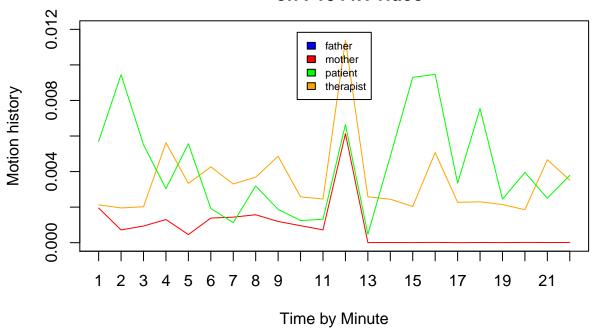
Mean motion history (non overlaping minute intervals) on F1044G video



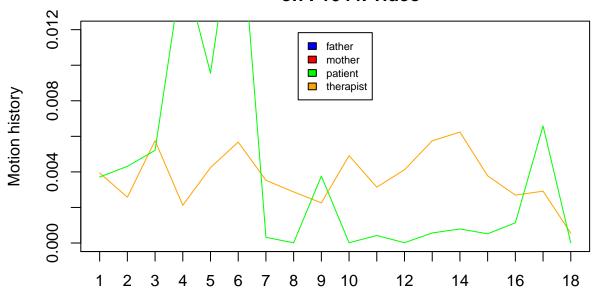
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044H video



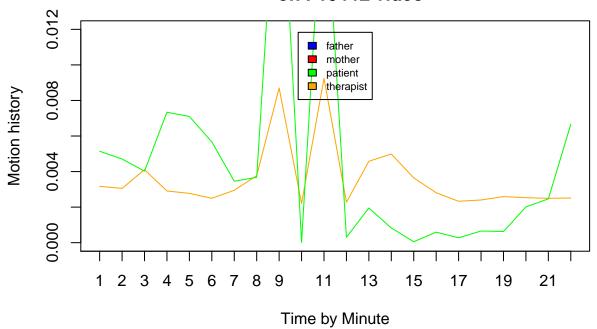
Mean motion history (non overlaping minute intervals) on F1044I video



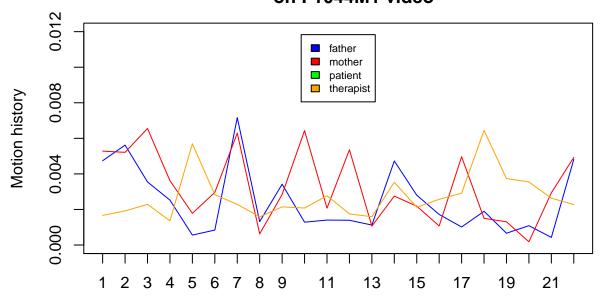
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044L video



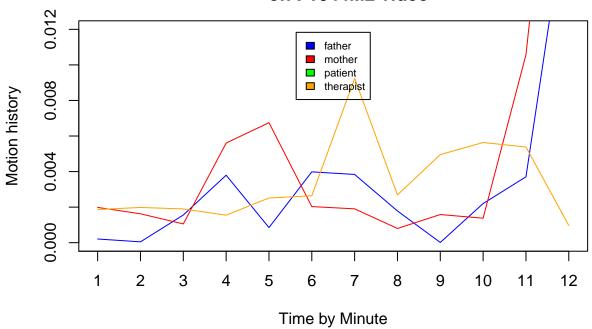
Mean motion history (non overlaping minute intervals) on F1044M1 video



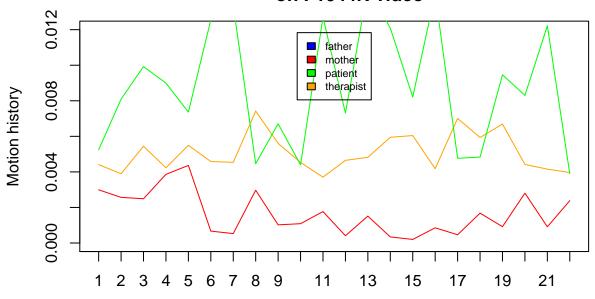
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044M2 video



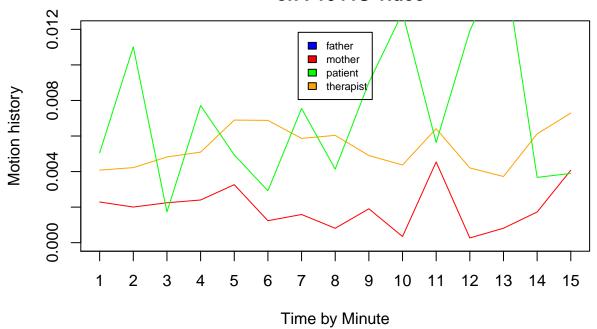
Mean motion history (non overlaping minute intervals) on F1044N video



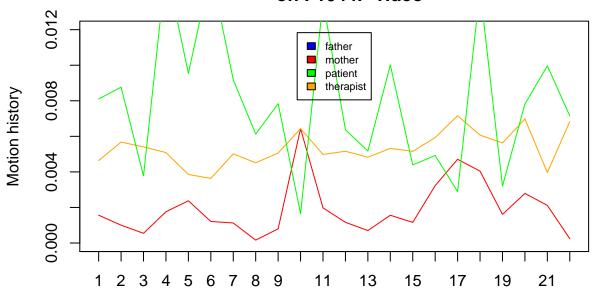
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044O video



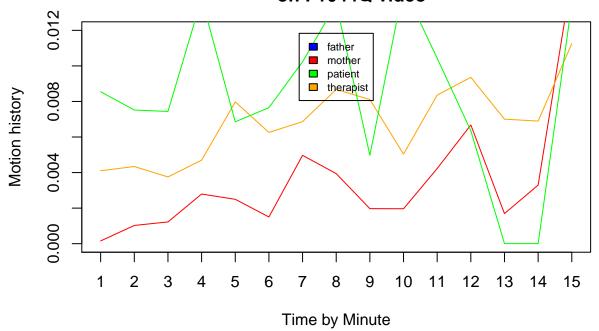
Mean motion history (non overlaping minute intervals) on F1044P video



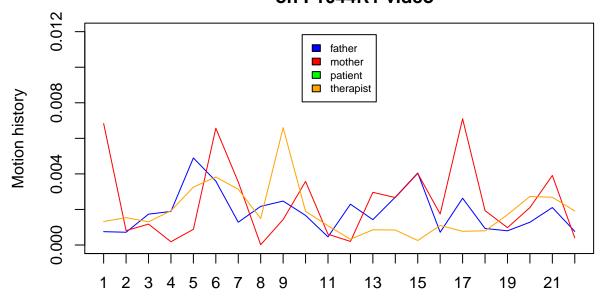
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044Q video



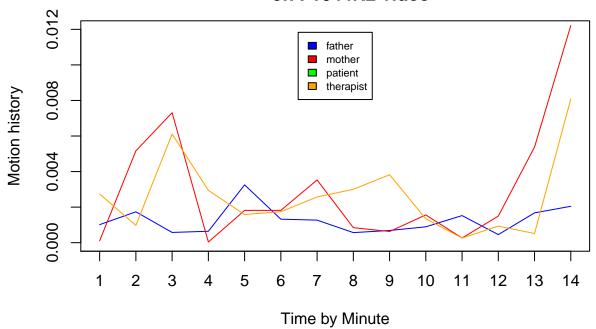
Mean motion history (non overlaping minute intervals) on F1044R1 video



Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044R2 video



```
# slidedfatherwoNA <- slidedfather[which(is.na(slidedfather)==FALSE)]
# slidedmotherwoNA <- slidedmother[which(is.na(slidedmother)==FALSE)]
# slidedtherapistwoNA <- slidedtherapist[which(is.na(slidedtherapist)==FALSE)]
# slidedpatientwoNA <- slidedpatient[which(is.na(slidedpatient)==FALSE)]</pre>
```

Export data in text files

```
## REMINDER:
#SlidingInterval <- function(subject, indexOfvideos=1:NumberOfvideos, interval, data) with :
# subject : subject studied (patient, mother, father or therapist)
# indexOfvideos : list of videos studied (element eq. 3 or list eq 1:3 or c(1,2,4))
# interval : number of frames in the studied interval
# data : data frame where there is data
#index de la vid?eo de 1ere a la length de indexvideo
videoIndex <- 1</pre>
# videoName est le nom de la video actuelle
for (videoName in indexlist){
# Compute slinding interval for each participant
           print(paste("Computing slidedFather", videoName))
           slidedFather <- SlidingInterval("father", videoIndex, 5, data)</pre>
           print(paste("Computing slidedMother", videoName))
           slidedMother <- SlidingInterval("mother", videoIndex, 5, data)</pre>
           print(paste("Computing slidedTherapist", videoName))
           slidedTherapist <- SlidingInterval("therapist", videoIndex, 5, data)</pre>
           print(paste("Computing slidedPatient", videoName))
           slidedPatient <- SlidingInterval("patient", videoIndex, 5, data)</pre>
# create a data frame to store temporarily this data with NA
            slidedVideo <- data.frame(slidedFather, slidedMother, slidedTherapist, slidedPatient)</pre>
##### Creating a data frame if the information is available ##########
           dataFrame <- FALSE
           dfSliding <- data.frame()</pre>
           for (participant in 1:4){
# If the colum is not empty, takes its length and begin a data frame with it
                       if (dataFrame==FALSE){
\#if\ (length(slidedVideo[participant][!is.na(slidedVideo[participant])]) > 0 \ \&\ dataFrame==FALSE)\{
                           dfSliding <- data.frame("video"=rep(indexlist[videoIndex],length(slidedVideo[participant]
                                           dataFrame <- TRUE}</pre>
 \# \ if \ (length(slidedVideo[participant][!is.na(slidedVideo[participant])]) > 0 \ \& \ dataFrame == TRUE) \{ (length(slidedVideo[participant]), (length(slidedVideo[participant]), (length(slidedVideo[participant)), (length(sl
                       if (dataFrame==TRUE){
                                           dfSliding <- cbind(dfSliding, slidedVideo[participant])}}</pre>
           print(str(dfSliding))
CCdfSliding <- complete.cases(dfSliding)</pre>
           emptyLine <- c()</pre>
           for (i in 1:nrow(dfSliding)){
                   dfLine <- dfSliding[i,3:6]</pre>
                   NaLine <- is.na(dfLine)</pre>
                   if (all(NaLine)){
                           emptyLine <- c(emptyLine, i)}}</pre>
           print (emptyLine)
```

```
if (length(emptyLine)>0){
    dfSliding <- dfSliding[-emptyLine,]}

write.csv(dfSliding, paste("/Users/Ofix/Documents/Fac/internat/Recherche/projets/synchro/synchroD
videoIndex <-(videoIndex+1)}</pre>
```

SyncPy utilisation for creating synchrony dataframe

After extracting filtered motion motion history with mean on sliding interval (overlapping interval) of 5 frames

And after puting this data on a CSV file slideddata.csv

We import this data on python Script with panda module Call_S_Estimator.py

This script will compute the synchrony between each dyad of the interaction and of the whole group

It will return a csv file for each video SSIXXXX.csv with XXXX the name of the video (F10044C, F1044D1, etc) that we can import with R with

this following function

```
## [1] "/Users/Ofix/Documents/Fac/internat/Recherche/projets/synchro/synchroData"
## [1] "SSI Files Directory"

## [1] "SS Files List"

## [1] "SSIF1044C.csv" "SSIF1044D1.csv" "SSIF1044D2.csv" "SSIF1044E.csv"
## [5] "SSIF1044F.csv" "SSIF1044G.csv" "SSIF1044H.csv" "SSIF1044I.csv"
## [9] "SSIF1044L.csv" "SSIF1044M1.csv" "SSIF1044M2.csv" "SSIF1044N.csv"
## [13] "SSIF10440.csv" "SSIF1044P.csv" "SSIF1044Q.csv" "SSIF1044R1.csv"
## [17] "SSIF1044R2.csv"
```

Description of SSI data frame

```
str(SSIDataFrames)
## 'data.frame':
                   1900 obs. of 13 variables:
   $ X
                 : int 0 1 2 3 4 5 6 7 8 9 ...
##
## $ Interval
                : int 1 2 3 4 5 6 7 8 9 10 ...
                 : num 0 0.167 0.333 0.5 0.667 ...
## $ Time_min
                 : Factor w/ 17 levels "F1044C.VOB", "F1044D1.VOB", ...: 1 1 1 1 1 1 1 1 1 1 1 ...
## $ video
                : num 3.55e-04 4.27e-03 4.76e-05 1.61e-02 5.53e-02 ...
   $ SSI_fa_mo
## $ SSI_fa_mo_th: num 0.01428 0.00538 0.02604 0.02166 0.02527 ...
## $ SSI fa th : num 0.014475 0.000896 0.01422 0.005114 0.004301 ...
## $ SSI mo th
                 : num 0.018606 0.007802 0.047529 0.031851 0.000883 ...
```

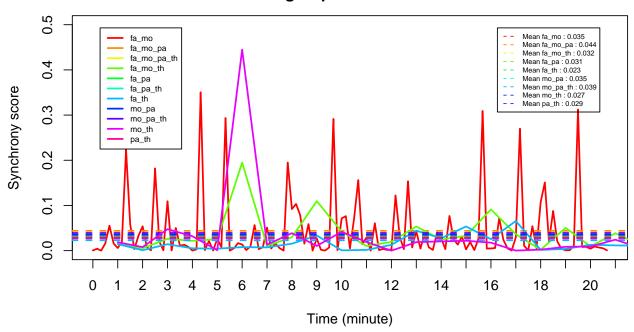
Synchrony scores for each dyad and for the whole group

```
for (i in unique(SSIDataFrames$video))
      {par(mar=c(4,4,4,3), mfrow=c(1,1))}
     plot(SSIDataFrames[which(SSIDataFrames$video==i),]$Time_min,
           SSIDataFrames[which(SSIDataFrames$video==i),]$SSI_fa_mo,
           type="l", col=rainbow(4)[1],
           main=paste("Synchrony scores for each dyad and for \n the whole group in", i, "video"),
           xlab = "Time (minute)", ylab="Synchrony score", lwd=2,
xaxp=c(0,length(SSIDataFrames$Time_min), length(SSIDataFrames$Time_min)), ylim=c(0,0.5))
      abline(h=mean(SSIDataFrames$SSI_fa_mo, na.rm=TRUE), col=rainbow(11)[1], lwd=2, lty=2)
      lines(SSIDataFrames[which(SSIDataFrames$video==i),]$SSI fa mo pa, col=rainbow(11)[2], lwd=2)
      abline(h= mean(SSIDataFrames$SSI_fa_mo_pa, na.rm=TRUE), col=rainbow(11)[2], lwd=2, lty=2)
       lines (SSIDataFrames [which (SSIDataFrames \$video == i),] \$SSI\_fa\_mo\_pa\_th, \ col = rainbow (11) [3], \ lwd = 2) \\
       abline(h= mean(SSIDataFrames$SSI_fa_mo_pa_th, na.rm=TRUE), col=rainbow(11)[3], lwd=2, lty=2)
      lines(SSIDataFrames[which(SSIDataFrames$video==i),]$SSI_fa_mo_th, col=rainbow(11)[4], lwd=2)
      abline(h= mean(SSIDataFrames$SSI_fa_mo_th, na.rm=TRUE), col=rainbow(11)[4], lwd=2, lty=2)
      lines(SSIDataFrames[which(SSIDataFrames$video==i),]$SSI_fa_pa, col=rainbow(11)[5], lwd=2)
      abline(h= mean(SSIDataFrames$SSI_fa_pa, na.rm=TRUE), col=rainbow(11)[5], lwd=2, lty=2)
       lines(SSIDataFrames[which(SSIDataFrames$video==i),]$SSI_fa_pa_th, col=rainbow(11)[6], lwd=2)
#
       abline(h= mean(SSIDataFrames$SSI_fa_pa_th, na.rm=TRUE), col=rainbow(11)[6], lwd=2, lty=2)
      lines(SSIDataFrames[which(SSIDataFrames$video==i),]$SSI_fa_th, col=rainbow(11)[7], lwd=2)
      abline(h= mean(SSIDataFrames$SSI_fa_th, na.rm=TRUE), col=rainbow(11)[7], lwd=2, lty=2)
      lines(SSIDataFrames[which(SSIDataFrames$video==i),]$SSI_mo_pa, col=rainbow(11)[8], lwd=2)
      abline(h= mean(SSIDataFrames$SSI_mo_pa, na.rm=TRUE), col=rainbow(11)[8], lwd=2, lty=2)
     lines(SSIDataFrames[which(SSIDataFrames$video==i),]$SSI_mo_pa_th, col=rainbow(11)[9], lwd=2)
      abline(h= mean(SSIDataFrames$SSI_mo_pa_th, na.rm=TRUE), col=rainbow(11)[9], lwd=2, lty=2)
      lines(SSIDataFrames[which(SSIDataFrames$video==i),]$SSI_mo_th, col=rainbow(11)[10], lwd=2)
      abline(h= mean(SSIDataFrames$SSI_mo_th, na.rm=TRUE), col=rainbow(11)[10], lwd=2, lty=2)
      lines(SSIDataFrames[which(SSIDataFrames$video==i),]$SSI_pa_th, col=rainbow(11)[11], lwd=2)
      abline(h= mean(SSIDataFrames$SSI_pa_th, na.rm=TRUE), col=rainbow(11)[11], lwd=2, lty=2)
legend("topleft", inset=.05, c("fa_mo", "fa_mo_pa", "fa_mo_pa_th",
                               "fa_mo_th", "fa_pa", "fa_pa_th", "fa_th",
                               "mo_pa", "mo_pa_th", "mo_th", "pa_th"), col=rainbow(11), cex=0.6, lwd=2)
legend("topright", inset=.05, c(paste ("Mean fa mo :",
```

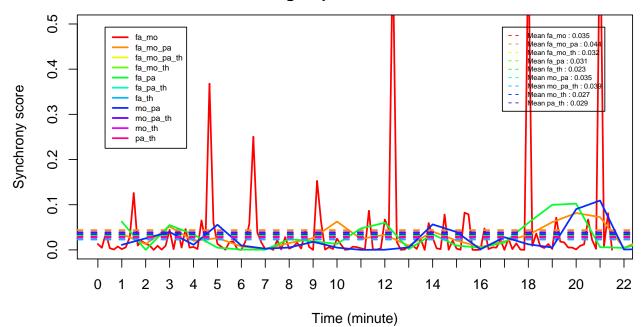
```
round(mean(SSIDataFrames$SSI_fa_mo, na.rm=TRUE),3)),
paste ("Mean fa_mo_pa :", round(mean(SSIDataFrames$SSI_fa_mo_pa,na.rm=TRUE),3)),

# paste ("Mean fa_mo_pa_th :", #round(mean(SSIDataFrames$SSI_fa_mo_pa_th),3)),
paste ("Mean fa_mo_th :", round(mean(SSIDataFrames$SSI_fa_mo_th,na.rm=TRUE),3)),
paste ("Mean fa_pa :", round(mean(SSIDataFrames$SSI_fa_pa, na.rm=TRUE),3)),
paste ("Mean fa_pa_th :", round(mean(SSIDataFrames$SSI_fa_pa_th,na.rm=TRUE),3)),
paste ("Mean mo_pa_th :", round(mean(SSIDataFrames$SSI_fa_th,na.rm=TRUE),3)),
paste ("Mean mo_pa_th :", round(mean(SSIDataFrames$SSI_mo_pa,na.rm=TRUE),3)),
paste ("Mean mo_th :", round(mean(SSIDataFrames$SSI_mo_th,na.rm=TRUE),3)),
paste ("Mean pa_th :", round(mean(SSIDataFrames$SSI_mo_th,na.rm=TRUE),3)),
paste ("Mean pa_th :", round(mean(SSIDataFrames$SSI_pa_th,na.rm=TRUE),3))),
col=rainbow(11), cex=0.5, lty=2, lwd=1)}
```

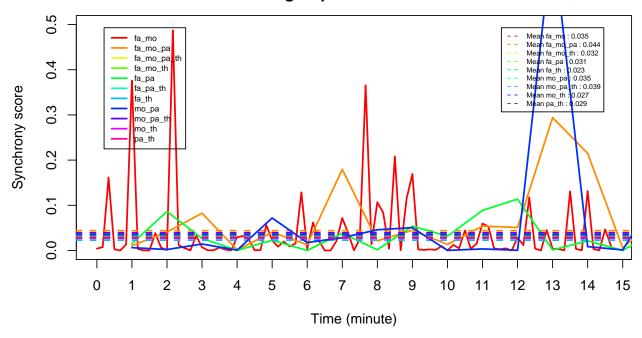
Synchrony scores for each dyad and for the whole group in F1044C.VOB video



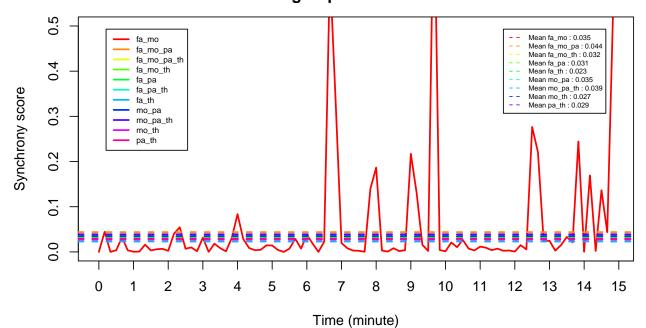
Synchrony scores for each dyad and for the whole group in F1044D1.VOB video



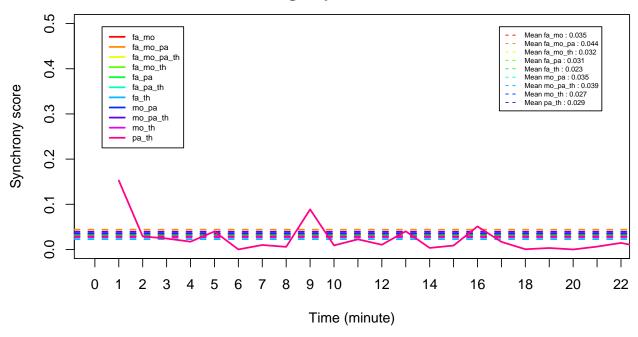
Synchrony scores for each dyad and for the whole group in F1044D2.VOB video



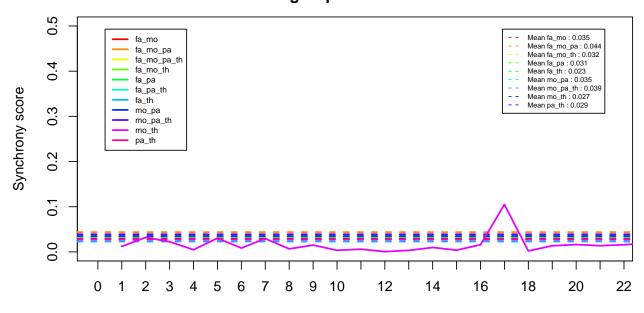
Synchrony scores for each dyad and for the whole group in F1044E.VOB video



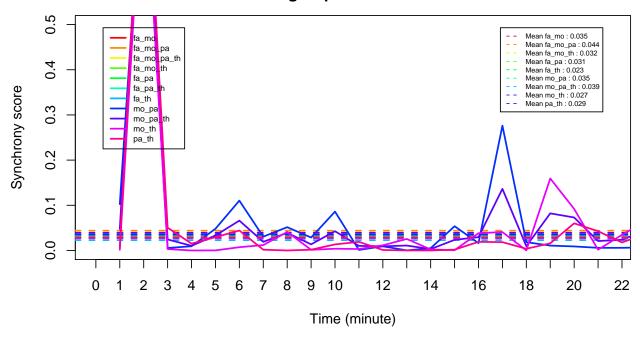
Synchrony scores for each dyad and for the whole group in F1044F.VOB video



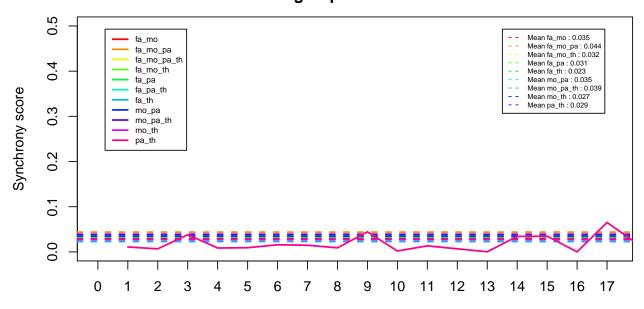
Synchrony scores for each dyad and for the whole group in F1044G.VOB video



Time (minute)
Synchrony scores for each dyad and for the whole group in F1044H.VOB video

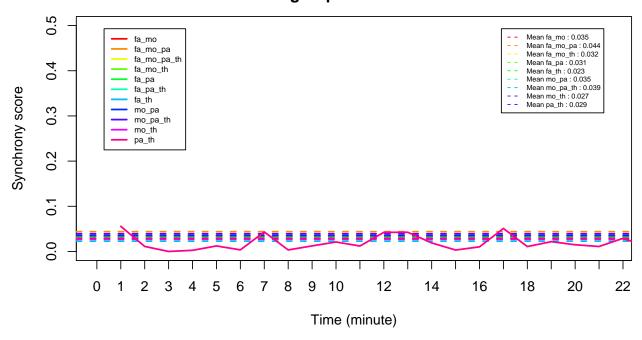


Synchrony scores for each dyad and for the whole group in F1044I.VOB video

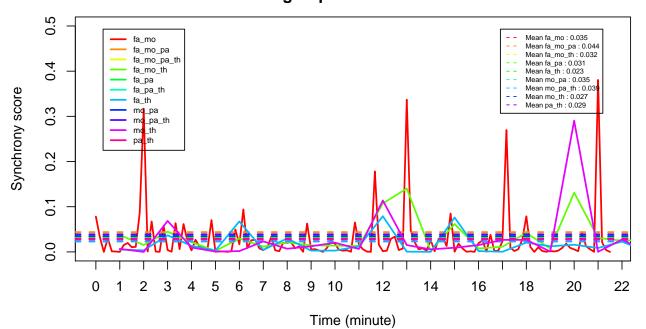


Time (minute)

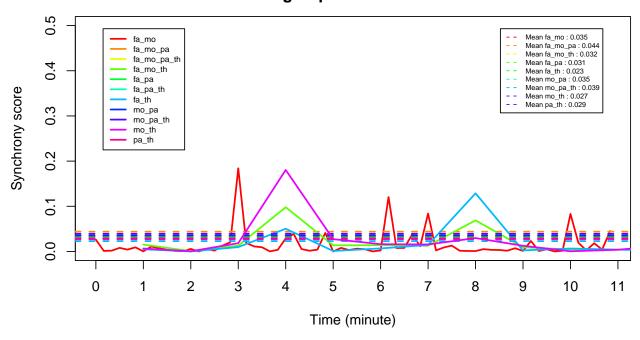
Synchrony scores for each dyad and for the whole group in F1044L.VOB video



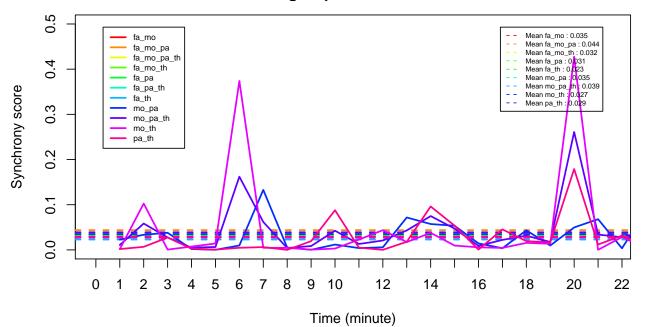
Synchrony scores for each dyad and for the whole group in F1044M1.VOB video



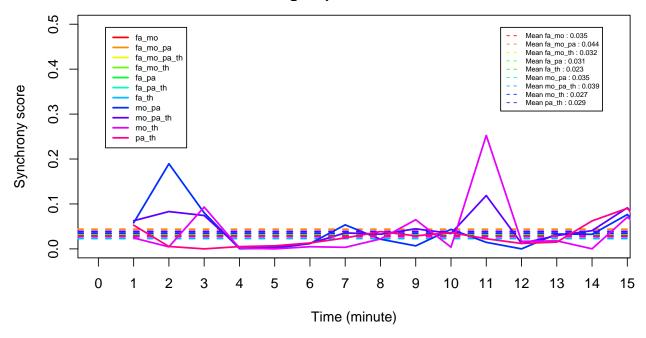
Synchrony scores for each dyad and for the whole group in F1044M2.VOB video



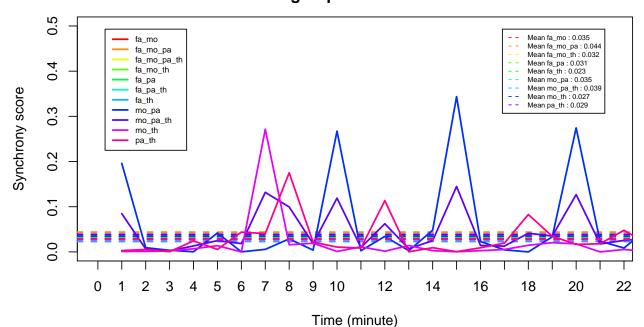
Synchrony scores for each dyad and for the whole group in F1044N.VOB video



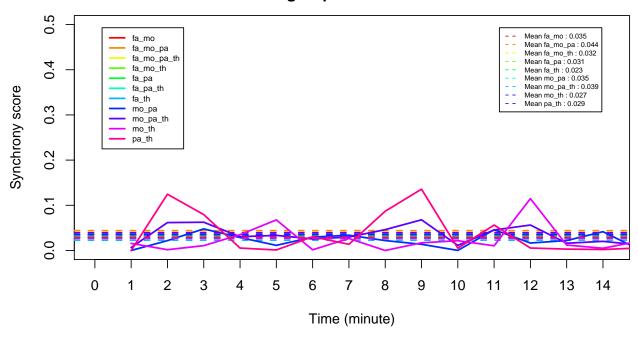
Synchrony scores for each dyad and for the whole group in F10440.VOB video



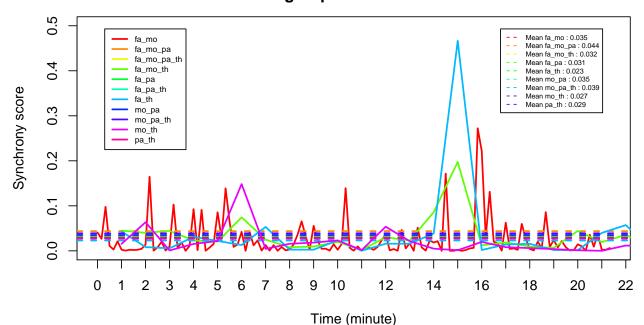
Synchrony scores for each dyad and for the whole group in F1044P.VOB video



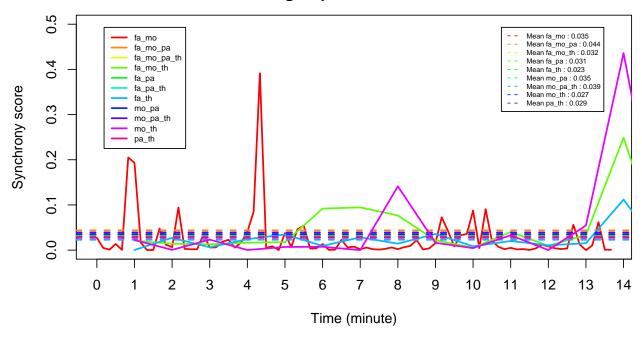
Synchrony scores for each dyad and for the whole group in F1044Q.VOB video



Synchrony scores for each dyad and for the whole group in F1044R1.VOB video



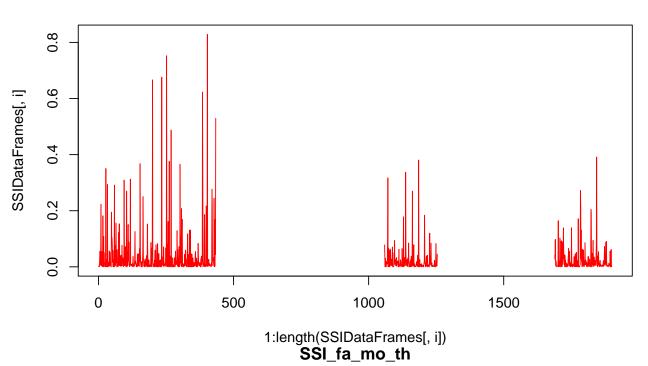
Synchrony scores for each dyad and for the whole group in F1044R2.VOB video

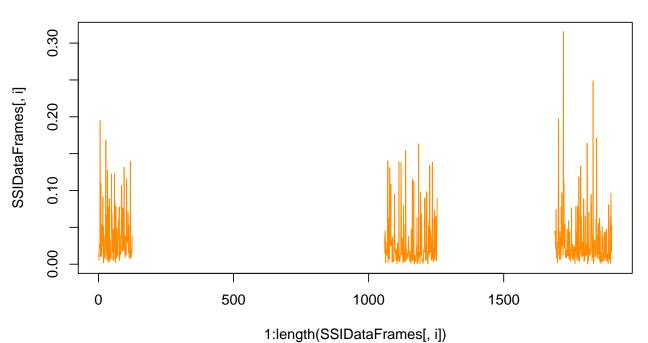


Evolution of synchrony through time, raw each second

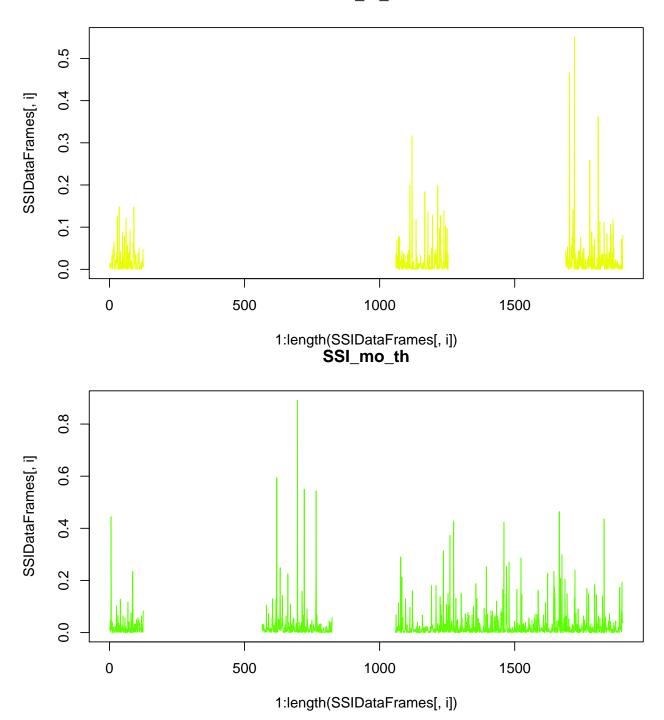
```
par(mar=c(4,4,4,4))
    col= 1
for (i in 5:13){
```

SSI_fa_mo

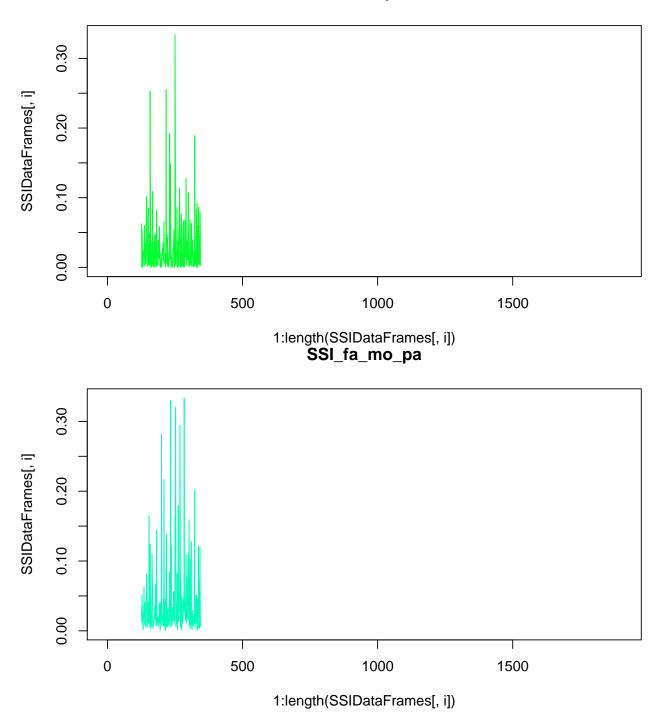




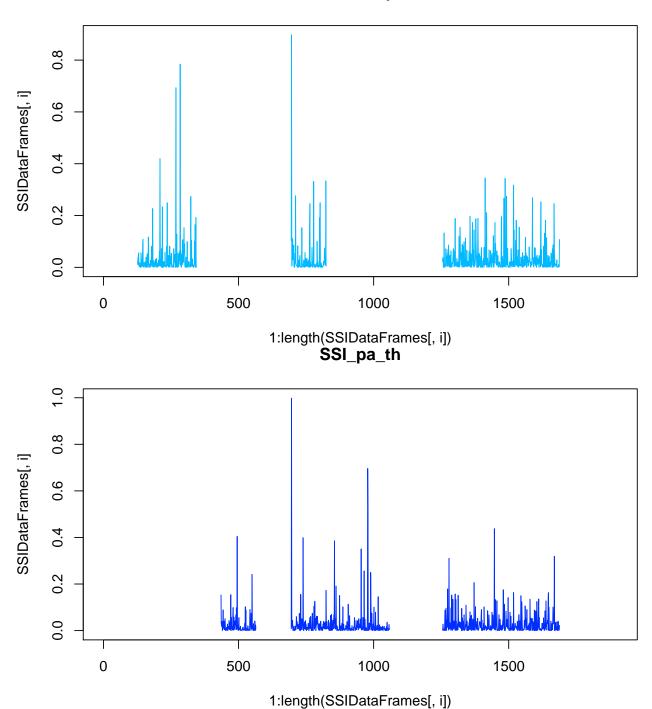
SSI_fa_th



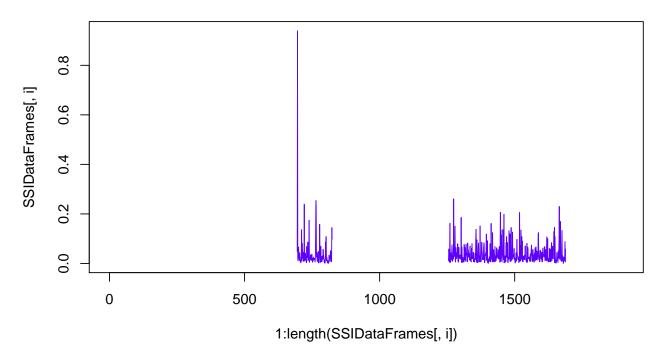
SSI_fa_pa



SSI_mo_pa



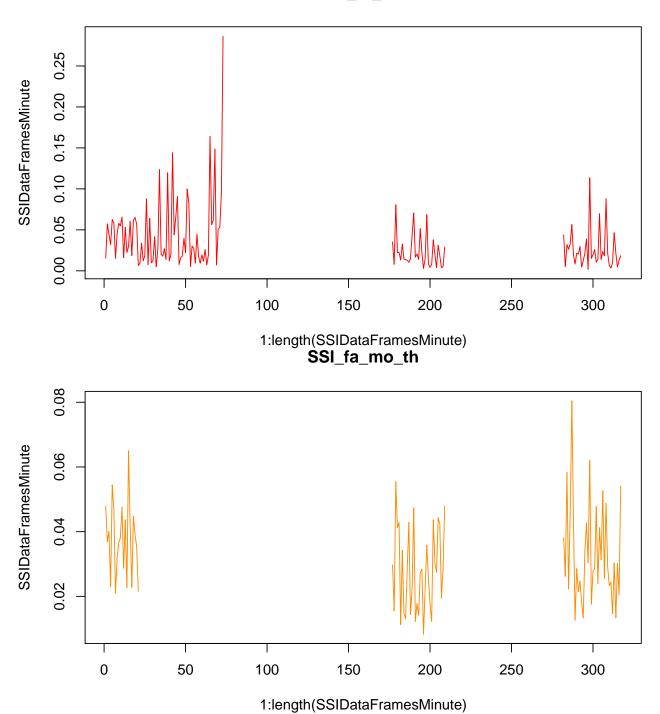
SSI_mo_pa_th



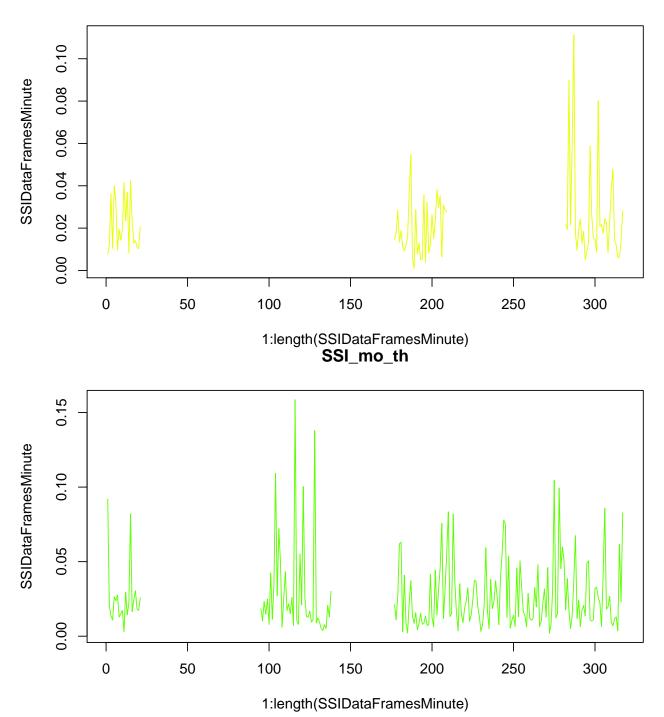
Evolution of synchrony through time, mean by minute

```
par(mar=c(4,4,4,4))
    col = 1
for (indexSSI in 5:13){
    IntervalNumbersVideo <- ceiling(length(SSIDataFrames[,indexSSI])/6)
    SSIColumn <- SSIDataFrames[,indexSSI]
    SSIDataFramesMinute <- c()
    for (i in 1:IntervalNumbersVideo){
        borneInf <- 1+(i-1)*6
        borneSup <- i * 6
        SSIVectorInterval <- SSIColumn[borneInf:borneSup]
        mean <- mean(SSIVectorInterval, na.rm=TRUE)
        SSIDataFramesMinute <- c(SSIDataFramesMinute, mean)}
    plot(1:length(SSIDataFramesMinute), SSIDataFramesMinute, type="l", col=rainbow(11)[col], main = nam col <- col+1}</pre>
```

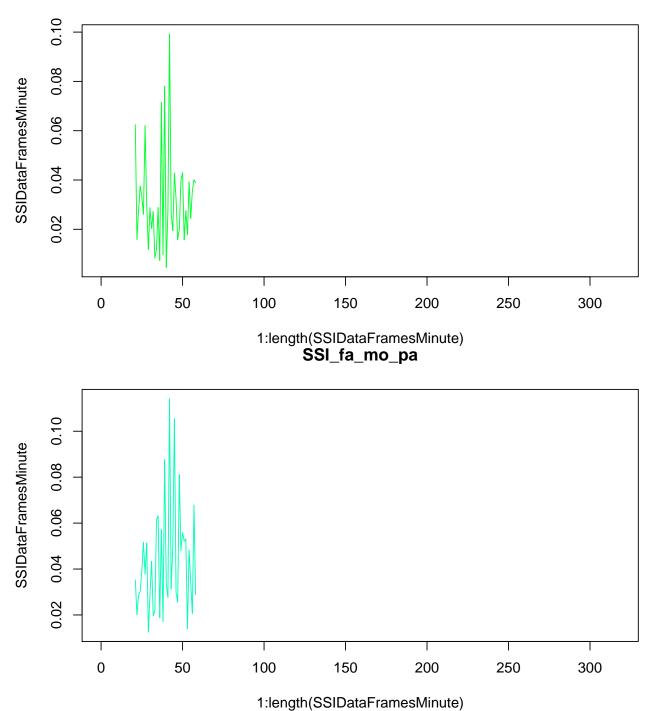
SSI_fa_mo

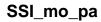


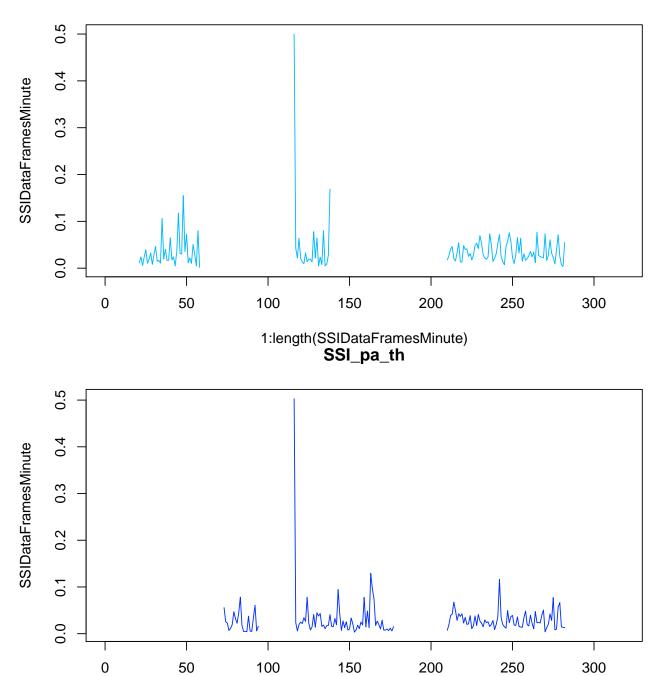
SSI_fa_th











1:length(SSIDataFramesMinute)

SSI_mo_pa_th

