Synchrony in Psychotherapy, Analysis example with F1044 patient data

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Description of the differents steps of data management

Lists

Functions list

MeanMotionByTime

Function that takes raw motion history data and compute the mean on a given interval. Intervals don't overlap, so the frequency of the data change (from 25 frames by second 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 compute the 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>
```

MeanSynchronyByTime (TODO)

File lists

Participants list

```
## [1] "father" "mother" "patient" "therapist"
```

Presentation of the data

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

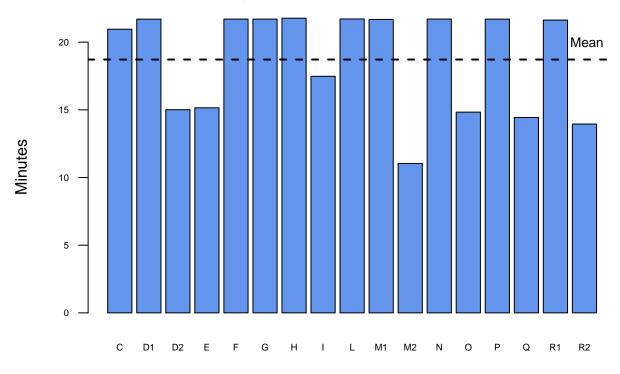
summary(data)

##	frame	father	mother	patient
##	Min. : 1	Min. :0.00	Min. :0.00	Min. :0.00
##	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
##	Mean :14576	Mean :0.00	Mean :0.00	Mean :0.01
##	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	
##	Min. :0.0	F1044H.VOB: 3265	6 Min. : 0.00	00667
##	1st Qu.:0.0	F1044L.VOB: 32570	0 1st Qu.: 4.6	79333
##	Median :0.0	F1044N.VOB: 3256	2 Median : 9.3	58333
##	Mean :0.0	F1044G.VOB: 32556	6 Mean : 9.7	17052
##	3rd Qu.:0.0	F1044F.VOB: 3255	5 3rd Qu.:14.24	42667
##	Max. :0.8	F1044P.VOB: 3255	4 Max. :21.7	70667
##	NA's :77972	(Other) :28180	5	

The time Min is calculated with a frame rate of $25/\mathrm{sec}.$

Length of the videos in minutes

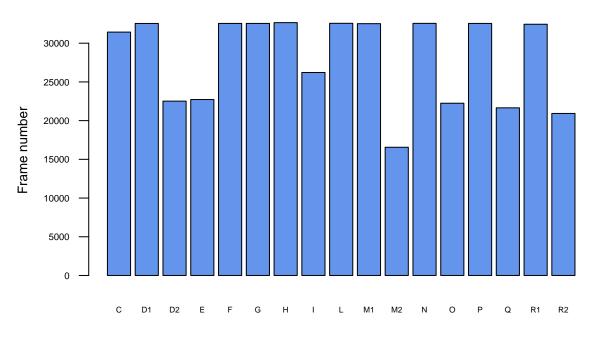
Length in each F1044 video (min)



Video Name

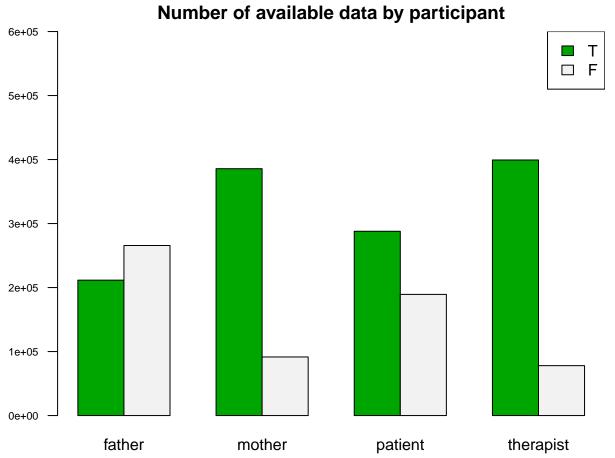
Length of the videos in number of frames

Number of frames in each F1044 video



Video Name

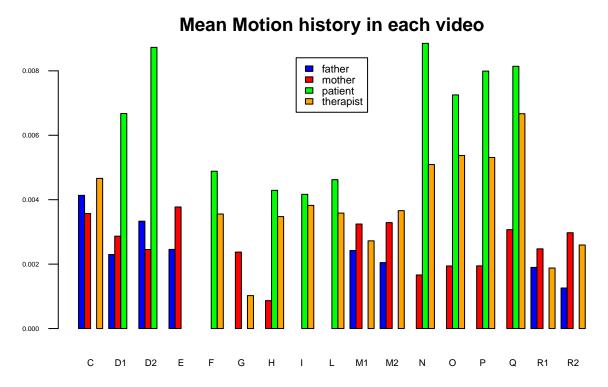
Number of Available (True) and Not Available (False) data for each participant



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 F1044C. The father, mother and the rapist are present. The patient is absent. ## Raw data

```
rawdatafather <- data[which(data$file=="F1044C.VOB"),]$father</pre>
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)
```

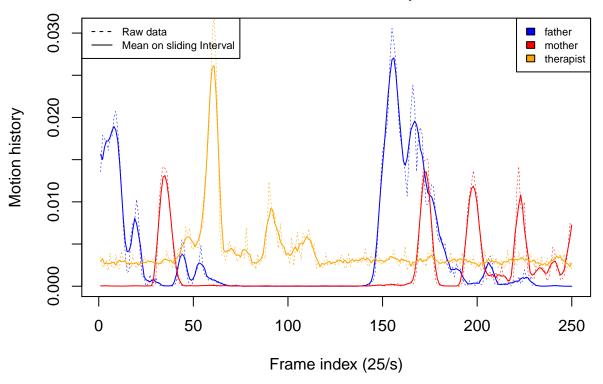
Non overlapping interval

```
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
                                                                      NA's
        Min.
                                       Mean
                                              3rd Qu.
                                                            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)

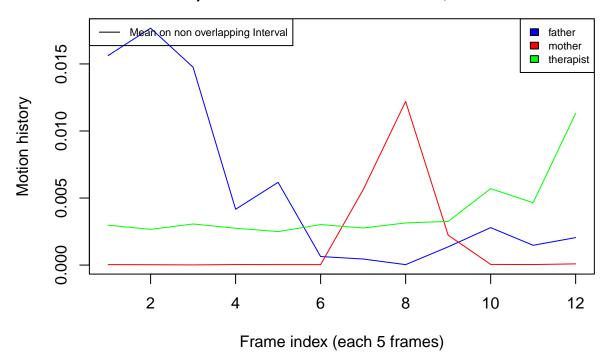
Sliding interval function on a 5 frames interval

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)

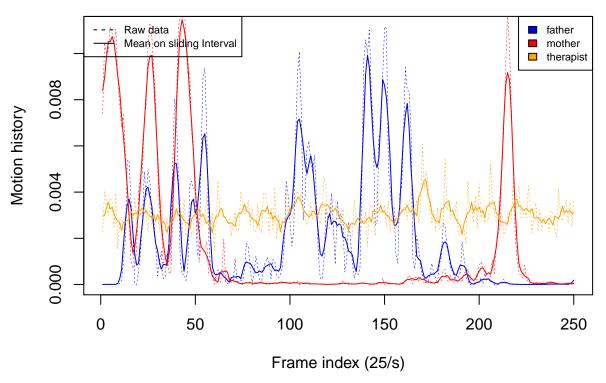
Motion history of the father during 10-20 seconds of the first video(C)

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

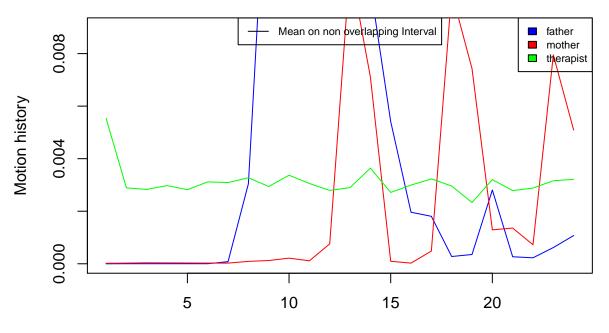


Non overlapping interval function on a 5 frames interval

```
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)

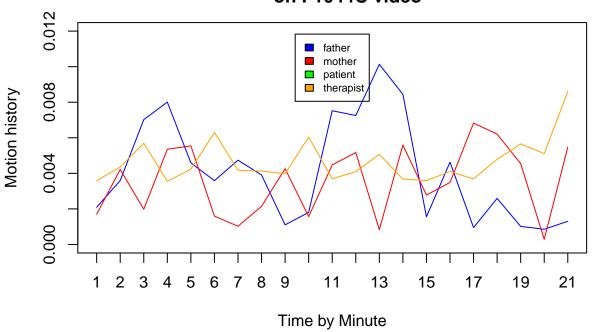
```
### Non overlapping interval function on a 5 frames interval with shifting of therapist (substraction o
plot (1:24, fatherFive[23:46], type="l", col="blue",
main="Mean motion history (non overlapping 5 frames intervals) for
father on F1044C video, between 10-20 seconds,
data therapist shifted",
ylab="Motion 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

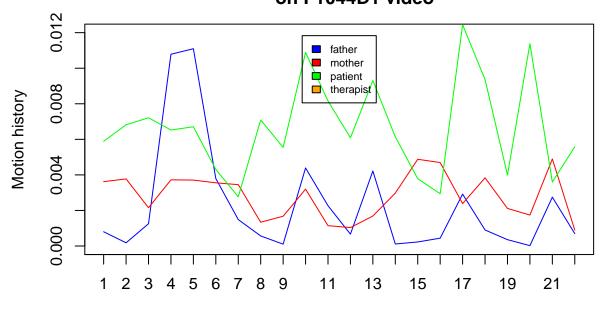
```
for (i in 1:NumberOfvideos){
  fatherMinute<- MeanMotionByTime("father", indexOfvideos=i, interval=1500, data)

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

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



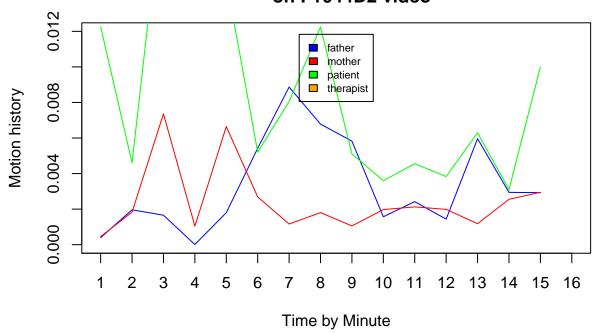
Mean motion history (non overlaping minute intervals) on F1044D1 video



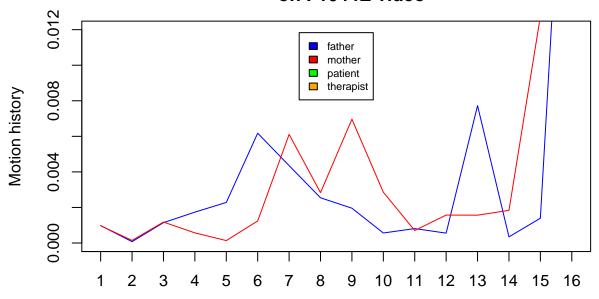
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044D2 video



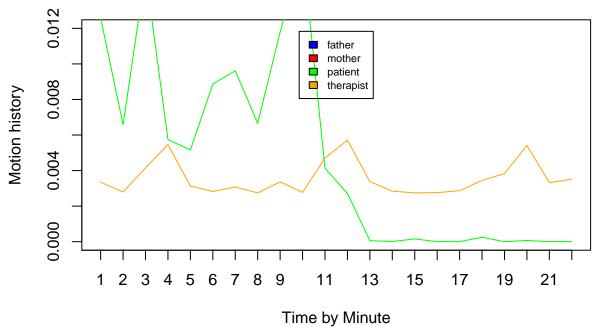
Mean motion history (non overlaping minute intervals) on F1044E video



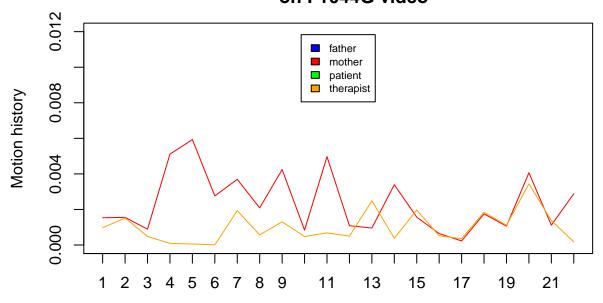
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044F video



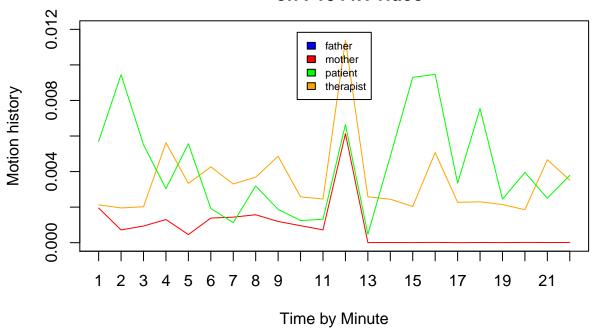
Mean motion history (non overlaping minute intervals) on F1044G video



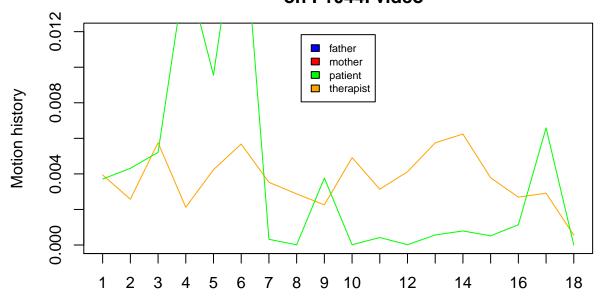
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044H video



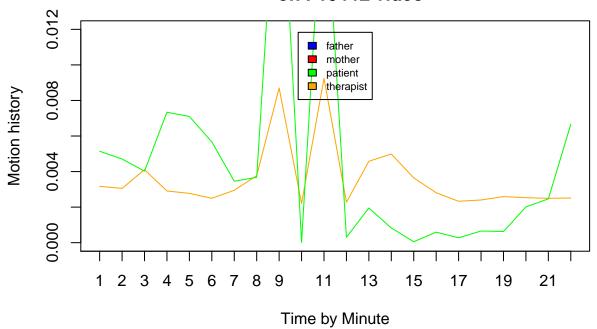
Mean motion history (non overlaping minute intervals) on F1044I video



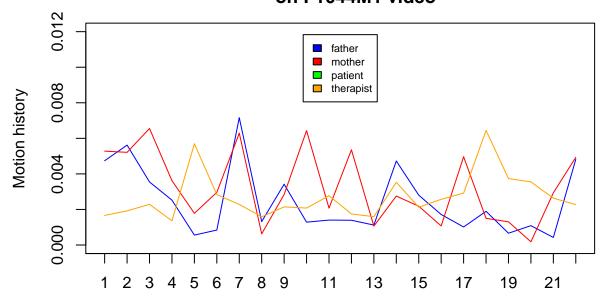
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044L video



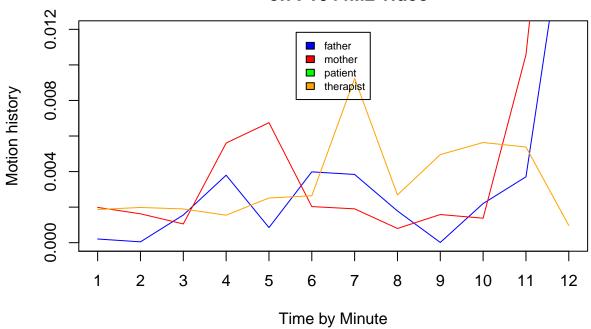
Mean motion history (non overlaping minute intervals) on F1044M1 video



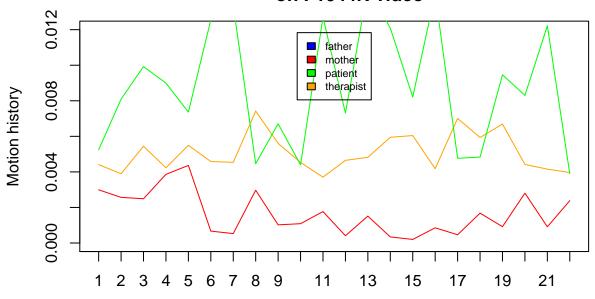
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044M2 video



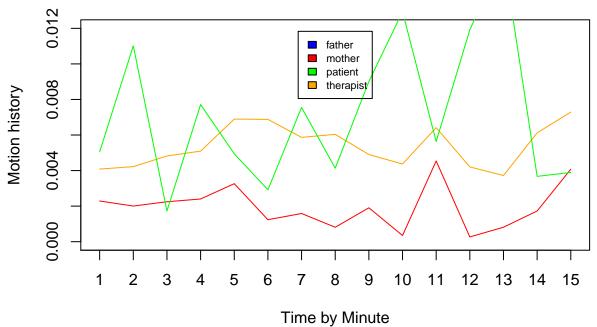
Mean motion history (non overlaping minute intervals) on F1044N video



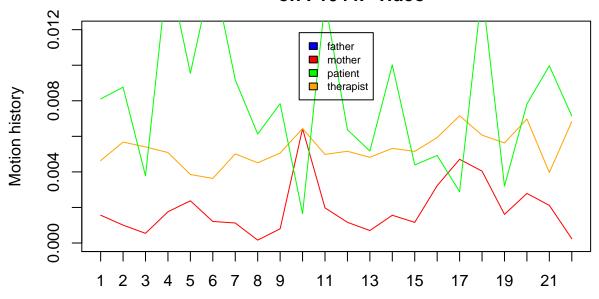
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044O video



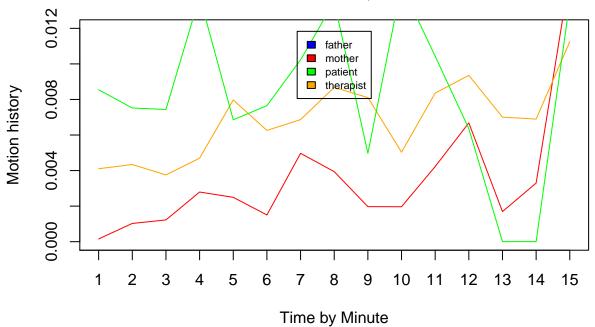
Mean motion history (non overlaping minute intervals) on F1044P video



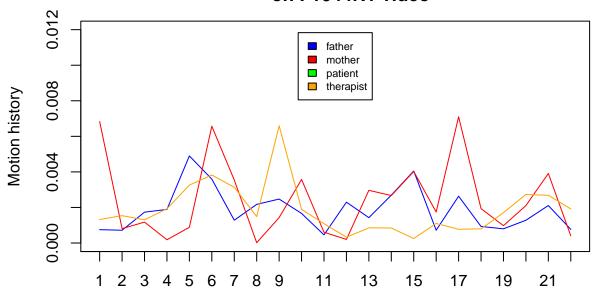
Time by Minute

Mean motion history (non overlaping minute intervals)

on F1044Q video



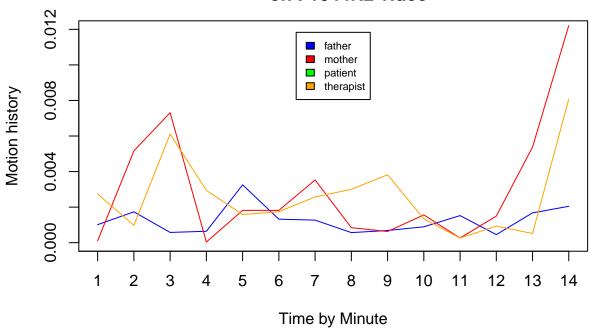
Mean motion history (non overlaping minute intervals) on F1044R1 video



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 video 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("./CSV/filtered/",videoName, ".slideddata.csv", sep=""))
  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 (F1044C, F1044D1, etc) that we can import with R with

this following function

```
## factor(0)
## 17 Levels: F1044C.VOB F1044D1.VOB F1044D2.VOB F1044E.VOB ... F1044R2.VOB
## [1] "SSI Files Directory"
## [1] "/Users/Ofix/Documents/Fac/internat/Recherche/projets/synchro/synchroData/Git"
## [1] "SS Files List"
   [1] "SSIF1044C.csv"
                        "SSIF1044D1.csv" "SSIF1044D2.csv" "SSIF1044E.csv"
                        "SSIF1044G.csv" "SSIF1044H.csv" "SSIF1044I.csv"
   [5] "SSIF1044F.csv"
  [9] "SSIF1044L.csv"
                        "SSIF1044M1.csv" "SSIF1044M2.csv" "SSIF1044N.csv"
## [13] "SSIF10440.csv"
                        "SSIF1044P.csv" "SSIF1044Q.csv" "SSIF1044R1.csv"
## [17] "SSIF1044R2.csv"
   [1] "SSIF1044C.csv" "SSIF1044D1.csv" "SSIF1044D2.csv" "SSIF1044E.csv"
##
                        "SSIF1044G.csv" "SSIF1044H.csv" "SSIF1044I.csv"
   [5] "SSIF1044F.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(SSIdataFrame)
```

```
1900 obs. of 14 variables:
## 'data.frame':
## $ X
                 : int 0 1 2 3 4 5 6 7 8 9 ...
                : int 1 2 3 4 5 6 7 8 9 10 ...
## $ Interval
## $ Time min : num 0 0.167 0.333 0.5 0.667 ...
## $ video
                : Factor w/ 17 levels "F1044C.VOB", "F1044D1.VOB", ...: 1 1 1 1 1 1 1 1 1 1 ...
## $ date
                 : chr "07/02/08" "07/02/08" "07/02/08" "07/02/08" ...
                : 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 ...
## $ SSI_fa_pa : num NA ...
## $ SSI_fa_mo_pa: num NA ...
## $ SSI_mo_pa
                : num NA NA NA NA NA NA NA NA NA ...
                : num NA NA NA NA NA NA NA NA NA ...
## $ SSI_pa_th
## $ SSI_mo_pa_th: num NA ...
```

Synchrony scores for each dyad and for the whole group

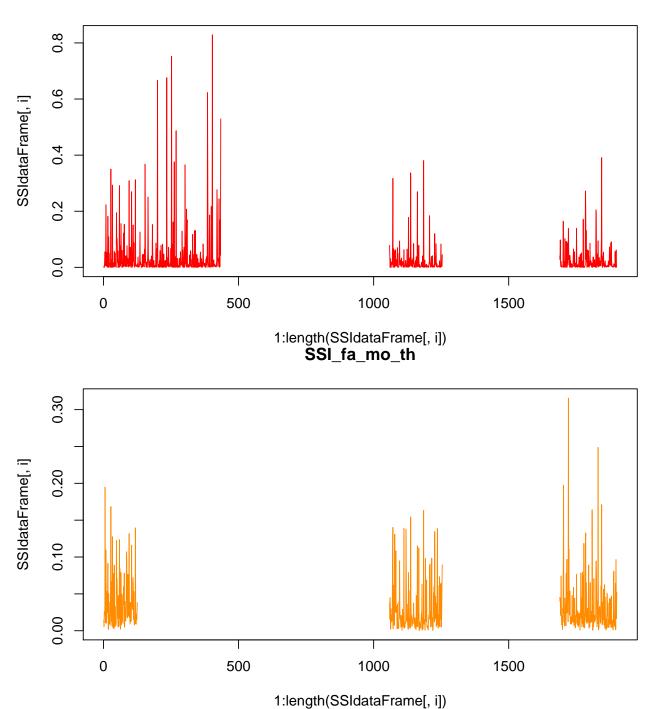
```
for (i in unique(SSIdataFrame$video))
      \{par(mar=c(4,4,4,3), mfrow=c(1,1))\}
      plot(SSIdataFrame[which(SSIdataFrame$video==i),]$Time min,
           SSIdataFrame[which(SSIdataFrame$video==i),]$SSI fa mo,
           type="1", col=rainbow(4)[1],
           main=paste("Synchrony scores for each dyad and for \n the whole group in", i, "video"),
           xlab = "Time (minute)", ylab="Synchrony score", lwd=2,
xaxp=c(0,length(SSIdataFrame$Time min), length(SSIdataFrame$Time min)))
\#, ylim=c(0,0.5))
      abline(h=mean(SSIdataFrame$SSI_fa_mo, na.rm=TRUE), col=rainbow(11)[1], lwd=2, lty=2)
      lines(SSIdataFrame[which(SSIdataFrame$video==i),]$SSI_fa_mo_pa, col=rainbow(11)[2], lwd=2)
      abline(h= mean(SSIdataFrame$SSI_fa_mo_pa, na.rm=TRUE), col=rainbow(11)[2], lwd=2, lty=2)
       lines(SSIdataFrame[which(SSIdataFrame$video==i),]$SSI_fa_mo_pa_th, col=rainbow(11)[3], lwd=2)
#
       abline(h= mean(SSIdataFrame$SSI_fa_mo_pa_th, na.rm=TRUE), col=rainbow(11)[3], lwd=2, lty=2)
      lines(SSIdataFrame[which(SSIdataFrame$video==i),]$SSI_fa_mo_th, col=rainbow(11)[4], lwd=2)
      abline(h= mean(SSIdataFrame$SSI_fa_mo_th, na.rm=TRUE), col=rainbow(11)[4], lwd=2, lty=2)
     lines(SSIdataFrame[which(SSIdataFrame$video==i),]$SSI_fa_pa, col=rainbow(11)[5], lwd=2)
      abline(h= mean(SSIdataFrame$SSI_fa_pa, na.rm=TRUE), col=rainbow(11)[5], lwd=2, lty=2)
       lines (SSIdataFrame[which(SSIdataFrame$video==i),] \$SSI\_fa\_pa\_th, \ col=rainbow(11)[6], \ lwd=2)
#
       abline(h= mean(SSIdataFrame$SSI fa pa th, na.rm=TRUE), col=rainbow(11)[6], lwd=2, lty=2)
lines(SSIdataFrame[which(SSIdataFrame$video==i),]$SSI_fa_th, col=rainbow(11)[7], lwd=2)
abline(h= mean(SSIdataFrame$SSI_fa_th, na.rm=TRUE), col=rainbow(11)[7], lwd=2, lty=2)
lines(SSIdataFrame[which(SSIdataFrame$video==i),]$SSI_mo_pa, col=rainbow(11)[8], lwd=2)
abline(h= mean(SSIdataFrame$SSI_mo_pa, na.rm=TRUE), col=rainbow(11)[8], lwd=2, lty=2)
lines(SSIdataFrame[which(SSIdataFrame$video==i),]$SSI_mo_pa_th, col=rainbow(11)[9], lwd=2)
```

```
abline(h= mean(SSIdataFrame$SSI_mo_pa_th, na.rm=TRUE), col=rainbow(11)[9], lwd=2, lty=2)
lines(SSIdataFrame[which(SSIdataFrame$video==i),]$SSI_mo_th, col=rainbow(11)[10], lwd=2)
abline(h= mean(SSIdataFrame$SSI_mo_th, na.rm=TRUE), col=rainbow(11)[10], lwd=2, lty=2)
      lines(SSIdataFrame[which(SSIdataFrame$video==i),]$SSI_pa_th, col=rainbow(11)[11], lwd=2)
      abline(h= mean(SSIdataFrame$SSI_pa_th, na.rm=TRUE), col=rainbow(11)[11], lwd=2, lty=2)
legend("topleft", inset=.05, c("fa mo", "fa mo pa", "fa mo pa th",
"fa_mo_th", "fa_pa", "fa_pa_th", "fa_th",
"mo_pa", "mo_pa_th", "mo_th", "pa_th"),
col=rainbow(11), cex=0.6, lwd=2)
legend("topright", inset=.05, c(paste ("Mean fa_mo :",
                                       round(mean(SSIdataFrame$SSI_fa_mo, na.rm=TRUE),3)),
      paste ("Mean fa_mo_pa :", round(mean(SSIdataFrame$SSI_fa_mo_pa,na.rm=TRUE),3)),
#
      paste \ ("Mean \ fa\_mo\_pa\_th \ :", \ \#round(mean(SSIdataFrame\$SSI\_fa\_mo\_pa\_th), 3)),
      paste ("Mean fa_mo_th :", round(mean(SSIdataFrame$SSI_fa_mo_th,na.rm=TRUE),3)),
      paste ("Mean fa_pa :", round(mean(SSIdataFrame$SSI_fa_pa, na.rm=TRUE),3)),
      paste ("Mean fa_pa_th :", round(mean(SSIdataFrame$SSI_fa_pa_th,na.rm=TRUE),3)),
      paste ("Mean fa_th :", round(mean(SSIdataFrame$SSI_fa_th,na.rm=TRUE),3)),
      paste ("Mean mo_pa :", round(mean(SSIdataFrame$SSI_mo_pa,na.rm=TRUE),3)),
      paste ("Mean mo_pa_th :", round(mean(SSIdataFrame$SSI_mo_pa_th,na.rm=TRUE),3)),
      paste ("Mean mo_th :", round(mean(SSIdataFrame$SSI_mo_th,na.rm=TRUE),3)),
      paste ("Mean pa_th :", round(mean(SSIdataFrame$SSI_pa_th,na.rm=TRUE),3))),
col=rainbow(11), cex=0.5, lty=2, lwd=1)}
```

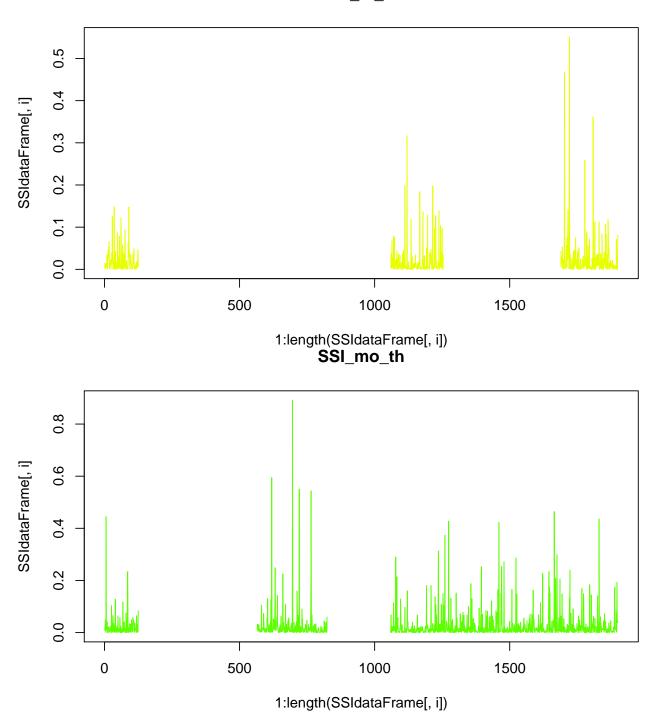
Evolution of synchrony through time, raw each second

```
par(mar=c(4,4,4,4))
    col <- 1
for (i in 6:length(SSIdataFrame)){
    plot(1:length(SSIdataFrame[,i]), SSIdataFrame[,i], type="l",
    col=rainbow(11)[col], main = names(SSIdataFrame)[i])
    col <- col+1}</pre>
```

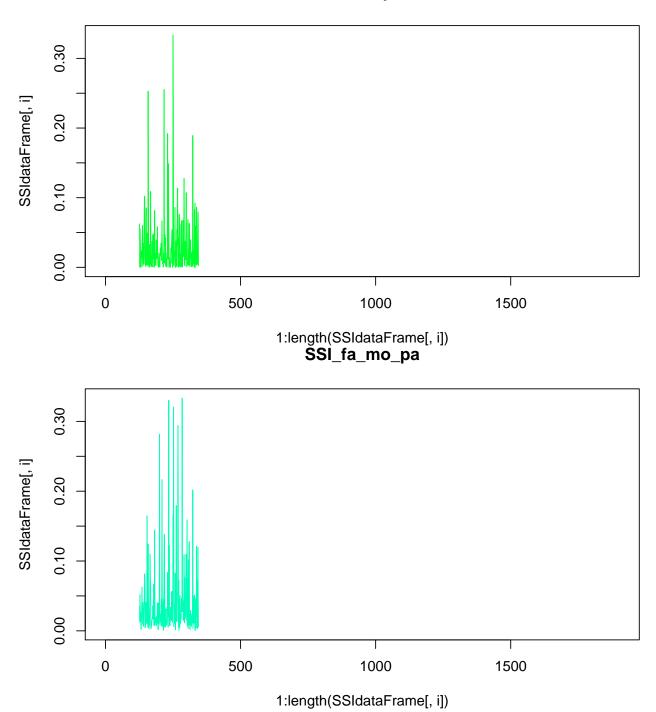




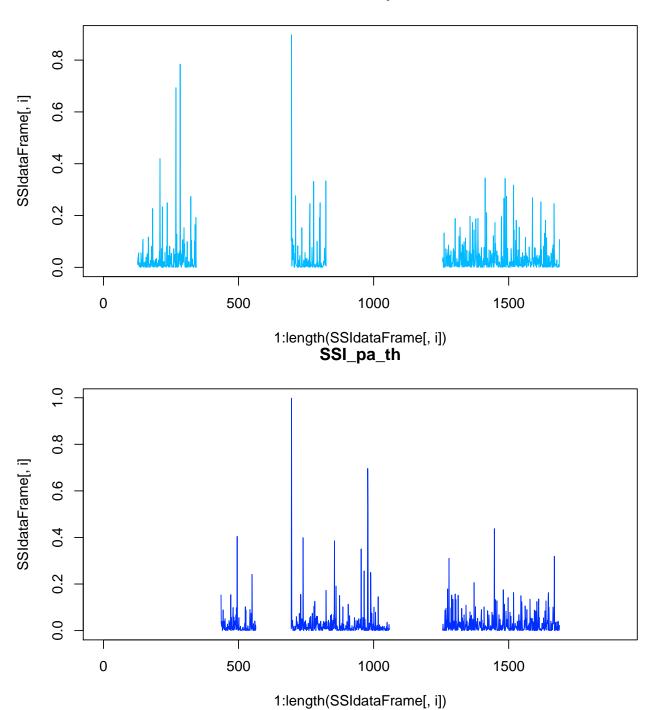
SSI_fa_th



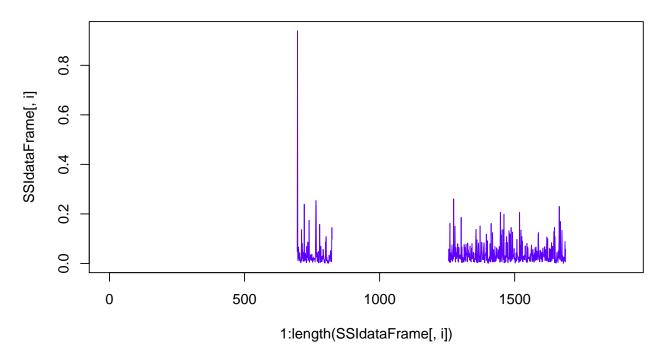
SSI_fa_pa



SSI_mo_pa



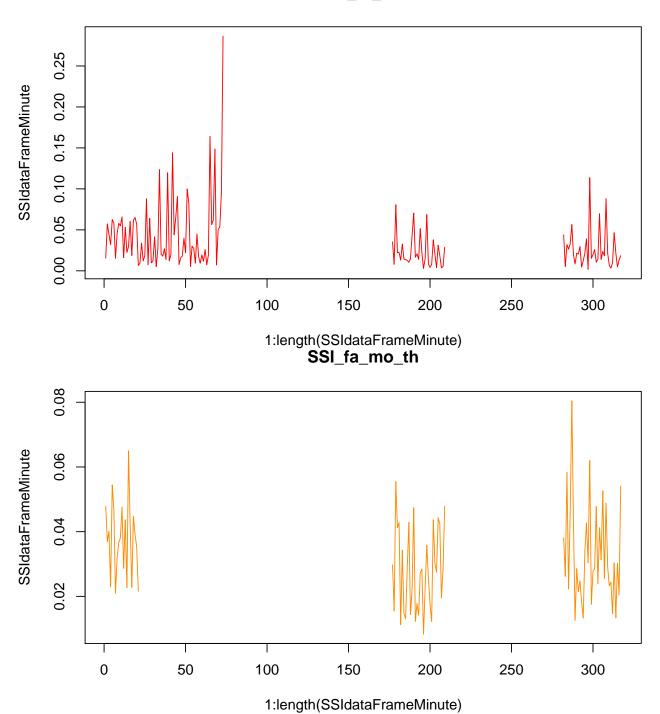
SSI_mo_pa_th



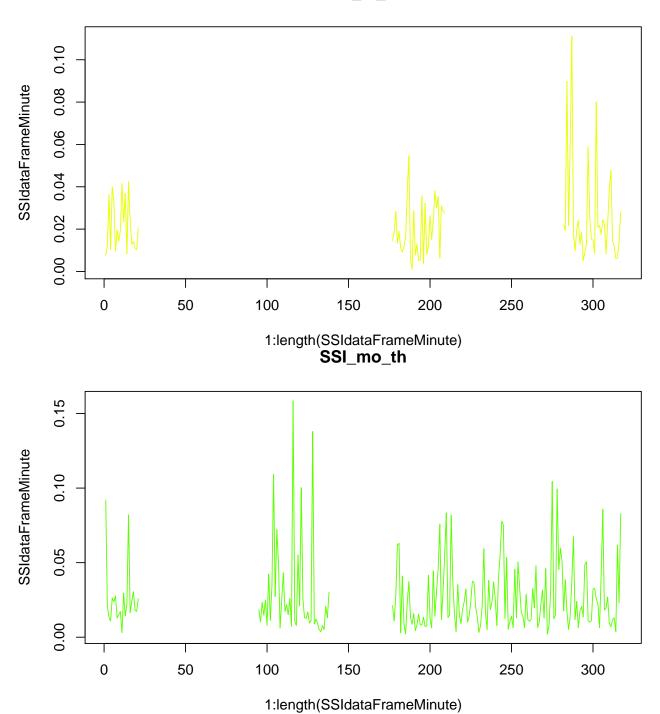
Evolution of synchrony through time, mean by minute

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

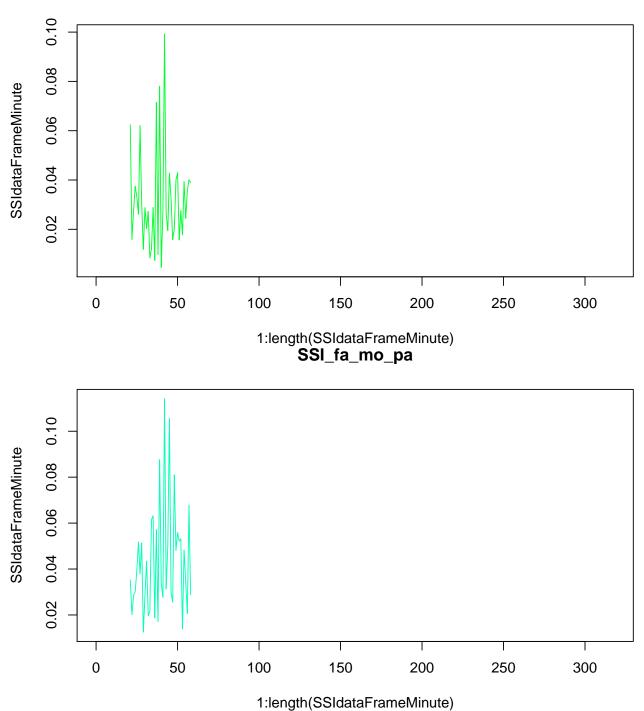
SSI_fa_mo

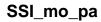


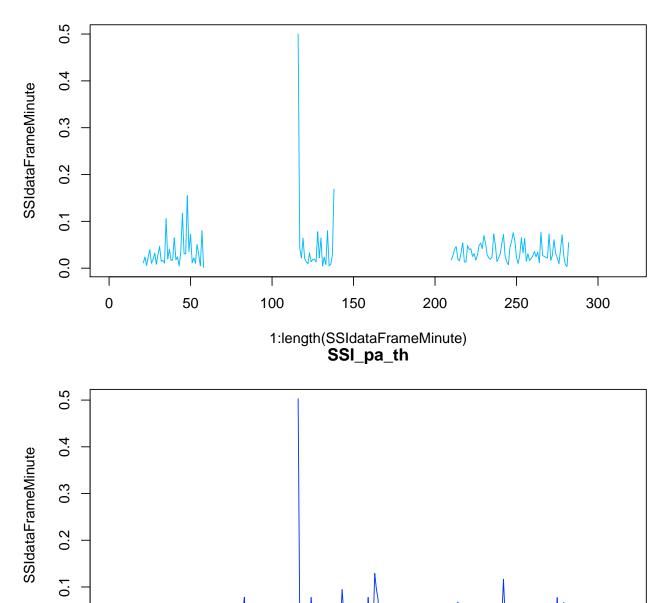
SSI_fa_th







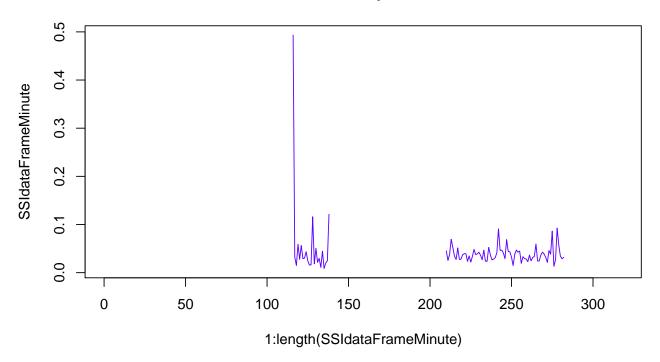




1:length(SSIdataFrameMinute)

0.0

SSI_mo_pa_th

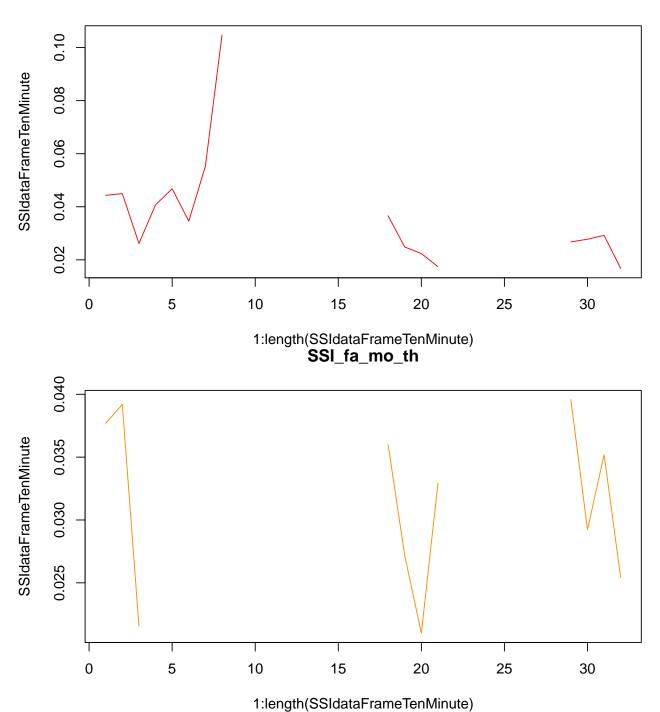


Evolution of synchrony through time, mean by 10 minutes

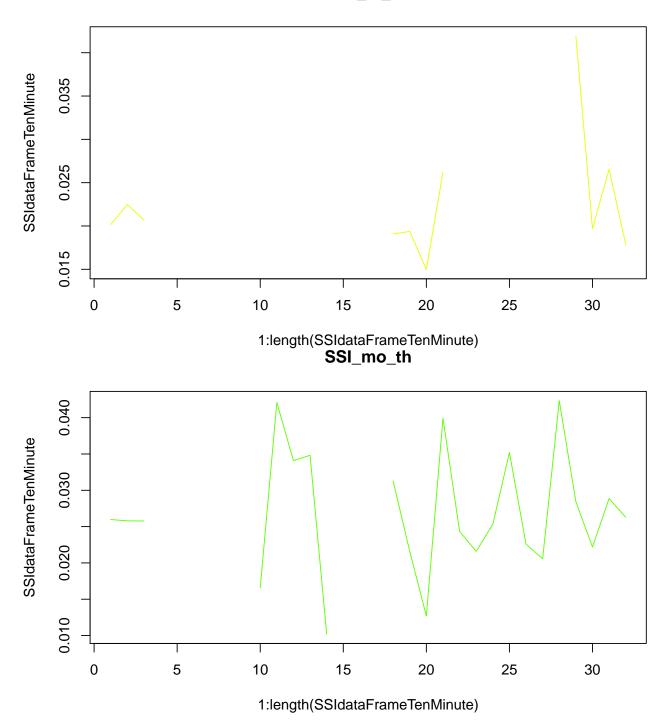
```
par(mar=c(4,4,4,4))
    col = 1

for (indexSSI in 6:length(SSIdataFrame)){
    IntervalNumbersVideo <- ceiling(length(SSIdataFrame[,indexSSI])/60)
    SSIColumn <- SSIdataFrame[,indexSSI]
    SSIdataFrameTenMinute <- c()
    for (i in 1:IntervalNumbersVideo){
        borneInf <- 1+(i-1)*60
        borneSup <- i * 60
        SSIVectorInterval <- SSIColumn[borneInf:borneSup]
        mean <- mean(SSIVectorInterval, na.rm=TRUE)
        SSIdataFrameTenMinute <- c(SSIdataFrameTenMinute, mean)}
    plot(1:length(SSIdataFrameTenMinute), SSIdataFrameTenMinute, type="l", col=rainbow(11)[col], main = col <- col+1}</pre>
```

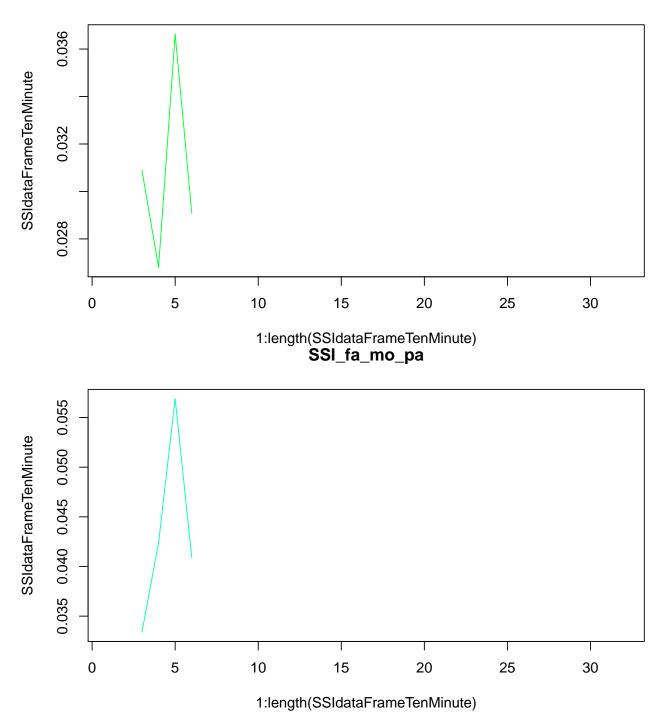


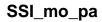


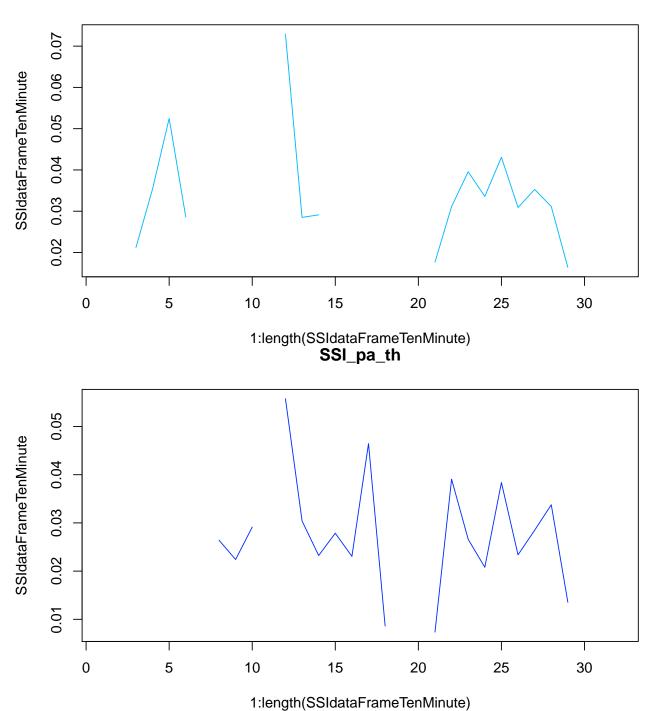
SSI_fa_th



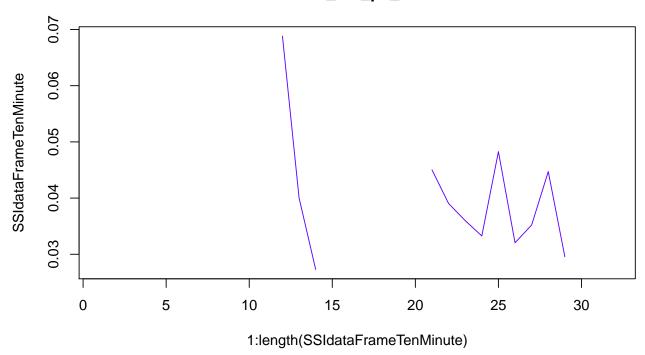






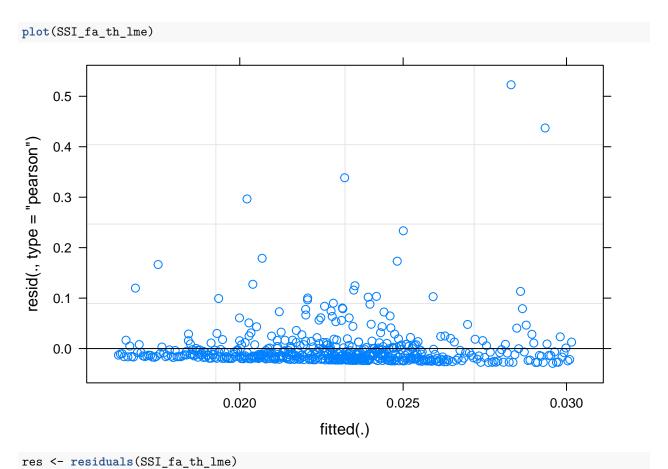


SSI_mo_pa_th



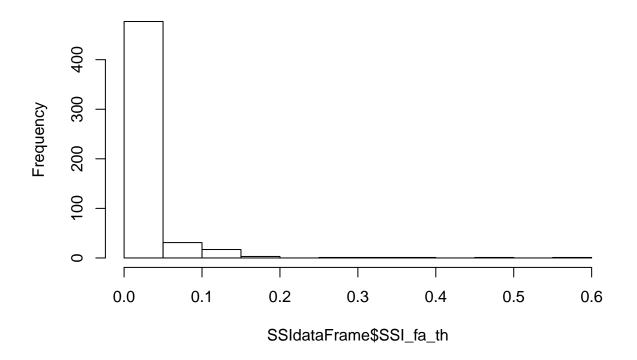
```
SSI_fa_th_lme <- lmer(SSI_fa_th ~ Time_min + (1|video), data=SSIdataFrame)
summary(SSI_fa_th_lme)</pre>
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: SSI_fa_th ~ Time_min + (1 | video)
      Data: SSIdataFrame
##
## REML criterion at convergence: -1708.1
##
## Scaled residuals:
       Min
                1Q Median
                                3Q
                                       Max
## -0.6153 -0.4189 -0.2830 0.0065 10.9778
##
## Random effects:
## Groups
                         Variance Std.Dev.
             Name
## video
             (Intercept) 1.514e-05 0.00389
  Residual
                         2.268e-03 0.04762
## Number of obs: 533, groups: video, 5
##
## Fixed effects:
##
                 Estimate Std. Error t value
## (Intercept) 0.0261037 0.0042446
## Time_min
               -0.0003472 0.0003557 -0.976
## Correlation of Fixed Effects:
            (Intr)
## Time_min -0.767
```



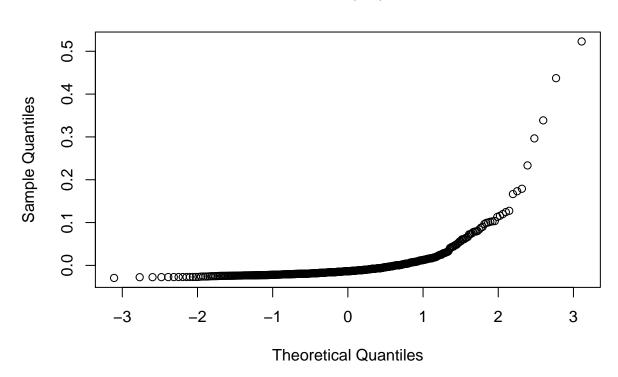
hist(SSIdataFrame\$SSI_fa_th)

Histogram of SSIdataFrame\$SSI_fa_th

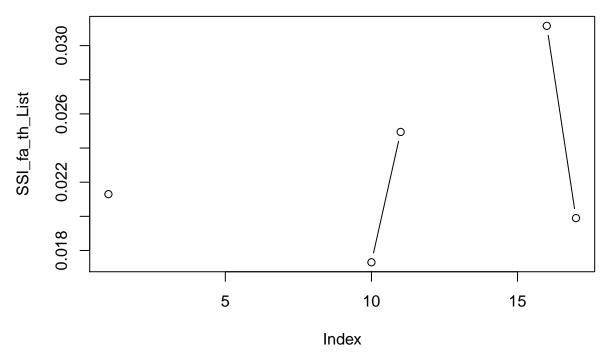


qqnorm(res)

Normal Q-Q Plot

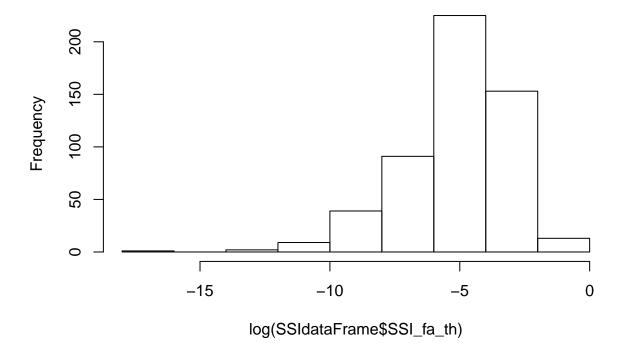


```
SSI_fa_th_List <- c()</pre>
for (i in indexlist){
  SSI_fa_th_List <- c(SSI_fa_th_List, mean(SSIdataFrame[which(SSIdataFrame$video==i),]$SSI_fa_th, na.rm
print(SSI_fa_th_List)
    [1] 0.02129936
##
                             {\tt NaN}
                                         NaN
                                                                              {\tt NaN}
                                                      NaN
                                                                  NaN
    [7]
                {\tt NaN}
                             NaN
                                         NaN 0.01730778 0.02494321
                                                                              NaN
## [13]
                {\tt NaN}
                             NaN
                                         NaN 0.03115848 0.01989619
plot(SSI_fa_th_List, type="b")
```



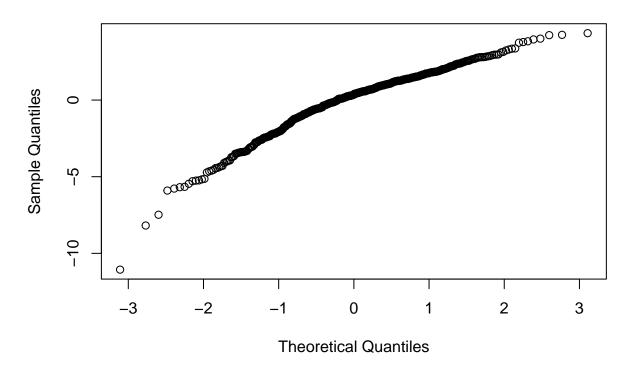
log of the data
log_SSI_fa_th <- hist(log(SSIdataFrame\$SSI_fa_th))</pre>

Histogram of log(SSIdataFrame\$SSI_fa_th)



SSI_fa_th_log_lme <- lmer(log(SSI_fa_th) ~ Time_min + (1|video), data=SSIdataFrame)
res_log <- residuals(SSI_fa_th_log_lme)
qqnorm(res_log)</pre>

Normal Q-Q Plot

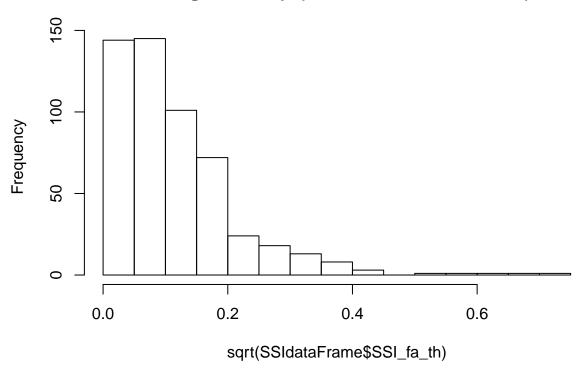


summary(SSI_fa_th_log_lme)

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: log(SSI_fa_th) ~ Time_min + (1 | video)
     Data: SSIdataFrame
##
##
## REML criterion at convergence: 2281.3
##
## Scaled residuals:
##
       Min
                1Q Median
                                3Q
                                       Max
  -5.4326 -0.4610 0.1755 0.6616 2.1427
##
##
## Random effects:
   Groups
##
             Name
                         Variance Std.Dev.
##
   video
             (Intercept) 0.04711 0.2171
   Residual
                         4.14617 2.0362
## Number of obs: 533, groups: video, 5
##
## Fixed effects:
               Estimate Std. Error t value
## (Intercept) -4.78183
                           0.19209 -24.893
                           0.01533 -2.451
## Time_min
               -0.03758
## Correlation of Fixed Effects:
            (Intr)
## Time_min -0.725
```

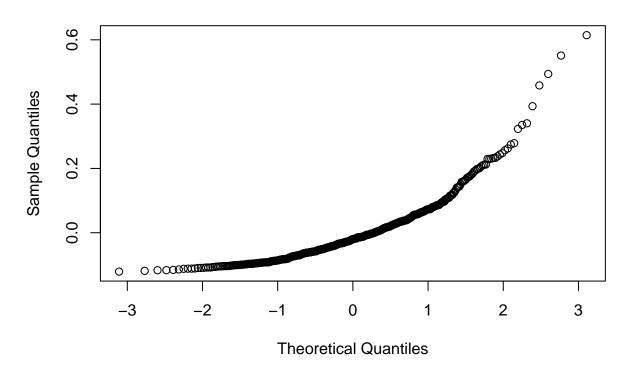
```
# root square of the data
sq_SSI_fa_th <- hist(sqrt(SSIdataFrame$SSI_fa_th))</pre>
```

Histogram of sqrt(SSIdataFrame\$SSI_fa_th)



```
SSI_fa_th_sq_lme <- lmer(sqrt(SSI_fa_th) ~ Time_min + (1|video), data=SSIdataFrame)
res_sq <- residuals(SSI_fa_th_sq_lme)
qqnorm(res_sq)</pre>
```

Normal Q-Q Plot



summary(SSI_fa_th_sq_lme)

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: sqrt(SSI_fa_th) ~ Time_min + (1 | video)
     Data: SSIdataFrame
##
##
## REML criterion at convergence: -937.9
##
## Scaled residuals:
                1Q Median
##
       Min
                                3Q
                                       Max
  -1.2301 -0.6704 -0.2018 0.3764 6.2496
##
##
## Random effects:
   Groups
##
             Name
                         Variance Std.Dev.
##
   video
             (Intercept) 8.351e-05 0.009139
  Residual
                         9.665e-03 0.098313
## Number of obs: 533, groups: video, 5
##
## Fixed effects:
                 Estimate Std. Error t value
## (Intercept) 0.1276072 0.0089814 14.208
## Time_min
               -0.0013773 0.0007371 -1.868
## Correlation of Fixed Effects:
            (Intr)
## Time_min -0.749
```