

SynchronyScoresAnalysis

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Clean environment

```
rm(list = ls(all.names = TRUE))
```

Session info

```
sessionInfo()

## R version 3.2.2 (2015-08-14)
## Platform: x86_64-apple-darwin13.4.0 (64-bit)
## Running under: OS X 10.12.3 (unknown)
##
## locale:
## [1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8
##
## attached base packages:
## [1] stats      graphics   grDevices  utils      datasets   methods    base
##
## loaded via a namespace (and not attached):
## [1] backports_1.0.5 magrittr_1.5     rprojroot_1.2   tools_3.2.2
## [5] htmltools_0.3.5 yaml_2.1.13    Rcpp_0.12.6    stringi_1.1.1
## [9] rmarkdown_1.3   knitr_1.15.1   stringr_1.0.0   digest_0.6.9
## [13] evaluate_0.10
```

Preparation of the session

Fixed variables

```
FileExtension <- ".MTS.avi_res.csv"

# working directory
# where this report is
#setwd("./Git/Monrado/Reports/")
setwd("/Users/Ofix/Documents/Fac/internat/Recherche/projets/synchro/synchroData/Git/Monrado/Reports")

# blue will refer to father
# red will refer to mother
# green to child
colOrderList <- c("blue", "red", "green")

ParticipantsList <- c("father", "mother", "child")
```

```

# Import data
cutFrames <- read.csv2("../Data/CSV/Cutframes.csv")
str(cutFrames)

## 'data.frame':   34 obs. of  7 variables:
##   $ family    : Factor w/ 34 levels "1606","BAJE059",...: 1 2 3 4 5 6 7 8 9 10 ...
##   $ CutBefore : Factor w/ 11 levels "00:06","00:07",...: 6 4 3 5 3 7 5 11 7 5 ...
##   $ CutMiddle1: Factor w/ 26 levels "04:43","04:53",...: 21 15 16 16 4 10 17 18 20 17 ...
##   $ CutMiddle2: Factor w/ 31 levels "05:02","05:08",...: 27 20 25 14 2 20 17 26 19 21 ...
##   $ CutFinal   : Factor w/ 28 levels "11:40","14:43",...: 25 26 21 14 6 11 23 22 24 20 ...
##   $ ChildSex   : Factor w/ 2 levels "Female","Male": 1 1 2 1 1 1 1 1 1 2 ...
##   $ ParentSex  : Factor w/ 2 levels "Female","Male": 2 2 1 1 1 1 1 1 1 1 ...

# Change the vector in character and cut the string in two parts minutes and second for the 4 time labels
cutFrames$CutBefore <- as.character(cutFrames$CutBefore)
cutFramesCB <- strsplit(cutFrames$CutBefore, split=":")

# Compute the time in minutes from time in minutes and seconds for each video for Cut Before
Cut <- c()
for (i in 1:nrow(cutFrames)){
  CutBeforeAlone <- (as.numeric(cutFramesCB[i][[1]][1]) + as.numeric(cutFramesCB[i][[1]][2])/60)
  Cut <- c(Cut, CutBeforeAlone)
}
cutFrames$CutBeforeMin <- Cut

# Compute the time in minutes from time in minutes and seconds for each video for Cut CutMiddle1
Cut <- c()
cutFrames$CutMiddle1 <- as.character(cutFrames$CutMiddle1)
cutMiddleSplit <- strsplit(cutFrames$CutMiddle1, split=":")
for (i in 1:nrow(cutFrames)){
  CutAlone <- (as.numeric(cutMiddleSplit[i][[1]][1]) + as.numeric(cutMiddleSplit[i][[1]][2])/60)
  Cut <- c(Cut, CutAlone)
}
cutFrames$CutMiddle1Min <- Cut

# Compute the time in minutes from time in minutes and seconds for each video for Cut CutMiddle2
Cut <- c()
cutFrames$CutMiddle2 <- as.character(cutFrames$CutMiddle2)
cutMiddleSplit <- strsplit(cutFrames$CutMiddle2, split=":")
for (i in 1:nrow(cutFrames)){
  CutAlone <- (as.numeric(cutMiddleSplit[i][[1]][1]) + as.numeric(cutMiddleSplit[i][[1]][2])/60)
  Cut <- c(Cut, CutAlone)
}
cutFrames$CutMiddle2Min <- Cut

# Compute the time in minutes from time in minutes and seconds for each video for Cut CutFinal
Cut <- c()
cutFrames$CutFinal <- as.character(cutFrames$CutFinal)
cutSplit <- strsplit(cutFrames$CutFinal, split=":")
for (i in 1:nrow(cutFrames)){
  CutAlone <- (as.numeric(cutSplit[i][[1]][1]) + as.numeric(cutSplit[i][[1]][2])/60)
  Cut <- c(Cut, CutAlone)
}
cutFrames$CutFinalMin <- Cut

```

```
Cut <- c()
```

Data dictionary of cutFrames dataframe

- **family** : code of the family
- **CutBefore** : when the experimenter leave the room and the interaction begin character string in the form min:sec
- **CutMiddle1** : when the experimenter come back to explain that the participants are asked to have a conflictual discussion, character string in the form min:sec
- **CutMiddle2** : when the experimenter leave and the conflictual discussion begin Between is the conflictual discussion, character string in the form min:sec
- **CutFinal** : when the experimenter come back to shut down the camera, character string in the form min:sec
- **ChildSex** : Factor variable : Male or Female
- **ParentSex** : Factor variable : Male or Female
- **CutBeforeMin** : when the experimenter leave the room and the interaction begin numeric variable in minutes
- **CutMiddle1Min** : when the experimenter come back to explain that the participants are asked to have a conflictual discussion, numeric variable in minutes
- **CutMiddle2Min** : when the experimenter leave and the conflictual discussion begin Between is the conflictual discussion, numeric variable in minutes
- **CutFinalMin** : when the experimenter come back to shut down the camera, numeric variable in minutes

SyncPy utilisation for creating synchrony dataframe

After extracting filtered motion history with mean on sliding interval (overlapping interval) of 5 frames And after putting 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 SSIXXX.csv with XXXX the name of the video (1606, BAJE059 etc) that we can import with R with this following function

```
# Create a list with the file names
print("SSI Files Directory")

## [1] "SSI Files Directory"
SSInoLogFilesList <- list.files("../Data/CSV/Synchrony/noLog/S_estimator/noShuffle", full.name=TRUE)
#SSInoLogFilesList

# Import and merge the data
SSInoLog <- data.frame(video="Name")
for (file in SSInoLogFilesList){
  # print(file)
  SSIAalone <- read.csv(file)
  # print(str(SSIAalone))
  SSInoLog<- rbind.fill(SSInoLog, SSIAalone)}
SSInoLog$video <-as.factor(SSInoLog$video)
SSInoLog <- SSInoLog[-which(SSInoLog$video=="Name"),]
SSInoLog$Interval <- NULL
SSInoLog <- rename (SSInoLog, c("video" = "family"))
```

```
SSIoLog <- rename (SSIoLog, c("X" = "SSI-interval"))
SSIoLog$SSI<-rowSums(SSIoLog[, c("SSI_fa_ch", "SSI_mo_ch")], na.rm=T)
```

Annotating data frames noLog

```
SSIoLog <- merge(SSIoLog, cutFrames, by.x="family", by.y="family")
SSIoLog$LabelVideo <- rep(NA, nrow(SSIoLog))
SSIoLog[which(SSIoLog$Time_min <= SSIoLog$CutBeforeMin),]$LabelVideo <- "Cut"

SSIoLog[which(SSIoLog$Time_min > SSIoLog$CutBeforeMin & SSIoLog$Time_min < SSIoLog$CutMiddle1Min),]$LabelVideo <- "No Conflict"

SSIoLog[which(SSIoLog$Time_min >= SSIoLog$CutMiddle1Min & SSIoLog$Time_min <= SSIoLog$CutMiddle2Min),]$LabelVideo <- "Conflict"

SSIoLog[which(SSIoLog$Time_min > SSIoLog$CutMiddle2Min & SSIoLog$Time_min < SSIoLog$CutFinalMin),]$LabelVideo <- "Cut"

SSIoLog[which(SSIoLog$Time_min >= SSIoLog$CutFinalMin),]$LabelVideo <- "Cut"
```

Import motion history (MH)

```
MH_list <- list.files("../Data/CSV/filtered/noLog/noshuffled", full.names = TRUE)

MH <- data.frame(video="Name")

for (file in MH_list){
  # print(file)
  MHalone <- read.csv(file)
  # print(str(SSIAalone))
  MH<- rbind.fill(MH, MHalone)}
MH <- MH[-which(MH$video=="Name"),]
MH$X <- NULL
MH <- rename (MH, c("video" = "family"))
MH$slidedParent<-rowSums(MH[, c("slidedFather", "slidedMother")], na.rm=T)
```

Annotating dataframe MH

```
#View(MH)
MH <- merge(MH, cutFrames, by.x="family", by.y="family")
MH$LabelVideo <- rep(NA, nrow(MH))
#for 25 frmaes by second and 60 seconds by min
MH$Time_min <- MH$frame_index/(25*60)
MH[which(MH$Time_min <= MH$CutBeforeMin),]$LabelVideo <- "Cut"

MH[which(MH$Time_min > MH$CutBeforeMin & MH$Time_min < MH$CutMiddle1Min),]$LabelVideo <- "No Conflict"

MH[which(MH$Time_min >= MH$CutMiddle1Min & MH$Time_min <= MH$CutMiddle2Min),]$LabelVideo <- "Conflict"

MH[which(MH$Time_min > MH$CutMiddle2Min & MH$Time_min < MH$CutFinalMin),]$LabelVideo <- "Conflict"
```

```
MH[which(MH$Time_min >= MH$CutFinalMin),]$LabelVideo <- "Cut"
```

Description of noLogSSI data frame

```
str(SSInoLog)
```

```
## 'data.frame': 6404 obs. of 17 variables:
## $ family : Factor w/ 35 levels "1606","BAJE059",...: 1 1 1 1 1 1 1 1 1 ...
## $ SSI-interval : int 0 1 2 3 4 5 6 7 8 9 ...
## $ Time_min : num 0 0.0833 0.1667 0.25 0.3333 ...
## $ SSI_fa_ch : num 0.06532 0.00806 0.1161 0.11668 0.18393 ...
## $ SSI_mo_ch : num NA NA NA NA NA NA NA NA NA ...
## $ SSI : num 0.06532 0.00806 0.1161 0.11668 0.18393 ...
## $ CutBefore : chr "00:11" "00:11" "00:11" "00:11" ...
## $ CutMiddle1 : chr "05:30" "05:30" "05:30" "05:30" ...
## $ CutMiddle2 : chr "06:16" "06:16" "06:16" "06:16" ...
## $ CutFinal : chr "16:27" "16:27" "16:27" "16:27" ...
## $ ChildSex : Factor w/ 2 levels "Female","Male": 1 1 1 1 1 1 1 1 1 ...
## $ ParentSex : Factor w/ 2 levels "Female","Male": 2 2 2 2 2 2 2 2 2 ...
## $ CutBeforeMin : num 0.183 0.183 0.183 0.183 0.183 ...
## $ CutMiddle1Min: num 5.5 5.5 5.5 5.5 5.5 5.5 5.5 5.5 5.5 ...
## $ CutMiddle2Min: num 6.27 6.27 6.27 6.27 6.27 ...
## $ CutFinalMin : num 16.4 16.4 16.4 16.4 16.4 ...
## $ LabelVideo : chr "Cut" "Cut" "Cut" "No Conflict" ...
```

```
#View(SSInoLog)
```

Data dictionary of SSInolog data frame

- **family** : code of the family
- **SSI-interval** : interval of SSI
- **Time_min** : Time in minutes
- **SSI_fa_ch** : SSI index of Synchrony between father and child
- **SSI_mo_ch** : SSI index of Synchrony between mother and child
- **SSI** : SSI index of the interaction (father-child or mother-child)
- **CutBefore** : when the experimenter leave the room and the interaction begin character string in the form min:sec
- **CutMiddle1** : when the experimenter come back to explain that the participants are asked to have a conflictual discussion, character string in the form min:sec
- **CutMiddle2** : when the experimenter leave and the conflictual discussion begin Between is the conflictual discussion, character string in the form min:sec
- **CutFinal** : when the experimenter come back to shut down the camera, character string in the form min:sec
- **ChildSex** : Factor variable : Male or Female
- **ParentSex** : Factor variable : Male or Female
- **CutBeforeMin** : when the experimenter leave the room and the interaction begin numeric variable in minutes
- **CutMiddle1Min** : when the experimenter come back to explain that the participants are asked to have a conflictual discussion, numeric variable in minutes
- **CutMiddle2Min** : when the experimenter leave and the conflictual discussion begin Between is the conflictual discussion, numeric variable in minutes

- **CutFinalMin** : when the experimenter come back to shut down the camera, numeric variable in minutes
- **LabelVideo** : label of the video if it is related to No Conflict period or Conflict Period

Description of MH data frame

```
str(MH)
```

```
## 'data.frame': 802834 obs. of 18 variables:
## $ family      : chr "1606" "1606" "1606" "1606" ...
## $ slidedFather : num 0.002084 0.001758 0.000729 0.000693 0.000537 ...
## $ slidedMother : num NA NA NA NA NA NA NA NA NA ...
## $ slidedChild  : num 2.57e-03 1.55e-03 2.54e-04 2.09e-04 5.28e-05 ...
## $ frame_index  : int 1 2 3 4 5 6 7 8 9 10 ...
## $ slidedParent : num 0.002084 0.001758 0.000729 0.000693 0.000537 ...
## $ CutBefore    : chr "00:11" "00:11" "00:11" "00:11" ...
## $ CutMiddle1   : chr "05:30" "05:30" "05:30" "05:30" ...
## $ CutMiddle2   : chr "06:16" "06:16" "06:16" "06:16" ...
## $ CutFinal     : chr "16:27" "16:27" "16:27" "16:27" ...
## $ ChildSex     : Factor w/ 2 levels "Female","Male": 1 1 1 1 1 1 1 1 1 ...
## $ ParentSex    : Factor w/ 2 levels "Female","Male": 2 2 2 2 2 2 2 2 2 ...
## $ CutBeforeMin : num 0.183 0.183 0.183 0.183 0.183 ...
## $ CutMiddle1Min: num 5.5 5.5 5.5 5.5 5.5 5.5 5.5 5.5 5.5 ...
## $ CutMiddle2Min: num 6.27 6.27 6.27 6.27 6.27 ...
## $ CutFinalMin  : num 16.4 16.4 16.4 16.4 16.4 ...
## $ LabelVideo   : chr "Cut" "Cut" "Cut" "Cut" ...
## $ Time_min     : num 0.000667 0.001333 0.002 0.002667 0.003333 ...
```

```
#View(SSInoLog)
```

Data dictionary of SSInolog data frame

- **family** : code of the family
- **SlidedFather** : Filtered motion History of the father
- **Time_min** : Time in minutes
- ****SlidedMother*** : Filtered Motion History of the mother
- **SlidedChild** : Filtered Motion History of the child
- ****SlidedParent*** : Filtered Motion History of the parent (either mother or father)
- ****frame index*** : index of the frame (from 1st to last frame of each video)
- **CutBefore** : when the experimenter leave the room and the interaction begin character string in the form min:sec
- **CutMiddle1** : when the experimenter come back to explain that the participants are asked to have a conflictual discussion, character string in the form min:sec
- **CutMiddle2** : when the experimenter leave and the conflictual discussion begin Between is the conflictual discussion, character string in the form min:sec
- **CutFinal** : when the experimenter come back to shut down the camera, character string in the form min:sec
- **ChildSex** : Factor variable : Male or Female
- **ParentSex** : Factor variable : Male or Female
- **CutBeforeMin** : when the experimenter leave the room and the interaction begin numeric variable in minutes
- **CutMiddle1Min** : when the experimenter come back to explain that the participants are asked to have a conflictual discussion, numeric variable in minutes

- **CutMiddle2Min** : when the experimenter leave and the conflictual discussion begin Between is the conflictual discussion, numeric variable in minutes
- **CutFinalMin** : when the experimenter come back to shut down the camera, numeric variable in minutes
- **LabelVideo** : label of the video if it is related to No Conflict period or Conflict Period

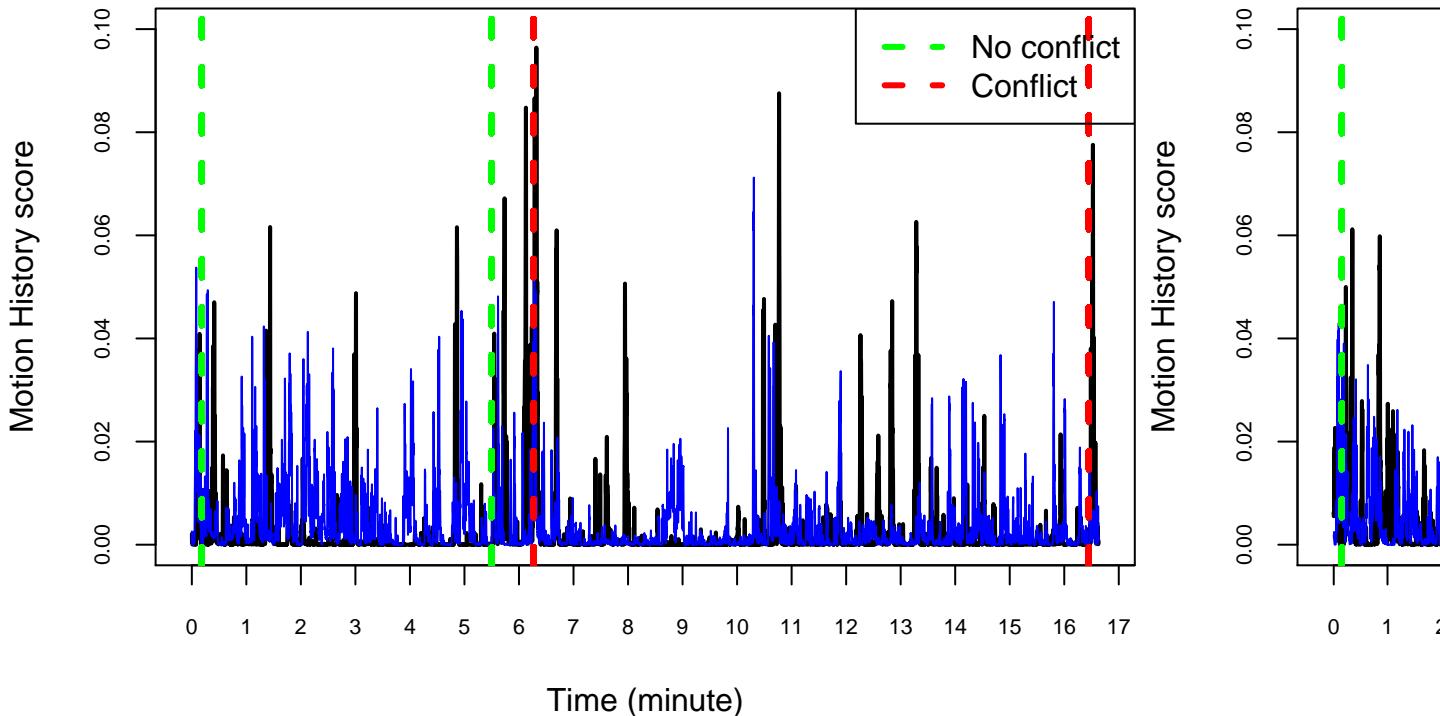
Motion history score for each dyad

```

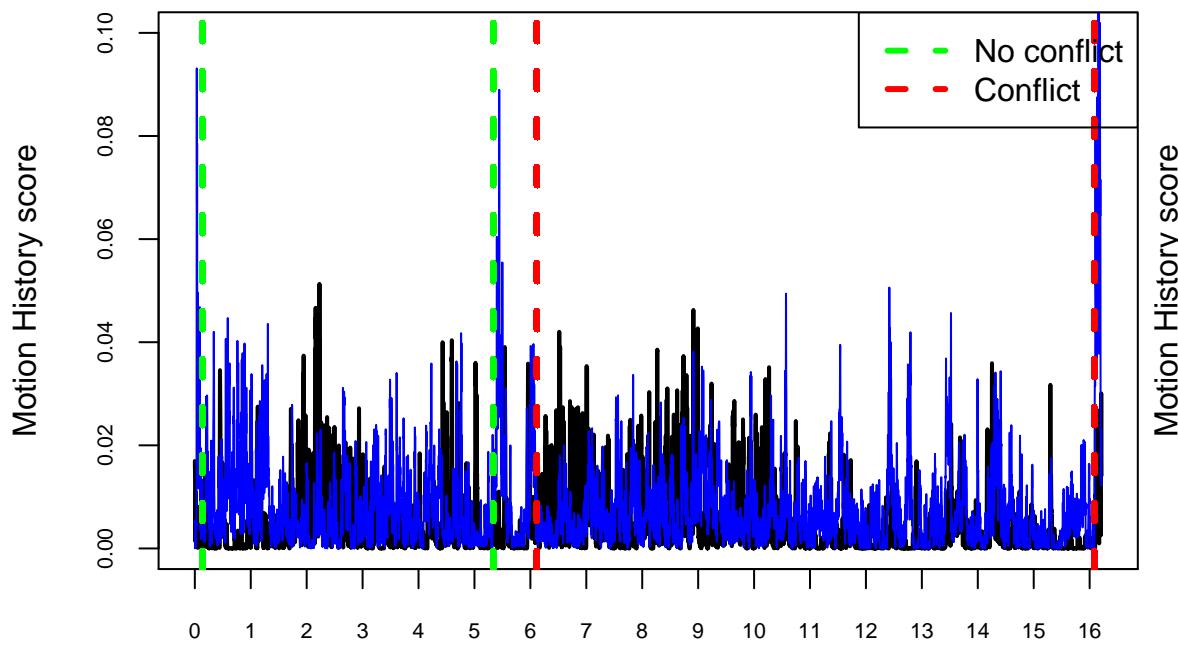
par(mar=c(4,4,4,3))
for (i in unique(MH$family)){
  plot(MH[which(MH$family==i),]$Time_min, (MH[which(MH$family==i),]$slidedParent),
       ylim=c(0, 0.1),
       main=paste("Motion History scores in", i, "family"),
       xlab = "Time (minute)", ylab="Motion History score",
       lwd=2, type="l", cex.axis=0.7, xaxp=c(0,length(MH[which(MH$family==i),]$Time_min)),
       lines(MH[which(MH$family==i),]$Time_min, MH[which(MH$family==i),]$slidedChild, col="blue")
       abline(v= MH[which(MH$family==i),]$CutBeforeMin, col="green", lty=2, lwd=3)
       abline(v= MH[which(MH$family==i),]$CutMiddle1Min, col="green", lty=2, lwd=3)
       abline(v= MH[which(MH$family==i),]$CutMiddle2Min
              , col="red", lty=2, lwd=3)
       abline(v= MH[which(MH$family==i),]$CutFinalMin, col="red", lty=2, lwd=3)
       legend("topright", c("No conflict", "Conflict"), lty=c(2,2), lwd=c(3,3), col=c("green", "red"))
#      a <- paste(as.character(i), ".jpeg", sep="")
#
#      jpeg(filename = a, width = 560, height = 360, units = "px")
}

```

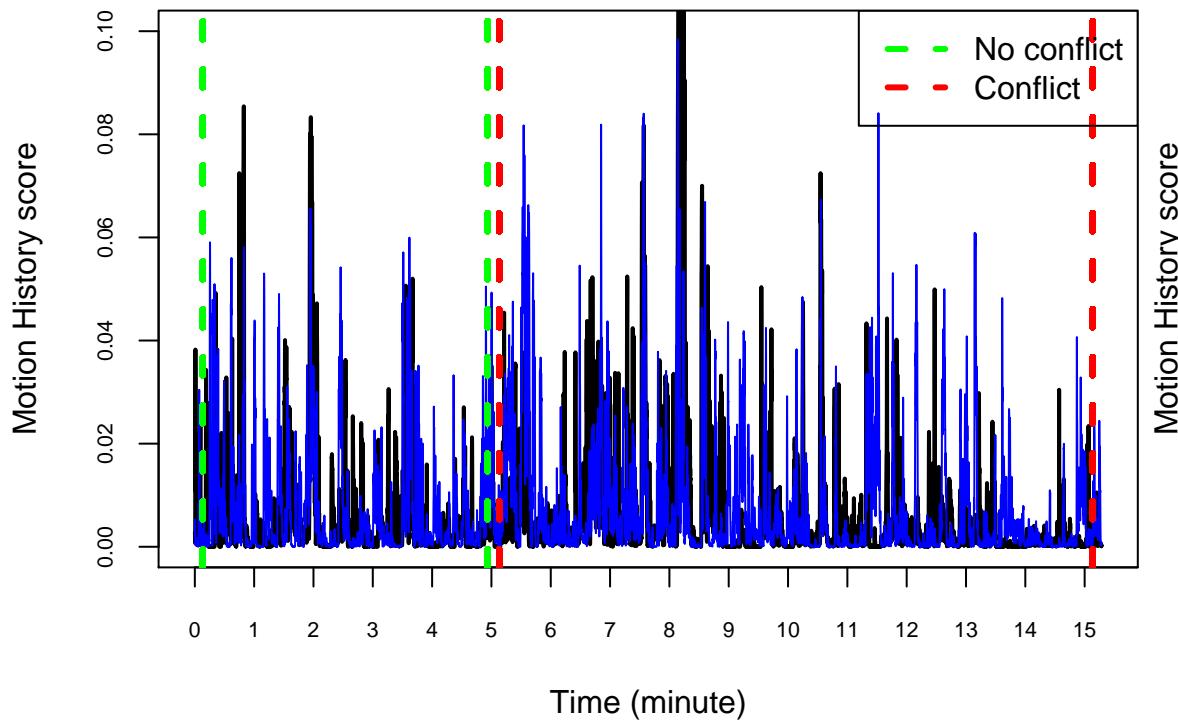
Motion History scores in 1606 family



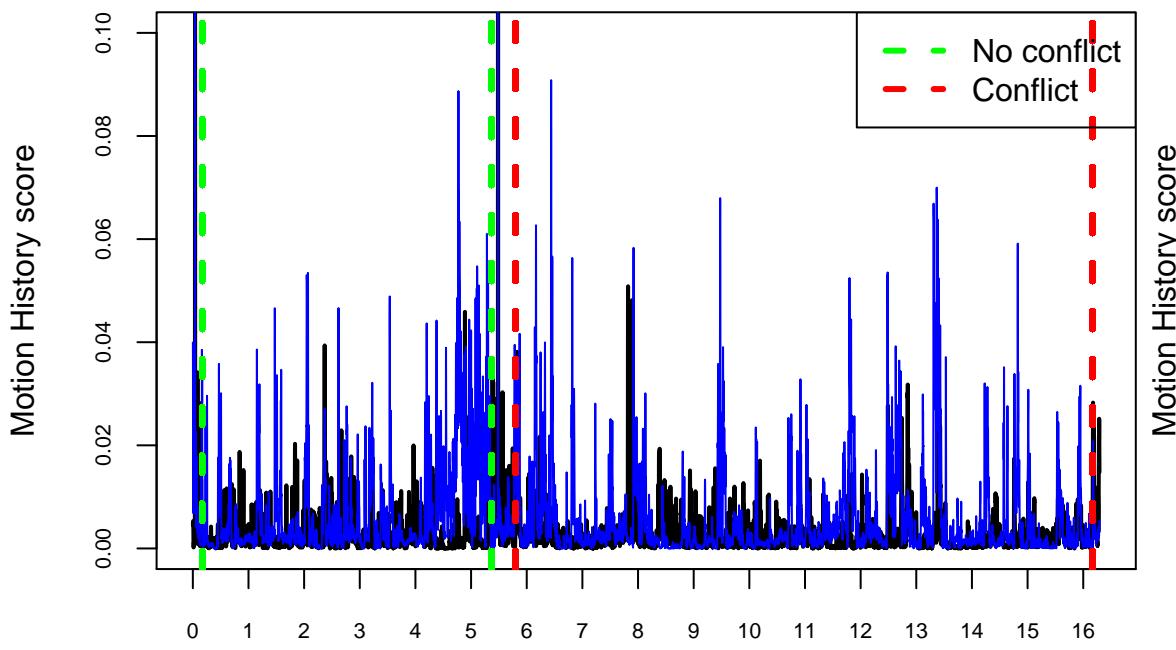
Motion History scores in BALU062 family



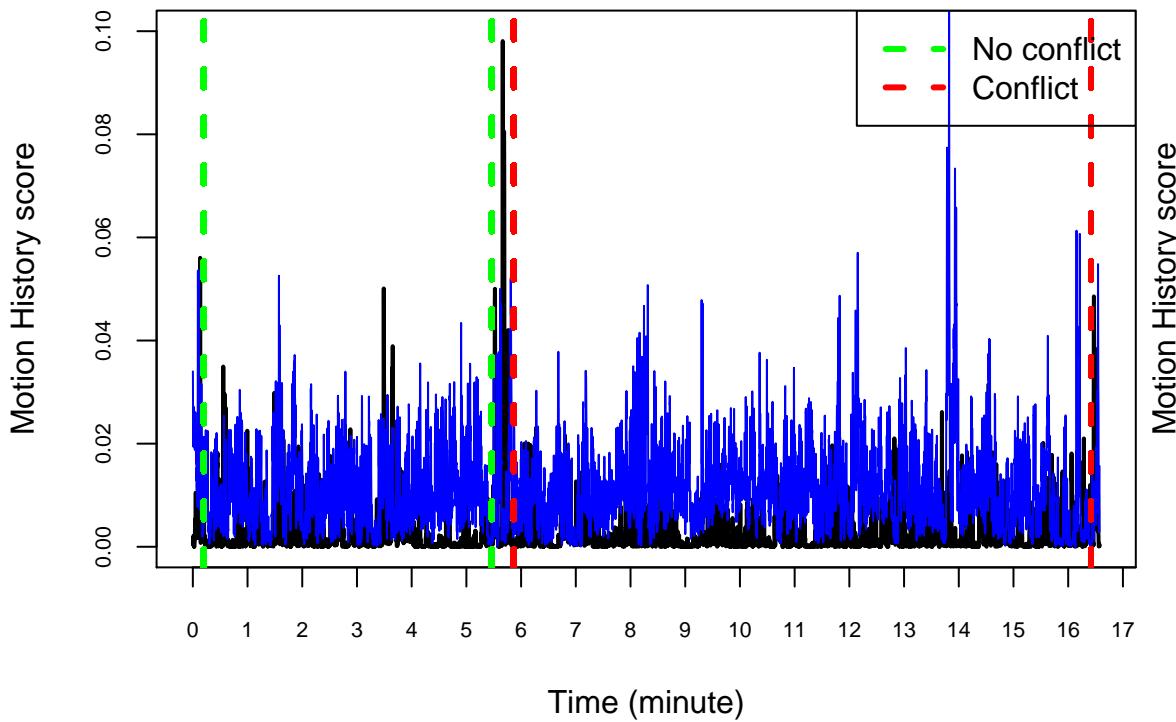
Motion History scores in BEAM031 family



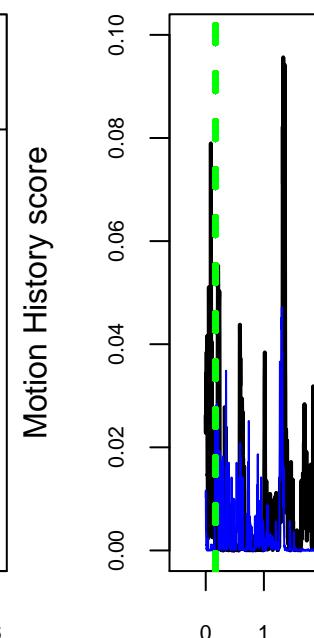
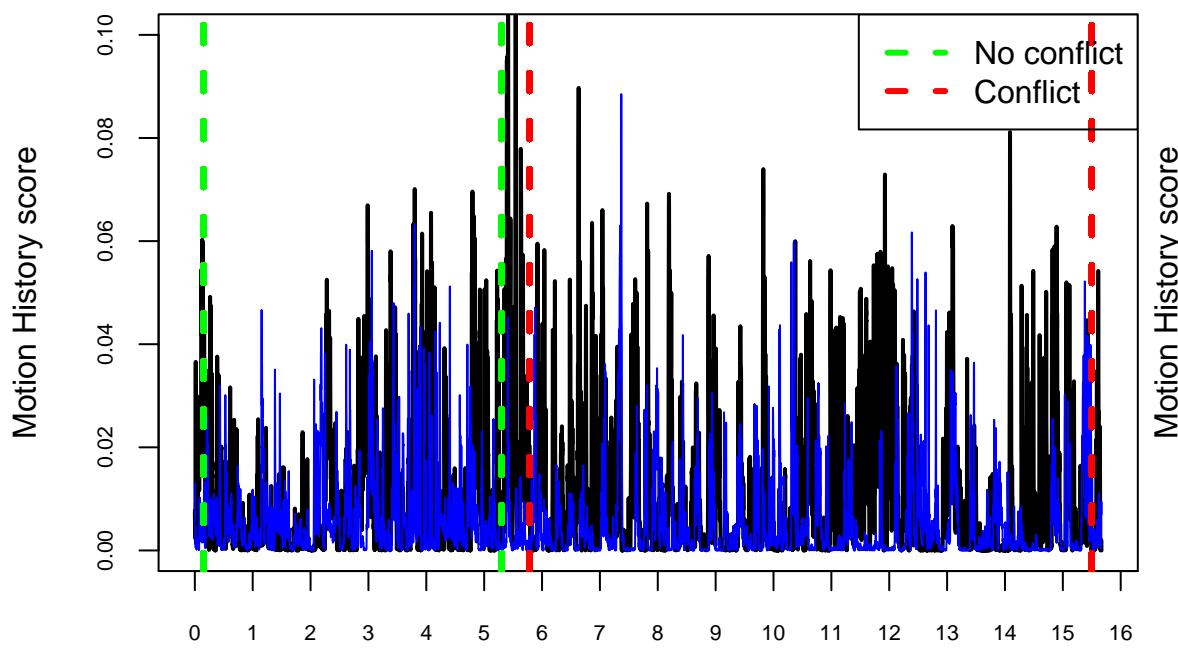
Motion History scores in COLO022 family



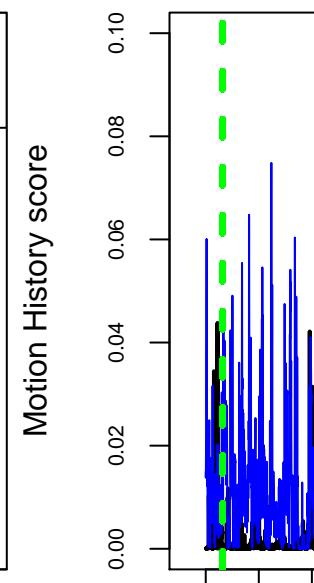
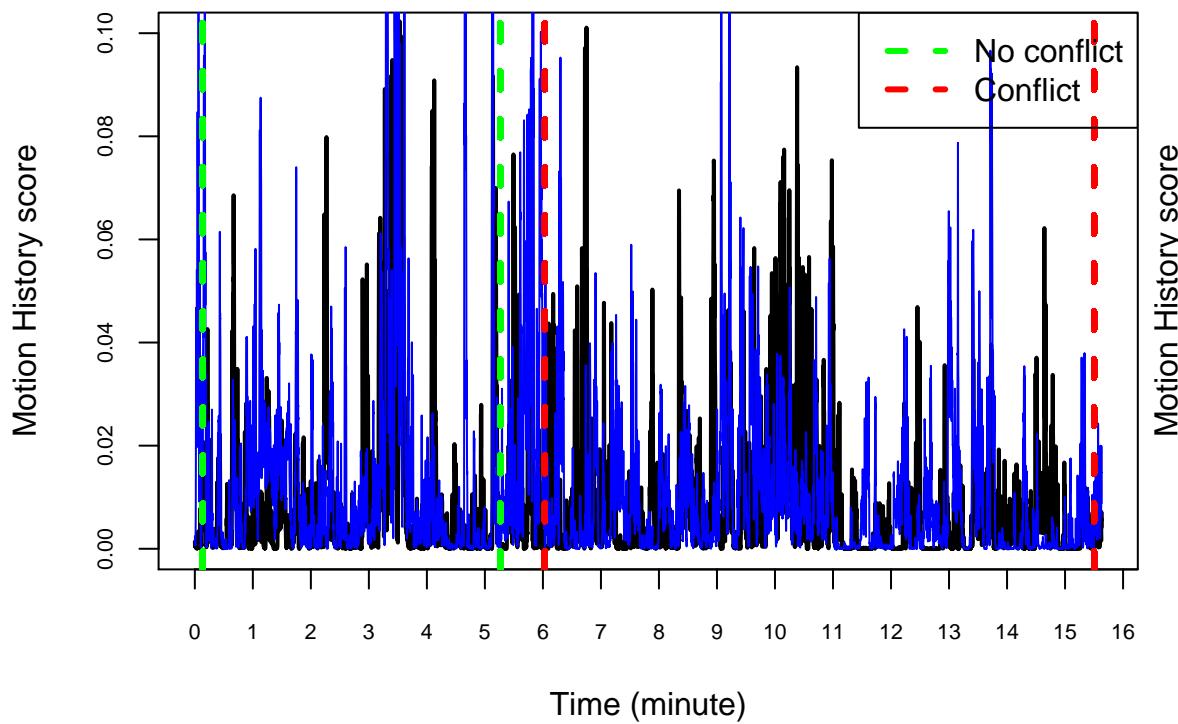
Time (minute)
Motion History scores in DOMA family



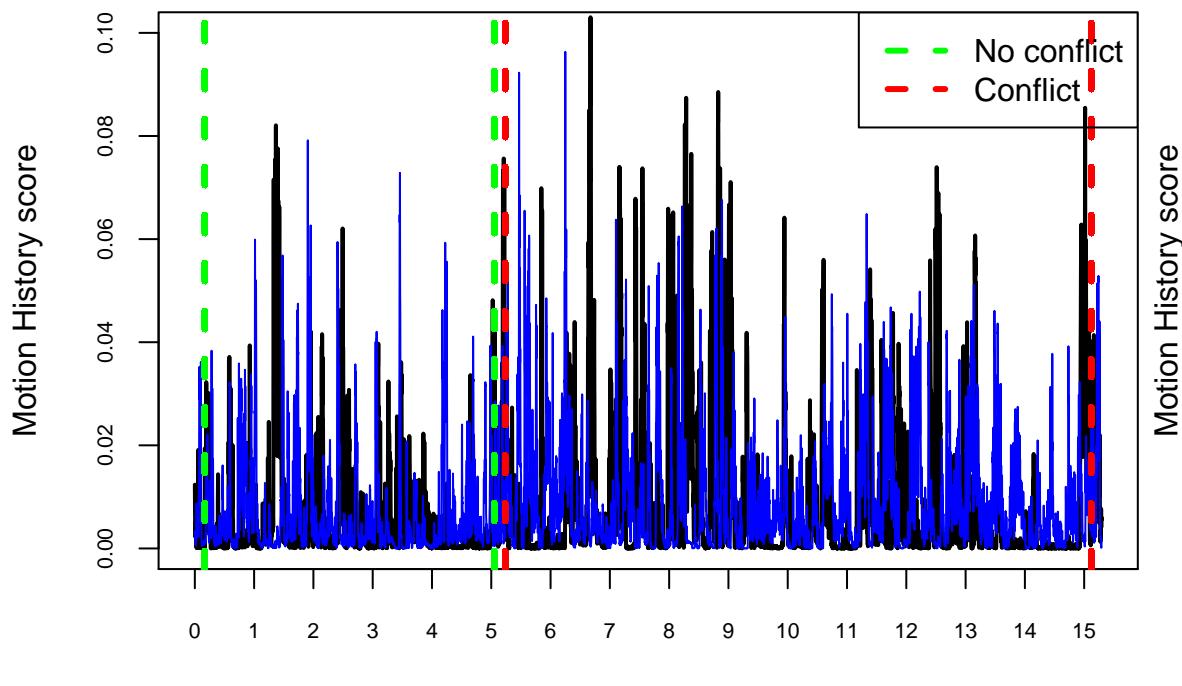
Motion History scores in FOMA057 family



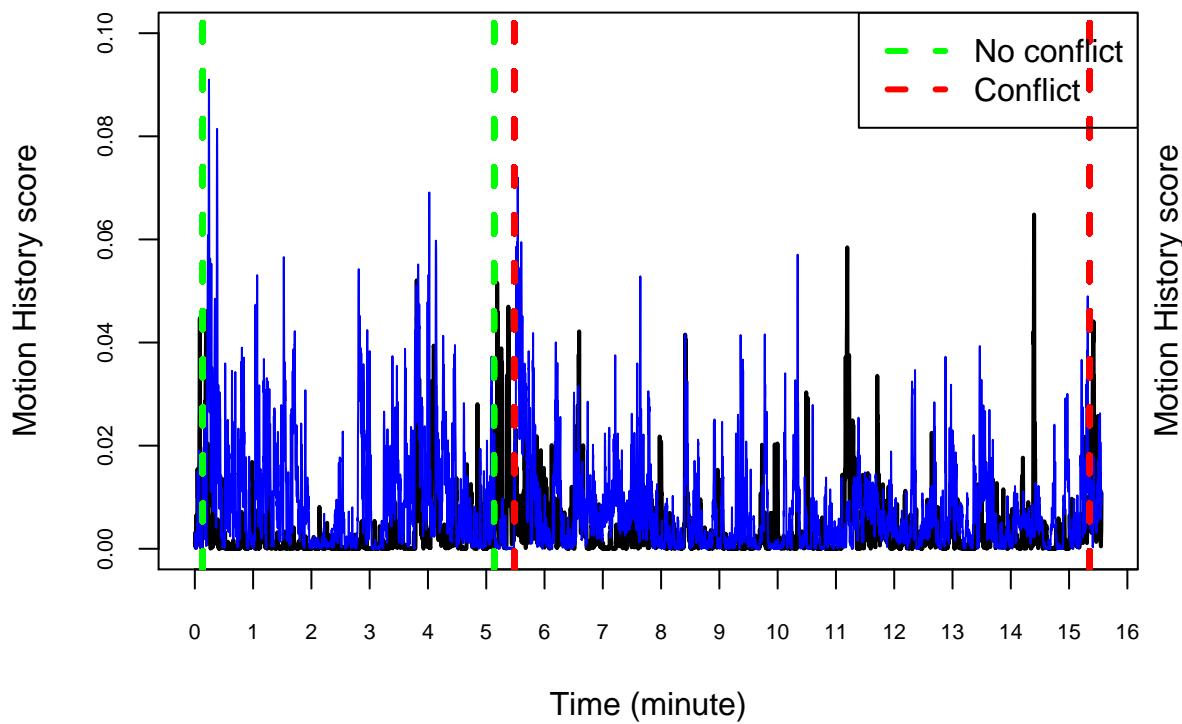
Motion History scores in HAJA052 family



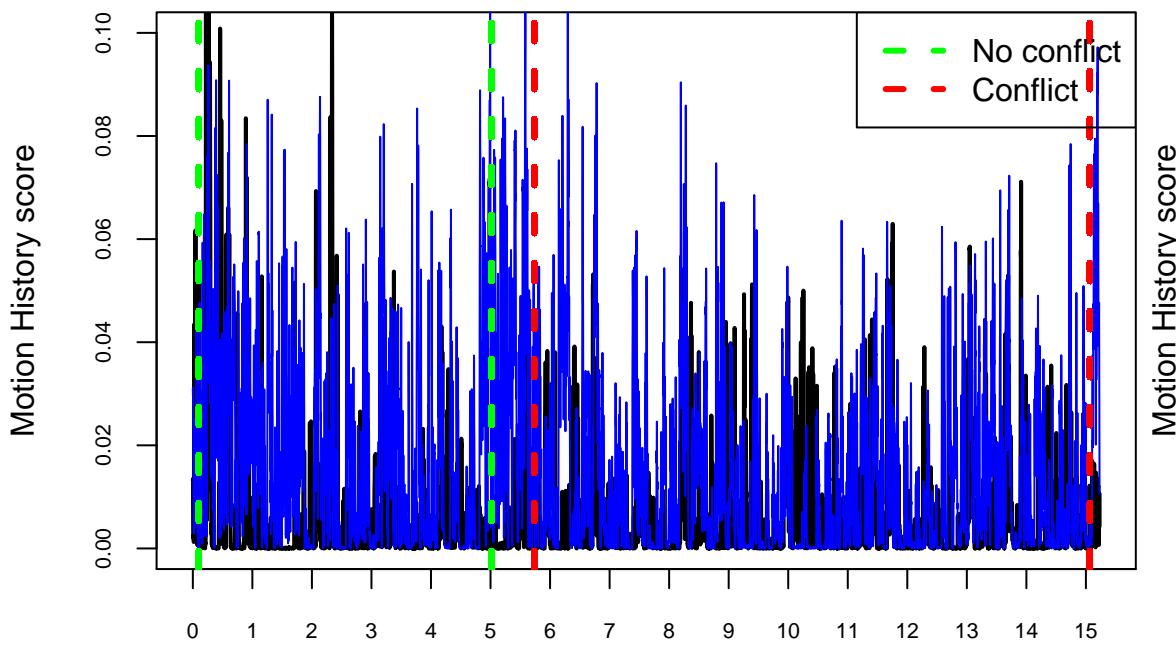
Motion History scores in JAEM046 family



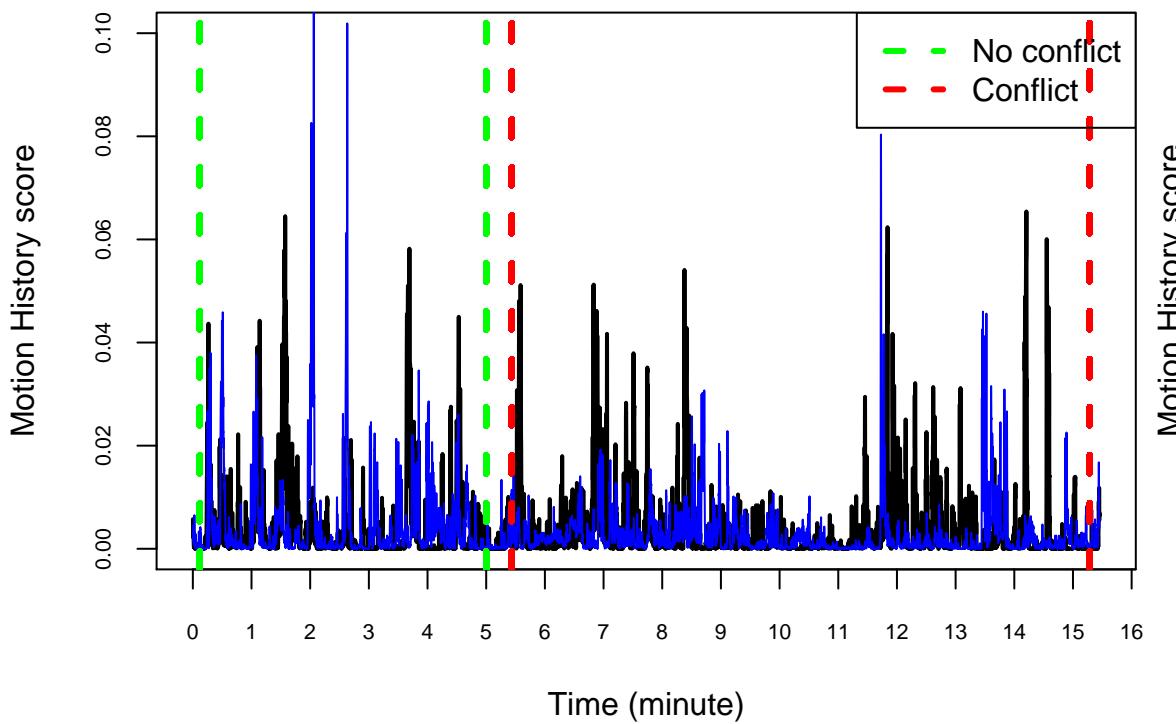
Motion History scores in JOCE014 family



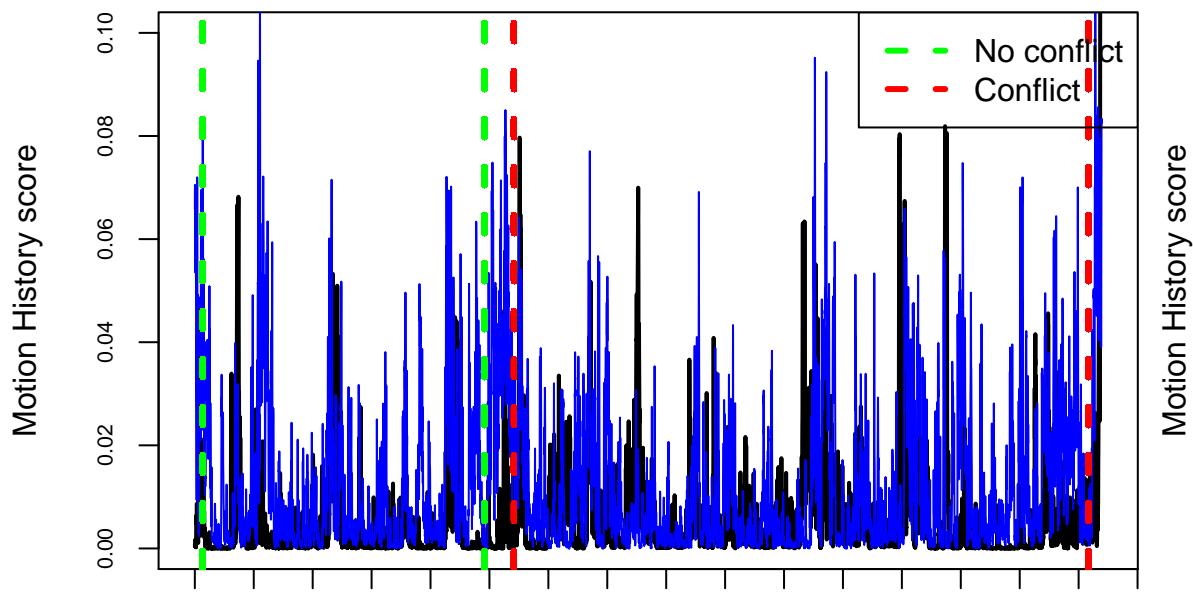
Motion History scores in MAEL048 family



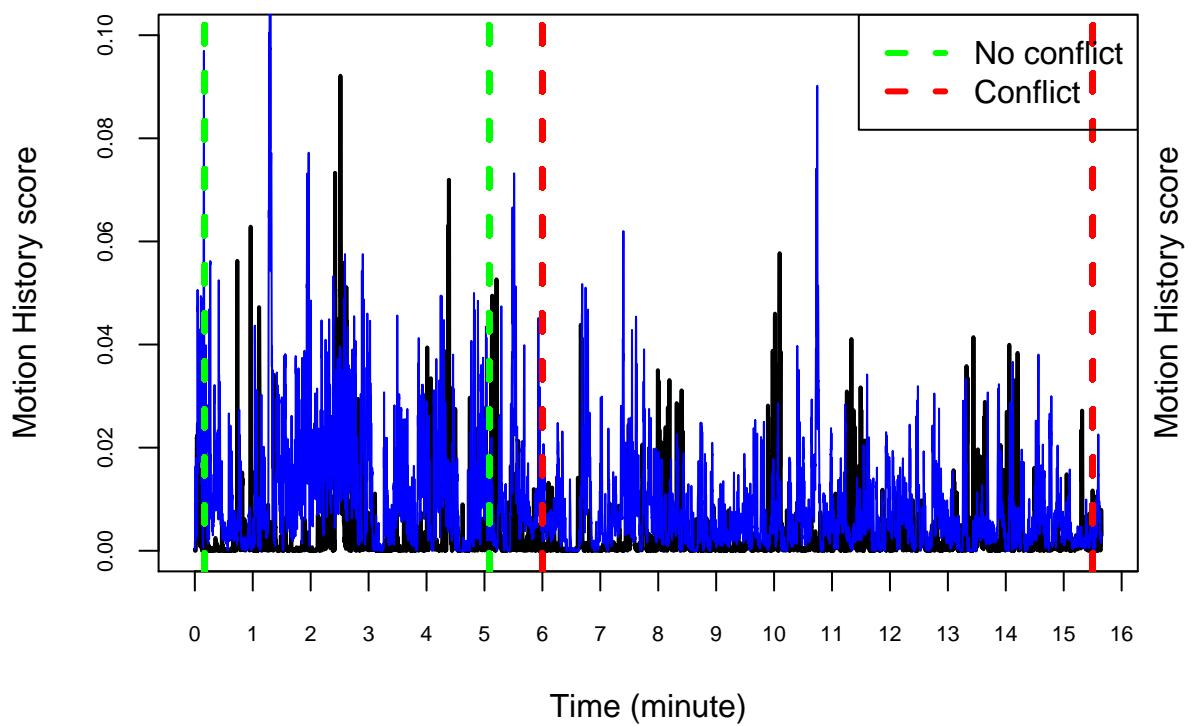
Motion History scores in MIPH043 family



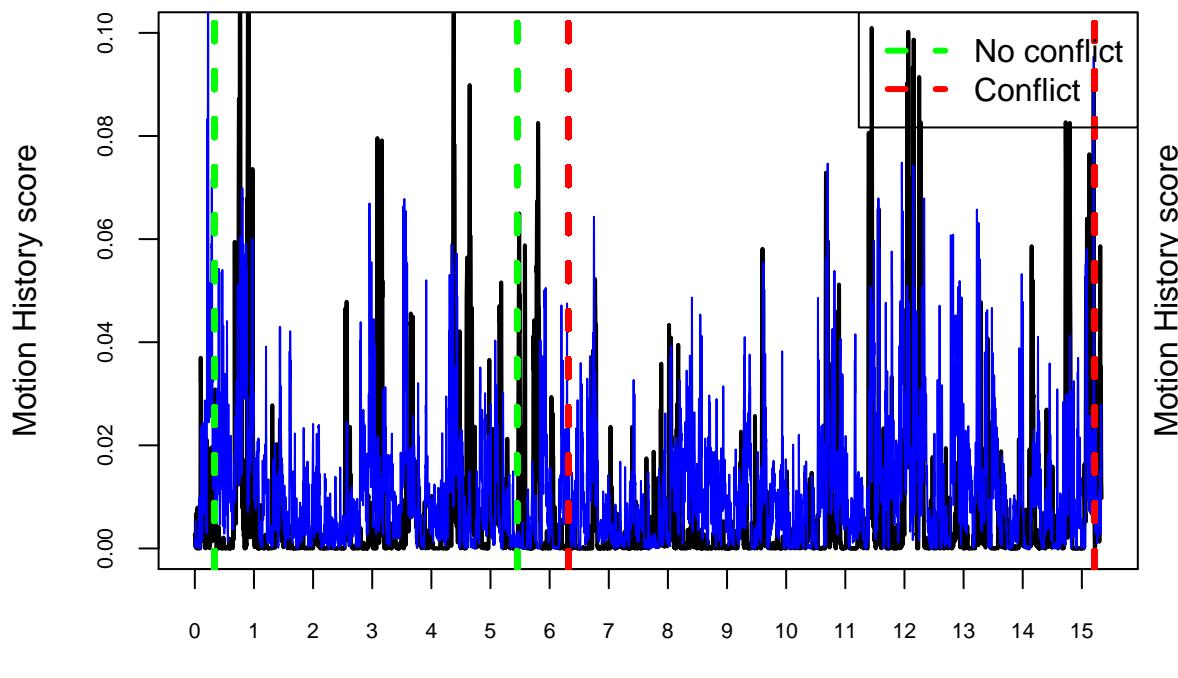
Motion History scores in NAMA045 family



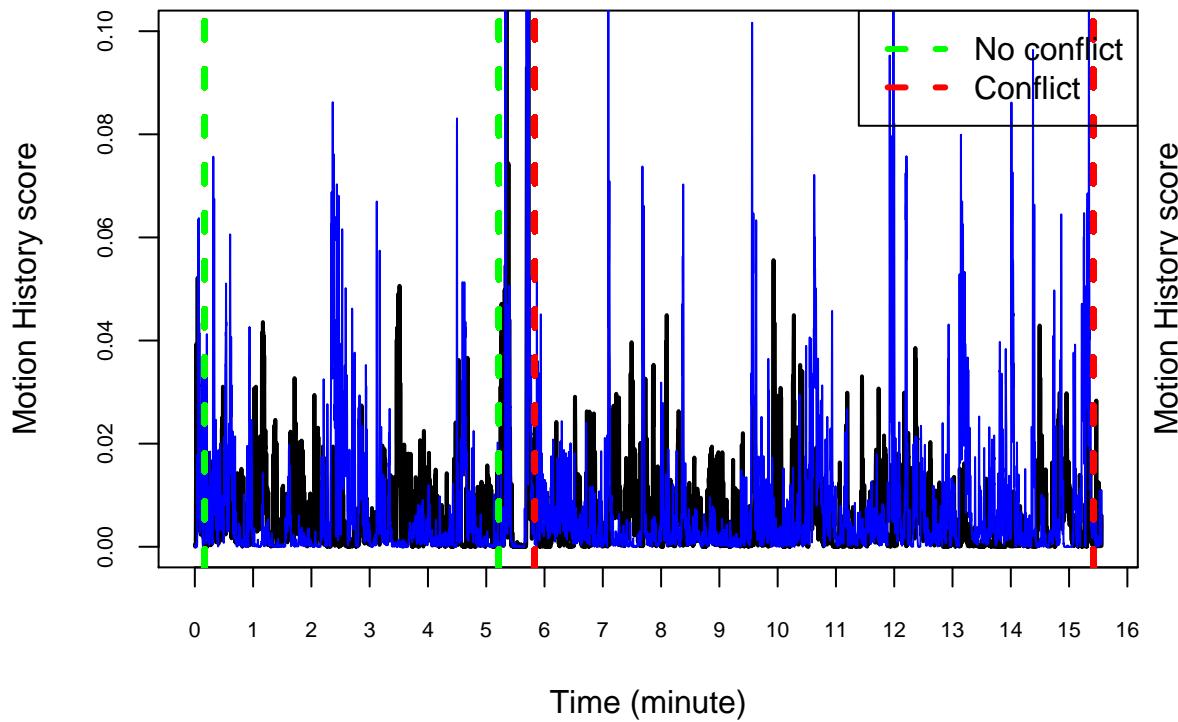
Motion History scores in OGGA034 family



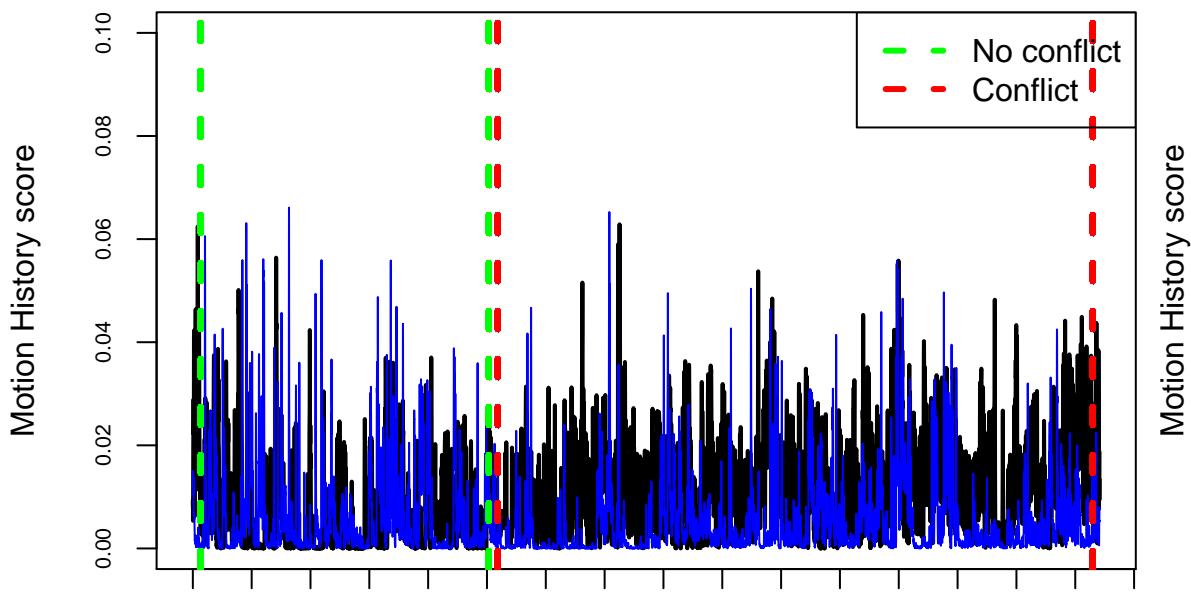
Motion History scores in PELI020 family



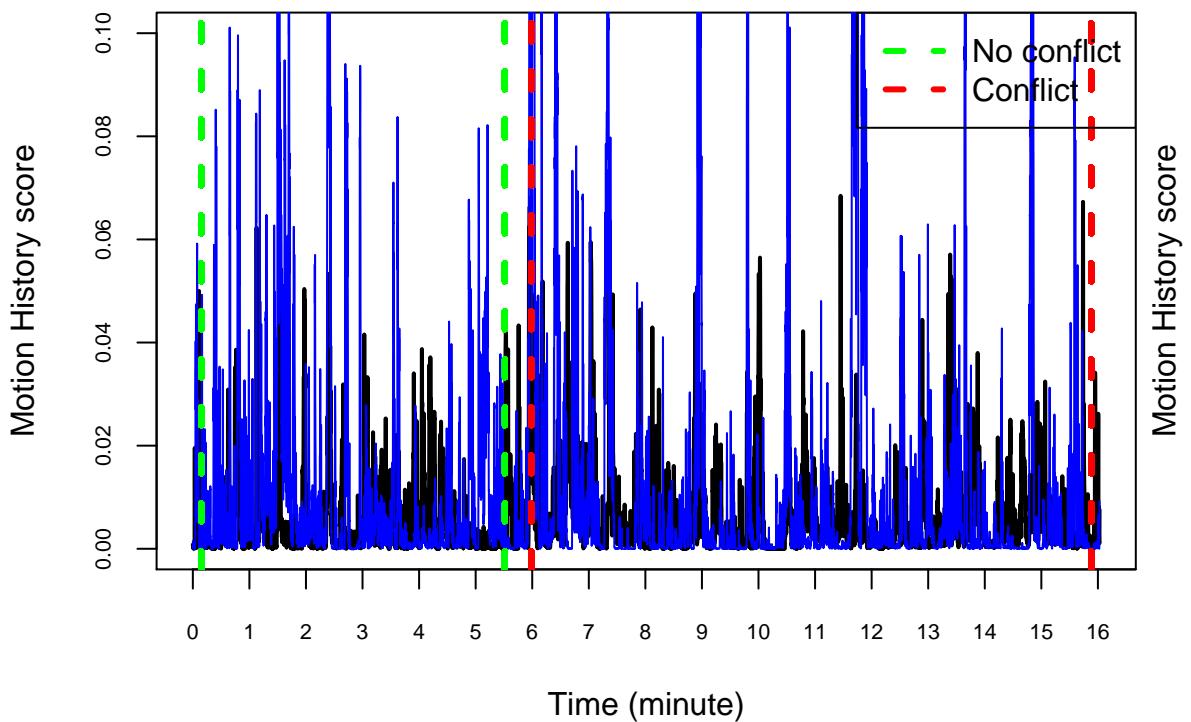
Motion History scores in RAMA054 family



Motion History scores in SHAN042 family



Motion History scores in TIUG032 family



Synchrony scores noLog for each dyad

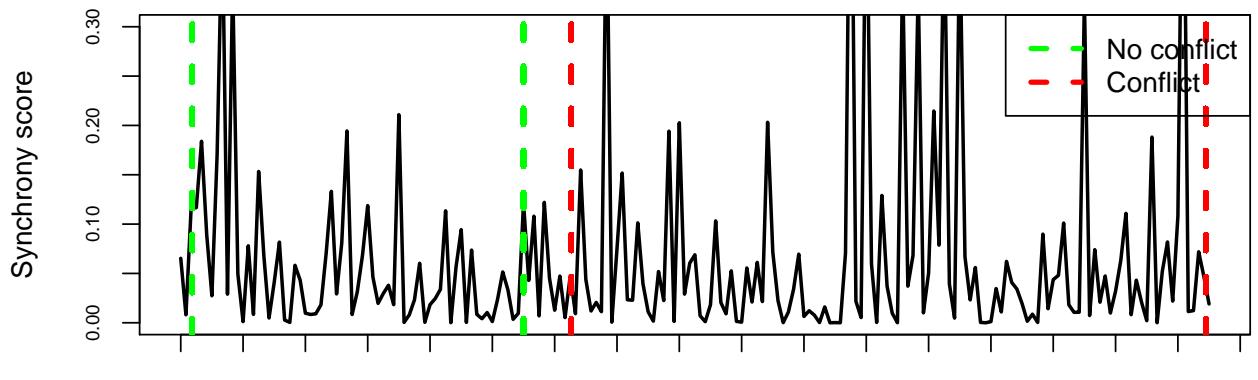
```

par(mar=c(4,4,4,3), mfrow=c(1,1))
for (i in unique(SSInoLog$family)){
  plot(SSInoLog[which(SSInoLog$family==i),]$Time_min,
       SSInoLog[which(SSInoLog$family==i),]$SSI,
       ylim=c(0, 0.3),
       main=paste("Synchrony scores in", i, "family"),
       xlab = "Time (minute)", ylab="Synchrony score",
       lwd=2, type="l", cex.axis=0.7, xaxp=c(0,length(SSInoLog[which(SSInoLog$family==i),]$Time_m
abline(v= SSInoLog[which(SSInoLog$family==i),]$CutBeforeMin, col="green", lty=2, lwd=3)
abline(v= SSInoLog[which(SSInoLog$family==i),]$CutMiddle1Min, col="green", lty=2, lwd=3)
abline(v= SSInoLog[which(SSInoLog$family==i),]$CutMiddle2Min
      , col="red", lty=2, lwd=3)
abline(v= SSInoLog[which(SSInoLog$family==i),]$CutFinalMin, col="red", lty=2, lwd=3)
legend("topright", c("No conflict", "Conflict"), lty=c(2,2), lwd=c(3,3), col=c("green", "red"))
#max(SSInoLog[which(SSInoLog$family==i),]$SSI), col="green")
}

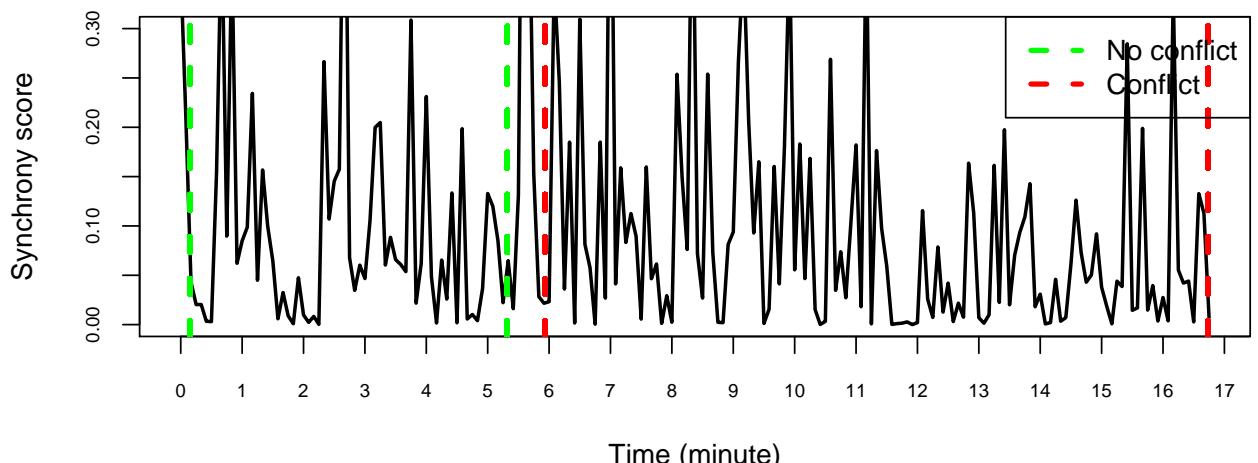
}

```

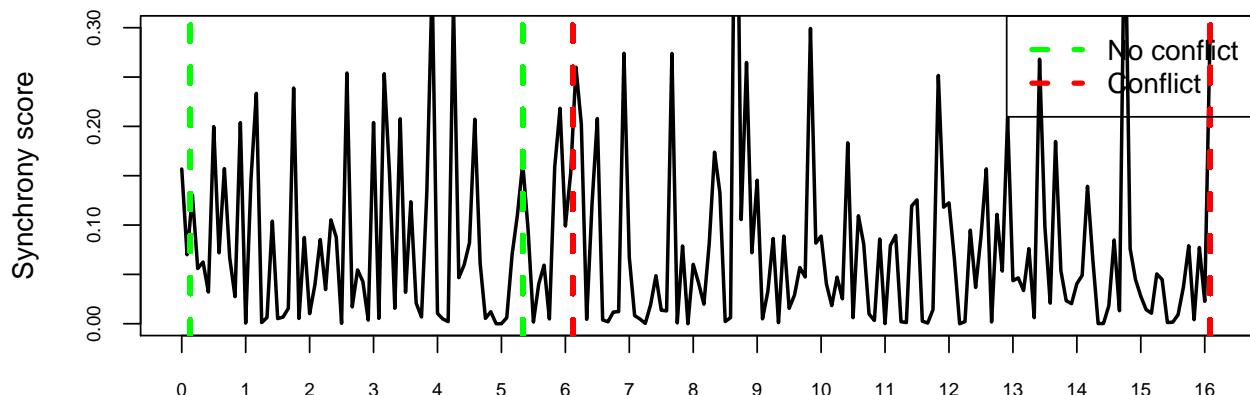
Synchrony scores in 1606 family



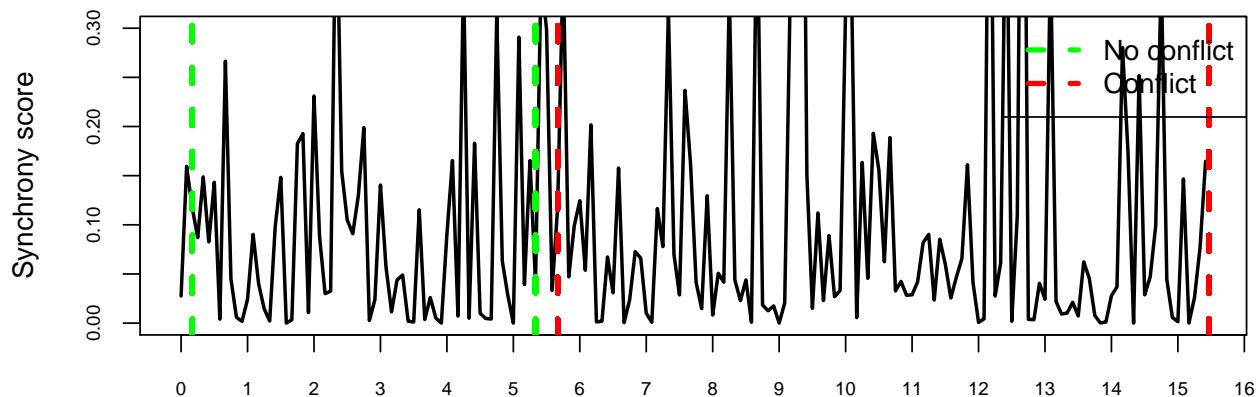
Synchrony scores in BAJE059 family



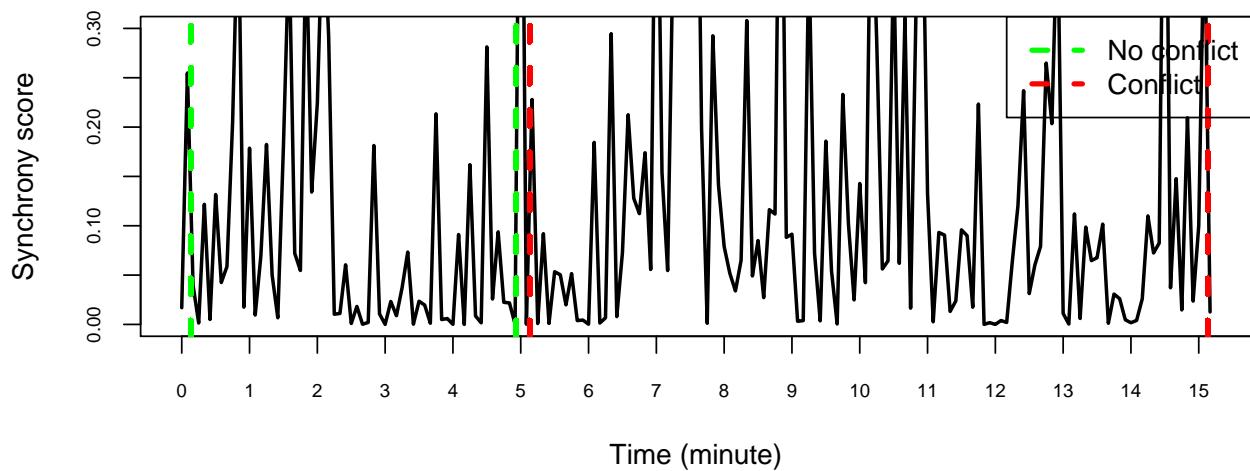
Synchrony scores in BALU062 family



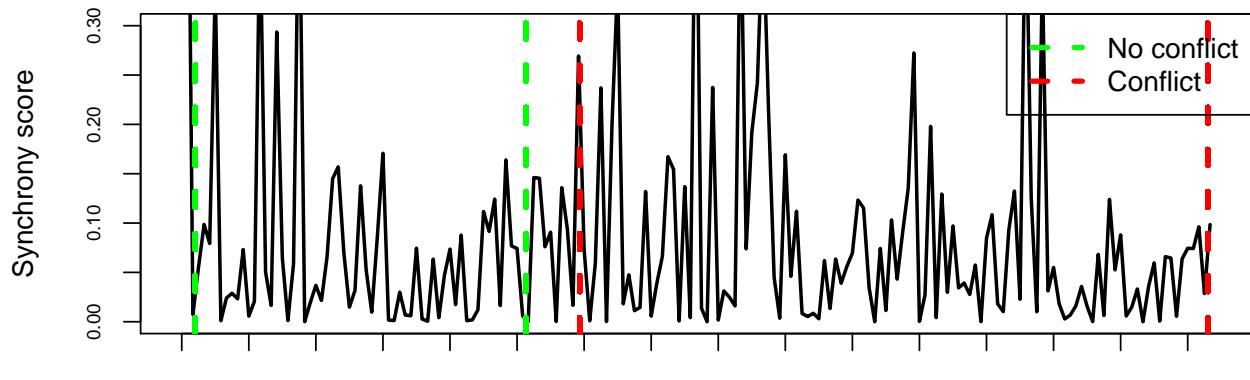
Synchrony scores in BEAL036 family



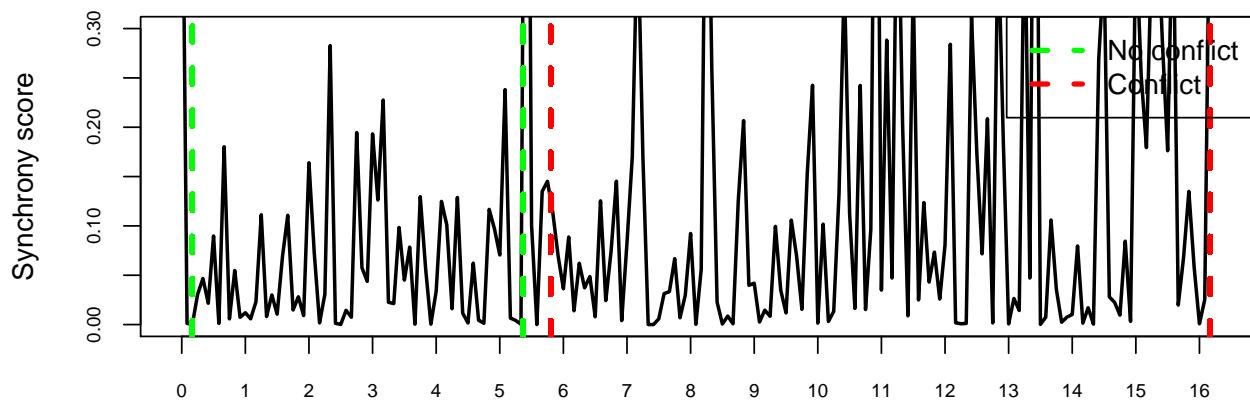
Synchrony scores in BEAM031 family



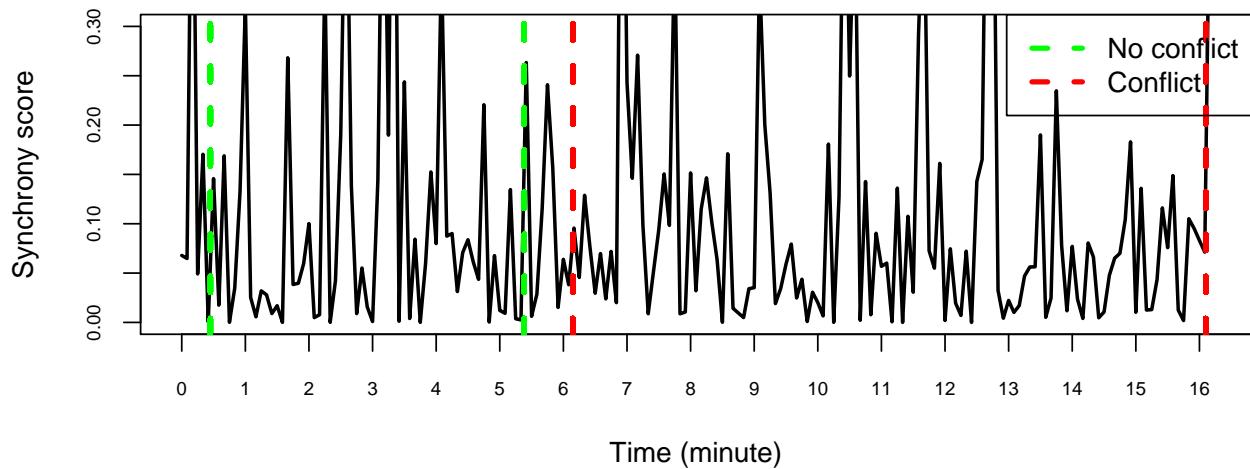
Synchrony scores in BRLO041 family



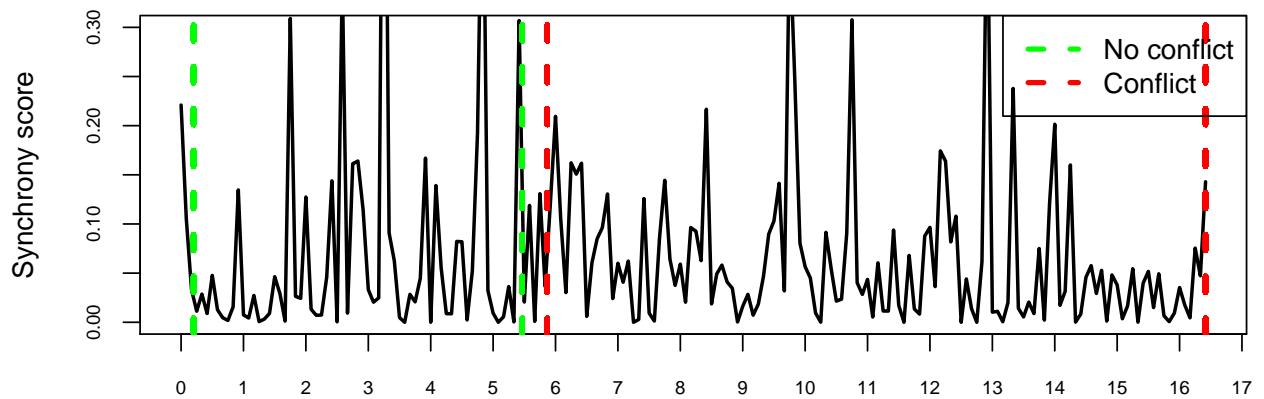
Synchrony scores in COLO022 family



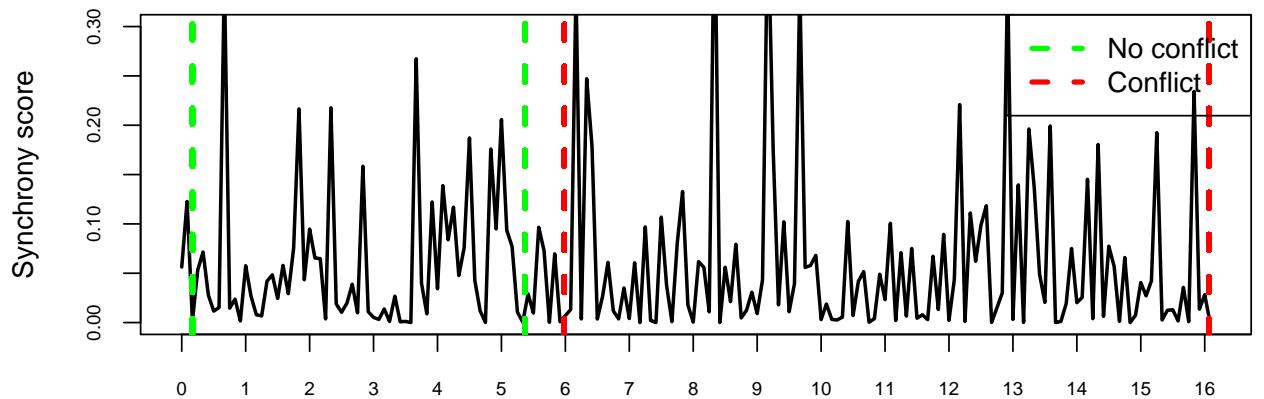
Synchrony scores in DIPE004 family



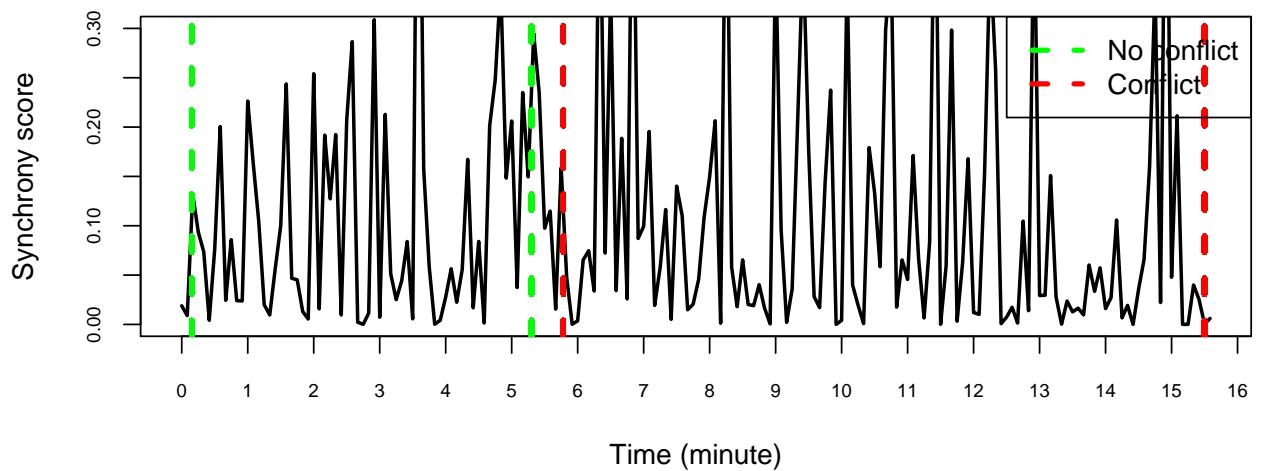
Synchrony scores in DOMA family



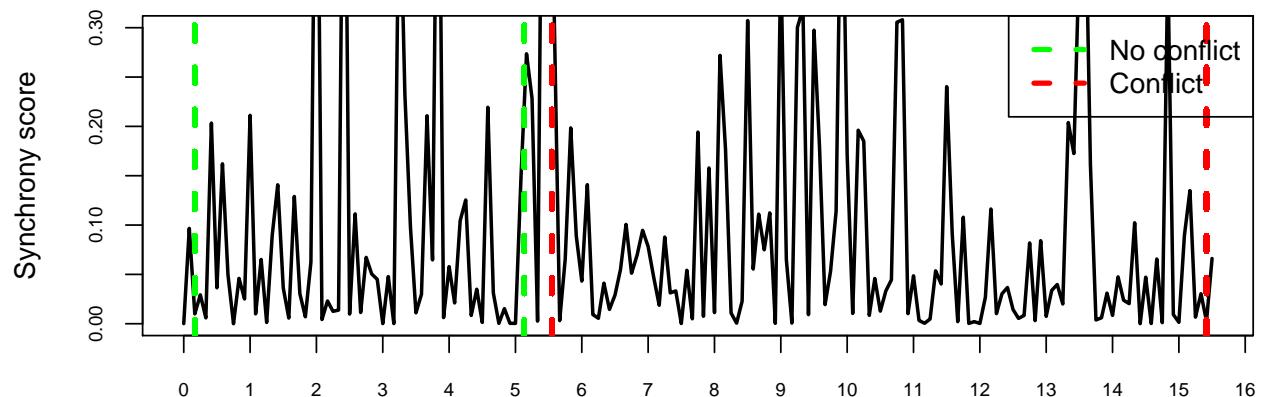
Synchrony scores in DRNE family



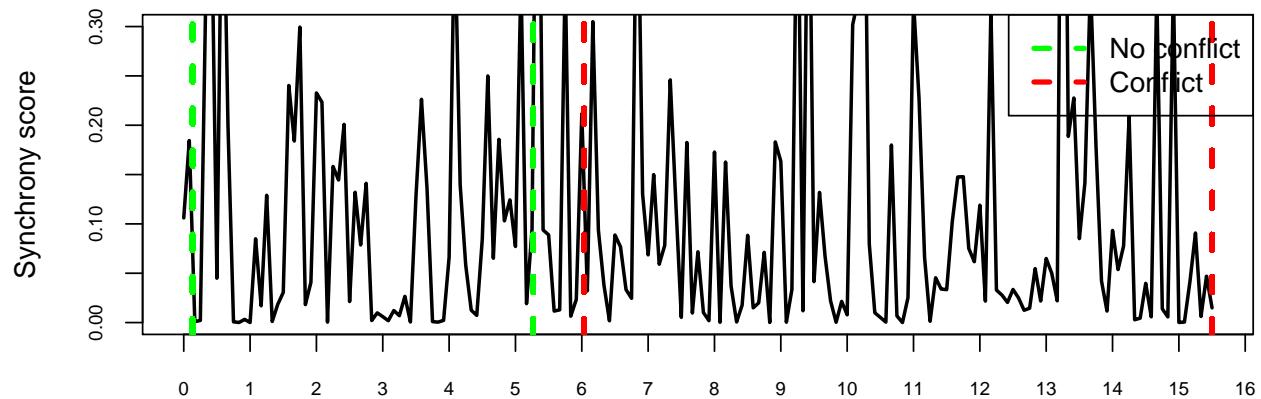
Synchrony scores in FOMA057 family



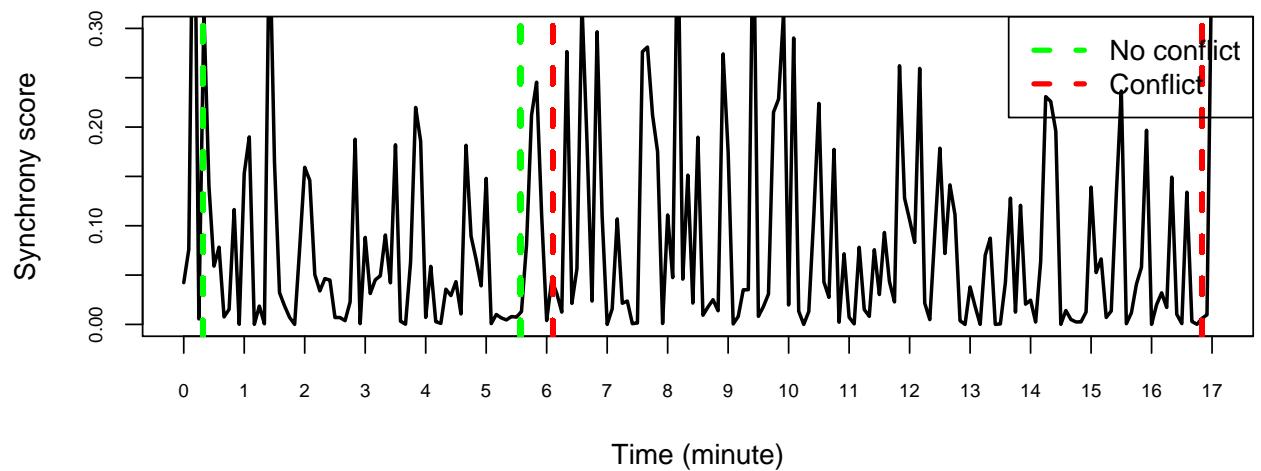
Synchrony scores in GROP039 family



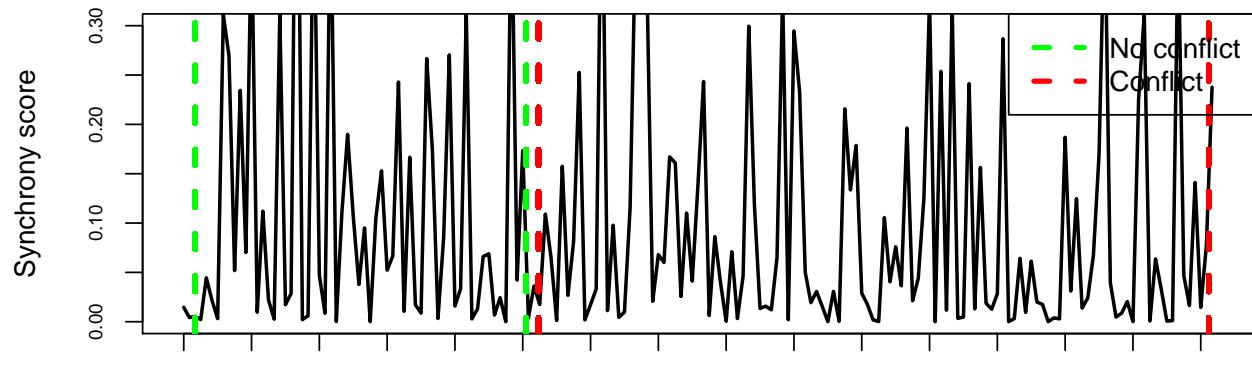
Synchrony scores in HAJA052 family



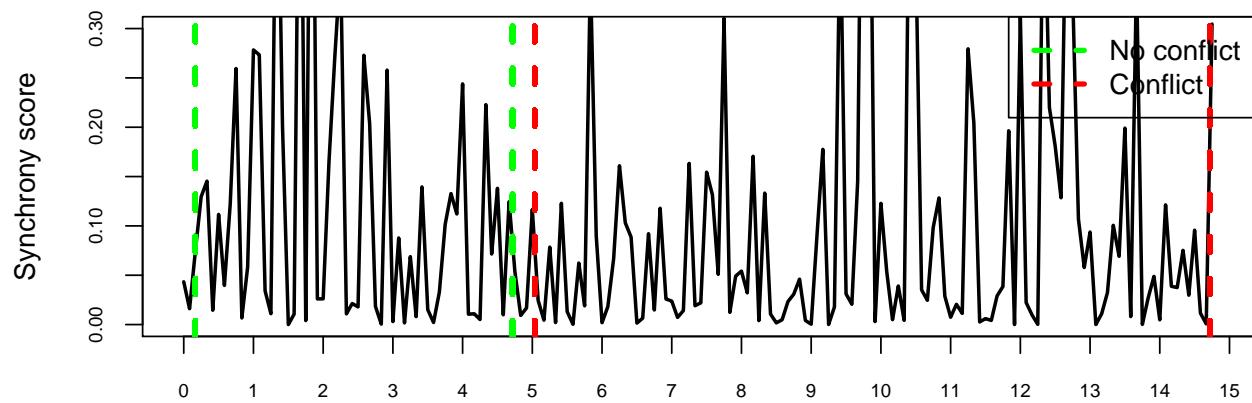
Synchrony scores in HUMA058 family



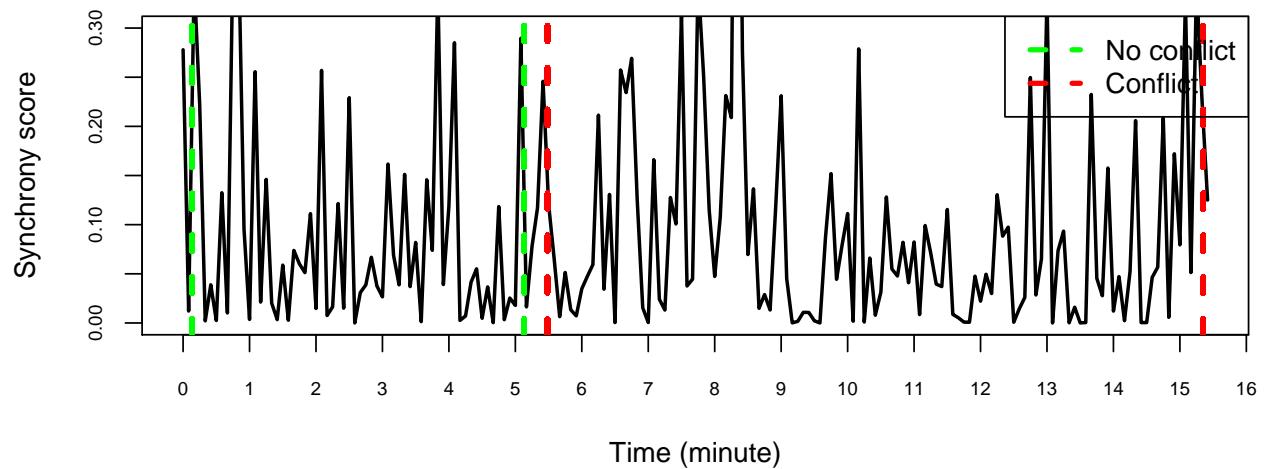
Synchrony scores in JAEM046 family



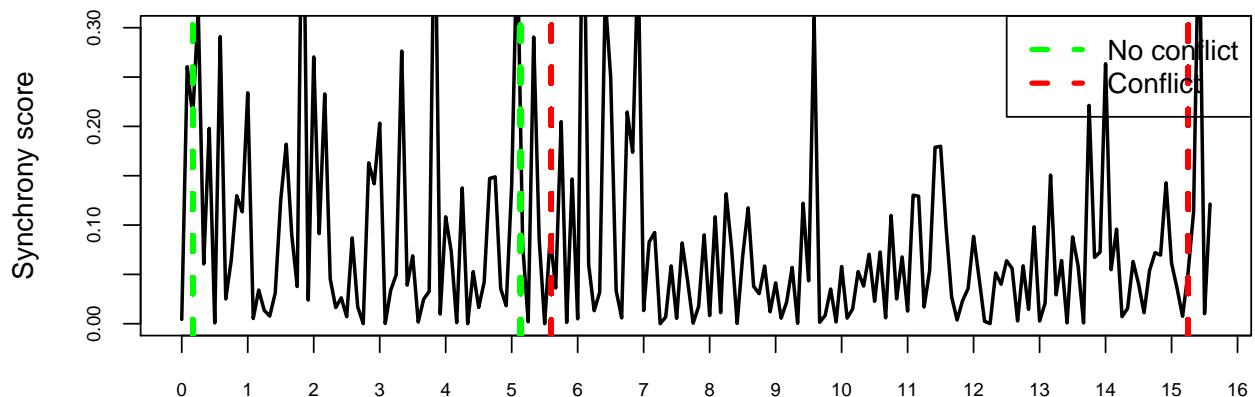
Synchrony scores in JEEO040 family



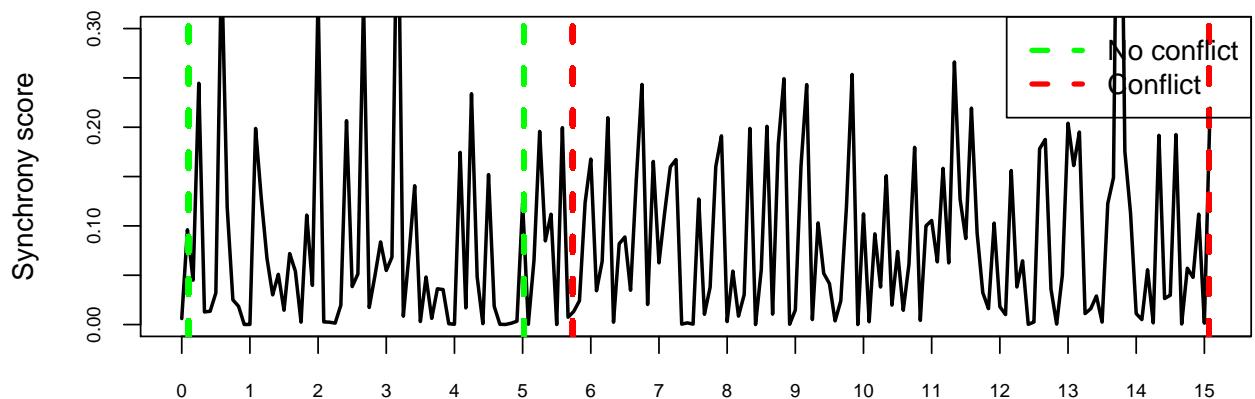
Synchrony scores in JOCE014 family



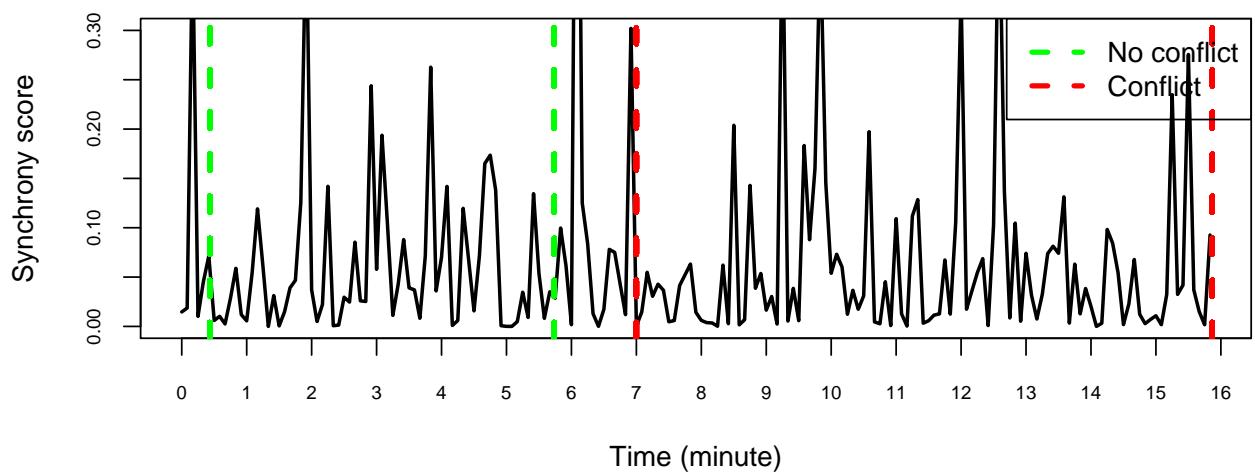
Synchrony scores in LACL family



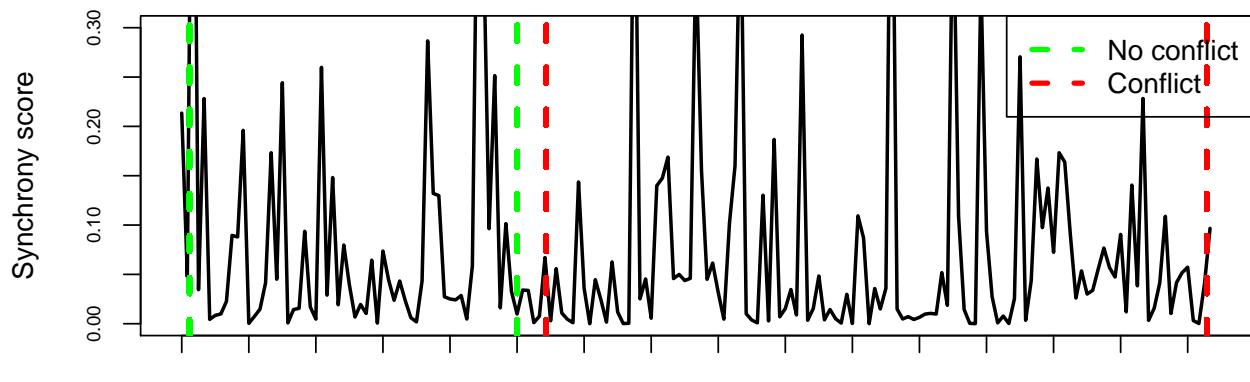
Time (minute) Synchrony scores in MAEL048 family



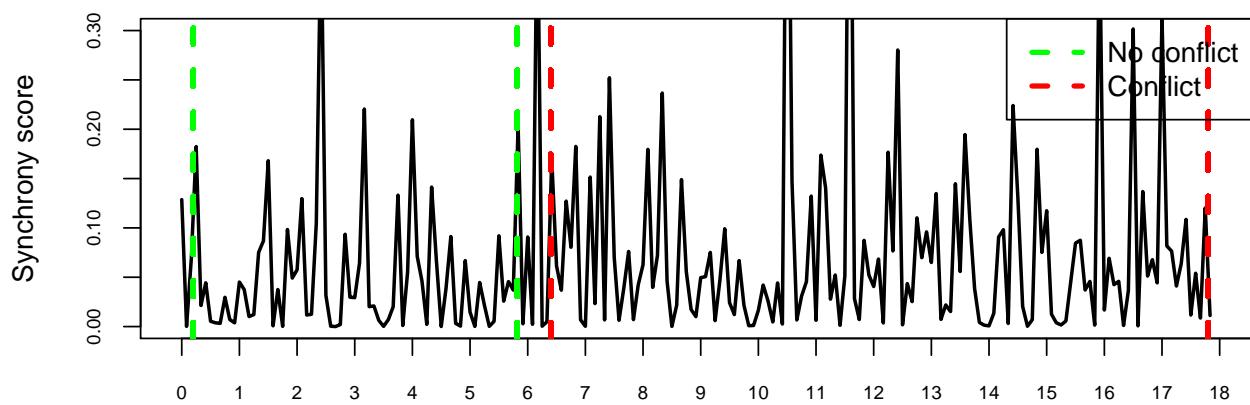
Time (minute) Synchrony scores in MAME20 family



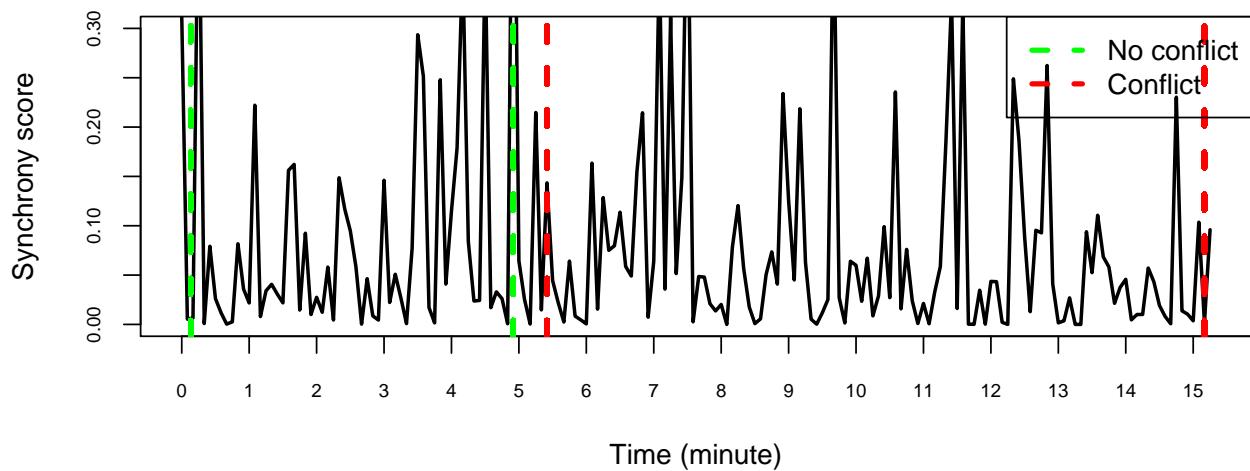
Synchrony scores in MIPH043 family



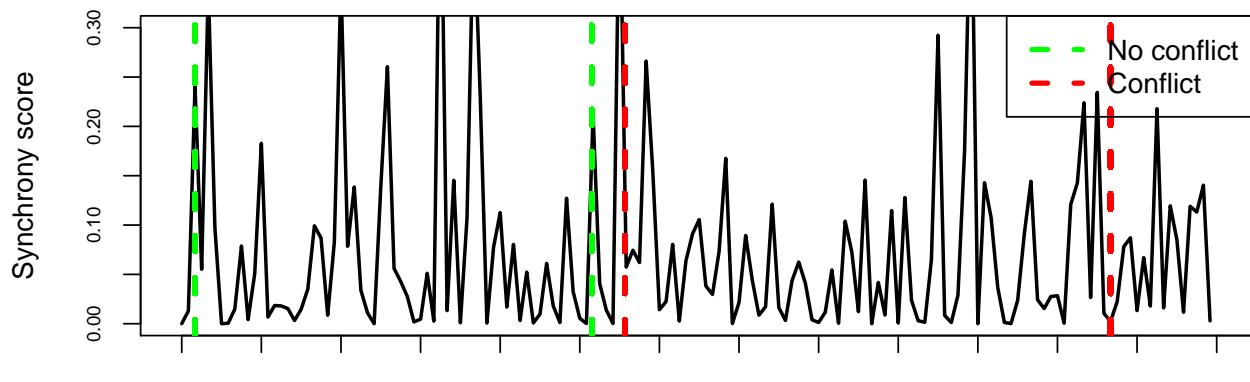
Synchrony scores in MOSA065 family



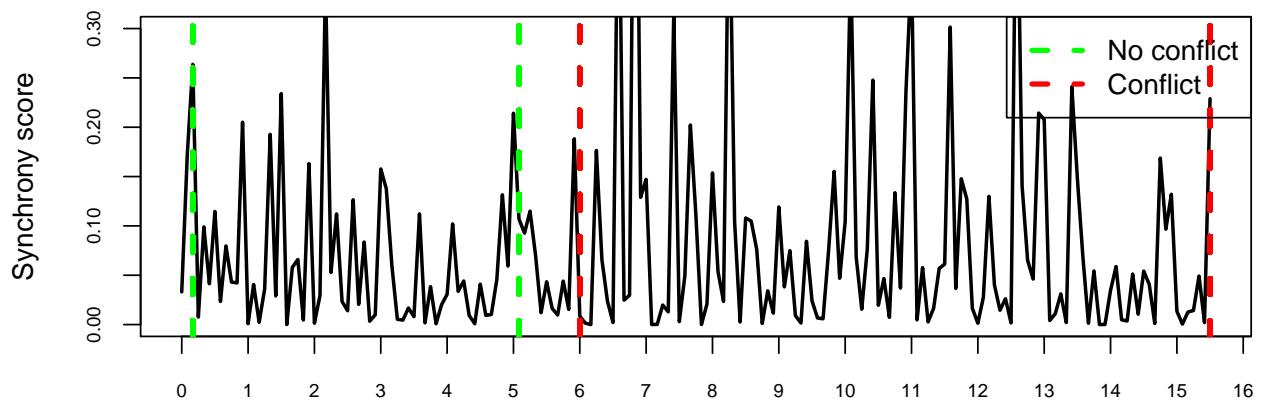
Synchrony scores in NAMA045 family



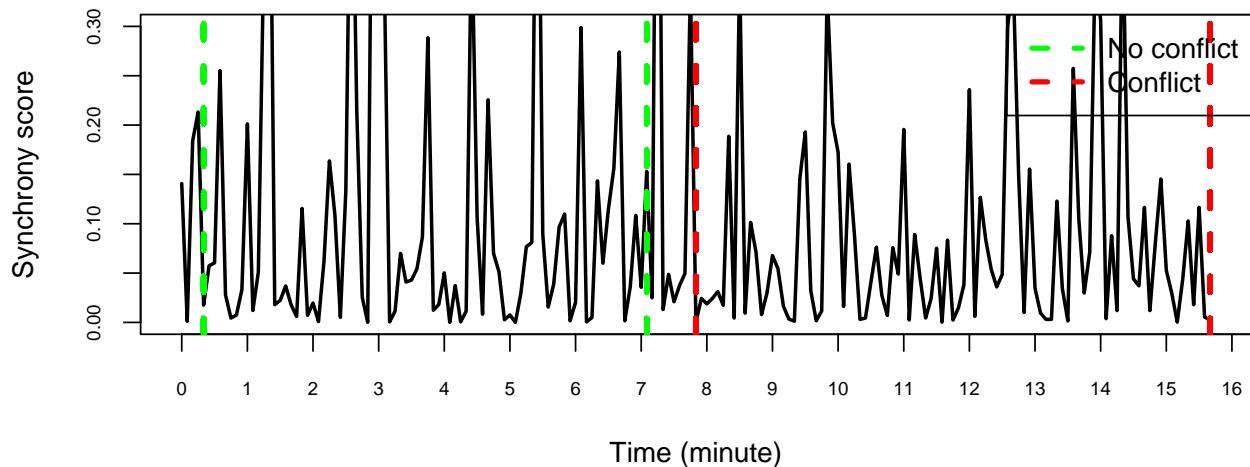
Synchrony scores in NUMA027 family



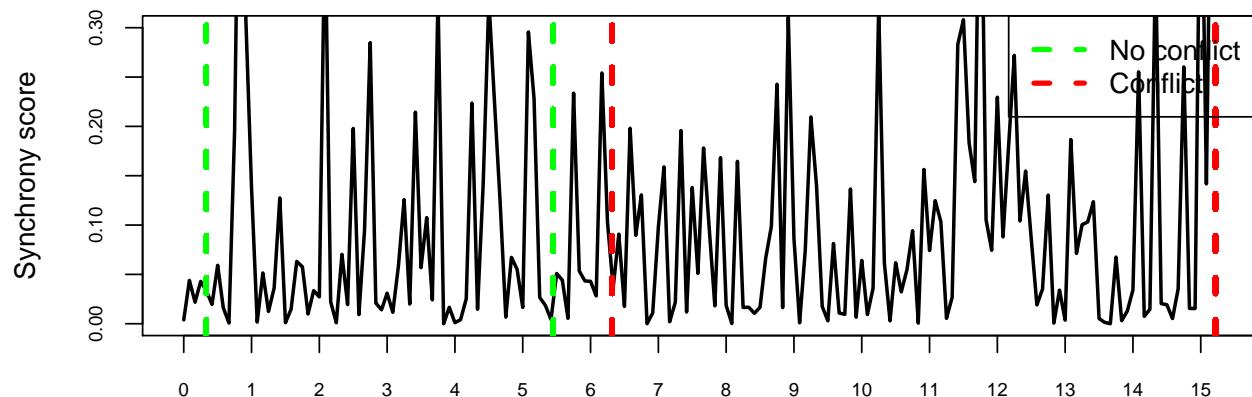
Synchrony scores in OGGA034 family



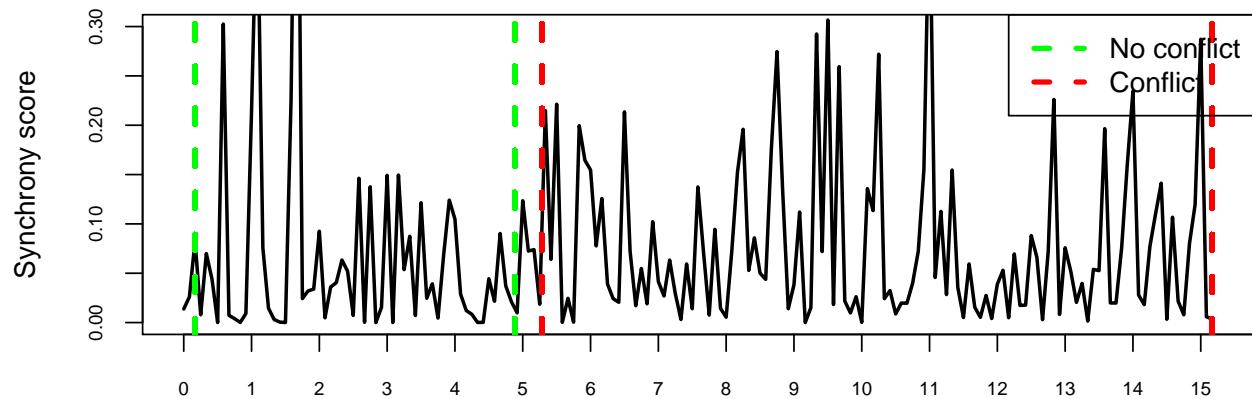
Synchrony scores in PAMA029 family



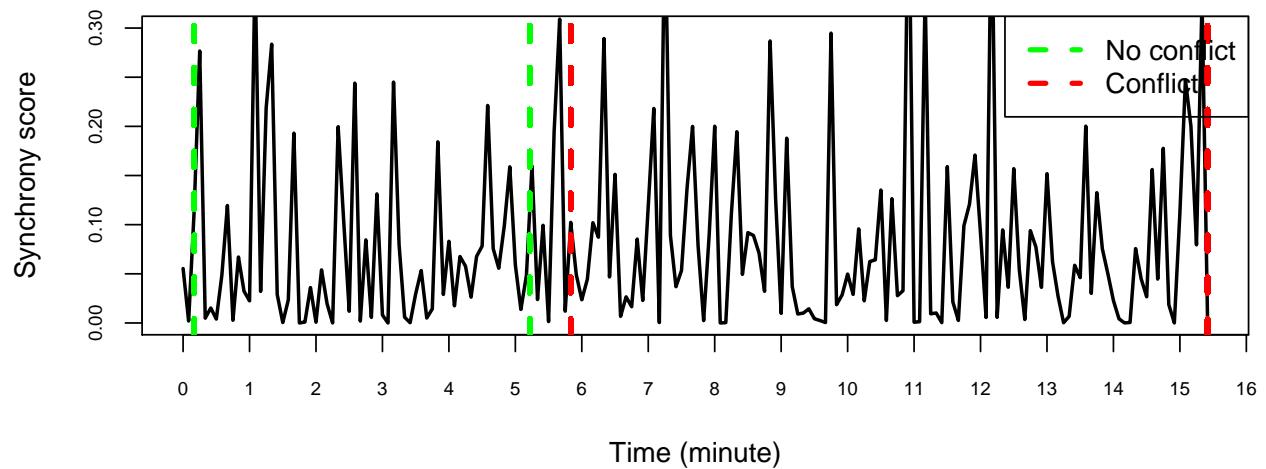
Synchrony scores in PELI020 family



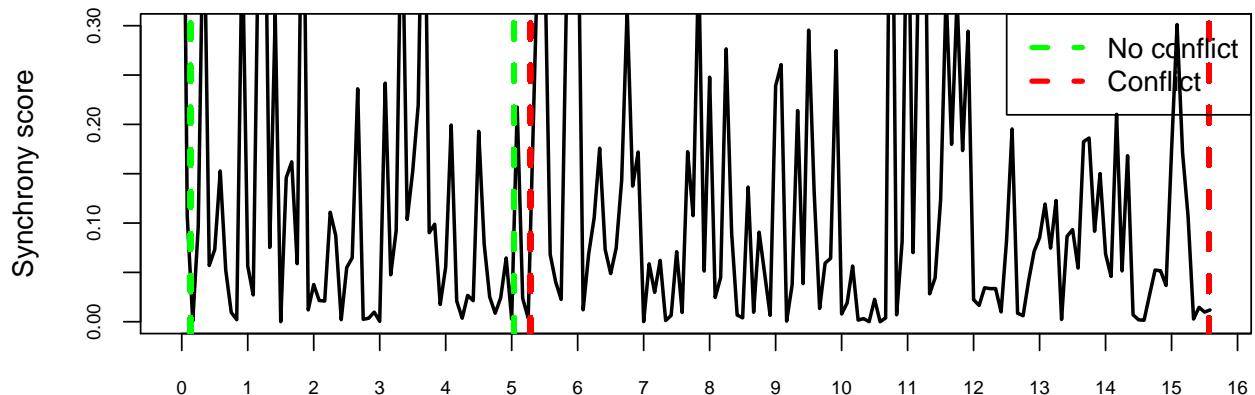
Synchrony scores in RAEM049 family



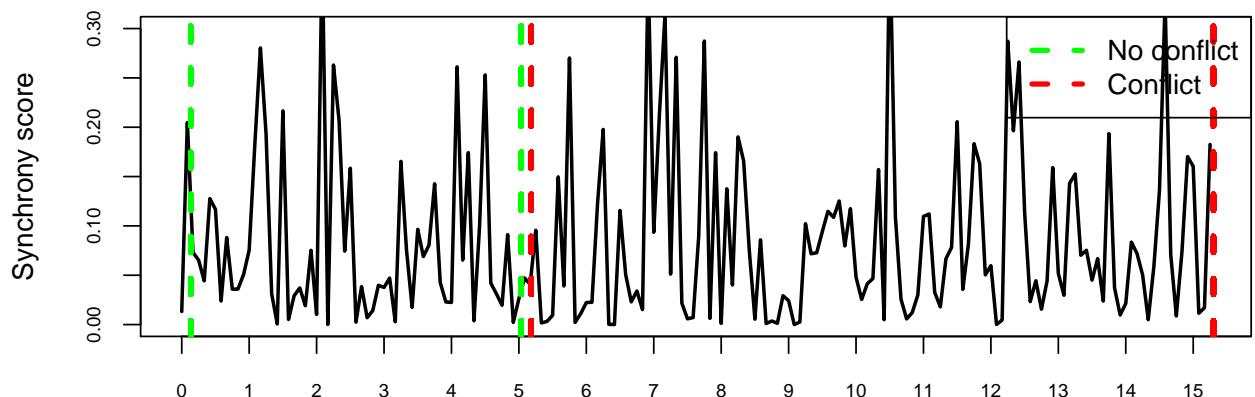
Synchrony scores in RAMA054 family



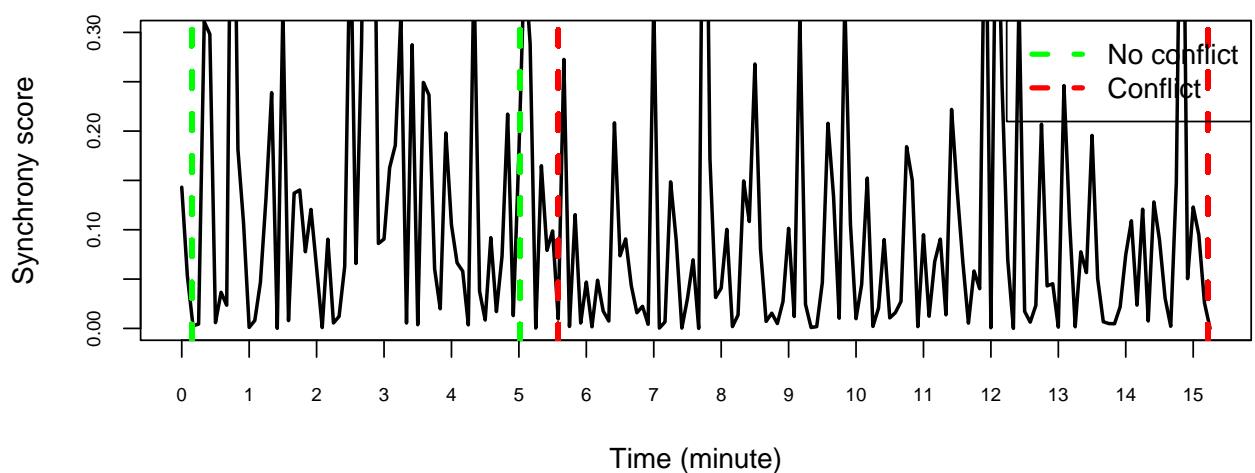
Synchrony scores in SEEM035 family



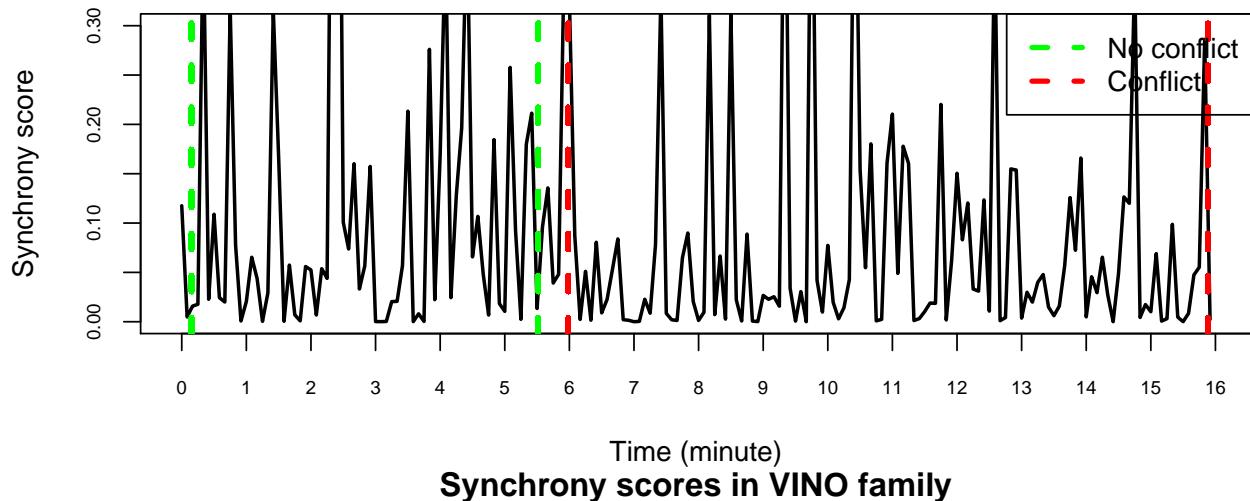
Synchrony scores in SHAN042 family



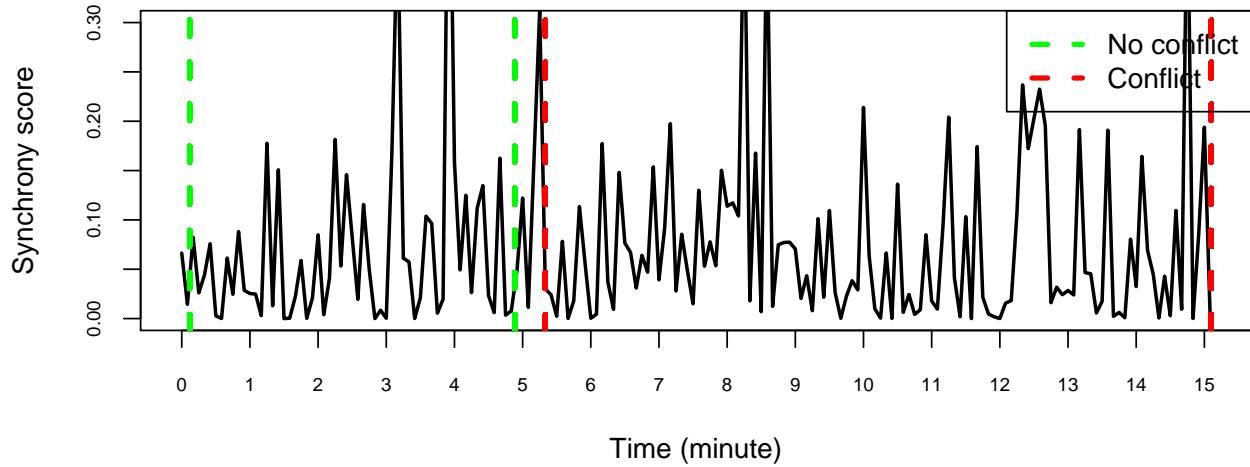
Synchrony scores in SOGA061 family



Synchrony scores in TIUG032 family



Synchrony scores in VINO family



```
#SSIInoLog$SSI <- as.ts(SSIInoLog$SSI)

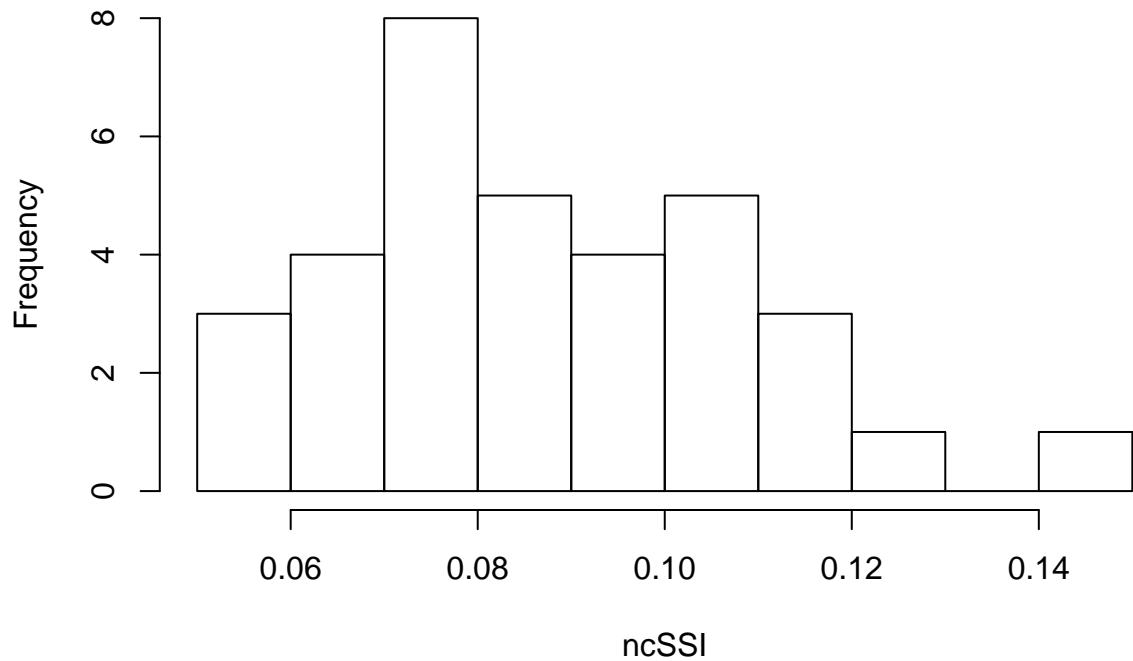
#for (i in unique(SSIInoLog$family)[5:35]){
#  print(i)
#  SSI.decompose <- #decompose(SSIInoLog[(which(SSIInoLog$family==i)),]$SSI, type="mult")
#}

ncSSI <- c()
cSSI <- c()

for (i in unique(SSIInoLog$family)){
#  print (i)
  nc <- mean(SSIInoLog[which(SSIInoLog$LabelVideo=="No Conflict" & SSIInoLog$family==i),]$SSI)
#  print(nc)
  ncSSI <- c(ncSSI, nc)
  co <-mean(SSIInoLog[which(SSIInoLog$LabelVideo=="Conflict" & SSIInoLog$family==i),]$SSI)
#  print (co)
  cSSI <- c(cSSI, co)
}
```

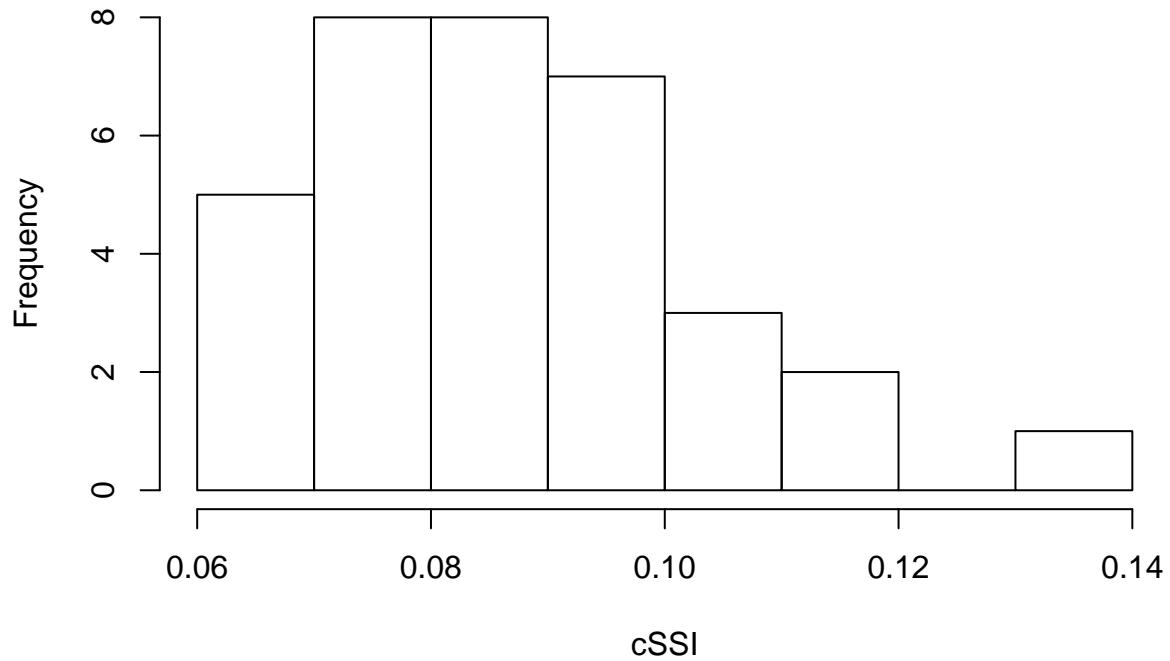
```
hist(ncSSI)
```

Histogram of ncSSI



```
hist(cSSI)
```

Histogram of cSSI



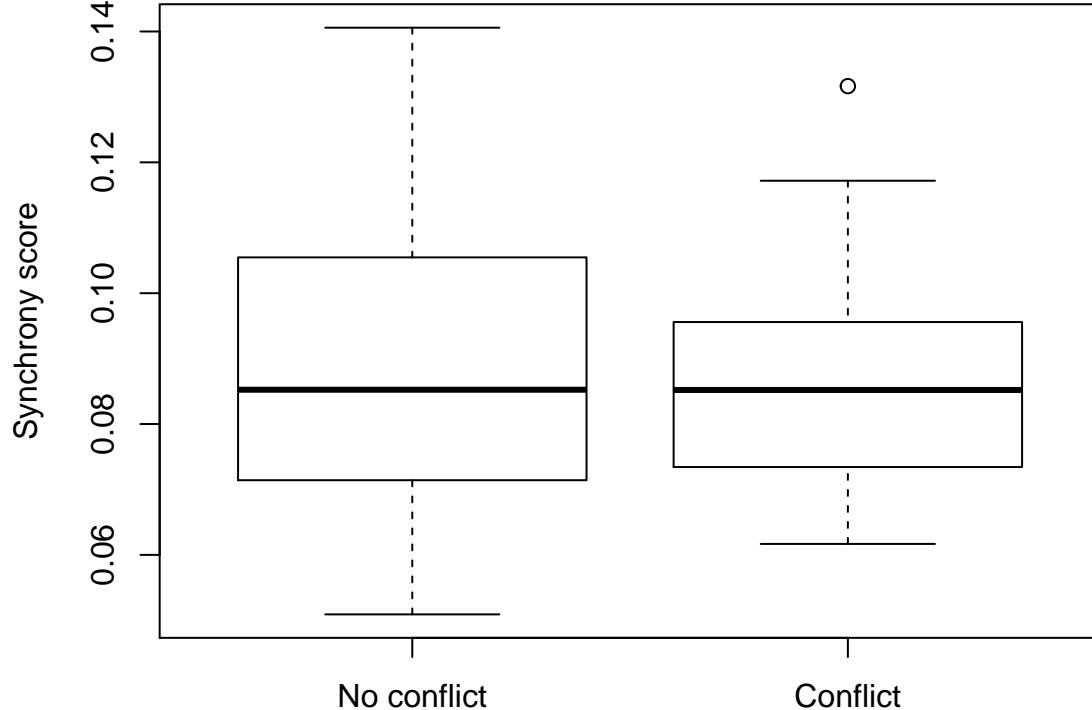
```
sd(ncSSI)/sd(cSSI)
```

```

## [1] 1.280335
SSImean <- data.frame(sort(ncSSI, decreasing = TRUE), sort(cSSI, decreasing = TRUE))
par(mar=c(3,5,3,3))

boxplot(SIImean, names= c("No conflict", "Conflict"), ylab="Synchrony score")

```



```

#effectifs <30 donc pas de t-test
#t.test(ncSSI, cSSI, alternative = c("two.sided"),
#paired=TRUE, conf.level = 0.95)

```

```
wilcox.test(ncSSI, cSSI)
```

```

##
## Wilcoxon rank sum test
##
## data: ncSSI and cSSI
## W = 578, p-value = 1
## alternative hypothesis: true location shift is not equal to 0

```

There isn't any difference between the mean SSI before and after the conflict.

```

SSI <- SSInoLog[-which(SSInoLog$LabelVideo=="Cut"),]
names(SI)

## [1] "family"          "SSI-interval"    "Time_min"        "SSI_fa_ch"
## [5] "SSI_mo_ch"       "SSI"            "CutBefore"       "CutMiddle1"
## [9] "CutMiddle2"      "CutFinal"        "ChildSex"        "ParentSex"
## [13] "CutBeforeMin"    "CutMiddle1Min"   "CutMiddle2Min"   "CutFinalMin"
## [17] "LabelVideo"

SSIagg <- subset(SI, select=c(family, SSI, LabelVideo))
#View(SSIagg)
names(SSIagg)

```

```

## [1] "family"      "SSI"          "LabelVideo"
str(SSIagg)

## 'data.frame':   6040 obs. of  3 variables:
##   $ family    : Factor w/ 35 levels "1606","BAJE059",...: 1 1 1 1 1 1 1 1 1 ...
##   $ SSI       : num  0.1167 0.1839 0.0893 0.0274 0.1701 ...
##   $ LabelVideo: chr  "No Conflict" "No Conflict" "No Conflict" "No Conflict" ...
SSIagg$LabelVideo <- as.factor(SSIagg$LabelVideo)
names(SSIagg)

## [1] "family"      "SSI"          "LabelVideo"
SSIagg <-aggregate(SSIagg, by=list(SSIagg$family, SSIagg$LabelVideo), FUN=mean)

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
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## returning NA

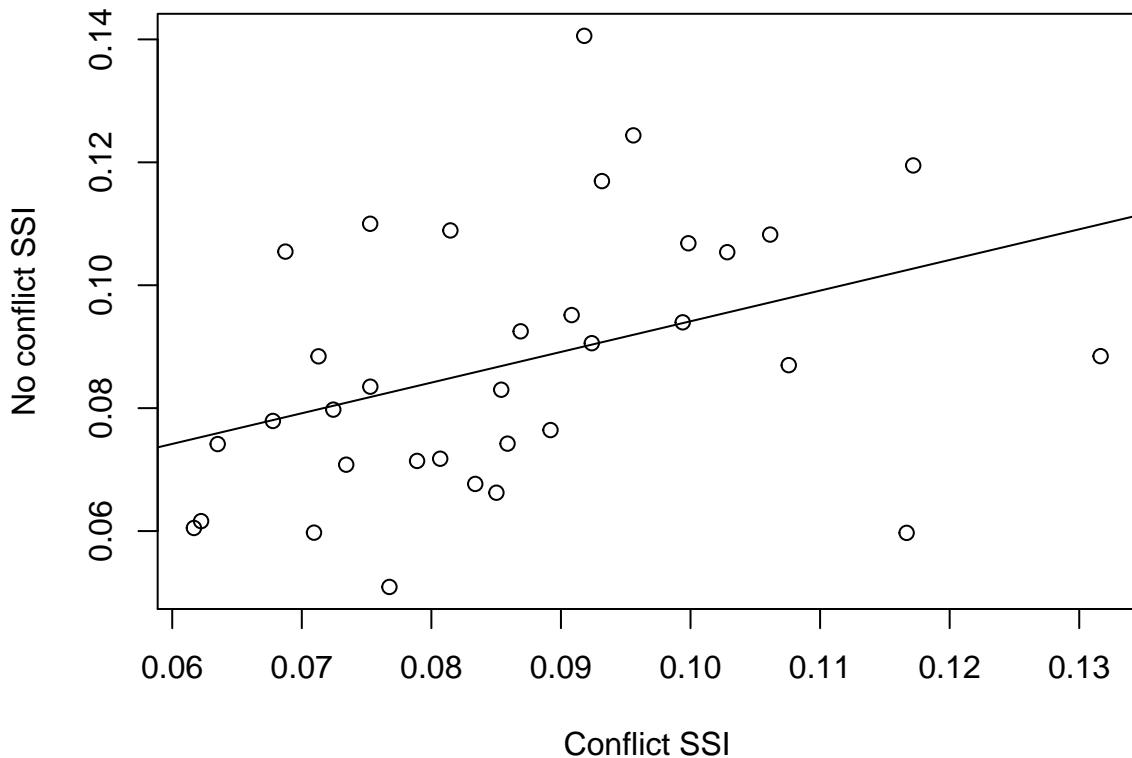
## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

```

```
cor.test(SSIagg[which(SSIagg$LabelVideo=="Conflict"),]$SSI,SSIagg[which(SSIagg$LabelVideo=="No Conflict")]

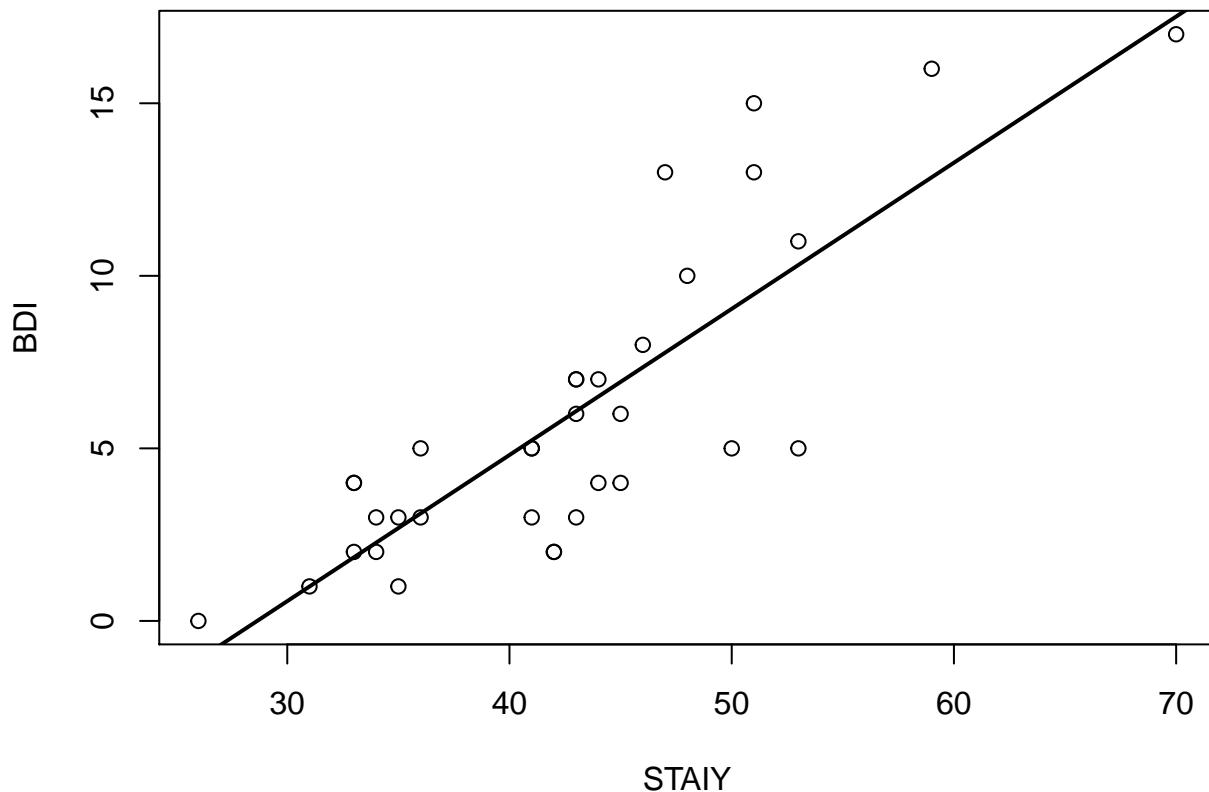
##
## Pearson's product-moment correlation
##
## data: SSIagg[which(SSIagg$LabelVideo == "Conflict"), ]$SSI and SSIagg[which(SSIagg$LabelVideo == "No Conflict")]
## t = 2.3934, df = 32, p-value = 0.02273
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.05929851 0.64307996
## sample estimates:
##        cor
## 0.3896506
#View(SSIagg)
```

There is a correlation between the synchrony before and after the conflict.

```
psycho<-read.csv2("/Users/Ofix/Documents/Fac/internat/Recherche/projets/synchro/synchroData/Monrado/Data")
#View(psycho)
psycho <- subset(psycho, select=c(Num._ident_videos, attachement_cluster, insecurite_level, TAS_total,
#View(psycho)

par(mar=c(4,4,2,1))
plot(psycho$STAIYB_total, psycho$BDI_total, main="BDI en fonction de la STAIY", xlab="STAIY", ylab="BDI")
model <- lm(psycho$BDI_total~psycho$STAIYB_total)
abline(model, lwd=2)
```

BDI en fonction de la STAIY



```
cor.test(psycho$BDI_total,psycho$STAIYB_total)
```

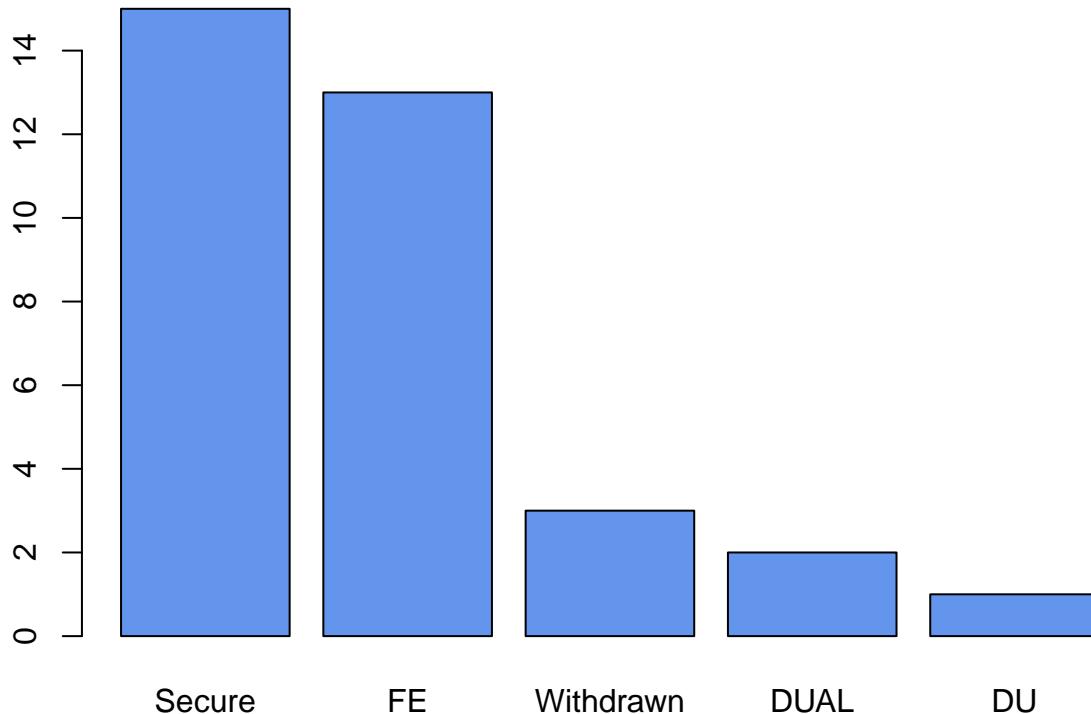
```
##  
## Pearson's product-moment correlation  
##  
## data: psycho$BDI_total and psycho$STAIYB_total  
## t = 8.4354, df = 32, p-value = 1.223e-09  
## alternative hypothesis: true correlation is not equal to 0  
## 95 percent confidence interval:  
## 0.6846660 0.9124366  
## sample estimates:  
## cor  
## 0.830537
```

```
table(psycho$attachement_cluster)
```

```
##  
## DU DUAL FE Secure Withdrawn  
## 1 2 13 15 3
```

```
barplot(sort(table(psycho$attachement_cluster), decreasing=TRUE), col="cornflower blue", main="Effectif")
```

Effectifs par style d'attachement



```
#View(SSInoLog)
```

```
SSI <- SSInoLog[-which(SSInoLog$LabelVideo=="Cut"),]  
SSIagg <- subset(SSI, select=c(family, SSI, LabelVideo))  
#View(SSIagg)
```

```
str(SSIagg)
```

```
## 'data.frame': 6040 obs. of 3 variables:  
## $ family : Factor w/ 35 levels "1606","BAJE059",...: 1 1 1 1 1 1 1 1 1 1 ...  
## $ SSI : num 0.1167 0.1839 0.0893 0.0274 0.1701 ...  
## $ LabelVideo: chr "No Conflict" "No Conflict" "No Conflict" "No Conflict" ...  
SSIagg$LabelVideo <- as.factor(SSIagg$LabelVideo)
```

```
SSIagg <- aggregate(SSIagg, by=list(SSIagg$family), FUN=mean)
```

```
## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:  
## returning NA
```

```
## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:  
## returning NA
```

```
## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:  
## returning NA
```

```
## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:  
## returning NA
```

```
## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:  
## returning NA
```



```

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
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## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

```

```

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

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## returning NA

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## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

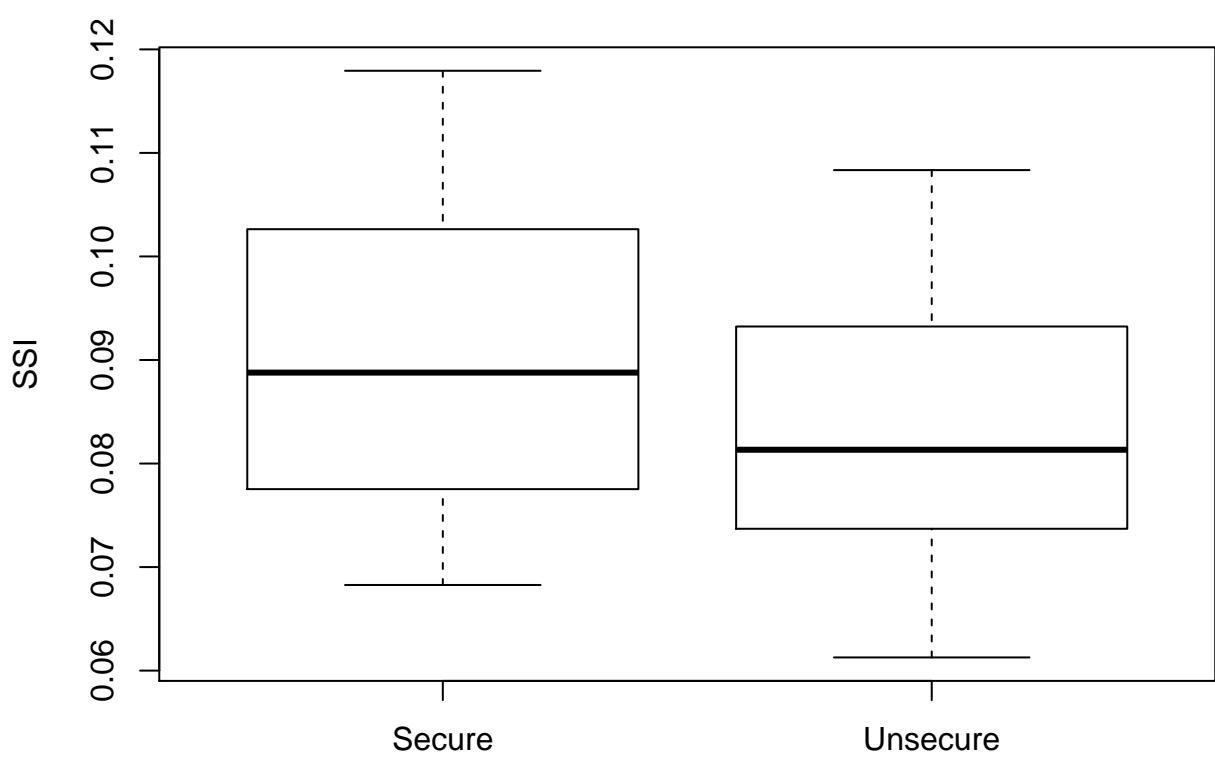
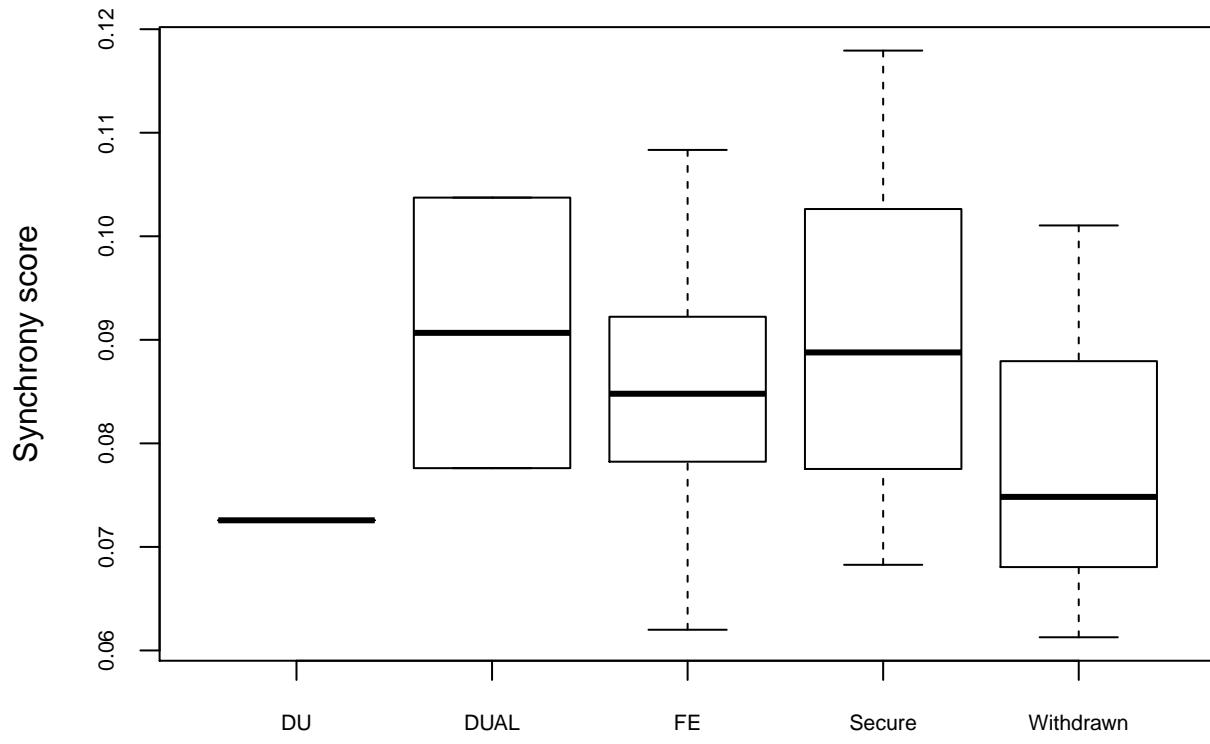
## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

#View(SSIagg)
SSIagg$family <-NULL
SSIagg$LabelVideo <-NULL
names(SSIagg) <-c("family", "SSI")
SSI <- merge(SSIagg, psycho, by.x= "family", by.y= "Num._ident_videos")

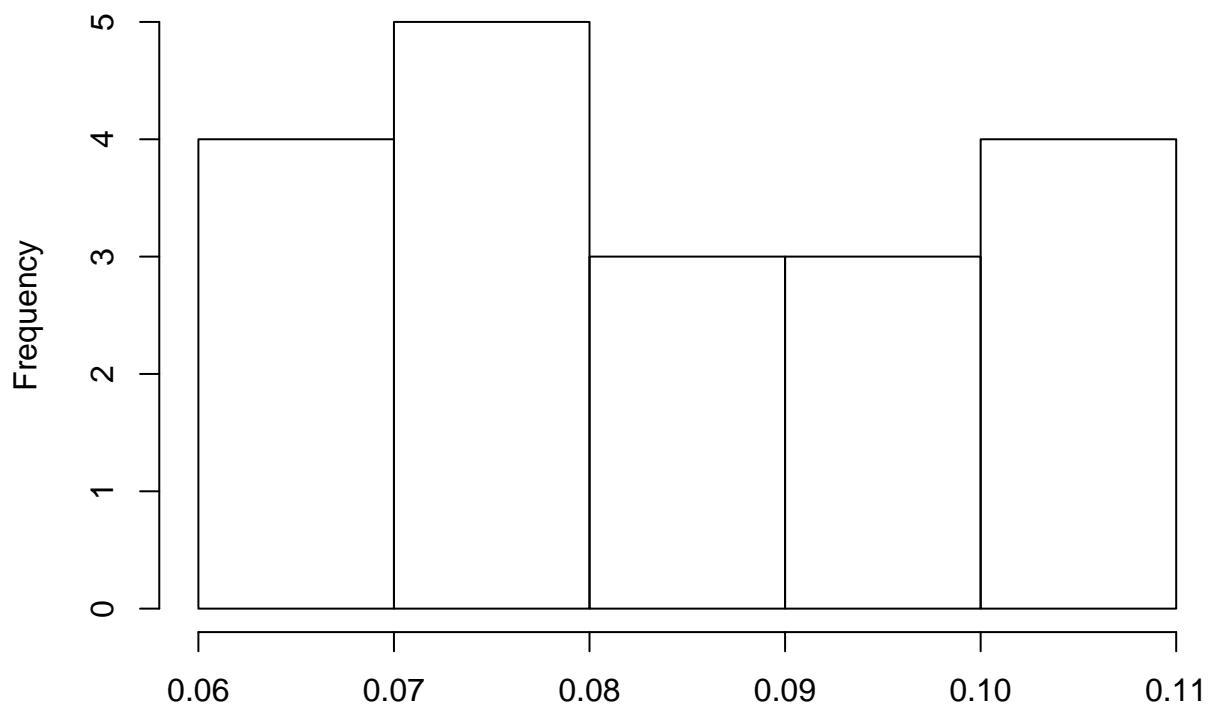
#View(SSI)
SSI$attachement_duo <- rep(NA)
SSI[which(SSI$attachement_cluster=="Secure"),]$attachement_duo <- "Secure"
SSI[which(SSI$attachement_cluster!="Secure"),]$attachement_duo <- "Unsecure"

boxplot(SSI ~ attachement_cluster, data = SSI, cex.axis=0.7, ylab="Synchrony score")

```



Histogram of SSI[which(SSI\$attachement_duo != "Secure"),]\$SS



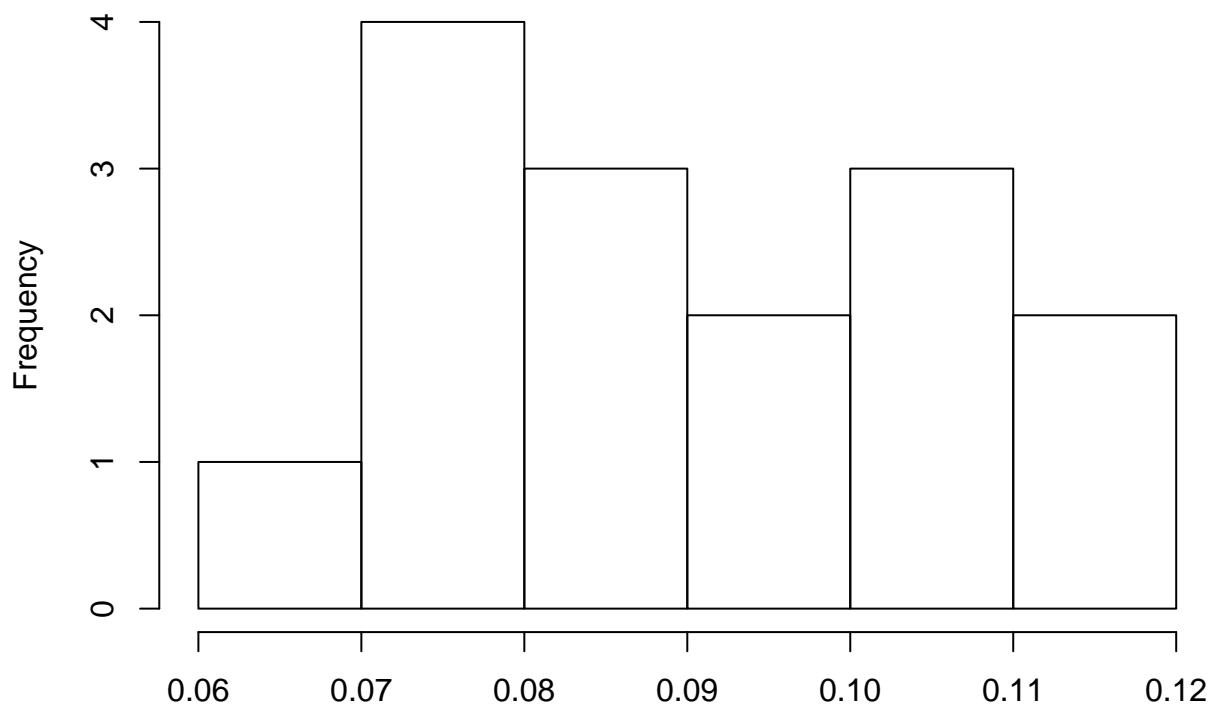
```
SSI[which(SSI$attachement_duo != "Secure"), ]$SSI
```

```
sd(SSI[which(SSI$attachement_duo != "Secure"), ]$SSI)
```

```
## [1] 0.01441683
```

```
hist(SSI[which(SSI$attachement_duo == "Secure"), ]$SSI)
```

Histogram of SSI[which(SSI\$attachement_duo == "Secure"),]\$SSI



```
SSI[which(SSI$attachement_duo == "Secure"), ]$SSI
```

```
sd(SSI[which(SSI$attachement_duo=="Secure"),]$SSI)
```

```
## [1] 0.01607062
```

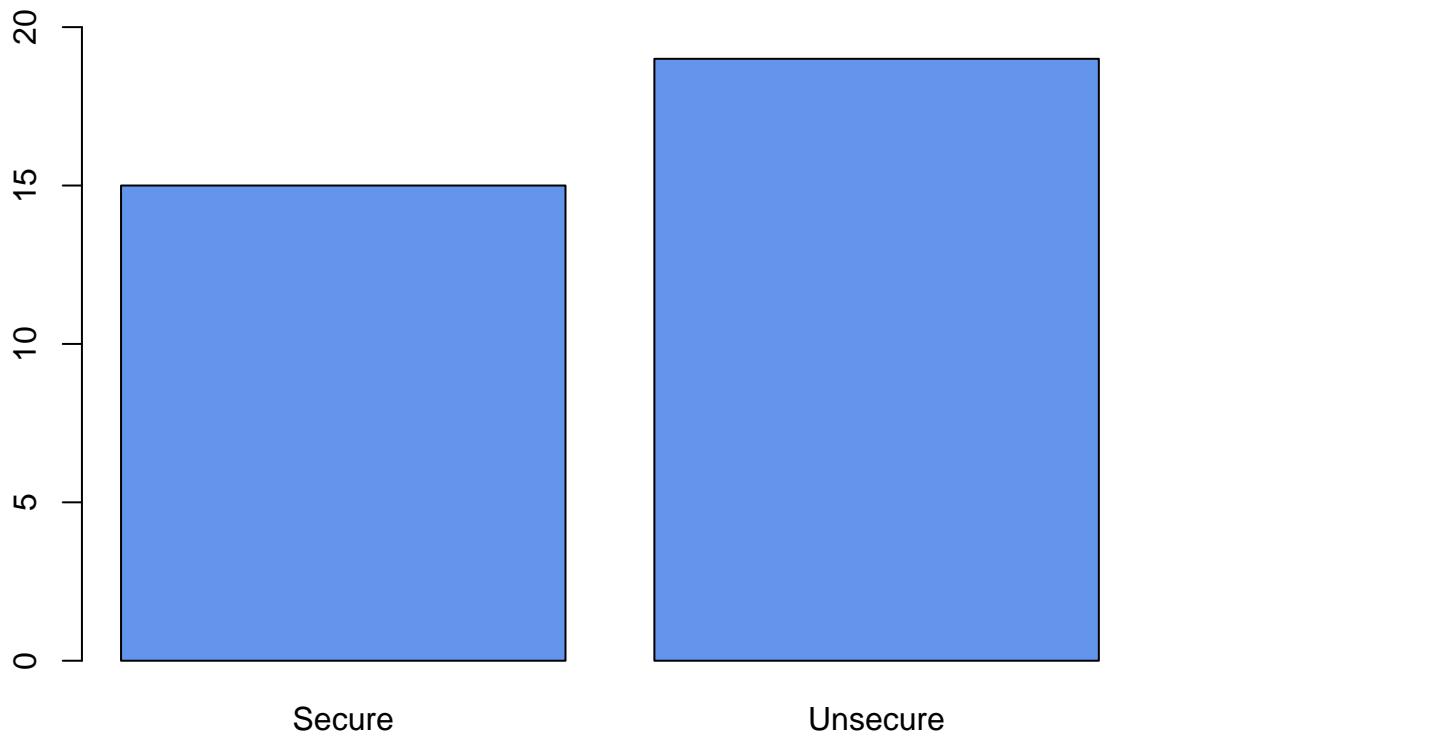
```
table(SSI$attachement_duo)
```

```
##
```

```
##   Secure Unsecure
```

```
##      15      19
```

```
barplot(table(SSI$attachement_duo), ylim=c(0,20), col="cornflowerblue")
```



```
#Effectifs inf à 30 donc pas possibles
#t.test(SSI[which(SSI$attachement_duo=="Unsecure") ,]$SSI, SSI[which(SSI$attachement_duo=="Secure") ,]$SSI)

wilcox.test(SSI[which(SSI$attachement_duo=="Unsecure") ,]$SSI, SSI[which(SSI$attachement_duo=="Secure") ,]$SSI)

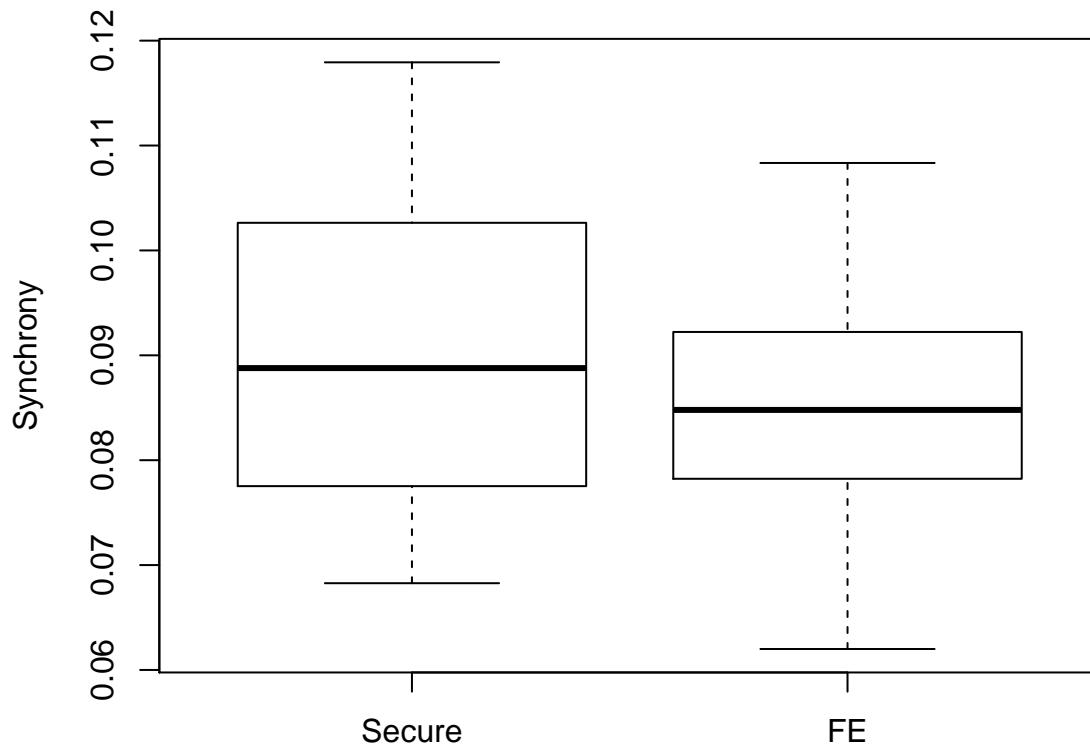
## Wilcoxon rank sum test
## data: SSI[which(SSI$attachement_duo == "Unsecure"), ]$SSI and SSI[which(SSI$attachement_duo == "Secure"), ]$SSI
## W = 107, p-value = 0.2281
## alternative hypothesis: true location shift is not equal to 0

SSIattach <- data.frame(c(sort(SSI[which(SSI$attachement_cluster=="FE") ,]$SSI, decreasing = TRUE), rep(mean(SSI[which(SSI$attachement_cluster=="FE") ,]$SSI), 107)))

#View(SSIattach)
par(mar=c(3,5,3,3))

FE <- c(sort(SSI[which(SSI$attachement_cluster=="FE") ,]$SSI, decreasing = TRUE), rep(mean(SSI[which(SSI$attachement_cluster=="FE") ,]$SSI), 107))

boxplot(SSI[which(SSI$attachement_cluster=="Secure") ,]$SSI, SSI[which(SSI$attachement_cluster=="FE") ,]$SSI)
```



```
t.test(SSI[which(SSI$attachement_cluster=="Secure"),]$SSI, SSI[which(SSI$attachement_cluster=="FE"),]$SSI)
##
##  Welch Two Sample t-test
##
## data: SSI[which(SSI$attachement_cluster == "Secure"), ]$SSI and SSI[which(SSI$attachement_cluster =
## t = 1.2518, df = 26, p-value = 0.2218
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.004540924 0.018684528
## sample estimates:
## mean of x mean of y
## 0.09132637 0.08425457
SSIM <- aggregate(SSI, by=list(SSI$family), FUN=mean)

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

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## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
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## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA
```



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## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

```



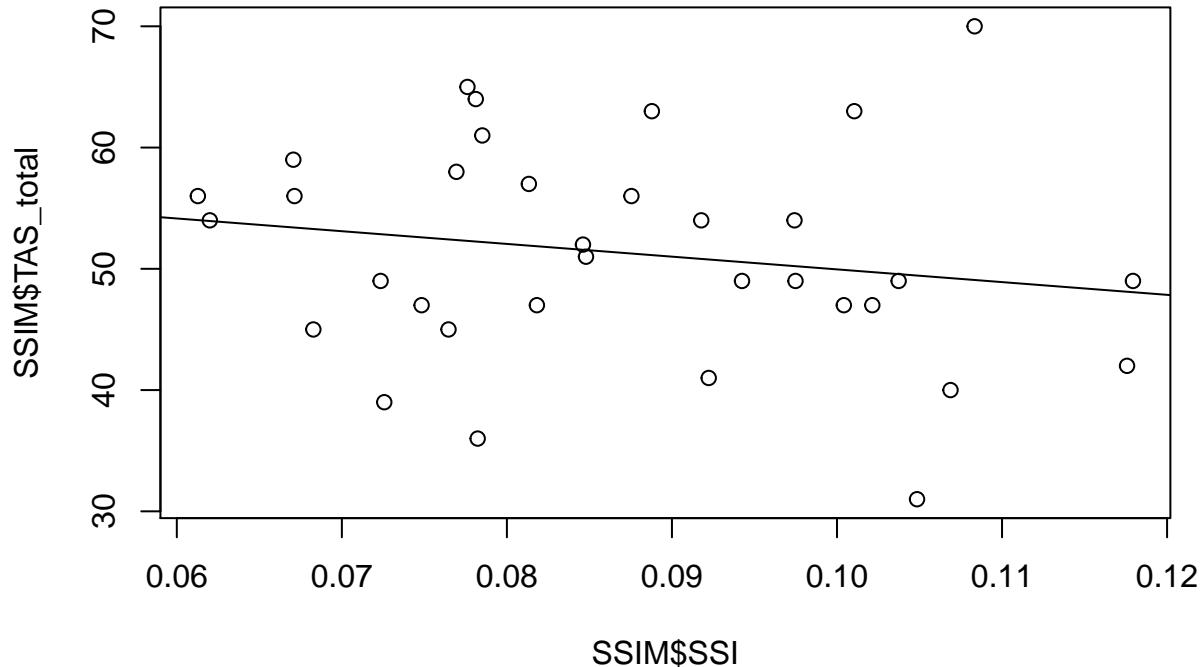
```

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA
plot(SSIM$SSI, SSIM$TAS_total)
model <- lm(SSIM$TAS_total~SSIM$SSI)
abline(model)

```

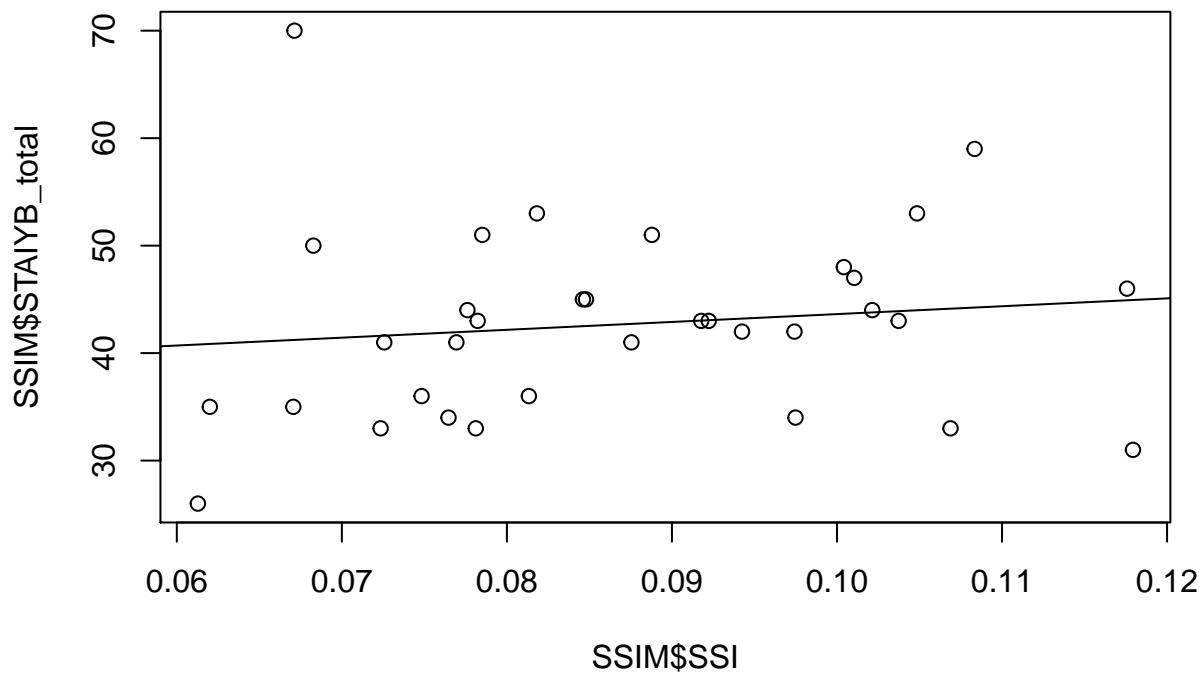


```

cor.test(SSIM$SSI, SSIM$TAS_total)

##
## Pearson's product-moment correlation
##
## data: SSIM$SSI and SSIM$TAS_total
## t = -1.0528, df = 32, p-value = 0.3003
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.4907738 0.1654264
## sample estimates:
## cor
## -0.1829752
plot(SSIM$SSI, SSIM$STAIYB_total)
model <- lm(SSIM$STAIYB_total~SSIM$SSI)
abline(model)

```



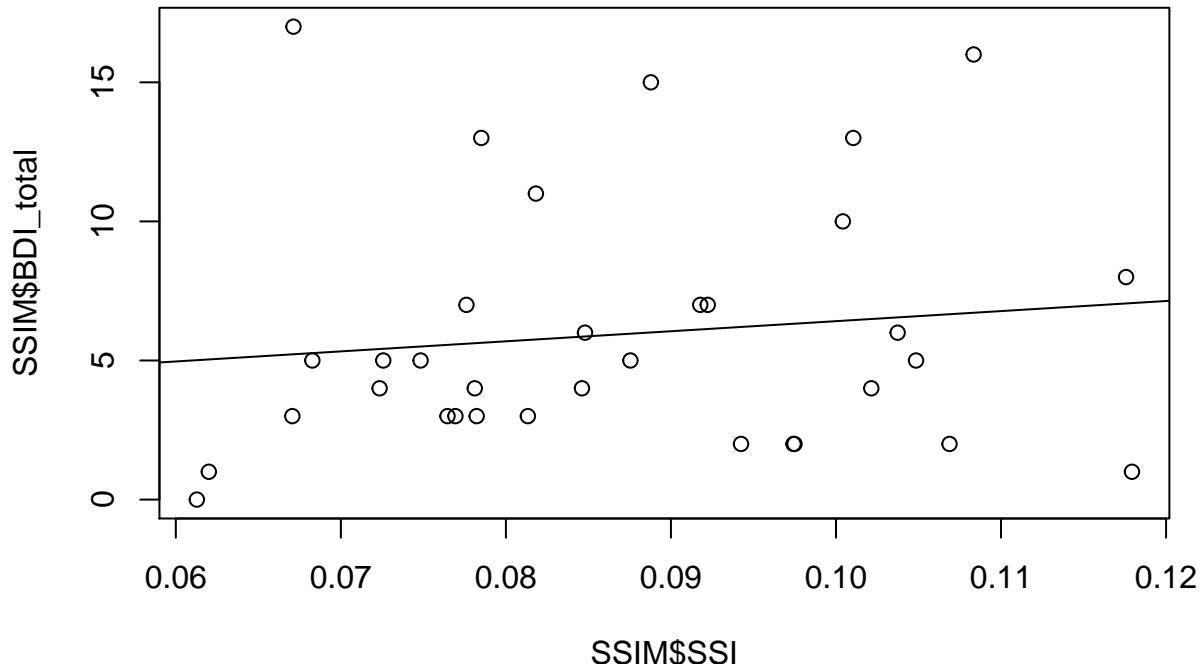
```

cor.test(SSIM$SSI, SSIM$STAIYB_total)

##
## Pearson's product-moment correlation
##
## data: SSIM$SSI and SSIM$STAIYB_total
## t = 0.73236, df = 32, p-value = 0.4693
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.2192940 0.4471445
## sample estimates:
##       cor
## 0.128393

plot(SSIM$SSI, SSIM$BDI_total)
model <- lm(SSIM$BDI_total~SSIM$SSI)
abline(model)

```



```
cor.test(SSIM$SSI, SSIM$BDI_total)
```

```
##
## Pearson's product-moment correlation
##
## data: SSIM$SSI and SSIM$BDI_total
## t = 0.7112, df = 32, p-value = 0.4821
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.2228239 0.4441704
## sample estimates:
## cor
## 0.1247412
```

#View(SSInoLog)

```
SSI <- SSInoLog[-which(SSInoLog$LabelVideo=="Cut"),]
SSIagg <- subset(SSI, select=c(family, SSI, LabelVideo))
#View(SSIagg)
names(SSIagg)
```

```
## [1] "family"      "SSI"          "LabelVideo"
str(SSIagg)
```

```
## 'data.frame': 6040 obs. of 3 variables:
## $ family      : Factor w/ 35 levels "1606","BAJE059",...: 1 1 1 1 1 1 1 1 1 ...
## $ SSI         : num  0.1167 0.1839 0.0893 0.0274 0.1701 ...
## $ LabelVideo: chr  "No Conflict" "No Conflict" "No Conflict" "No Conflict" ...
SSIagg$LabelVideo <- as.factor(SSIagg$LabelVideo)
names(SSIagg)
```

```
## [1] "family"      "SSI"          "LabelVideo"
```

```

SSIagg <-aggregate(SSIagg, by=list(SSIagg$family, SSIagg$LabelVideo), FUN=mean)

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

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## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA
```



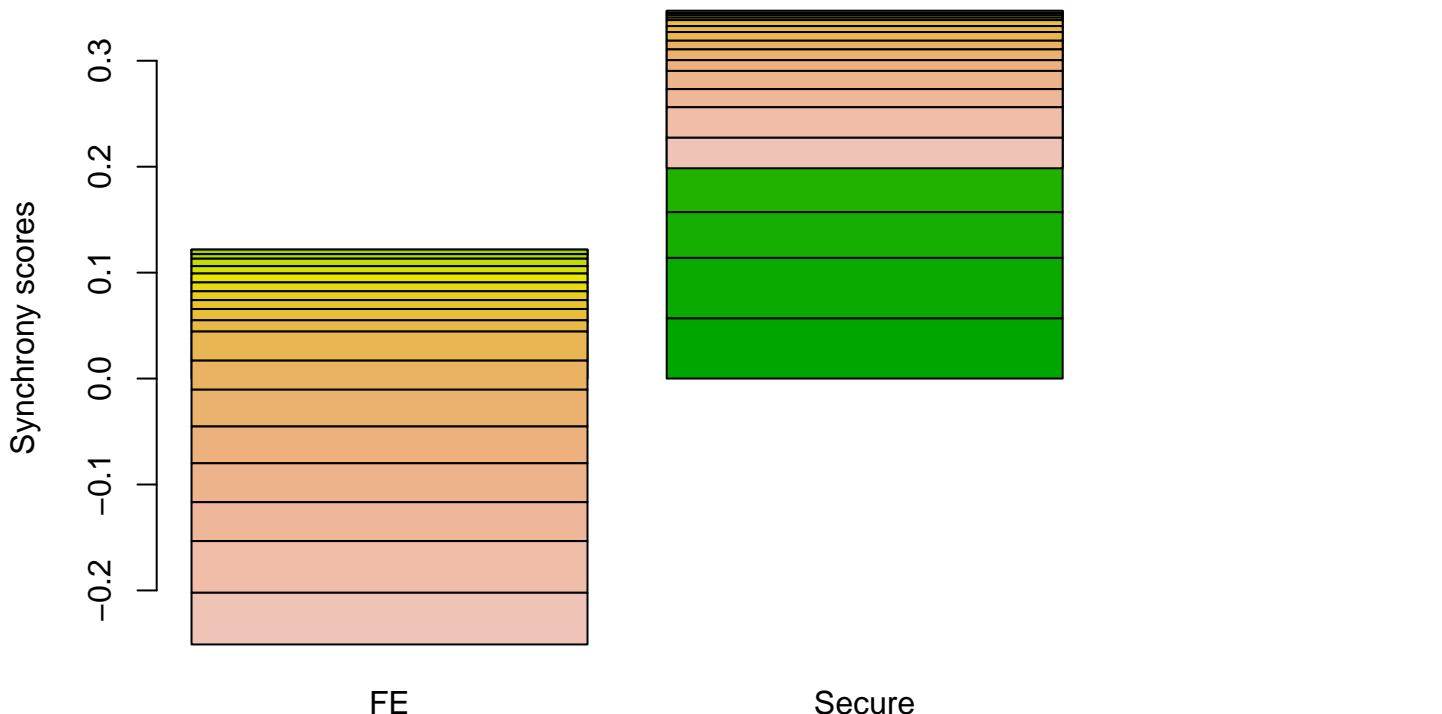
```

## [1] 0.001173828
## [1] -0.002096179
## [1] -0.005612297
## [1] -0.002542164
## [1] 0.01571619
## [1] -0.0237813
## [1] -0.02880027
## [1] 0.001811938
## [1] -0.03673722
## [1] 0.01275727
## [1] 0.0006123615
## [1] -0.01714405
## [1] 0.02585718
## [1] -0.007329623
## [1] -0.01014714
## [1] 0.01877874
## [1] -0.02744715
## [1] 0.00543184
## [1] 0.007484757
## [1] 0.01165041
## [1] -0.002303032
## [1] 0.002393419
## [1] -0.04878173
## [1] -0.03472463
## [1] 0.002624708

SSIdiff <- data.frame(sort(c(SSI[which(SSI$attachement_cluster=="FE"),]$diff, rep(mean(SSI[which(SSI$at
par(mar=c(3,5,3,3))
barplot(as.matrix(SSIdiff), col=terrain.colors(34), ylab="Synchrony scores", main="Sum of mean synchrony

```

Sum of mean synchrony scores for all videos



```

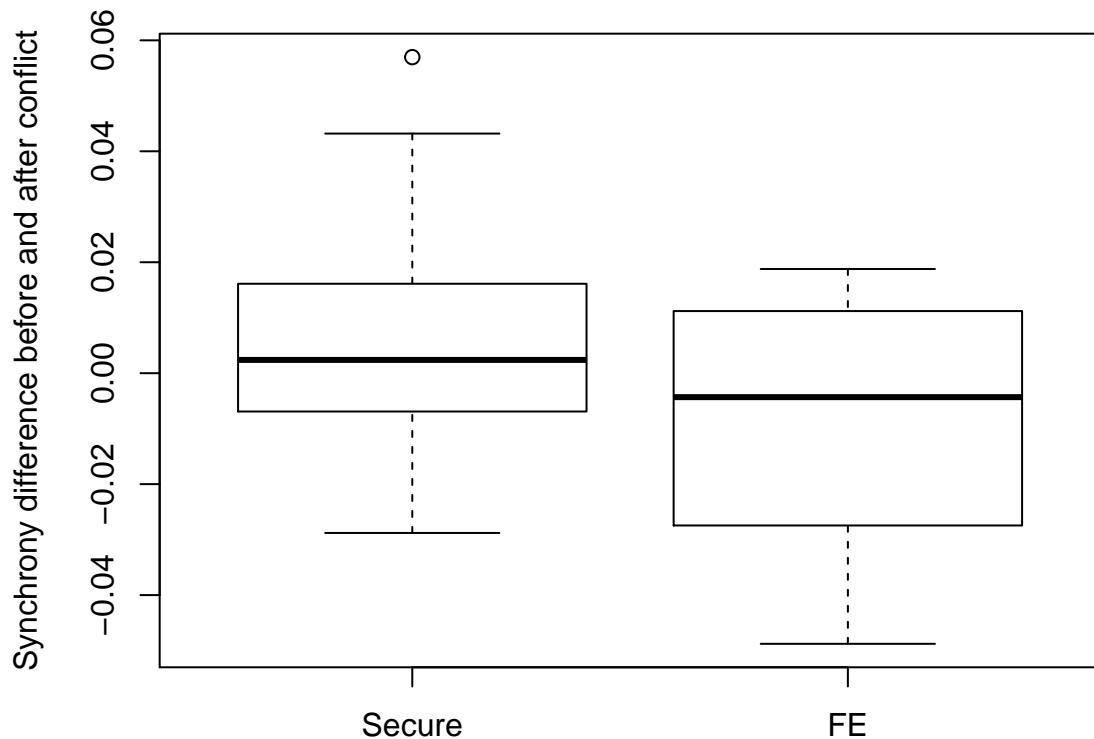
FE <- c(sort(SSI[which(SSI$attachement_cluster=="FE")],]$SSI, decreasing = TRUE), rep(mean(SSI[which(SSI$attachement_cluster=="FE")],]$SSI, times=10))

t.test(SSI[which(SSI$attachement_cluster=="Secure" & SSI$LabelVideo=="Conflict")],]$diff, SSI[which(SSI$attachement_cluster=="FE")],]$diff)

##
##  Welch Two Sample t-test
##
## data:  SSI[which(SSI$attachement_cluster == "Secure" & SSI$LabelVideo == "Conflict")], SSI[which(SSI$attachement_cluster == "FE")]
## t = 1.7738, df = 25.578, p-value = 0.088
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.002393389  0.032361620
## sample estimates:
##   mean of x   mean of y
## 0.006617209 -0.008366906

boxplot(SSI[which(SSI$attachement_cluster=="Secure" & SSI$LabelVideo=="Conflict")],]$diff, SSI[which(SSI$attachement_cluster=="FE")],]$diff)

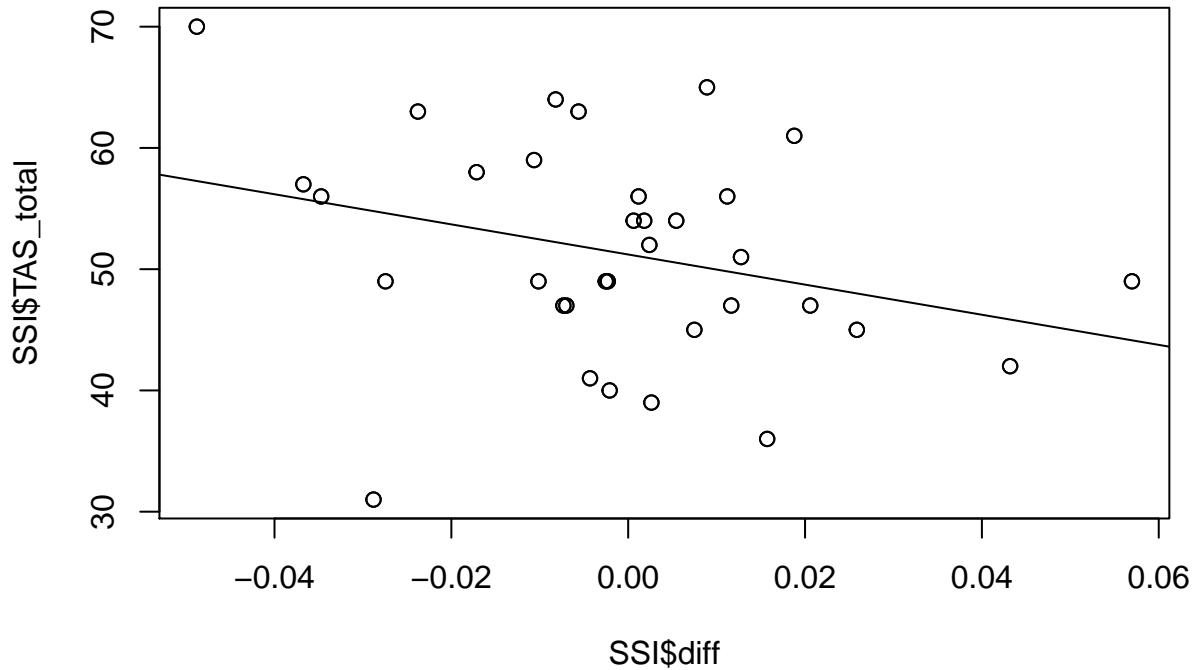
```



```

#View(SSI)
plot(SSI$diff, SSI$TAS_total)
model <- lm(SSI$TAS_total~SSI$diff)
abline(model)

```



```

cor.test(ssi[which(ssi$LabelVideo=="Conflict"),]$diff, ssi[which(ssi$LabelVideo=="Conflict"),]$TAS_total)

##
## Pearson's product-moment correlation
##
## data: SSI[which(ssi$LabelVideo == "Conflict"), ]$diff and SSI[which(ssi$LabelVideo == "Conflict"), ]
## t = -1.7862, df = 32, p-value = 0.08355
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.58018843 0.04126969
## sample estimates:
## cor
## -0.3010981

table(ssi$insecurite_level)

##
##          Clearly      Markedly      Midly
##        4           28           8           2
## Mildly Mildly insecure Mildly Insecure   Moderately
##       10            2            2            4
## Moderately
##       8

#Quantite de mut et BDI
str(MH)

## 'data.frame': 802834 obs. of 18 variables:
## $ family      : chr "1606" "1606" "1606" "1606" ...
## $ slidedFather : num 0.002084 0.001758 0.000729 0.000693 0.000537 ...
## $ slidedMother : num NA NA NA NA NA NA NA NA NA ...
## $ slidedChild  : num 2.57e-03 1.55e-03 2.54e-04 2.09e-04 5.28e-05 ...
## $ frame_index  : int 1 2 3 4 5 6 7 8 9 10 ...
## $ slidedParent : num 0.002084 0.001758 0.000729 0.000693 0.000537 ...

```

```

## $ CutBefore      : chr  "00:11" "00:11" "00:11" "00:11" ...
## $ CutMiddle1    : chr  "05:30" "05:30" "05:30" "05:30" ...
## $ CutMiddle2    : chr  "06:16" "06:16" "06:16" "06:16" ...
## $ CutFinal       : chr  "16:27" "16:27" "16:27" "16:27" ...
## $ ChildSex       : Factor w/ 2 levels "Female","Male": 1 1 1 1 1 1 1 1 1 ...
## $ ParentSex      : Factor w/ 2 levels "Female","Male": 2 2 2 2 2 2 2 2 2 ...
## $ CutBeforeMin   : num  0.183 0.183 0.183 0.183 0.183 ...
## $ CutMiddle1Min : num  5.5 5.5 5.5 5.5 5.5 5.5 5.5 5.5 5.5 ...
## $ CutMiddle2Min : num  6.27 6.27 6.27 6.27 6.27 ...
## $ CutFinalMin   : num  16.4 16.4 16.4 16.4 16.4 ...
## $ LabelVideo     : chr  "Cut" "Cut" "Cut" "Cut" ...
## $ Time_min       : num  0.000667 0.001333 0.002 0.002667 0.003333 ...

str(psycho)

## 'data.frame': 34 obs. of 7 variables:
## $ Num._ident_videos : Factor w/ 34 levels "1606","BAJE059",...: 1 2 3 4 5 6 7 8 9 10 ...
## $ attachement_cluster: Factor w/ 5 levels "DU","DUAL","FE",...: 3 3 4 4 4 2 4 3 3 5 ...
## $ insecurite_level  : Factor w/ 9 levels "","Clearly","Markedly",...: 3 5 2 2 2 3 2 7 4 5 ...
## $ TAS_total        : int  56 41 64 47 42 65 49 47 59 56 ...
## $ global_score     : Factor w/ 14 levels "","10 Midly Fearful",...: 6 10 5 5 5 13 12 11 2 3 ...
## $ STAIYB_total    : int  70 43 33 48 46 44 34 44 35 26 ...
## $ BDI_total        : int  17 7 4 10 8 7 2 4 3 0 ...

MH1 <- merge(MH, psycho, by.x="family", by.y="Num._ident_videos")
MH1 <- MH1[-which(MH1$LabelVideo=="Cut"),]
str(MH1)

## 'data.frame': 755257 obs. of 24 variables:
## $ family          : chr  "1606" "1606" "1606" "1606" ...
## $ slidedFather    : num  0.000116 0.000327 0.000807 0.001484 0.002285 ...
## $ slidedMother    : num  NA NA NA NA NA NA NA NA NA ...
## $ slidedChild     : num  0.00671 0.00784 0.00867 0.00907 0.00888 ...
## $ frame_index     : int  276 277 278 279 280 281 282 283 284 285 ...
## $ slidedParent    : num  0.000116 0.000327 0.000807 0.001484 0.002285 ...
## $ CutBefore       : chr  "00:11" "00:11" "00:11" "00:11" ...
## $ CutMiddle1      : chr  "05:30" "05:30" "05:30" "05:30" ...
## $ CutMiddle2      : chr  "06:16" "06:16" "06:16" "06:16" ...
## $ CutFinal        : chr  "16:27" "16:27" "16:27" "16:27" ...
## $ ChildSex        : Factor w/ 2 levels "Female","Male": 1 1 1 1 1 1 1 1 1 ...
## $ ParentSex       : Factor w/ 2 levels "Female","Male": 2 2 2 2 2 2 2 2 2 ...
## $ CutBeforeMin    : num  0.183 0.183 0.183 0.183 0.183 ...
## $ CutMiddle1Min  : num  5.5 5.5 5.5 5.5 5.5 5.5 5.5 5.5 5.5 ...
## $ CutMiddle2Min  : num  6.27 6.27 6.27 6.27 6.27 ...
## $ CutFinalMin    : num  16.4 16.4 16.4 16.4 16.4 ...
## $ LabelVideo      : chr  "No Conflict" "No Conflict" "No Conflict" "No Conflict" ...
## $ Time_min        : num  0.184 0.185 0.185 0.186 0.187 ...
## $ attachement_cluster: Factor w/ 5 levels "DU","DUAL","FE",...: 3 3 3 3 3 3 3 3 3 ...
## $ insecurite_level : Factor w/ 9 levels "","Clearly","Markedly",...: 3 3 3 3 3 3 3 3 3 ...
## $ TAS_total        : int  56 56 56 56 56 56 56 56 56 ...
## $ global_score     : Factor w/ 14 levels "","10 Midly Fearful",...: 6 6 6 6 6 6 6 6 6 ...
## $ STAIYB_total    : int  70 70 70 70 70 70 70 70 70 ...
## $ BDI_total        : int  17 17 17 17 17 17 17 17 17 ...

```

```

MHagg <- aggregate(MH1, by=list(MH1$family, MH1$LabelVideo), FUN=mean)

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
## returning NA

## Warning in mean.default(X[[i]], ...): argument is not numeric or logical:
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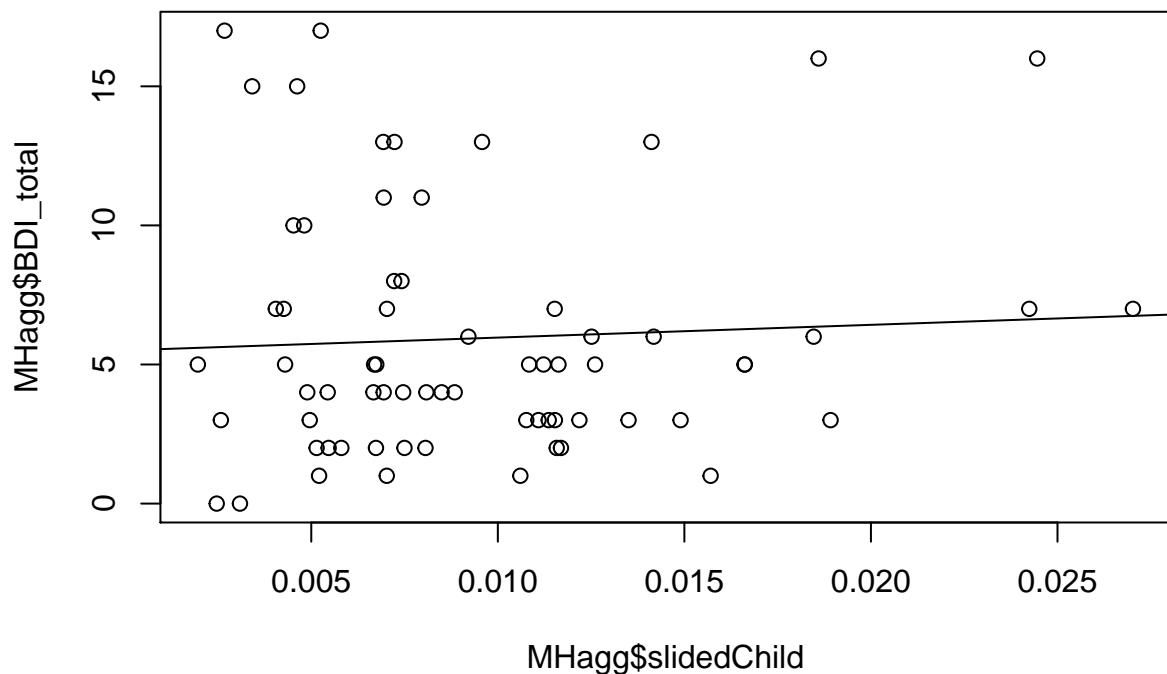
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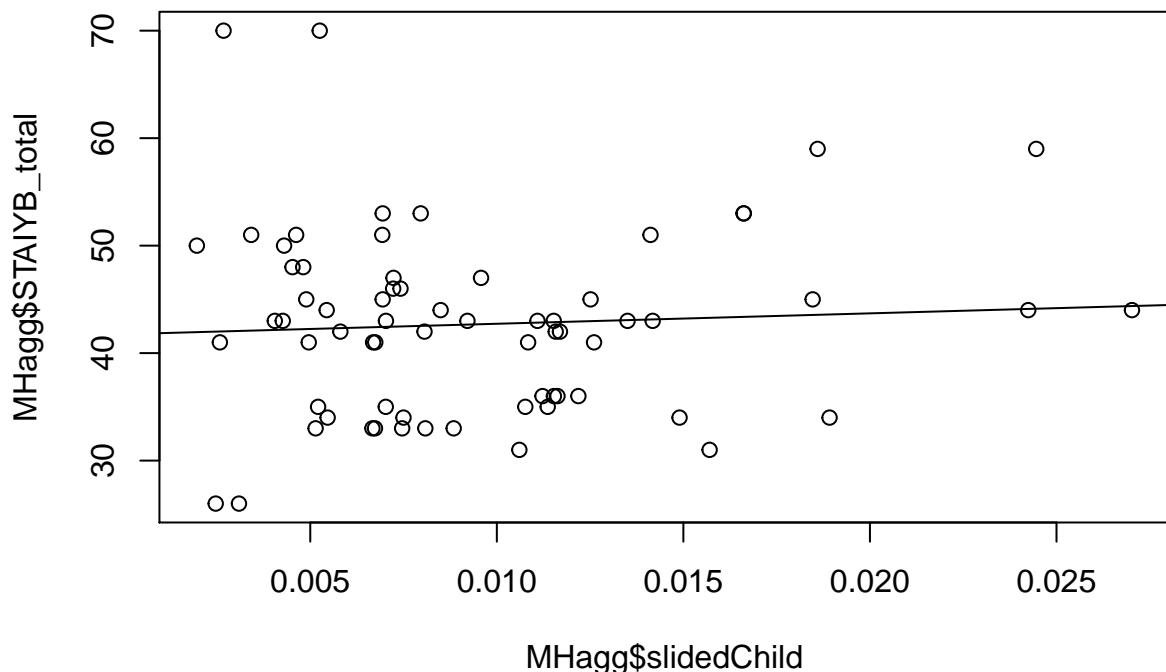



```
cor.test(MHagg$slidedChild, MHagg$BDI_total)
```

```
##
## Pearson's product-moment correlation
##
## data: MHagg$slidedChild and MHagg$BDI_total
## t = 0.45377, df = 66, p-value = 0.6515
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.1851177 0.2903334
## sample estimates:
## cor
## 0.05576882
```

#View(MH1)

```
plot(MHagg$slidedChild, MHagg$STAIYB_total)
lm <- lm(MHagg$STAIYB_total ~ MHagg$slidedChild)
abline(lm)
```



```

cor.test(MHagg$slidedChild, MHagg$STAIYB_total)

##
## Pearson's product-moment correlation
##
## data: MHagg$slidedChild and MHagg$STAIYB_total
## t = 0.48863, df = 66, p-value = 0.6267
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.1809776 0.2942512
## sample estimates:
##       cor
## 0.06003815

library(reshape2)
SSIPsycho <- merge(SSInoLog, psycho, by.x="family", by.y="Num._ident_videos")
data_long <- melt(SSInoLog, id.vars = c('family', 'ChildSex', "ParentSex"), measure.vars = c(1,2,3,4))

## Warning: attributes are not identical across measure variables; they will
## be dropped

SSIPsycho$CutBefore <-NULL
SSIPsycho$SSI_fa_ch <-NULL
SSIPsycho$SSI_mo_ch <-NULL
SSIPsycho$CutMiddle1 <-NULL
SSIPsycho$CutMiddle2 <-NULL
SSIPsycho$CutFinal <-NULL
SSIPsycho$CutBeforeMin <-NULL
SSIPsycho$CutMiddle1Min <-NULL
SSIPsycho$CutMiddle2Min <-NULL
SSIPsycho$CutFinalMin <-NULL
SSIPsycho$global_score <-NULL
SSIPsycho <- SSIPsycho[-which(SSIPsycho$LabelVideo=="Cut"),]
#View(data_long)

```

```

write.csv(SSIPsycho, file="SSIPsycho.csv")

SSIPsycho$LabelVideo <- as.factor(SSIPsycho$LabelVideo)

mod1 <- glmer(LabelVideo ~ SSI + (1|family), data=SSIPsycho, family=binomial, nAGQ=20)
summary(mod1)

## Generalized linear mixed model fit by maximum likelihood (Adaptive
## Gauss-Hermite Quadrature, nAGQ = 20) [glmerMod]
## Family: binomial ( logit )
## Formula: LabelVideo ~ SSI + (1 | family)
## Data: SSIPsycho
##
##      AIC      BIC  logLik deviance df.resid
##    7784.2  7804.3 -3889.1    7778.2     6037
##
## Scaled residuals:
##      Min      1Q  Median      3Q      Max
## -0.7363 -0.7245 -0.7236  1.3793  1.3823
##
## Random effects:
##   Groups Name      Variance Std.Dev.
##   family (Intercept) 0        0
## Number of obs: 6040, groups: family, 34
##
## Fixed effects:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.64745   0.03400 -19.043  <2e-16 ***
## SSI          0.04105   0.23601   0.174    0.862
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##      (Intr)
## SSI -0.605

mod <- glmer(LabelVideo ~ SSI + (1|family) + ChildSex + ParentSex + attachement_cluster, data=SSIPsycho)
summary(mod)

## Generalized linear mixed model fit by maximum likelihood (Adaptive
## Gauss-Hermite Quadrature, nAGQ = 20) [glmerMod]
## Family: binomial ( logit )
## Formula:
## LabelVideo ~ SSI + (1 | family) + ChildSex + ParentSex + attachement_cluster
## Data: SSIPsycho
##
##      AIC      BIC  logLik deviance df.resid
##    7794.6  7855.0 -3888.3    7776.6     6031
##
## Scaled residuals:
##      Min      1Q  Median      3Q      Max
## -0.7553 -0.7292 -0.7181  1.3571  1.4347
##
## Random effects:

```

```

## Groups Name      Variance Std.Dev.
## family (Intercept) 0        0
## Number of obs: 6040, groups: family, 34
##
## Fixed effects:
##                               Estimate Std. Error z value Pr(>|z|)
## (Intercept)                -0.72196   0.16244 -4.445 8.81e-06 ***
## SSI                      0.03907   0.23641  0.165  0.869
## ChildSexMale               0.02778   0.07215  0.385  0.700
## ParentSexMale              -0.05291   0.07342 -0.721  0.471
## attachement_clusterDUAL    0.09810   0.19702  0.498  0.619
## attachement_clusterFE      0.11124   0.16917  0.658  0.511
## attachement_clusterSecure   0.06600   0.16770  0.394  0.694
## attachement_clusterWithdrawn 0.01896   0.18736  0.101  0.919
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##             (Intr) SSI     ChldSM PrntSM a_DUAL att_FE attc_S
## SSI          -0.106
## ChildSexMal -0.004  0.034
## ParentSexMl -0.002  0.020  0.013
## attchm_DUAL -0.813 -0.022 -0.001  0.000
## attchmnt_FE -0.947 -0.021 -0.068 -0.134  0.783
## attchmnt_cS -0.955 -0.031 -0.089 -0.064  0.790  0.934
## attchmnt_cW -0.856 -0.012 -0.133 -0.002  0.707  0.832  0.842
mod <- glmer(attachement_cluster ~ SSI + (1|family) + LabelVideo, data=SSIPsycho, family=binomial , nAGQ=20)
summary(mod)

## Generalized linear mixed model fit by maximum likelihood (Adaptive
## Gauss-Hermite Quadrature, nAGQ = 20) [glmerMod]
## Family: binomial ( logit )
## Formula: attachement_cluster ~ SSI + (1 | family) + LabelVideo
## Data: SSIPsycho
##
##      AIC      BIC  logLik deviance df.resid
## 28.8    55.6   -10.4    20.8      6036
##
## Scaled residuals:
##      Min      1Q  Median      3Q     Max
## -0.007148  0.000000  0.000000  0.000000  0.000000
##
## Random effects:
## Groups Name      Variance Std.Dev.
## family (Intercept) 9036    95.06
## Number of obs: 6040, groups: family, 34
##
## Fixed effects:
##                               Estimate Std. Error z value Pr(>|z|)
## (Intercept)                44.8007   49.3331  0.908  0.364
## SSI                      0.7120    59.2376  0.012  0.990
## LabelVideoNo Conflict    0.0394   12.7557  0.003  0.998
##
## Correlation of Fixed Effects:

```

```

##          (Intr) SSI
## SSI      -0.094
## LblVdNCnflc -0.086  0.002

mod <- glmer(attachement_cluster ~ SSI + (1|family) + LabelVideo, data=SSIPsycho, family=binomial , nAGQ=20)
summary(mod)

## Generalized linear mixed model fit by maximum likelihood (Adaptive
## Gauss-Hermite Quadrature, nAGQ = 20) [glmerMod]
## Family: binomial ( logit )
## Formula: attachement_cluster ~ SSI + (1 | family) + LabelVideo
## Data: SSIPsycho
##
##      AIC      BIC  logLik deviance df.resid
##      28.8     55.6    -10.4     20.8     6036
##
## Scaled residuals:
##      Min      1Q  Median      3Q      Max
## -0.007148  0.000000  0.000000  0.000000  0.000000
##
## Random effects:
##   Groups Name      Variance Std.Dev.
##   family (Intercept) 9036     95.06
## Number of obs: 6040, groups: family, 34
##
## Fixed effects:
##                  Estimate Std. Error z value Pr(>|z|)
## (Intercept)      44.8007   49.3331  0.908   0.364
## SSI             0.7120   59.2376  0.012   0.990
## LabelVideoNo Conflict  0.0394   12.7557  0.003   0.998
##
## Correlation of Fixed Effects:
##          (Intr) SSI
## SSI      -0.094
## LblVdNCnflc -0.086  0.002

mod <- glmer(attachement_cluster ~ SSI + (1|family) + LabelVideo, data=SSIPsycho, family=binomial , nAGQ=20)
summary(mod)

## Generalized linear mixed model fit by maximum likelihood (Adaptive
## Gauss-Hermite Quadrature, nAGQ = 20) [glmerMod]
## Family: binomial ( logit )
## Formula: attachement_cluster ~ SSI + (1 | family) + LabelVideo
## Data: SSIPsycho
##
##      AIC      BIC  logLik deviance df.resid
##      28.8     55.6    -10.4     20.8     6036
##
## Scaled residuals:
##      Min      1Q  Median      3Q      Max
## -0.007148  0.000000  0.000000  0.000000  0.000000
##
## Random effects:
##   Groups Name      Variance Std.Dev.
##   family (Intercept) 9036     95.06

```

```

## Number of obs: 6040, groups: family, 34
##
## Fixed effects:
##                               Estimate Std. Error z value Pr(>|z|)
## (Intercept)             44.8007   49.3331  0.908   0.364
## SSI                   0.7120   59.2376  0.012   0.990
## LabelVideoNo Conflict  0.0394   12.7557  0.003   0.998
##
## Correlation of Fixed Effects:
##          (Intr) SSI
## SSI      -0.094
## LblVdNCnflc -0.086  0.002

SSIPsycho$attachement_duo <- rep(NA)
SSIPsycho[which(SSSIPsycho$attachement_cluster=="Secure"),]$attachement_duo <- "Secure"
SSIPsycho[which(SSSIPsycho$attachement_cluster!="Secure"),]$attachement_duo <- "Unsecure"
SSIPsycho$attachement_duo <- as.factor(SSSIPsycho$attachement_duo)

mod <- glmer(attachement_duo ~ SSI + (1|family) + LabelVideo, data=SSIPsycho, family=binomial , nAGQ=20)
summary(mod)

## Generalized linear mixed model fit by maximum likelihood (Adaptive
## Gauss-Hermite Quadrature, nAGQ = 20) [glmerMod]
## Family: binomial ( logit )
## Formula: attachement_duo ~ SSI + (1 | family) + LabelVideo
## Data: SSIPsycho
##
##      AIC      BIC  logLik deviance df.resid
## 61.7    88.6   -26.9     53.7     6036
##
## Scaled residuals:
##      Min      1Q  Median      3Q      Max
## -0.006484 -0.005592  0.004850  0.005008  0.005612
##
## Random effects:
## Groups Name        Variance Std.Dev.
## family (Intercept) 2090     45.72
## Number of obs: 6040, groups: family, 34
##
## Fixed effects:
##                               Estimate Std. Error z value Pr(>|z|)
## (Intercept)             1.313203   4.507116  0.291   0.771
## SSI                  -0.305572  11.498227 -0.027   0.979
## LabelVideoNo Conflict  0.006044   2.761918  0.002   0.998
##
## Correlation of Fixed Effects:
##          (Intr) SSI
## SSI      -0.222
## LblVdNCnflc -0.211 -0.003

mod1 <- glmer(LabelVideo ~ SSI + (1|family) + attachement_duo, data=SSIPsycho, family=binomial, nAGQ=20)
summary(mod1)

## Generalized linear mixed model fit by maximum likelihood (Adaptive
## Gauss-Hermite Quadrature, nAGQ = 20) [glmerMod]

```

```

## Family: binomial ( logit )
## Formula: LabelVideo ~ SSI + (1 | family) + attachement_duo
## Data: SSIPsycho
##
##      AIC      BIC  logLik deviance df.resid
## 7786.1 7812.9 -3889.0    7778.1     6036
##
## Scaled residuals:
##      Min      1Q  Median      3Q      Max
## -0.7401 -0.7267 -0.7204  1.3753  1.3898
##
## Random effects:
## Groups Name        Variance Std.Dev.
## family (Intercept) 0         0
## Number of obs: 6040, groups: family, 34
##
## Fixed effects:
##                               Estimate Std. Error z value Pr(>|z|)
## (Intercept)              -0.65834   0.04632 -14.212 <2e-16 ***
## SSI                      0.04390   0.23615   0.186   0.853
## attachement_duoUnsecure  0.01894   0.05463   0.347   0.729
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##          (Intr) SSI
## SSI      -0.467
## attachmnt_dU -0.679  0.035

```