

## Supplementary Material

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## Hypothesis 1

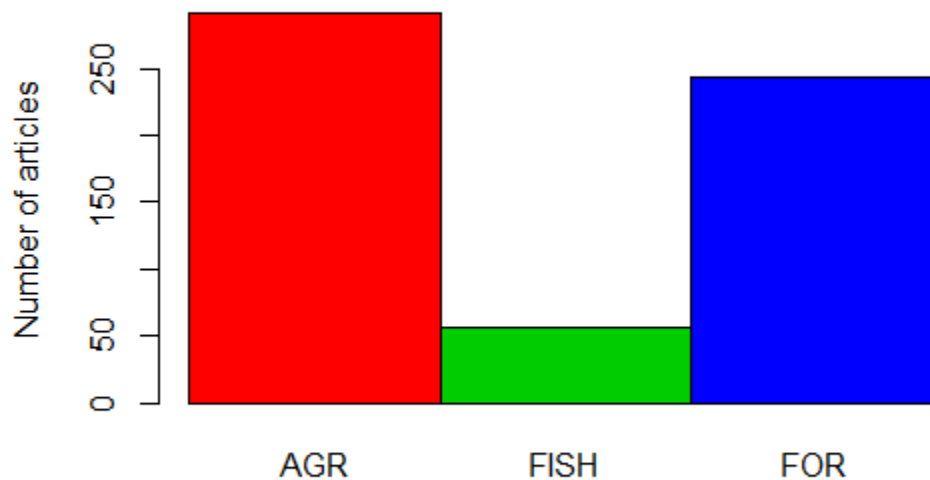
1. The application of the Multi-Criteria Decision-Making (MCDM) techniques does not significantly differ in the three fields (Agriculture, Forestry and Fishery), given that we have not seen comparative studies that support the opposite.

==> We reject the Hypothesis

==> With DEA: We reject the null hypothesis that they are independent. They significantly differ when the field is changed.

==> Without DEA: We cannot reject the null hypothesis that they are independent. They do not significantly differ when the field is changed.

```
db$Campo <- NA; db$Campo[db$FOR > 0 ] <- "FOR"; db$Campo[db$AGR > 0 ] <- "AGR"; db$Campo[db$FISH > 0 ] <- "FISH"
db$Campo[(db$FOR > 0 & db$AGR > 0) | (db$FISH > 0 & db$AGR > 0) | (db$FISH > 0 & db$FOR > 0)] <- NA
tbl <- table(db$Campo)
barplot(as.matrix(tbl), beside = TRUE, col=c(2:4), ylab="Número de artículos",names.arg = rownames(tbl)
)
```

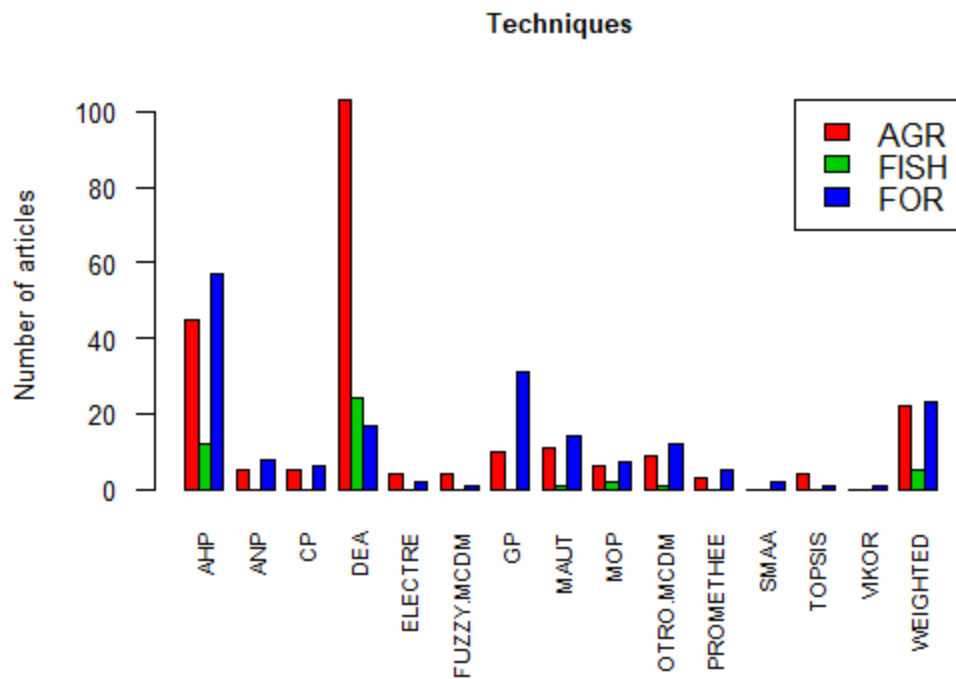


```
chisq.test(tbl) # The number of items differs significantly when we change the field.
```

```
##
## Chi-squared test for given probabilities
##
```

```
## data: tbl
## X-squared = 155.09, df = 2, p-value < 2.2e-16

Tecnicas_name <- c("GP","CP","TOPSIS","MOP","VIKOR","ELECTRE","PROMETHEE","AHP","ANP","DE
A",
  "MAUT","WEIGHTED","SMAA","FUZZY.MCDM","OTRO.MCDM")
db$Tecnicas <- NA
for(i in 1:length(Tecnicas_name)){
  db$Tecnicas[db[,Tecnicas_name[i]] > 0 ] <- Tecnicas_name[i]}
db$Tecnicas[rowSums(db[,Tecnicas_name], na.rm=TRUE) > 1] <- NA
tbl <- table(db$Campo, db$Tecnicas)
barplot(as.matrix(tbl), beside = TRUE, las=2, col=c(2:4), ylab="Número de artículos",
  main = "Técnicas", cex.main=0.8, cex.axis=0.8, cex.names = 0.7, cex.lab=0.8)
legend("topright",rownames(tbl), fill=c(2:4))
```



```
fisher.test(tbl,simulate.p.value=TRUE)#,B=1e7)

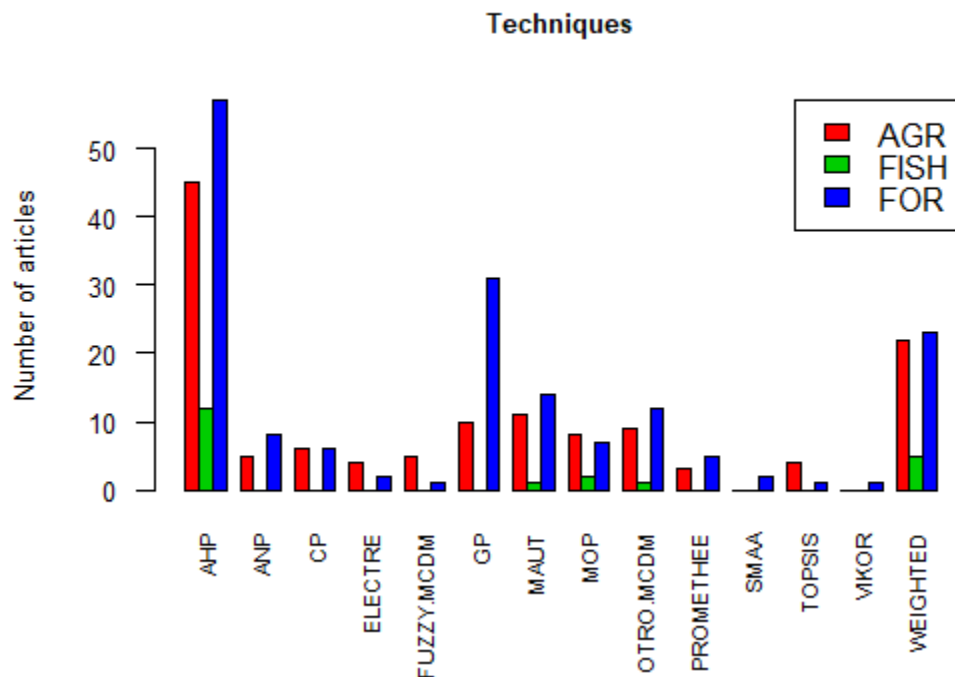
##
## Fisher's Exact Test for Count Data with simulated p-value (based
## on 2000 replicates)
##
## data: tbl
## p-value = 0.0004998
## alternative hypothesis: two.sided

Tecnicas_sinDEA_name <- c("GP","CP","TOPSIS","MOP","VIKOR","ELECTRE","PROMETHEE","AHP","A
NP",
```

```

"MAUT","WEIGHTED","SMAA","FUZZY.MCDM","OTRO.MCDM")
db$Tecnicas_sinDEA <- NA
for(i in 1:length(Tecnicas_sinDEA_name)){
  db$Tecnicas_sinDEA[db[,Tecnicas_sinDEA_name[i]] > 0 ] <- Tecnicas_sinDEA_name[i]}
db$Tecnicas_sinDEA[rowSums(db[,Tecnicas_sinDEA_name], na.rm=TRUE) > 1] <- NA
tbl <- table(db$Campo, db$Tecnicas_sinDEA)
barplot(as.matrix(tbl), beside = TRUE, las=2, col=c(2:4), ylab="Número de artículos",
  main = "Técnicas", cex.main=0.8, cex.axis=0.8, cex.names = 0.7, cex.lab=0.8)
legend("topright",rownames(tbl), fill=c(2:4))

```



```

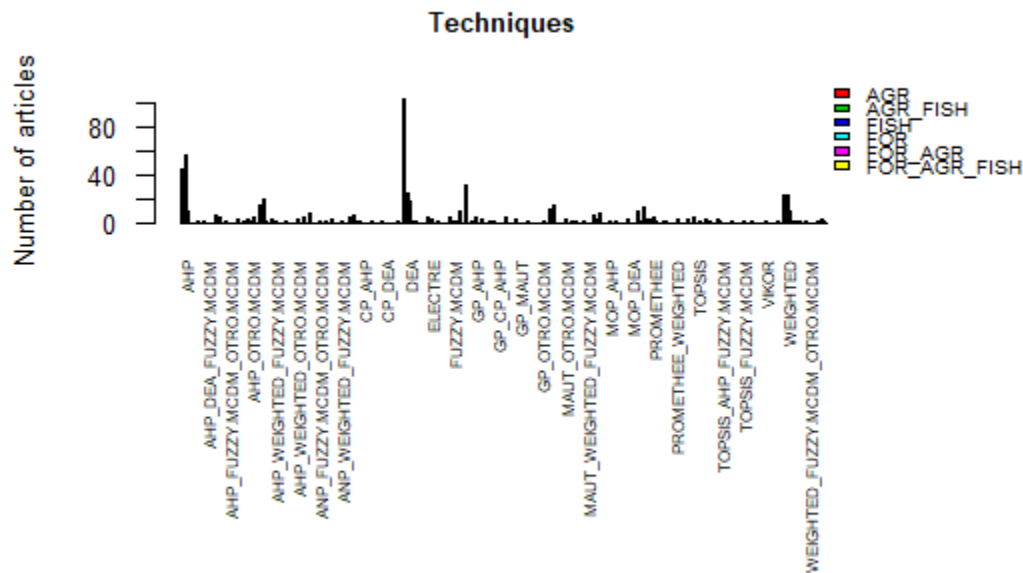
fisher.test(tbl,simulate.p.value=TRUE)#,B=1e7)
##
## Fisher's Exact Test for Count Data with simulated p-value (based
## on 2000 replicates)
##
## data: tbl
## p-value = 0.2484
## alternative hypothesis: two.sided

db$Campo_mixtos <- db$Campo
db$Campo_mixtos[(db$FOR > 0 & db$AGR > 0) ] <- "FOR_AGR"
db$Campo_mixtos[(db$FOR > 0 & db$FISH > 0) ] <- "FOR_FISH"
db$Campo_mixtos[(db$AGR > 0 & db$FISH > 0) ] <- "AGR_FISH"
db$Campo_mixtos[(db$FOR > 0 & db$AGR > 0 & db$FISH > 0) ] <- "FOR_AGR_FISH"
table(db$Campo_mixtos)

```

```
##
##   AGR  AGR_FISH  FISH  FOR  FOR_AGR
##   291    1    57   243   30
## FOR_AGR_FISH
##    1

db$Tecnicas_mixto <- db$Tecnicas
a <- db[,Tecnicas_name ]
obs_greater1_tech <- rowSums(a) > 1 # to identify observations with more than 1 technique
for(i in 1:nrow(a)){
  if(obs_greater1_tech[i] == TRUE){
    db$Tecnicas_mixto[i] <- paste(colnames(a)[which(a[i,] %in% 1)], collapse = "_")
  }
}
tbl <- table(db$Campo_mixtos, db$Tecnicas_mixto)
par(mar=c(12.1, 4.1, 4.1, 5.1))
barplot(as.matrix(tbl), beside = TRUE, las=2, col=c(2:7), ylab="Número de artículos",
        main = "Técnicas", cex.main=0.8, cex.axis=0.8, cex.names = 0.5, cex.lab=0.8)
legend(400,120,rownames(tbl), fill=c(2:7), cex=0.6,bty = "n", xpd=TRUE)
```



```
fisher.test(tbl,simulate.p.value=TRUE)#,B=1e7)

##
## Fisher's Exact Test for Count Data with simulated p-value (based
## on 2000 replicates)
##
## data: tbl
```

```
## p-value = 0.0004998
## alternative hypothesis: two.sided
par(mar=c(5, 4, 4, 2) + 0.1)
```

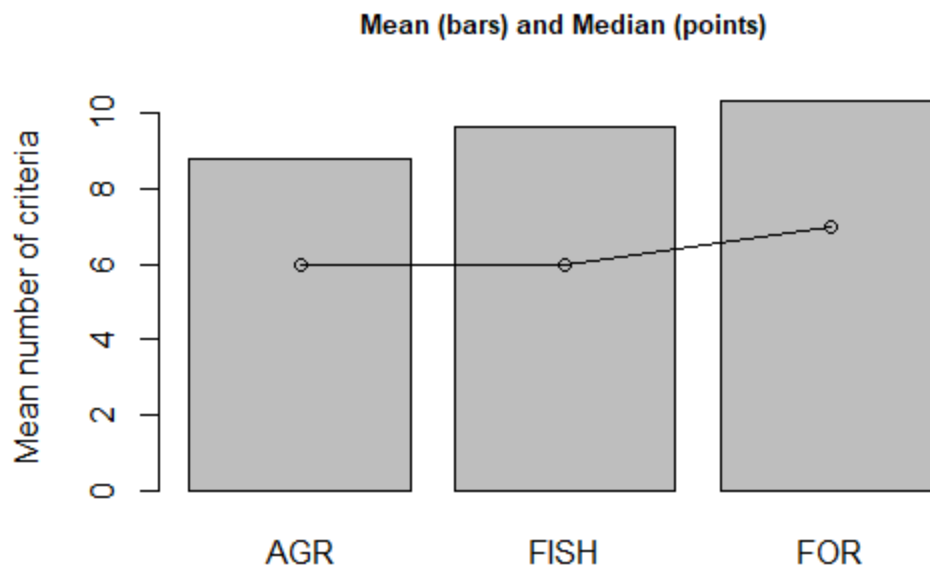
## Hypothesis 2

- The number of criteria used in MCDM problems is similar in the three fields (Agriculture, Forestry and Fishery)

==> We accept the Hypothesis

==> We cannot reject the null hypothesis that they are equal.

```
tbl_mean <- aggregate(db[, c("nº.criterios")], list(db$Campo), mean, na.rm=TRUE)
tbl_median <- aggregate(db[, c("nº.criterios")], list(db$Campo), median, na.rm=TRUE)
df.bar <- barplot(tbl_mean$x, names.arg = tbl_mean$Group.1, ylab="Número medio de criterios",
  main="Media (barras) y Mediana (puntos)", cex.main=0.8)
lines(x = df.bar, y = tbl_median$x); points(x = df.bar, y = tbl_median$x)
```



```
summary(aov(db$nº.criterios ~ db$Campo))
##           Df Sum Sq Mean Sq F value Pr(>F)
## db$Campo  2   297   148.6   1.255 0.286
## Residuals 569 67358   118.4
## 56 observations deleted due to missingness
```

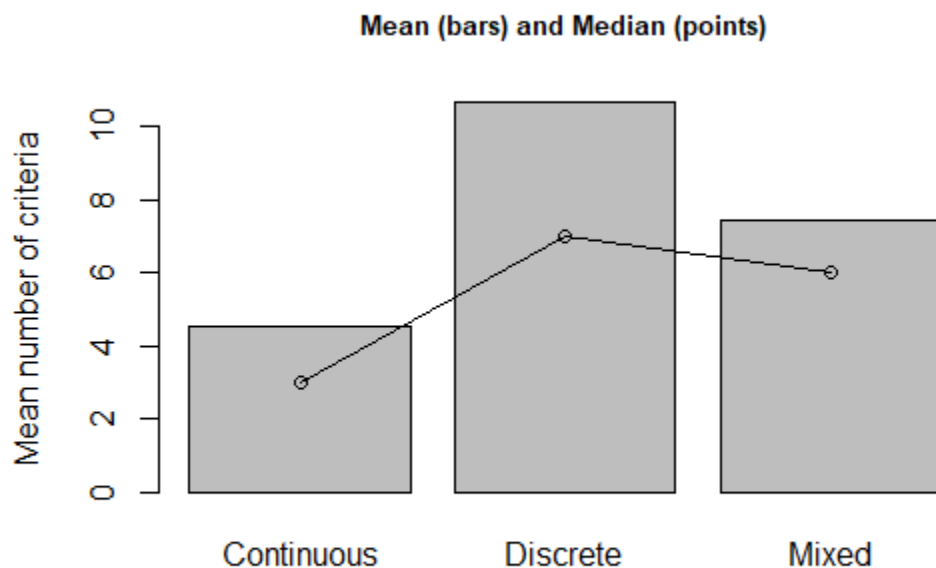
### Hypothesis 3

3. The number of criteria is lower in continuous problems than in discrete problems.

==> We accept the Hypothesis

==> We reject the null hypothesis the fact that they are equal. The number of criteria in continuous is less than in discrete.

```
Continuos <- c("GP","CP","TOPSIS","MOP","VIKOR")
Discretos <-
c("ELECTRE","PROMETHEE","AHP","ANP","DEA","MAUT","WEIGHTED","SMAA","FUZZY.MCDM","
OTRO.MCDM")
Continuos0 <- rowSums(db[,Continuos]); Continuos0[Continuos0 >0] <- 1; db$Continuos <- Continuos0
Discretos0 <- rowSums(db[,Discretos]); Discretos0[Discretos0 >0] <- 1; db$Discretos <- Discretos0
db$Cont_Disc <- NA; db$Cont_Disc[db$Continuos > 0] <- "Continuo"; db$Cont_Disc[db$Discretos > 0] <-
"Discreto";
db$Cont_Disc[db$Continuos > 0 & db$Discretos > 0] <- "Mixto"
tbl_mean <- aggregate(db[, c("nº.criterios")], list(db$Cont_Disc), mean, na.rm=TRUE)
tbl_median <- aggregate(db[, c("nº.criterios")], list(db$Cont_Disc), median, na.rm=TRUE)
df.bar <- barplot(tbl_mean$x, names.arg = tbl_mean$Group.1, ylab="Número medio de criterios",
main="Media (barras) y Mediana (puntos)", cex.main=0.8)
lines(x = df.bar, y = tbl_median$x); points(x = df.bar, y = tbl_median$x)
```



```
a <- db$Cont_Disc; a[a=="Mixto"] <- NA
t.test(db$nº.criterios ~ a)
##
## Welch Two Sample t-test
```



```

##
## data: db$nº.criterios by a
## t = -9.1647, df = 442.38, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -7.460490 -4.825736
## sample estimates:
## mean in group Continuo mean in group Discreto
##      4.513514      10.656627

t.test(db$nº.criterios ~ a, alternative = "less")

##
## Welch Two Sample t-test
##
## data: db$nº.criterios by a
## t = -9.1647, df = 442.38, p-value < 2.2e-16
## alternative hypothesis: true difference in means is less than 0
## 95 percent confidence interval:
## -Inf -5.038247
## sample estimates:
## mean in group Continuo mean in group Discreto
##      4.513514      10.656627

summary(aov(db$nº.criterios ~ db$Cont_Disc))

##      Df Sum Sq Mean Sq F value Pr(>F)
## db$Cont_Disc  2  2609 1304.7  9.993 5.38e-05 ***
## Residuals    601 78467  130.6
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## 24 observations deleted due to missingness

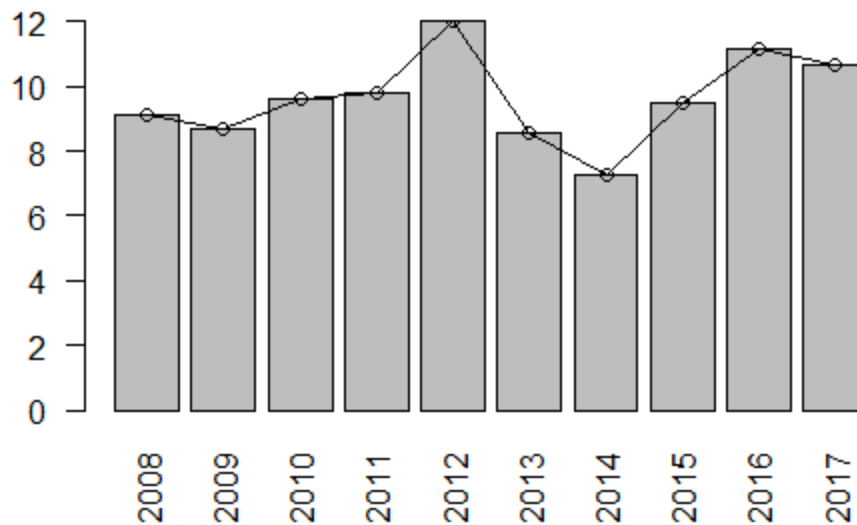
summary(db$nº.criterios)

##  Min. 1st Qu.  Median  Mean 3rd Qu.  Max.  NA's
## 1.000  4.000  6.000  9.724 11.000 133.000  22

tbl_mean <- aggregate(db[, c("nº.criterios")], list(db$year), mean, na.rm=TRUE)
df.bar <- barplot(tbl_mean$x, las=2, main="Evolución del número total de criterios", cex.main=0.8,
names.arg = tbl_mean$Group.1)
lines(x = df.bar, y = tbl_mean$x); points(x = df.bar, y = tbl_mean$x)

```

Evolution of the number of criteria



```
tbl1.2008_2012 <- tbl_mean [1:5,];tbl1.2013_2017 <- tbl_mean [6:10,]
mean(tbl1.2008_2012[,2]); mean(tbl1.2013_2017[,2])

## [1] 9.834254
## [1] 9.421236

t.test(tbl1.2008_2012[,2], tbl1.2013_2017[,2]) # No hay un cambio significativo en el número de criterios

##
## Welch Two Sample t-test
##
## data: tbl1.2008_2012[, 2] and tbl1.2013_2017[, 2]
## t = 0.4532, df = 7.6871, p-value = 0.6629
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -1.703506 2.529543
## sample estimates:
## mean of x mean of y
## 9.834254 9.421236
```

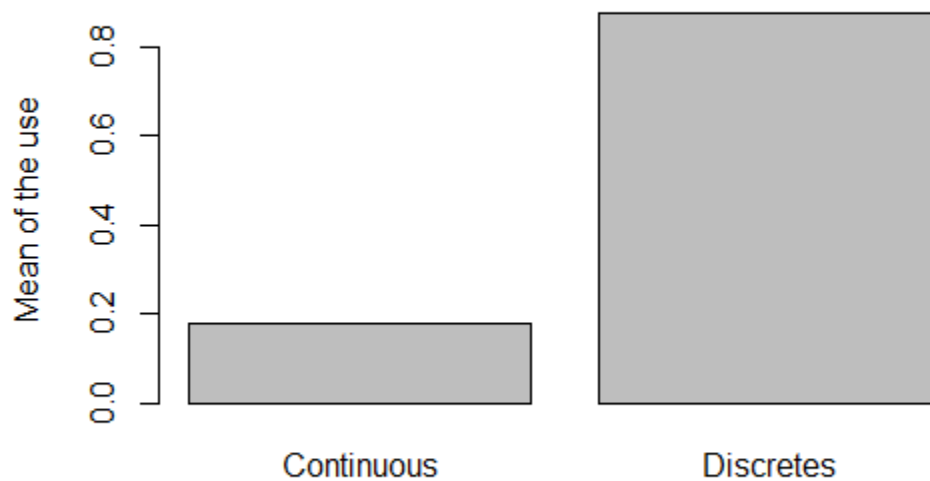
#### Hypothesis 4

4. There seems to be less use of MCDM techniques that apply to continuous problems than others that can only be applied to discrete problems.

==> We accept the Hypothesis.

==> There is less use of techniques that apply to Continuous than Discrete problems

```
barplot(c(mean(db$Continuos), mean(db$Discretos)), names.arg = c("Continuos", "Discretos"), ylab="Me  
dia de uso")
```



```
t.test(db$Continuos, db$Discretos)
```

```
##  
## Welch Two Sample t-test  
##  
## data: db$Continuos and db$Discretos  
## t = -34.541, df = 1233.1, p-value < 2.2e-16  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## -0.7370668 -0.6578377  
## sample estimates:  
## mean of x mean of y  
## 0.1751592 0.8726115
```

## Hypothesis 5

5. There is no relationship between the use of a particular MCDM technique and the fact that the case studies are from one or from several countries.

==> **We reject the Hypothesis.**

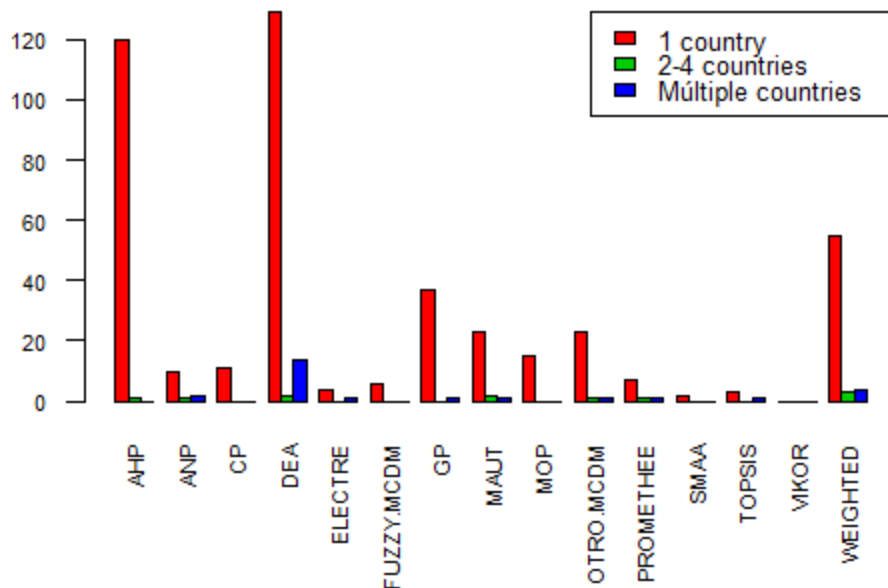
==> There exists a significant difference between the use of techniques and the number of countries. This may be because there are many techniques or groups of techniques and because in some techniques, the proportion of the number of countries is quite different.

```
db$Numero_paises[db$Numero_paises==0] <- NA; db$Numero_paises <- as.character(db$Numero_paises)
db$Numero_paises_3Cat <- NA
db$Numero_paises_3Cat[db$Numero_paises == "1"] <- "1 country"
db$Numero_paises_3Cat[db$Numero_paises == "2" | db$Numero_paises == "3" | db$Numero_paises == "4"] <- "2-4 countries"
db$Numero_paises_3Cat[db$Numero_paises == "Multiple countries"] <- "Multiple countries"
table(db$Tecnicas, db$Numero_paises_3Cat)

##
##      1 country 2-4 countries Multiple countries
## AHP          120          1           0
## ANP           10          1           2
## CP            11          0           0
## DEA           129          2          14
## ELECTRE        4          0           1
## FUZZY.MCDM     6          0           0
## GP             37          0           1
## MAUT           23          2           1
## MOP            14          0           0
## OTRO.MCDM     23          1           1
## PROMETHEE      7          1           1
## SMAA           2          0           0
## TOPSIS         3          0           1
## VIKOR          0          0           0
## WEIGHTED       54          3           4

barplot(as.matrix(t(table(db$Tecnicas, db$Numero_paises_3Cat))), beside=T, las=2, col=(2:4), cex.names=0.7, cex.axis = 0.7,
        main = "Técnicas por countries", cex.main=0.8)
legend("topright",c("1 country", "2-4 countries", "Multiple countries"), fill=c(2:4), cex=0.8)
```

Techniques by countries

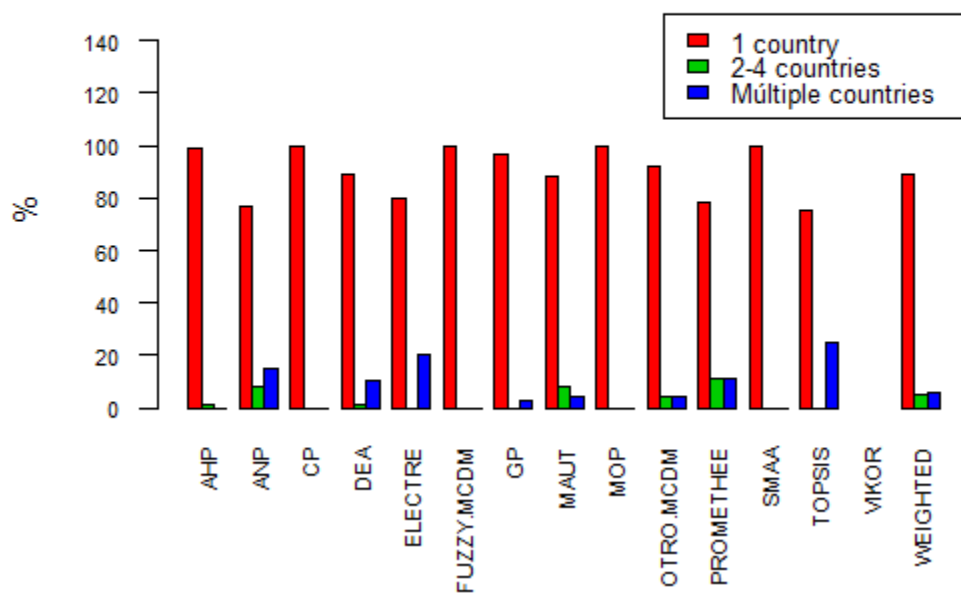


```

barplot(as.matrix(t(round(table(db$Técnicas, db$Numero_paises_3Cat) / rowSums(table(db$Técnicas, db$Numero_paises_3Cat)), 2) * 100)), beside=T, las=2, col=(2:4), cex.names= 0.7, cex.axis = 0.7,
main = "Proporciones de Técnicas por countries", cex.main=0.8, ylab="%", ylim=c(0,150))
legend("topright",c("1 country", "2-4 countries", "Multiple countries"), fill=c(2:4), cex=0.8)

```

Proportions of Techniques by countries



```

fisher.test(table(db$Tecnicas, db$Numero_paises_3Cat),simulate.p.value=TRUE)

##
## Fisher's Exact Test for Count Data with simulated p-value (based
## on 2000 replicates)
##
## data: table(db$Tecnicas, db$Numero_paises_3Cat)
## p-value = 0.007996
## alternative hypothesis: two.sided

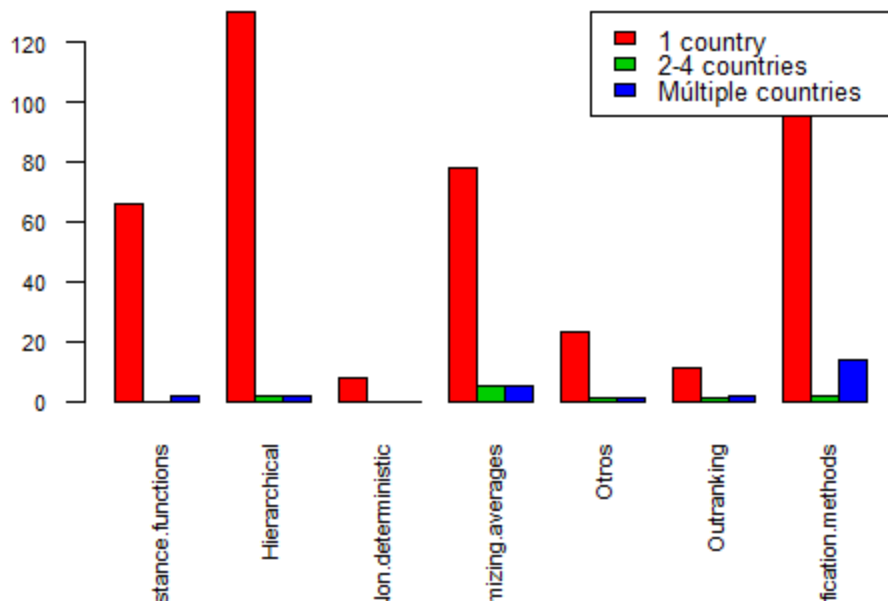
Distance.functions <- c("GP","CP","TOPSIS","MOP","VIKOR")
Outranking <- c("ELECTRE","PROMETHEE")
Hierarchical <- c("AHP","ANP")
Ranking.and.classification.methods <- "DEA"
Optimizing.averages <- c("MAUT","WEIGHTED")
Non.deterministic <- c("SMAA","FUZZY.MCDM")
Otros <- c("OTRO.MCDM")
db$Tecnicas_Grupos <- NA
db$Tecnicas_Grupos[which(db$Tecnicas %in% Distance.functions)] <- "Distance.functions"
db$Tecnicas_Grupos[which(db$Tecnicas %in% Outranking)] <- "Outranking"
db$Tecnicas_Grupos[which(db$Tecnicas %in% Hierarchical)] <- "Hierarchical"
db$Tecnicas_Grupos[which(db$Tecnicas %in% Ranking.and.classification.methods)] <- "Ranking.and.classification.methods"
db$Tecnicas_Grupos[which(db$Tecnicas %in% Optimizing.averages)] <- "Optimizing.averages"
db$Tecnicas_Grupos[which(db$Tecnicas %in% Non.deterministic)] <- "Non.deterministic"
db$Tecnicas_Grupos[which(db$Tecnicas %in% Otros)] <- "Otros"
table(db$Tecnicas_Grupos, db$Numero_paises_3Cat)

##
##           1 country 2-4 countries Multiple countries
## Distance.functions           65      0           2
## Hierarchical                 130      2           2
## Non.deterministic             8       0           0
## Optimizing.averages           77      5           5
## Otros                        23       1           1
## Outranking                    11      1           2
## Ranking.and.classification.methods 129      2          14

barplot(as.matrix(t(table(db$Tecnicas_Grupos, db$Numero_paises_3Cat))), beside=T, las=2, col=(2:4), cex
.names= 0.7, cex.axis = 0.7,
      main = "Grupos de Técnicas por countries", cex.main=0.8)
legend("topright",c("1 country", "2-4 countries", "Multiple countries"), fill=c(2:4), cex=0.8)

```

Group of techniques by countries

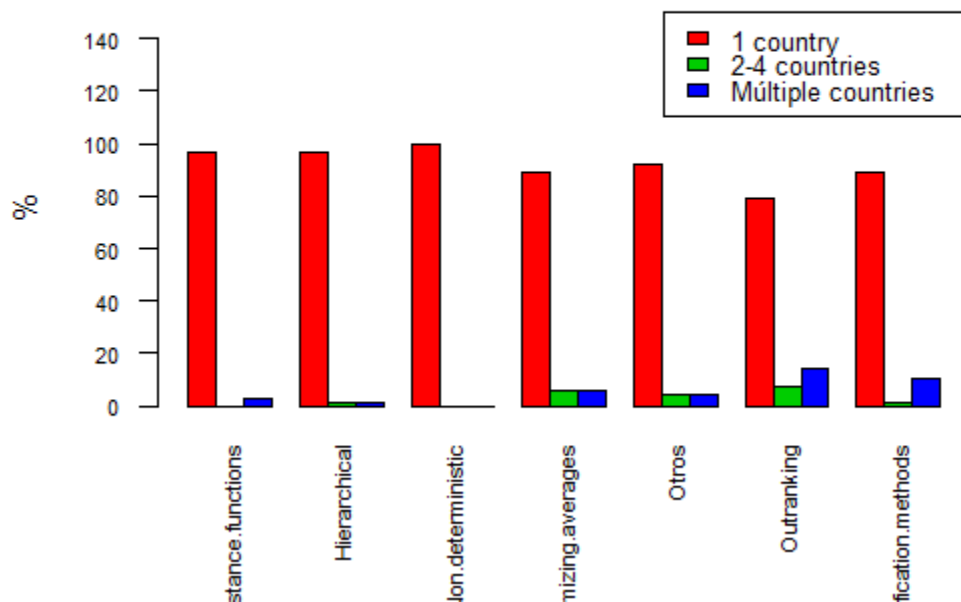


```
barplot(as.matrix(t(round(table(db$Tecnicas_Grupos, db$Numero_paises_3Cat) / rowSums(table(db$Tecnicas_Grupos, db$Numero_paises_3Cat)), 2)*100)), beside=T, las=2, col=(2:4), cex.names= 0.7, cex.axis = 0.7,
```

```
main = "Proporciones de Grupos de Técnicas por countries", cex.main=0.8, ylab="%", ylim=c(0,150))
```

```
legend("topright",c("1 country", "2-4 countries", "Multiple countries"), fill=c(2:4), cex=0.8)
```

Proportions of Group of techniques by countries



```
fisher.test(table(db$Tecnicas_Grupos, db$Numero_paises_3Cat),simulate.p.value=TRUE)
```

```
##  
## Fisher's Exact Test for Count Data with simulated p-value (based  
## on 2000 replicates)  
##  
## data: table(db$Tecnicas_Grupos, db$Numero_paises_3Cat)  
## p-value = 0.02149  
## alternative hypothesis: two.sided
```

```
fit <- multinom(Tecnicas_Grupos ~ Numero_paises_3Cat, data = db)
```

```
## # weights: 28 (18 variable)  
## initial value 934.036872  
## iter 10 value 774.855185  
## iter 20 value 769.965454  
## iter 30 value 769.933996  
## final value 769.933907  
## converged
```

```
summary(fit)
```

```
## Call:  
## multinom(formula = Tecnicas_Grupos ~ Numero_paises_3Cat, data = db)  
##  
## Coefficients:  
##  
## (Intercept)  
## Hierarchical 0.6931180  
## Non.deterministic -2.0953305  
## Optimizing.averages 0.1694081  
## Otros -1.0388386  
## Outranking -1.7764343  
## Ranking.and.classification.methods 0.6854186  
##  
## Numero_paises_3Cat2-4 paÃses  
## Hierarchical 11.927648  
## Non.deterministic -4.379108  
## Optimizing.averages 13.367685  
## Otros 12.966480  
## Outranking 13.704184  
## Ranking.and.classification.methods 11.935356  
##  
## Numero_paises_3CatMÃltiples paÃses  
## Hierarchical -0.6941672  
## Non.deterministic -11.6008832  
## Optimizing.averages 0.7462232  
## Otros 0.3447363  
## Outranking 1.7758651  
## Ranking.and.classification.methods 1.2598160  
##  
## Std. Errors:
```



```

## (Intercept)
## Hierarchical 0.1519105
## Non.deterministic 0.3747407
## Optimizing.averages 0.1684379
## Otros 0.2426100
## Outranking 0.3260167
## Ranking.and.classification.methods 0.1521061
## Numero_paises_3Cat2-4 paÃses
## Hierarchical 6.626728e-01
## Non.deterministic 7.742127e-07
## Optimizing.averages 5.121540e-01
## Otros 8.721243e-01
## Outranking 8.882524e-01
## Ranking.and.classification.methods 6.626978e-01
## Numero_paises_3CatMÃltiples paÃses
## Hierarchical 1.0114098
## Non.deterministic 665.9457003
## Optimizing.averages 0.8532568
## Otros 1.2485341
## Outranking 1.0516272
## Ranking.and.classification.methods 0.7708701
##
## Residual Deviance: 1539.868
## AIC: 1575.868

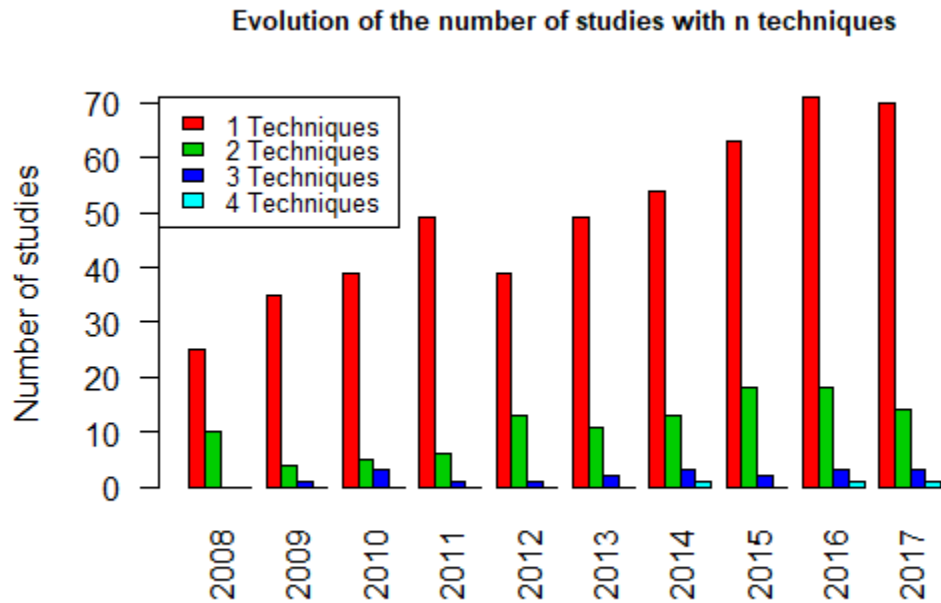
```

### Hypothesis 6

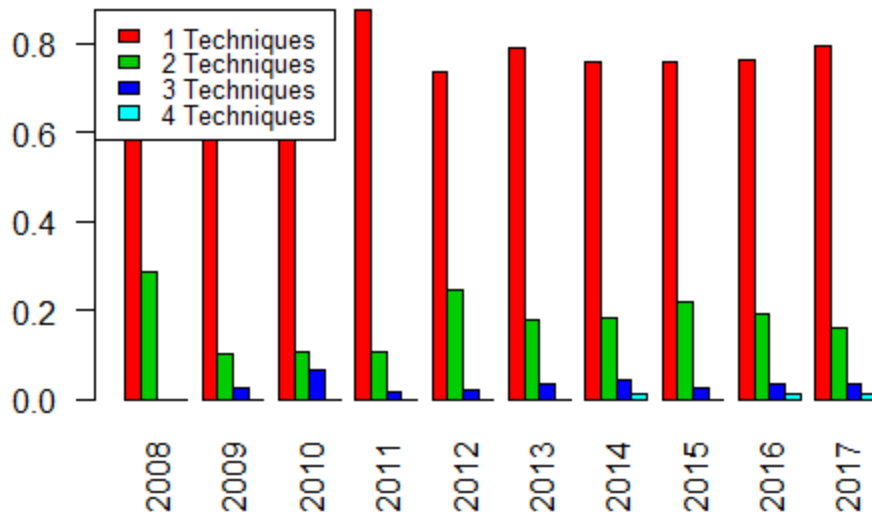
- The use in the same problem of several MCDM techniques simultaneously has increased over time

==> **The hypothesis is accepted with some nuances.**

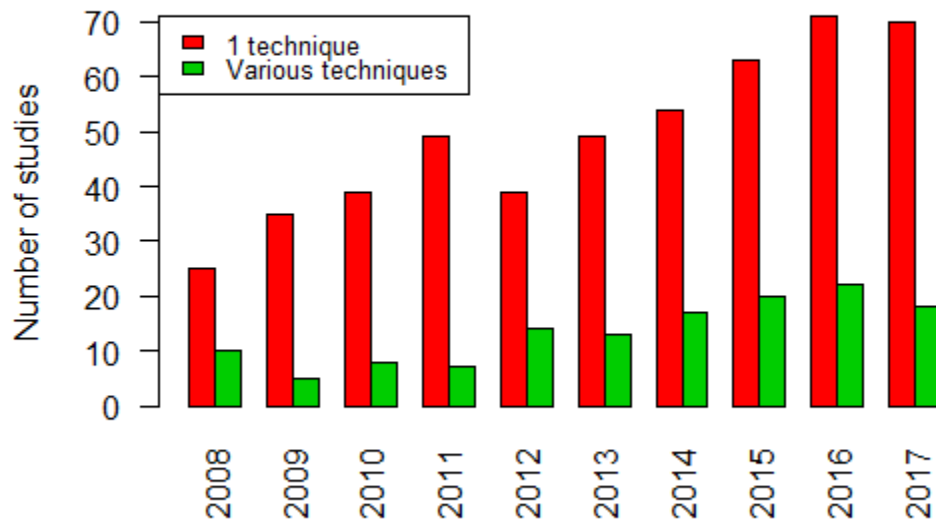
The number of total studies has increased. So has the number of studies with GDM and MCDM (significantly), but the proportion of papers with GDM and MCDM has decreased (not significant)



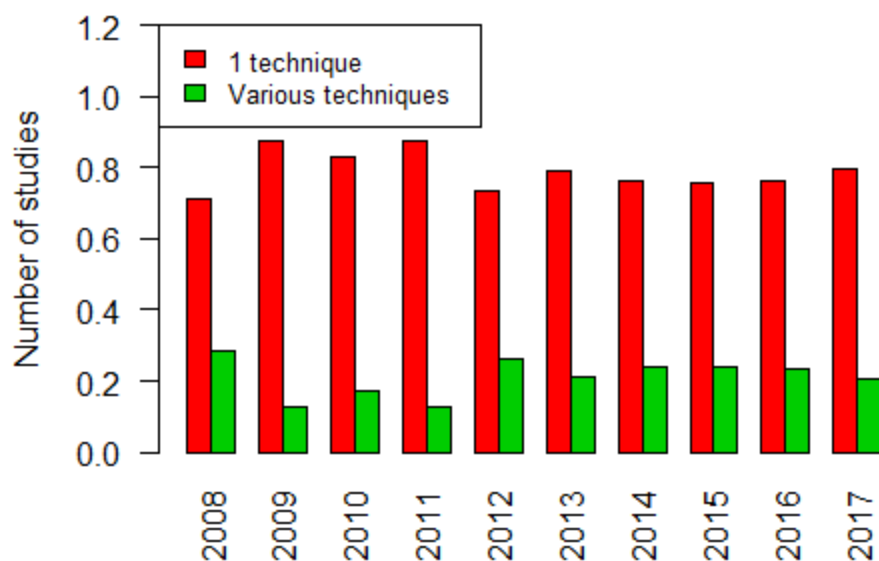
Evolution of the percentage of studies with n techniques



Evolution of the number of studies with n techniques



**Evolution of the percentage of studies with n techniques**



```
> mean(tbl2.2008_2012[1,]); mean(tbl2.2013_2017[1,])
[1] 37.4
[1] 61.4
> t.test(tbl2.2008_2012[1,], tbl2.2013_2017[1,])
```

Welch Two Sample t-test

```
data: tbl2.2008_2012[1, ] and tbl2.2013_2017[1, ]
t = -4.1269, df = 7.895, p-value = 0.003408
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
-37.44166 -10.55834
sample estimates:
mean of x mean of y
 37.4    61.4
```

```
> mean(tbl2.2008_2012[2,]); mean(tbl2.2013_2017[2,])
[1] 8.8
[1] 18
> t.test(tbl2.2008_2012[2,], tbl2.2013_2017[2,])
```

Welch Two Sample t-test

```
data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]
t = -4.271, df = 7.9994, p-value = 0.002721
```

alternative hypothesis: true difference in means is not equal to 0

95 percent confidence interval:

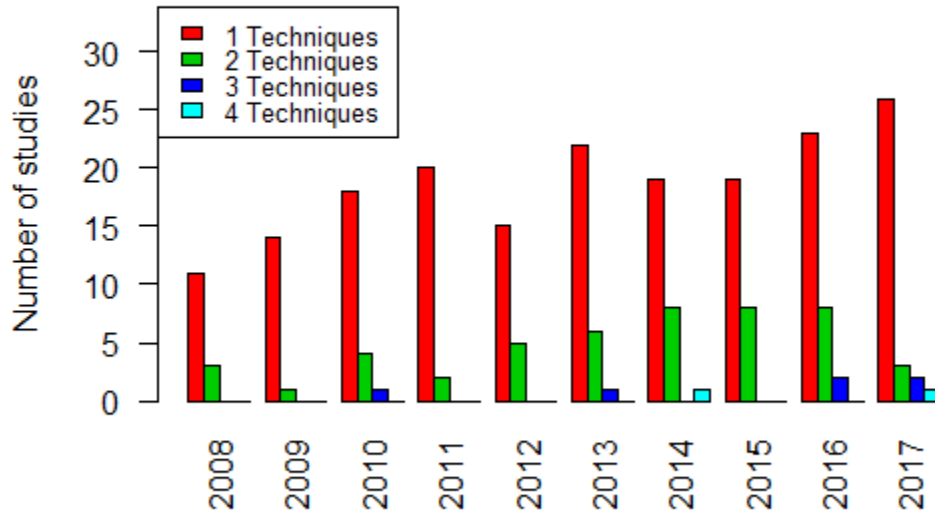
-14.167349 -4.232651

sample estimates:

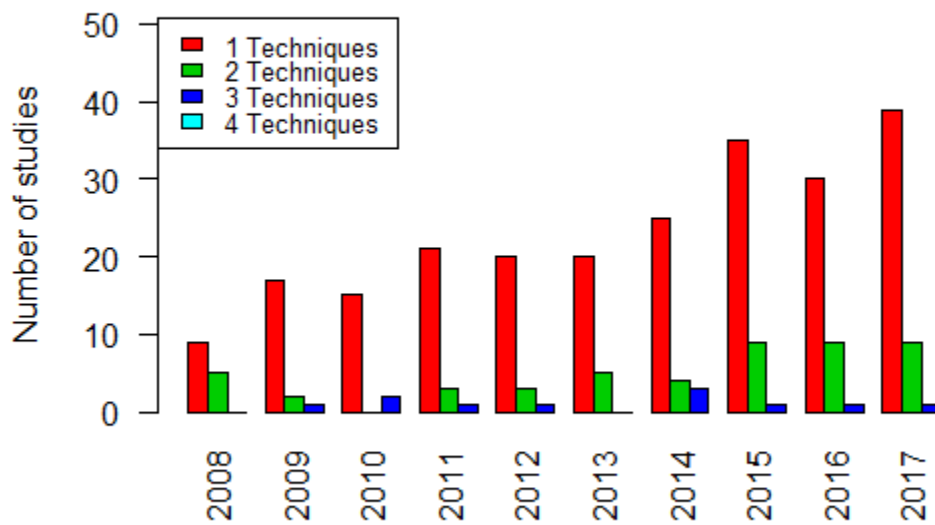
mean of x mean of y

8.8 18.0

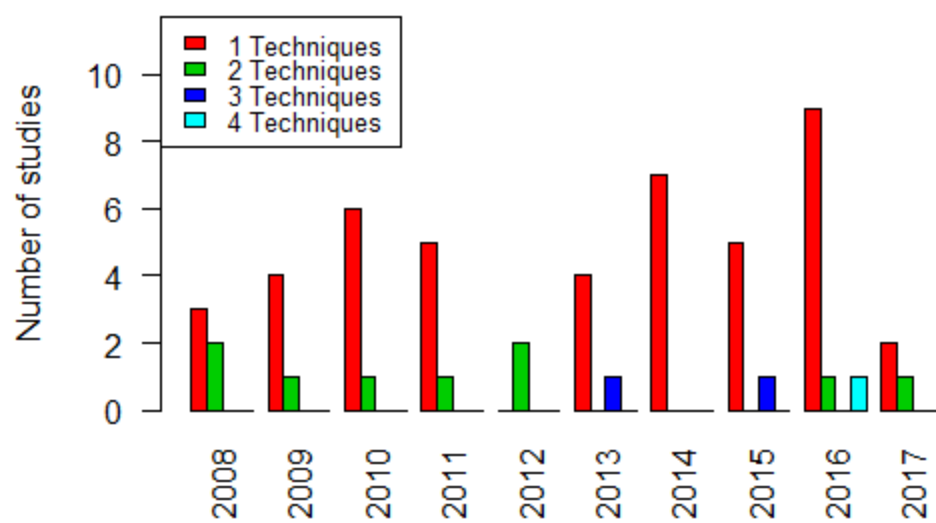
**Evolution of the number of studies (FOR) with n techniques**



**Evolution of the number of studies (AGR) with n techniques**



Evolution of the number of studies (FISH) with n techniques



## Hypothesis 7

7. A) The works are usually normalized, independently of the field to be used.
7. B) The works are usually normalized, independently of the multicriteria technique to be used.

==> We accept the Hypothesis in regards to the field (18 A). In regards to the techniques (18 B), we reject the hypothesis.

==> Regarding the field, the normalization does not vary significantly. Regarding the techniques, the normalization varies significantly. This may be due to the high number of techniques or group of techniques.

```
table(db$normalización, db$Campo)

##
##  AGR FISH FOR
##  0 156  32 112
##  1 135  25 131

chisq.test(table(db$normalización, db$Campo))

##
## Pearson's Chi-squared test
##
## data: table(db$normalización, db$Campo)
## X-squared = 3.7245, df = 2, p-value = 0.1553

table(db$normalización, db$Tecnicas)

##
##  AHP ANP CP DEA ELECTRE FUZZY.MCDM GP MAUT MOP OTRO.MCDM PROMETHEE
##  0 41  3  4 126  3  3 19 14 9  16  4
##  1 83 10  8 19  3  3 22 12 6  10  5
##
##  SMAA TOPSIS VIKOR WEIGHTED
##  0 1  1  1  25
##  1 1  4  0  36

fisher.test(table(db$normalización, db$Tecnicas),simulate.p.value=TRUE)

##
## Fisher's Exact Test for Count Data with simulated p-value (based
## on 2000 replicates)
##
## data: table(db$normalización, db$Tecnicas)
## p-value = 0.0004998
## alternative hypothesis: two.sided

table(db$normalización, db$Tecnicas_Grupos)

##
## Distance.functions Hierarchical Non.deterministic Optimizing.averages
```

```

## 0      34      44      4      39
## 1      40      93      4      48
##
## Otros Outranking Ranking.and.classification.methods
## 0  16      7      126
## 1  10      8      19

fisher.test(table(db$normalización, db$Tecnicas_Grupos),simulate.p.value=TRUE)

##
## Fisher's Exact Test for Count Data with simulated p-value (based
## on 2000 replicates)
##
## data: table(db$normalización, db$Tecnicas_Grupos)
## p-value = 0.0004998
## alternative hypothesis: two.sided

```



## Hypothesis 8

8. A) The justification of why the method is chosen is not usually given, whatever the method used.
8. B) The justification of why the method is chosen is not usually given, whatever the area where it is applied.

==> We reject the Hypothesis for the method used (significant differences) and we accept it for the country (there are significant differences).

==> 7 percent of studies justify the chosen method. By techniques or groups of techniques there are significant differences. By country there are no significant differences.

```
table(db$X.justificación.técnica.elección.)

##
## 0 1
## 583 45

mydf <- data.frame(Sin_justification = table(db$lugar.caso.de.estudio.1, db$X.justificación.técnica.elección.),[1],
  With_justification = table(db$lugar.caso.de.estudio.1, db$X.justificación.técnica.elección.))[2])
mydf$Prop_With_Just <- as.numeric(as.character(mydf$With_justification)) /
  (as.numeric(as.character(mydf$With_justification)) + as.numeric(as.character(mydf$Sin_justification)))
# Para ver el orden de los countries por Número de Estudios Con Justificación
data.frame(Country= rownames(mydf[order(mydf$With_justification),]), With_justification =mydf[order
(mydf$With_justification),"With_justification"])

##
## Country With_justification
## 1 10 case study regions across Europe 0
## 2 10 countries distintos 0
## 3 12 casos de estudio en countries Europeos 0
## 4 14 countries de Europa 0
## 5 15 EU Countries 0
## 6 180 countries 0
## 7 19 EU Member States 0
## 8 23 OECD Countries and Turkey 0
## 9 25 European Union (EU) countries 0
## 10 25 countries de la UE 0
## 11 27 countries of the European Union 0
## 12 31 countries de Europa 0
## 13 33 countries of the world 0
## 14 39 SubSaharian countries 0
## 15 4 countries de Europa 0
## 16 5 countries de Centroamérica 0
## 17 5 countries de la UE yUSA 0
## 18 5 countries UE 0
## 19 8 countries de Europa 0
## 20 88 countries 0
## 21 Algeria 0
## 22 Austria 0
```

## 23	Belgium	0	
## 24	Bulgaria	0	
## 25	Chile	0	
## 26	Colombia	0	
## 27	Costa Rica	0	
## 28	Croatia	0	
## 29	Cuba	0	
## 30	Czech Republic	0	
## 31	Denmark	0	
## 32	Ecuador	0	
## 33	Egypt	0	
## 34	Ethiopia	0	
## 35	Finland	0	
## 36	Ghana	0	
## 37	Indonesia	0	
## 38	Ireland	0	
## 39	Israel	0	
## 40	Jamaica	0	
## 41	Japan	0	
## 42	Kenya	0	
## 43	Kyrgyzstan	0	
## 44	Laos	0	
## 45	Madagascar	0	
## 46	Malaysia	0	
## 47	Mexico	0	
## 48	Morocco	0	
## 49	Nepal	0	
## 50	New Zealand	0	
## 51	Nigeria	0	
## 52	Norway	0	
## 53	Oman	0	
## 54	Papua Nueva Guinea y muchas islas de la Polinesia	0	
## 55	Portugal	0	
## 56	Reunion Island	0	
## 57	Romania	0	
## 58	Russia	0	
## 59	Rwanda	0	
## 60	Saudi Arabia	0	
## 61	Serbia	0	
## 62	Sierra Leone	0	
## 63	six European case study locations	0	
## 64	Slovenia	0	
## 65	South Africa	0	
## 66	South Korea	0	
## 67	Sri Lanka	0	
## 68	Sweden	0	
## 69	Tanzania	0	
## 70	Thailand	0	

```

## 71          todo el mundo          0
## 72             Tunisia          0
## 73             Turkey          0
## 74             Uruguay          0
## 75             Uzbekistan        0
## 76             Vietnam          0
## 77             Zimbabwe          0
## 78                                1
## 79      27 European Union (EU) countries      1
## 80      33 African countries          1
## 81             Argentina          1
## 82             Australia          1
## 83             Brazil            1
## 84             Chipre            1
## 85             Finlandia         1
## 86             France            1
## 87             Georgia           1
## 88             Germany           1
## 89             Greece            1
## 90             India             1
## 91             Lithuania         1
## 92             Netherlands        1
## 93             Pakistan          1
## 94             Slovakia          1
## 95             Switzerland        1
## 96             Syria             1
## 97             Taiwan            1
## 98             UK                1
## 99             USA               1
## 100            Bangladesh         2
## 101            Canada            2
## 102            Spain             3
## 103            China             4
## 104            Italy              4
## 105            Iran              8

```

*# Para ver el orden de los countries por Proporción de Estudios Con Justificación*

```

data.frame(Country= rownames(mydf[order(mydf$Prop_With_Just),]), Prop_With_Just =round(mydf[order(mydf$Prop_With_Just),"Prop_With_Just"],3))

```

```

##          Country Prop_With_Just
## 1      10 case study regions across Europe      0.000
## 2          10 countries distintos      0.000
## 3      12 casos de estudio en countries Europeos      0.000
## 4          14 countries de Europa      0.000
## 5          15 EU Countries      0.000
## 6          180 countries      0.000
## 7          19 EU Member States      0.000

```

## 8	23 OECD Countries and Turkey	0.000
## 9	25 European Union (EU) countries	0.000
## 10	25 countries de la UE	0.000
## 11	27 countries of the European Union	0.000
## 12	31 countries de Europa	0.000
## 13	33 countries of the world	0.000
## 14	39 SubSaharian countries	0.000
## 15	4 countries de Europa	0.000
## 16	5 countries de Centroamérica	0.000
## 17	5 countries de la UE yUSA	0.000
## 18	5 countries UE	0.000
## 19	8 countries de Europa	0.000
## 20	88 countries	0.000
## 21	Algeria	0.000
## 22	Austria	0.000
## 23	Belgium	0.000
## 24	Bulgaria	0.000
## 25	Chile	0.000
## 26	Colombia	0.000
## 27	Costa Rica	0.000
## 28	Croatia	0.000
## 29	Cuba	0.000
## 30	Czech Republic	0.000
## 31	Denmark	0.000
## 32	Ecuador	0.000
## 33	Egypt	0.000
## 34	Ethiopia	0.000
## 35	Finland	0.000
## 36	Ghana	0.000
## 37	Indonesia	0.000
## 38	Ireland	0.000
## 39	Israel	0.000
## 40	Jamaica	0.000
## 41	Japan	0.000
## 42	Kenya	0.000
## 43	Kyrgyzstan	0.000
## 44	Laos	0.000
## 45	Madagascar	0.000
## 46	Malaysia	0.000
## 47	Mexico	0.000
## 48	Morocco	0.000
## 49	Nepal	0.000
## 50	New Zealand	0.000
## 51	Nigeria	0.000
## 52	Norway	0.000
## 53	Oman	0.000
## 54	Papua Nueva Guinea y muchas islas de la Polinesia	0.000
## 55	Portugal	0.000

## 56	Reunion Island	0.000	
## 57	Romania	0.000	
## 58	Russia	0.000	
## 59	Rwanda	0.000	
## 60	Saudi Arabia	0.000	
## 61	Serbia	0.000	
## 62	Sierra Leone	0.000	
## 63	six European case study locations	0.000	0.000
## 64	Slovenia	0.000	
## 65	South Africa	0.000	
## 66	South Korea	0.000	
## 67	Sri Lanka	0.000	
## 68	Sweden	0.000	
## 69	Tanzania	0.000	
## 70	Thailand	0.000	
## 71	todo el mundo	0.000	
## 72	Tunisia	0.000	
## 73	Turkey	0.000	
## 74	Uruguay	0.000	
## 75	Uzbekistan	0.000	
## 76	Vietnam	0.000	
## 77	Zimbabwe	0.000	
## 78	USA	0.021	
## 79	India	0.042	
## 80	Brazil	0.050	
## 81		0.062	
## 82	Greece	0.062	
## 83	Spain	0.064	
## 84	Australia	0.067	
## 85	France	0.083	
## 86	China	0.087	
## 87	Netherlands	0.091	
## 88	UK	0.100	
## 89	Canada	0.118	
## 90	Iran	0.138	
## 91	Italy	0.154	
## 92	Taiwan	0.167	
## 93	Bangladesh	0.182	
## 94	Argentina	0.200	
## 95	Germany	0.200	
## 96	Finlandia	0.250	
## 97	Lithuania	0.333	
## 98	Pakistan	0.333	
## 99	Switzerland	0.333	
## 100	27 European Union (EU) countries	0.500	0.500
## 101	33 African countries	1.000	
## 102	Chipre	1.000	
## 103	Georgia	1.000	

```

## 104          Slovakia    1.000
## 105          Syria      1.000

fisher.test(table(db$lugar.caso.de.estudio.1, db$X.justificación.técnica.elección.),simulate.p.value=TRUE)

##
## Fisher's Exact Test for Count Data with simulated p-value (based
## on 2000 replicates)
##
## data: table(db$lugar.caso.de.estudio.1, db$X.justificación.técnica.elección.)
## p-value = 0.4168
## alternative hypothesis: two.sided

mydf <- data.frame(Sin_justification = table(db$Tecnicas, db$X.justificación.técnica.elección.)[,1],
                  With_justification = table(db$Tecnicas, db$X.justificación.técnica.elección.)[,2])
mydf$Prop_With_Just <- as.numeric(as.character(mydf$With_justification)) /
  (as.numeric(as.character(mydf$With_justification)) + as.numeric(as.character(mydf$Sin_justification)))
data.frame(Country= rownames(mydf[order(mydf$With_justification),]), With_justification =mydf[order
(mydf$With_justification), "With_justification"])

##   Country With_justification
## 1   ANP          0
## 2   CP           0
## 3 FUZZY.MCDM      0
## 4   MOP          0
## 5   SMAA         0
## 6   VIKOR        0
## 7 WEIGHTED        0
## 8 ELECTRE         1
## 9   MAUT         1
## 10 TOPSIS         1
## 11  GP           3
## 12 OTRO.MCDM      3
## 13 PROMETHEE      3
## 14  AHP           8
## 15  DEA          13

data.frame(Country= rownames(mydf[order(mydf$Prop_With_Just),]), Prop_With_Just =round(mydf[or
der(mydf$Prop_With_Just), "Prop_With_Just"],3))

##   Country Prop_With_Just
## 1   ANP      0.000
## 2   CP       0.000
## 3 FUZZY.MCDM  0.000
## 4   MOP      0.000
## 5   SMAA     0.000
## 6   VIKOR    0.000
## 7 WEIGHTED   0.000

```

```
## 8   MAUT    0.038
## 9   AHP     0.065
## 10  GP      0.073
## 11  DEA     0.090
## 12  OTRO.MCDM  0.115
## 13  ELECTRE  0.167
## 14  TOPSIS  0.200
## 15  PROMETHEE  0.333
```

```
table(db$Tecnicas, db$X.justificación.técnica.elección.)
```

```
##
##      0 1
## AHP   116 8
## ANP    13 0
## CP     12 0
## DEA   132 13
## ELECTRE 5 1
## FUZZY.MCDM 6 0
## GP     38 3
## MAUT   25 1
## MOP    15 0
## OTRO.MCDM 23 3
## PROMETHEE 6 3
## SMAA    2 0
## TOPSIS  4 1
## VIKOR   1 0
## WEIGHTED 61 0
```

```
fisher.test(table(db$Tecnicas, db$X.justificación.técnica.elección.),simulate.p.value=TRUE)
```

```
##
## Fisher's Exact Test for Count Data with simulated p-value (based
## on 2000 replicates)
##
## data: table(db$Tecnicas, db$X.justificación.técnica.elección.)
## p-value = 0.04898
## alternative hypothesis: two.sided
```

```
mydf <- data.frame(Sin_justification = table(db$Tecnicas_Grupos, db$X.justificación.técnica.elección.)[,1],
                  With_justification = table(db$Tecnicas_Grupos, db$X.justificación.técnica.elección.)[,2])
mydf$Prop_With_Just <- as.numeric(as.character(mydf$With_justification)) /
  (as.numeric(as.character(mydf$With_justification)) + as.numeric(as.character(mydf$Sin_justification)))
data.frame(Country= rownames(mydf[order(mydf$With_justification),]), With_justification =mydf[order
(mydf$With_justification),"With_justification"])
```

```
##           Country With_justification
## 1   Non.deterministic           0
```

```
## 2      Optimizing.averages      1
## 3          Otros      3
## 4      Distance.functions      4
## 5          Outranking      4
## 6          Hierarchical      8
## 7 Ranking.and.classification.methods      13
```

```
data.frame(Country= rownames(mydf[order(mydf$Prop_With_Just),]), Prop_With_Just =round(mydf[order(mydf$Prop_With_Just),"Prop_With_Just"],3))
```

```
##          Country Prop_With_Just
## 1      Non.deterministic      0.000
## 2      Optimizing.averages      0.011
## 3      Distance.functions      0.054
## 4          Hierarchical      0.058
## 5 Ranking.and.classification.methods      0.090
## 6          Otros      0.115
## 7          Outranking      0.267
```

```
table(db$Tecnicas_Grupos, db$X.justificación.técnica.elección.)
```

```
##
##          0  1
## Distance.functions      70  4
## Hierarchical      129  8
## Non.deterministic      8  0
## Optimizing.averages      86  1
## Otros      23  3
## Outranking      11  4
## Ranking.and.classification.methods 132 13
```

```
fisher.test(table(db$Tecnicas_Grupos, db$X.justificación.técnica.elección.),simulate.p.value=TRUE)
```

```
##
## Fisher's Exact Test for Count Data with simulated p-value (based
## on 2000 replicates)
##
## data: table(db$Tecnicas_Grupos, db$X.justificación.técnica.elección.)
## p-value = 0.01599
## alternative hypothesis: two.sided
```



## Hypothesis 9

9. AHP and weighted MCDM always go together.

==> We reject the Hypothesis.

==> There is no significant relationship between the use of AHP and WEIGHTED

```
summary(glm(db$WEIGHTED ~ db$AHP, family="binomial"))

##
## Call:
## glm(formula = db$WEIGHTED ~ db$AHP, family = "binomial")
##
## Deviance Residuals:
##   Min     1Q   Median     3Q      Max
## -0.7053 -0.7053 -0.6450 -0.6450  1.8289
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.4643    0.1265 -11.580 <2e-16 ***
## db$AHP       0.1997    0.2067  0.967  0.334
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##   Null deviance: 626.84  on 627  degrees of freedom
## Residual deviance: 625.91  on 626  degrees of freedom
## AIC: 629.91
##
## Number of Fisher Scoring iterations: 4

summary(glm(db$AHP ~ db$WEIGHTED, family="binomial"))

##
## Call:
## glm(formula = db$AHP ~ db$WEIGHTED, family = "binomial")
##
## Deviance Residuals:
##   Min     1Q   Median     3Q      Max
## -0.9844 -0.9082 -0.9082  1.4729  1.4729
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.67234    0.09426  -7.133 9.84e-13 ***
## db$WEIGHTED  0.19974    0.20665  0.967  0.334
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

```
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 810.94 on 627 degrees of freedom
## Residual deviance: 810.02 on 626 degrees of freedom
## AIC: 814.02
##
## Number of Fisher Scoring iterations: 4

table(db$WEIGHTED,db$AHP); chisq.test(table(db$WEIGHTED,db$AHP))

##
##      0 1
## 0 333 170
## 1  77  48

##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: table(db$WEIGHTED, db$AHP)
## X-squared = 0.74384, df = 1, p-value = 0.3884
```

## Hypothesis 10

10. Given the inclusion of methods such as AHP and weighted MCDM in different GIS packages, it seems sensible to point out that there is a positive relationship between the use of AHP and its application to spatial problems.

==> We accept the Hypothesis.

==> Those studies that use AHP or WEIGHTED have a greater chance of using GIS. We reject the null hypothesis that they are equal.

```
db$AHP_WEIGHTED <- db$AHP; db$AHP_WEIGHTED[db$WEIGHTED > 0] <- 1
summary(glm(db$GIS ~ db$AHP, family="binomial"))
```

```
##
## Call:
## glm(formula = db$GIS ~ db$AHP, family = "binomial")
##
## Deviance Residuals:
##   Min     1Q   Median     3Q      Max
## -1.1696 -0.6862 -0.6862  1.1852  1.7674
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.3264    0.1213 -10.935 < 2e-16 ***
## db$AHP       1.3080    0.1818  7.194 6.31e-13 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##   Null deviance: 776.50  on 627  degrees of freedom
## Residual deviance: 723.37  on 626  degrees of freedom
## AIC: 727.37
##
## Number of Fisher Scoring iterations: 4
```

```
summary(glm(db$GIS ~ db$WEIGHTED, family="binomial"))
```

```
##
## Call:
## glm(formula = db$GIS ~ db$WEIGHTED, family = "binomial")
##
## Deviance Residuals:
##   Min     1Q   Median     3Q      Max
## -1.3987 -0.7241 -0.7241  0.9712  1.7129
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.2048    0.1059 -11.382 < 2e-16 ***
```

```

## db$WEIGHTED 1.7114 0.2128 8.041 8.93e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 776.50 on 627 degrees of freedom
## Residual deviance: 708.78 on 626 degrees of freedom
## AIC: 712.78
##
## Number of Fisher Scoring iterations: 4

summary(glm(db$GIS ~ db$AHP_WEIGHTED, family="binomial"))

##
## Call:
## glm(formula = db$GIS ~ db$AHP_WEIGHTED, family = "binomial")
##
## Deviance Residuals:
## Min 1Q Median 3Q Max
## -1.1745 -0.5516 -0.5516 1.1803 1.9789
##
## Coefficients:
## Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.8058 0.1574 -11.473 <2e-16 ***
## db$AHP_WEIGHTED 1.7991 0.1958 9.189 <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 776.50 on 627 degrees of freedom
## Residual deviance: 680.04 on 626 degrees of freedom
## AIC: 684.04
##
## Number of Fisher Scoring iterations: 4

table(db$GIS,db$AHP); chisq.test(table(db$GIS,db$AHP))

##
## 0 1
## 0 324 110
## 1 86 108
##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: table(db$GIS, db$AHP)
## X-squared = 53.07, df = 1, p-value = 3.219e-13

```

```
table(db$GIS,db$WEIGHTED); chisq.test(table(db$GIS,db$WEIGHTED))

##
##    0 1
## 0 387 47
## 1 116 78

##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: table(db$GIS, db$WEIGHTED)
## X-squared = 70.743, df = 1, p-value < 2.2e-16

table(db$GIS,db$AHP_WEIGHTED); chisq.test(table(db$GIS,db$AHP_WEIGHTED))

##
##    0 1
## 0 286 148
## 1 47 147

##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: table(db$GIS, db$AHP_WEIGHTED)
## X-squared = 91.804, df = 1, p-value < 2.2e-16
```

## Hypothesis 11

11. AHP is used mostly to obtain weights from a set of stakeholders and / or experts and apply these weights to solve the problem in question.

==> We accept the Hypothesis.

==> Most times that AHP is used it is used to obtain weights

```
table(db$AHP); table(db$pesos.AHP.ANP); table(db$AHP, db$pesos.AHP.ANP)
```

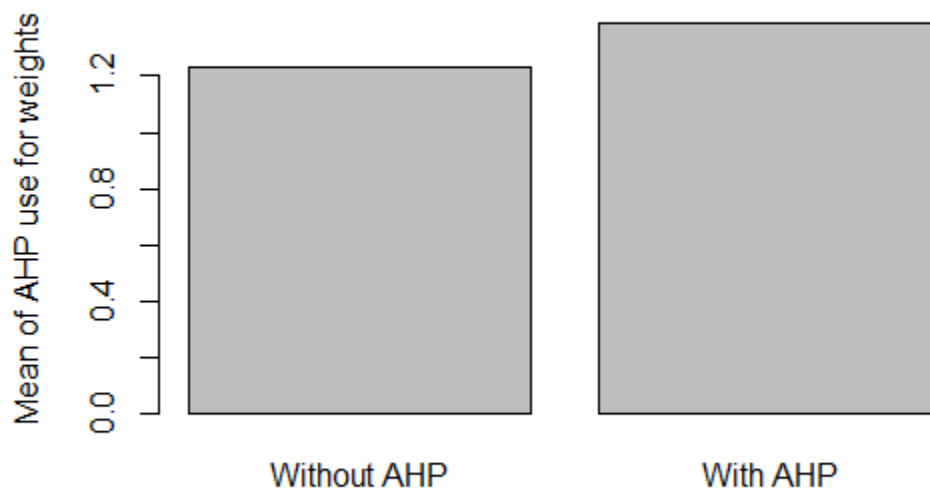
```
##  
## 0 1  
## 410 218
```

```
##  
## 0 1  
## 415 213
```

```
##  
## 0 1  
## 0 396 14  
## 1 19 199
```

```
tbl_mean <- aggregate(db[, c("AHP")], list(db$pesos.AHP.ANP), mean, na.rm=TRUE)
```

```
df.bar <- barplot(tbl_mean$x, names.arg = c("Sin AHP", "Con AHP"), ylab="Media de uso de AHP para pesos")
```



```
chisq.test(table(db$pesos.AHP.ANP, db$AHP))
```

```

##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: table(db$pesos.AHP.ANP, db$AHP)
## X-squared = 486.38, df = 1, p-value < 2.2e-16

summary(glm(db$AHP ~ db$pesos.AHP.ANP, family="binomial"))

##
## Call:
## glm(formula = db$AHP ~ db$pesos.AHP.ANP, family = "binomial")
##
## Deviance Residuals:
##   Min     1Q   Median     3Q      Max
## -2.3333 -0.3061 -0.3061  0.3688  2.4835
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept)   -3.0370    0.2348  -12.93 <2e-16 ***
## db$pesos.AHP.ANP  5.6912    0.3628  15.69 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##   Null deviance: 810.94  on 627  degrees of freedom
## Residual deviance: 257.58  on 626  degrees of freedom
## AIC: 261.58
##
## Number of Fisher Scoring iterations: 5

```

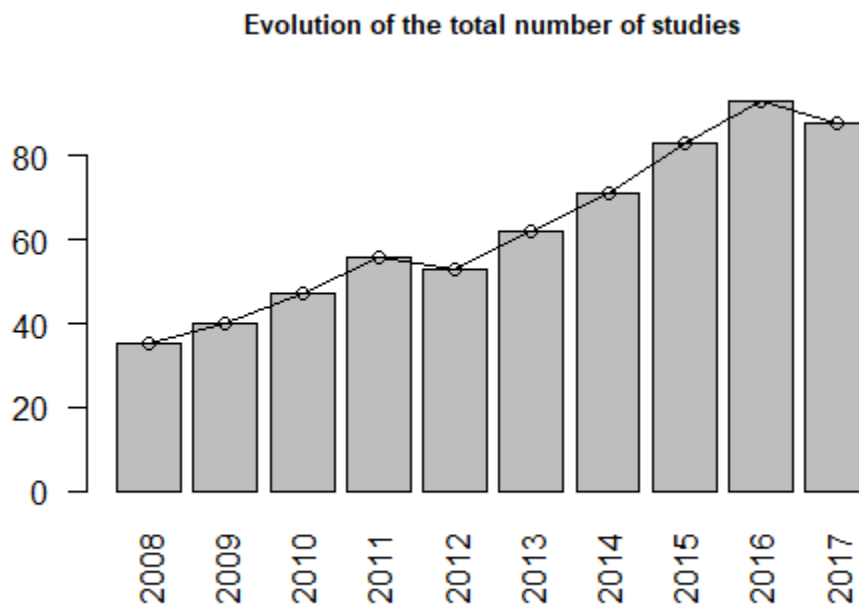
## Hypothesis 12

12. Hybridization of MCDM and GDM is increasing over time.

==> We accept the Hypothesis with some nuances.

==> Yes that is growing but with some nuances. The number of total studies has increased. The number of studies with GDM and MCDM has also (significantly) but the proportion of studies with GDM and MCDM has decreased (it is not significant)

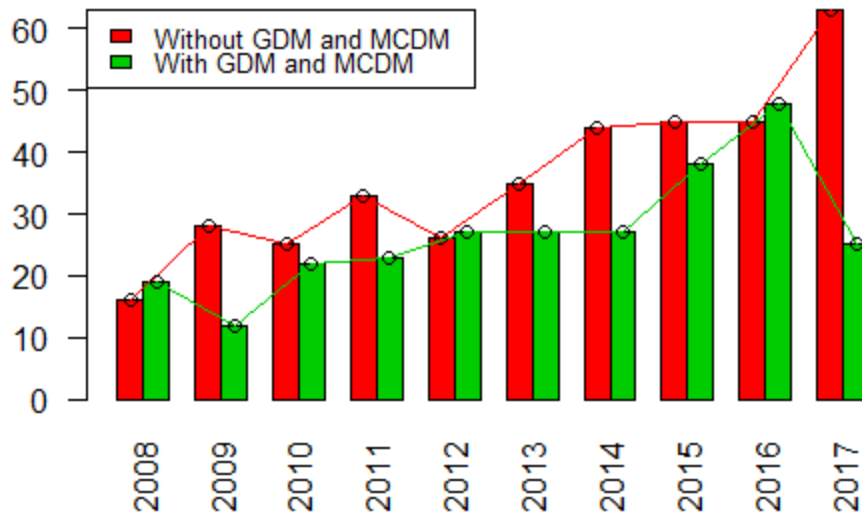
```
df.bar <- barplot(table(db$year), las=2, main="Evolución del número total de trabajos", cex.main=0.8)
lines(x = df.bar, y = table(db$year)); points(x = df.bar, y = table(db$year))
```



```
tbl1 <- as.matrix(table(db$GDM_MCDM, db$year))
df.bar <- barplot(tbl1, beside=T, col=c(2:3), las=2, main="Evolución del número de trabajos con GDM y MCDM", cex.main=0.8)
lines(x = df.bar[1,], y = tbl1[1,], col=2); points(x = df.bar[1,], y = tbl1[1,])
lines(x = df.bar[2,], y = tbl1[2,], col=3); points(x = df.bar[2,], y = tbl1[2,])
legend("topleft", c("Sin GDM y MCDM", "Con GDM y MCDM"), fill=c(2:3), cex=0.8)
```

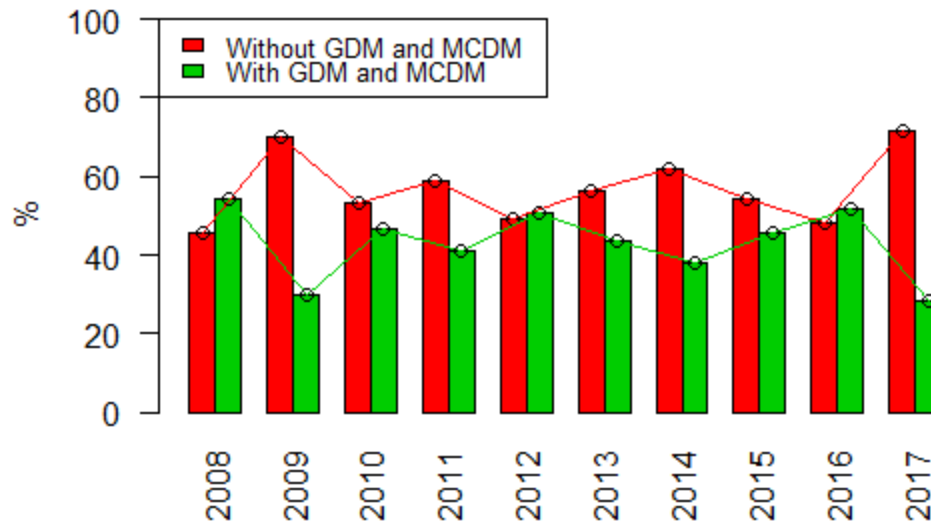


Evolution of the number of studies with GDM and MCDM



```
tbl2 <- rbind(table(db$GDM_MCDM, db$year)[1,] / (colSums(table(db$GDM_MCDM, db$year))),
             table(db$GDM_MCDM, db$year)[2,] / (colSums(table(db$GDM_MCDM, db$year))))
df.bar <- barplot(tbl2*100, beside=T, col=c(2:3), las=2, main="Evolución de la proporción de trabajos con GDM y MCDM",
                 ylim=c(0,100), ylab="%", cex.main=0.8)
lines(x = df.bar[1,], y = tbl2[1,]*100, col=2);points(x = df.bar[1,], y = tbl2[1,]*100)
lines(x = df.bar[2,], y = tbl2[2,]*100, col=3);points(x = df.bar[2,], y = tbl2[2,]*100)
legend("topleft",c("Sin GDM y MCDM", "Con GDM y MCDM"), fill=c(2:3), cex=0.8)
```

Evolution of the proportion of studies with GDM and MCDM



```
tbl1.2008_2012 <- tbl1 [,1:5];tbl1.2013_2017 <- tbl1 [,6:10]
tbl2.2008_2012 <- tbl2 [,1:5];tbl2.2013_2017 <- tbl2 [,6:10]
mean(tbl1.2008_2012[2,]); mean(tbl1.2013_2017[2,])

## [1] 20.6
## [1] 33

t.test(tbl1.2008_2012[2,], tbl1.2013_2017[2,])

##
## Welch Two Sample t-test
##
## data: tbl1.2008_2012[2, ] and tbl1.2013_2017[2, ]
## t = -2.4527, df = 6.3478, p-value = 0.04742
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -24.6083607 -0.1916393
## sample estimates:
## mean of x mean of y
## 20.6 33.0

mean(tbl2.2008_2012[2,]); mean(tbl2.2013_2017[2,])

## [1] 0.4462181
## [1] 0.4147634
```

```
t.test(tbl2.2008_2012[2,], tbl2.2013_2017[2,])  
  
##  
## Welch Two Sample t-test  
##  
## data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]  
## t = 0.54233, df = 7.9444, p-value = 0.6025  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## -0.1024540 0.1653634  
## sample estimates:  
## mean of x mean of y  
## 0.4462181 0.4147634
```

### Hypothesis 13

13. Given the nature of the forestry problems, it seems that the hybridization of GDM + MCDM techniques could be more frequent in forestry.

==> We accept the Hypothesis

==> We reject the null hypothesis that they are equal.

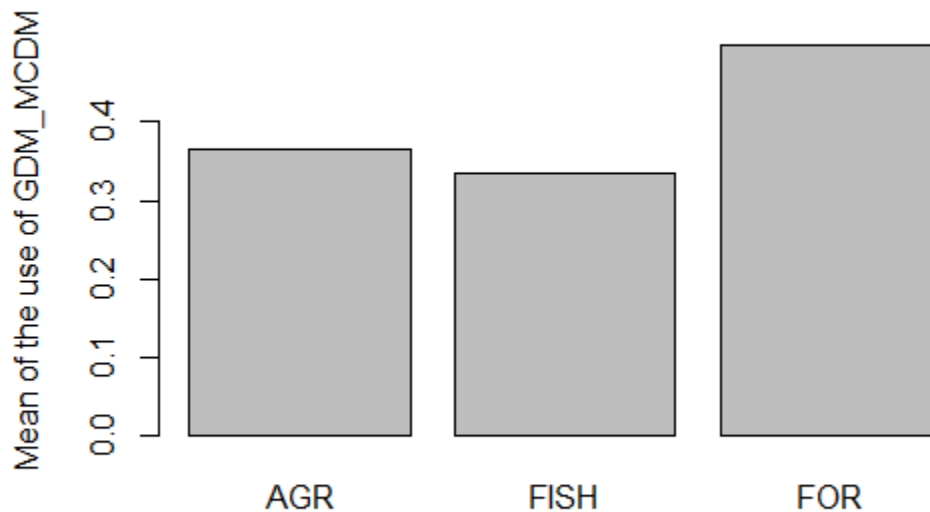
```
db$MCDM_n <- rowSums(db[,Tecnicas_name])
db$GDM_MCDM <- 0; db$GDM_MCDM[db$GDM > 0 & db$MCDM_n > 0] <- 1
table(db$GDM); table(db$GDM_MCDM); table(db$Campo,db$GDM_MCDM)

##
## 0 1
## 360 268

##
## 0 1
## 360 268

##
##    0 1
## AGR 185 106
## FISH 38 19
## FOR 122 121

tbl_mean <- aggregate(db[, c("GDM_MCDM")], list(db$Campo), mean, na.rm=TRUE)
df.bar <- barplot(tbl_mean$x, names.arg = tbl_mean$Group.1, ylab="Media de uso de GDM_MCDM")
```



```
chisq.test(table(db$Campo,db$GDM_MCDM))  
##  
## Pearson's Chi-squared test  
##  
## data: table(db$Campo, db$GDM_MCDM)  
## X-squared = 11.524, df = 2, p-value = 0.003145
```

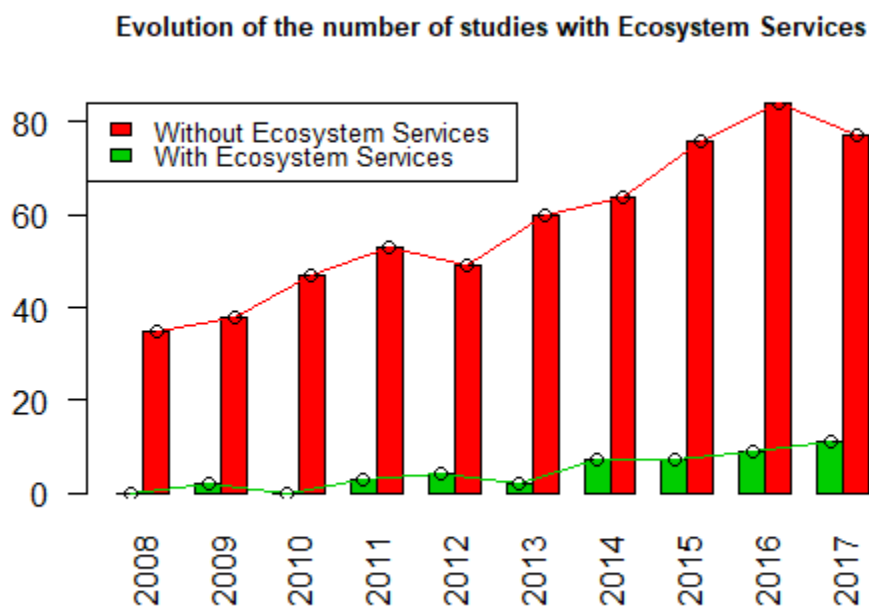
#### Hypothesis 14

14. The concepts of ecosystem services, climate change and sustainability are recent and has only become important in recent years.

==> We accept the Hypothesis for Ecosystem Services and Climate Change. We cannot accept it for Sustainability.

==> We accept the Hypothesis for Ecosystem Services and Climate Change (for both the number and the proportion). For sustainability, despite having increased the total number significantly, the proportion has decreased.

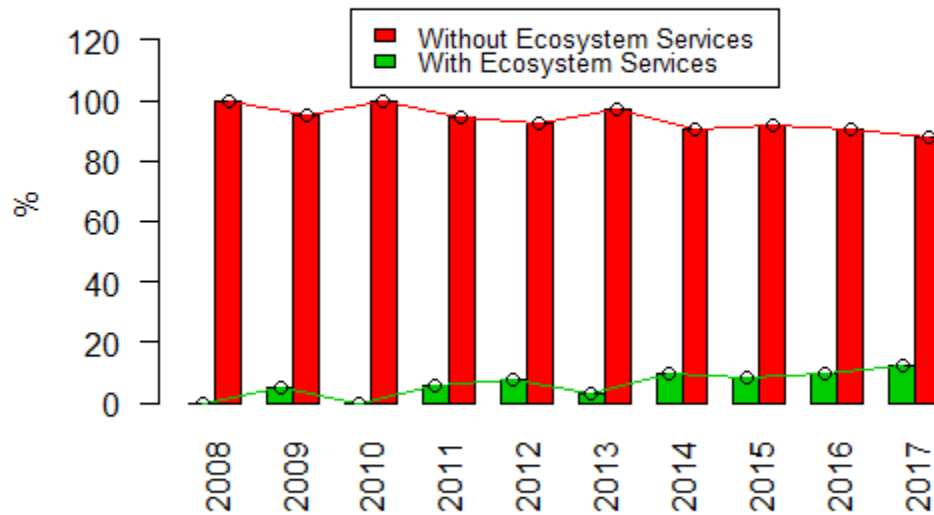
```
db$ecosystem.services <- "Sin Ecosystem Services"; db$ecosystem.services[db$nº.ecosystem.serv. > 0] <- "Con Ecosystem Services"
tbl1 <- as.matrix(table(db$ecosystem.services, db$year))
df.bar <- barplot(tbl1, beside=T, col=c(3,2),las=2, main="Evolución del número de trabajos con Ecosystem Services", cex.main=0.8)
lines(x = df.bar[1,], y = tbl1[1,], col=3);points(x = df.bar[1,], y = tbl1[1,])
lines(x = df.bar[2,], y = tbl1[2,], col=2);points(x = df.bar[2,], y = tbl1[2,])
legend("topleft",c("Sin Ecosystem Services","Con Ecosystem Services"), fill=c(2:3), cex=0.8)
```



```
tbl2 <- rbind(table(db$ecosystem.services, db$year)[1,] / (colSums(table(db$ecosystem.services, db$year))),
             table(db$ecosystem.services, db$year)[2,] / (colSums(table(db$ecosystem.services, db$year))))
df.bar <- barplot(tbl2*100, beside=T, col=c(3,2),las=2, main="Evolución de la proporción de trabajos con Ecosystem Services",
                 ylim=c(0,130), ylab="%", cex.main=0.8)
lines(x = df.bar[1,], y = tbl2[1,]*100, col=3);points(x = df.bar[1,], y = tbl2[1,]*100)
```

```
lines(x = df.bar[2,], y = tbl2[2,]*100, col=2);points(x = df.bar[2,], y = tbl2[2,]*100)
legend("top",c("Sin Ecosystem Services","Con Ecosystem Services"), fill=c(2:3), cex=0.8)
```

Evolution of the proportion of studies with Ecosystem Services



```
tbl1.2008_2012 <- tbl1 [,1:5];tbl1.2013_2017 <- tbl1 [,6:10]
tbl2.2008_2012 <- tbl2 [,1:5];tbl2.2013_2017 <- tbl2 [,6:10]
mean(tbl1.2008_2012[1,]); mean(tbl1.2013_2017[1,])
```

```
## [1] 1.8
```

```
## [1] 7.2
```

```
t.test(tbl1.2008_2012[1,], tbl1.2013_2017[1,])
```

```
##
```

```
## Welch Two Sample t-test
```

```
##
```

```
## data: tbl1.2008_2012[1, ] and tbl1.2013_2017[1, ]
```

```
## t = -3.182, df = 6.1132, p-value = 0.01855
```

```
## alternative hypothesis: true difference in means is not equal to 0
```

```
## 95 percent confidence interval:
```

```
## -9.533978 -1.266022
```

```
## sample estimates:
```

```
## mean of x mean of y
```

```
## 1.8 7.2
```

```
mean(tbl2.2008_2012[1,]); mean(tbl2.2013_2017[1,])
```

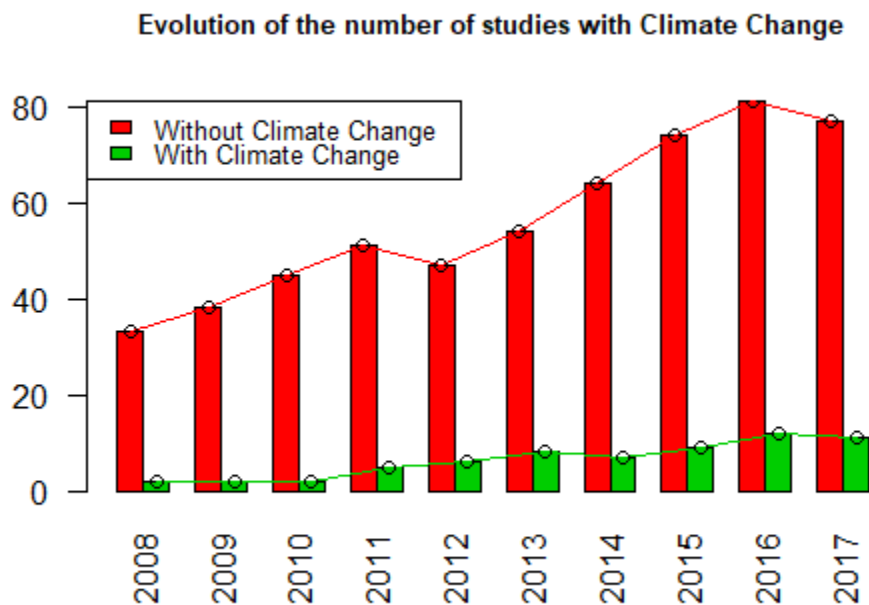
```
## [1] 0.03580863
```

```
## [1] 0.08739223

t.test(tbl2.2008_2012[1,], tbl2.2013_2017[1,])

##
## Welch Two Sample t-test
##
## data: tbl2.2008_2012[1, ] and tbl2.2013_2017[1, ]
## t = -2.3881, df = 8, p-value = 0.04399
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.101394108 -0.001773104
## sample estimates:
## mean of x mean of y
## 0.03580863 0.08739223

tbl1 <- as.matrix(table(db$cambio.climático, db$year))
df.bar <- barplot(tbl1, beside=T, col=c(2:3), las=2, main="Evolución del número de trabajos con Climate Change", cex.main=0.8)
lines(x = df.bar[1,], y = tbl1[1,], col=2);points(x = df.bar[1,], y = tbl1[1,])
lines(x = df.bar[2,], y = tbl1[2,], col=3);points(x = df.bar[2,], y = tbl1[2,])
legend("topleft",c("Sin Climate Change","Con Climate Change"), fill=c(2:3), cex=0.8)
```



```
tbl2 <- rbind(table(db$cambio.climático, db$year)[1,] / (colSums(table(db$cambio.climático, db$year))),
             table(db$cambio.climático, db$year)[2,] / (colSums(table(db$cambio.climático, db$year))))
df.bar <- barplot(tbl2*100, beside=T, col=c(2:3), las=2, main="Evolución de la proporción de trabajos con Climate Change",
```

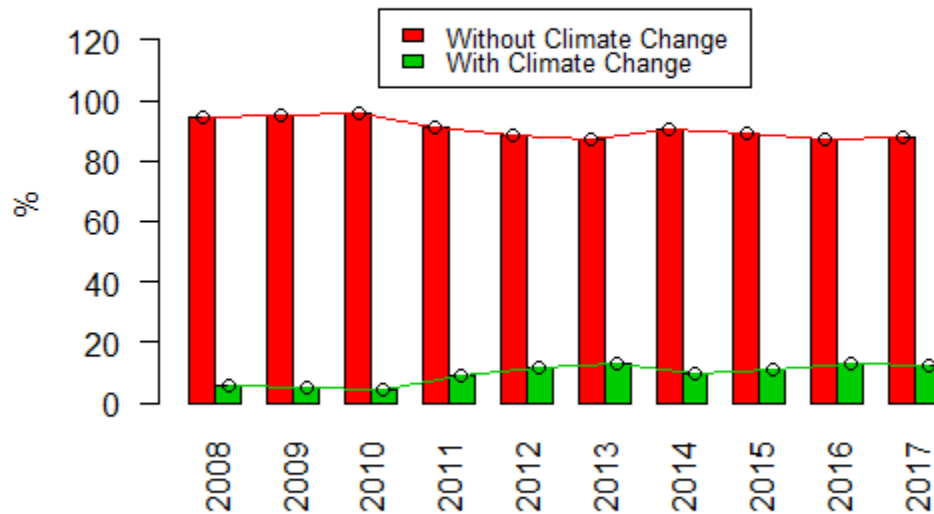


```

ylim=c(0,130), ylab="%", cex.main=0.8)
lines(x = df.bar[1,], y = tbl2[1,]*100, col=2);points(x = df.bar[1,], y = tbl2[1,]*100)
lines(x = df.bar[2,], y = tbl2[2,]*100, col=3);points(x = df.bar[2,], y = tbl2[2,]*100)
legend("top",c("Sin Climate Change","Con Climate Change"), fill=c(2:3), cex=0.8)

```

Evolution of the proportion of studies with Climate Change



```

tbl1.2008_2012 <- tbl1 [,1:5];tbl1.2013_2017 <- tbl1 [,6:10]
tbl2.2008_2012 <- tbl2 [,1:5];tbl2.2013_2017 <- tbl2 [,6:10]
mean(tbl1.2008_2012[2,]); mean(tbl1.2013_2017[2,])

## [1] 3.4
## [1] 9.4

t.test(tbl1.2008_2012[2,], tbl1.2013_2017[2,])

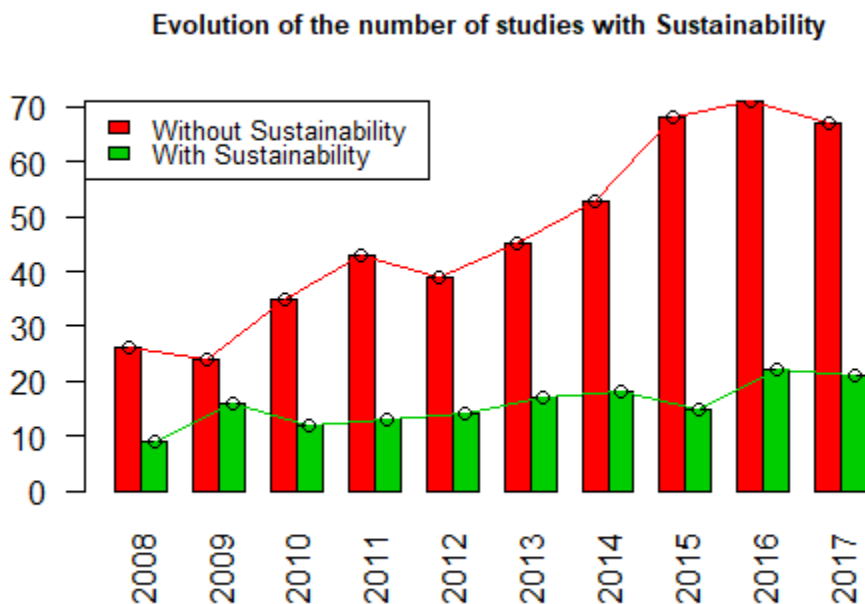
##
## Welch Two Sample t-test
##
## data: tbl1.2008_2012[2, ] and tbl1.2013_2017[2, ]
## t = -4.714, df = 7.9696, p-value = 0.001529
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -8.937012 -3.062988
## sample estimates:
## mean of x mean of y
## 3.4 9.4

mean(tbl2.2008_2012[2,]); mean(tbl2.2013_2017[2,])

```

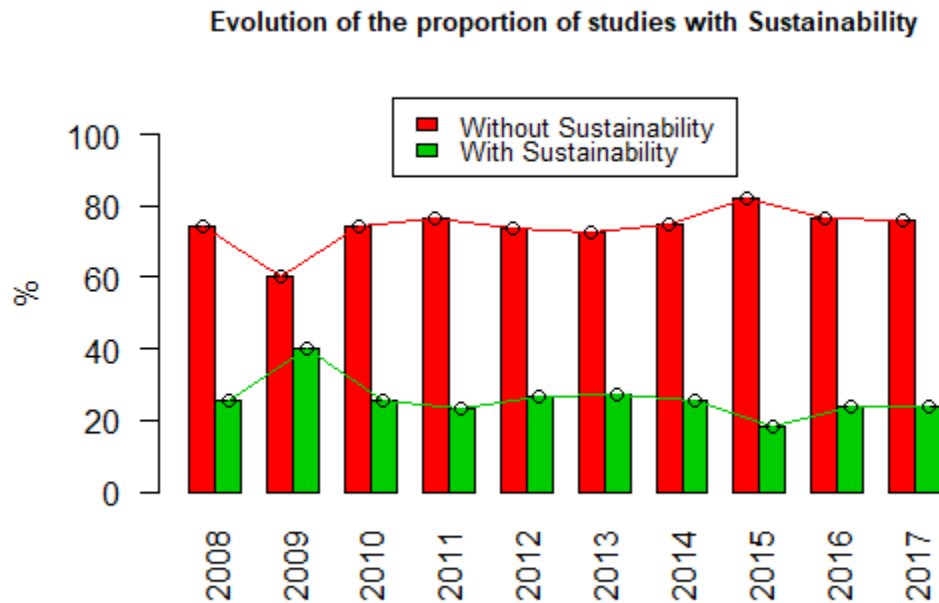
```
## [1] 0.07043786
## [1] 0.118018
t.test(tbl2.2008_2012[2,], tbl2.2013_2017[2,])
##
## Welch Two Sample t-test
##
## data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]
## t = -3.2385, df = 5.6361, p-value = 0.01937
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.08410046 -0.01105973
## sample estimates:
## mean of x mean of y
## 0.07043786 0.11801796

tbl1 <- as.matrix(table(db$sustainability, db$year))
df.bar <- barplot(tbl1, beside=T, col=c(2:3), las=2, main="Evolución del número de trabajos con Sustainability", cex.main=0.8)
lines(x = df.bar[1,], y = tbl1[1,], col=2);points(x = df.bar[1,], y = tbl1[1,])
lines(x = df.bar[2,], y = tbl1[2,], col=3);points(x = df.bar[2,], y = tbl1[2,])
legend("topleft",c("Sin Sustainability", "Con Sustainability"), fill=c(2:3), cex=0.8)
```



```
tbl2 <- rbind(table(db$sustainability, db$year)[1,] / (colSums(table(db$sustainability, db$year))),
table(db$sustainability, db$year)[2,] / (colSums(table(db$sustainability, db$year))))
```

```
df.bar <- barplot(tbl2*100, beside=T, col=c(2:3),las=2, main="Evolución de la proporción de trabajos con Sustainability",
  ylim=c(0,110), ylab="%", cex.main=0.8)
lines(x = df.bar[1,], y = tbl2[1,]*100, col=2);points(x = df.bar[1,], y = tbl2[1,]*100)
lines(x = df.bar[2,], y = tbl2[2,]*100, col=3);points(x = df.bar[2,], y = tbl2[2,]*100)
legend("top",c("Sin Sustainability","Con Sustainability"), fill=c(2:3), cex=0.8)
```



```
tbl1.2008_2012 <- tbl1 [,1:5];tbl1.2013_2017 <- tbl1 [,6:10]
tbl2.2008_2012 <- tbl2 [,1:5];tbl2.2013_2017 <- tbl2 [,6:10]
mean(tbl1.2008_2012[2,]); mean(tbl1.2013_2017[2,])

## [1] 12.8

## [1] 18.6

t.test(tbl1.2008_2012[2,], tbl1.2013_2017[2,])

##
## Welch Two Sample t-test
##
## data: tbl1.2008_2012[2, ] and tbl1.2013_2017[2, ]
## t = -3.3486, df = 7.91, p-value = 0.01027
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -9.80204 -1.79796
## sample estimates:
```

```
## mean of x mean of y
## 12.8 18.6

mean(tbl2.2008_2012[2,]); mean(tbl2.2013_2017[2,])

## [1] 0.2817512
## [1] 0.2367266

t.test(tbl2.2008_2012[2,], tbl2.2013_2017[2,])

##
## Welch Two Sample t-test
##
## data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]
## t = 1.3311, df = 5.9955, p-value = 0.2315
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.03775933 0.12780842
## sample estimates:
## mean of x mean of y
## 0.2817512 0.2367266
```

## Hypothesis 15

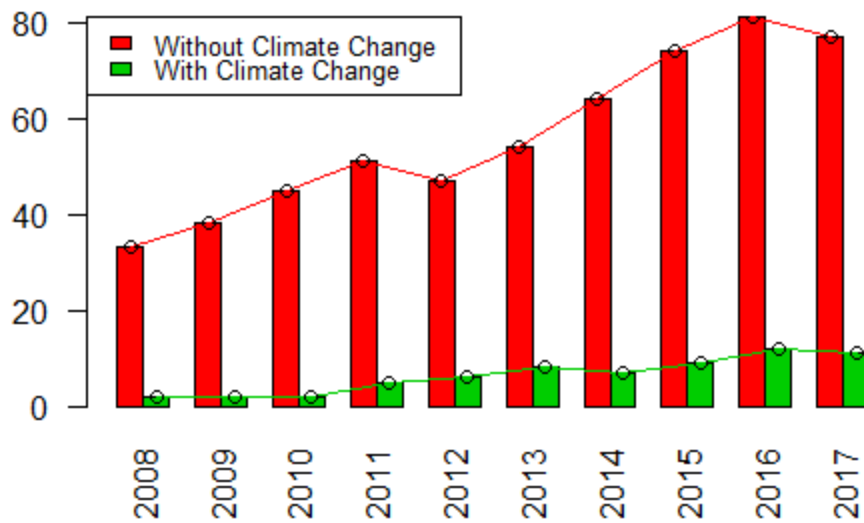
15. The topics that have been included to be relevant, are only relevant in the last years of the period.

==> In general, we reject the Hypothesis.

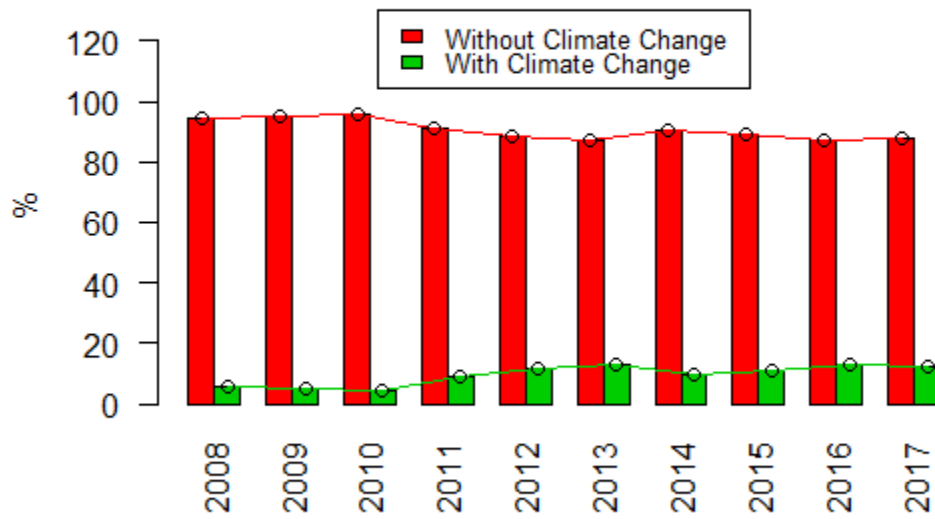
==> In most cases, we reject the Hypothesis. See graphs and test results one by one.

```
topics <- c("cambio.climático","multifunctionality","multiple.use","sustainability","bioeconomy","CSR.respons..Social.corp..",
           "certification","protected.areas","life.cycle")
for (i in 1:length(topics)){
  my_topic <- topics[i]
  tbl1 <- as.matrix(table(db[,my_topic], db$year))
  df.bar <- barplot(tbl1, beside=T, col=c(2:3),las=2, main=paste("Evolución del número de trabajos con", my_topic), cex.main=0.8)
  lines(x = df.bar[1,], y = tbl1[1,], col=2);points(x = df.bar[1,], y = tbl1[1,])
  lines(x = df.bar[2,], y = tbl1[2,], col=3);points(x = df.bar[2,], y = tbl1[2,])
  legend("topleft",legend = c(paste("Sin", my_topic),paste("Con", my_topic)), fill=c(2:3), cex=0.8)
  tbl2 <- rbind(table(db[,my_topic], db$year)[1,] / (colSums(table(db[,my_topic], db$year))),
               table(db[,my_topic], db$year)[2,] / (colSums(table(db[,my_topic], db$year))))
  df.bar <- barplot(tbl2*100, beside=T, col=c(2:3),las=2, main=paste("Evolución de la proporción de trabajos con", my_topic),
                   ylim=c(0,130), ylab="%", cex.main=0.8)
  lines(x = df.bar[1,], y = tbl2[1,]*100, col=2);points(x = df.bar[1,], y = tbl2[1,]*100)
  lines(x = df.bar[2,], y = tbl2[2,]*100, col=3);points(x = df.bar[2,], y = tbl2[2,]*100)
  legend("top",legend = c(paste("Sin", my_topic),paste("Con", my_topic)), fill=c(2:3), cex=0.8)
  tbl1.2008_2012 <- tbl1 [,1:5];tbl1.2013_2017 <- tbl1 [,6:10]
  tbl2.2008_2012 <- tbl2 [,1:5];tbl2.2013_2017 <- tbl2 [,6:10]
  mean(tbl1.2008_2012[2,]); mean(tbl1.2013_2017[2,])
  print(t.test(tbl1.2008_2012[2,], tbl1.2013_2017[2,]))
  mean(tbl2.2008_2012[2,]); mean(tbl2.2013_2017[2,])
  print(t.test(tbl2.2008_2012[2,], tbl2.2013_2017[2,]))
}
```

Evolution of the number of studies with Climate Change



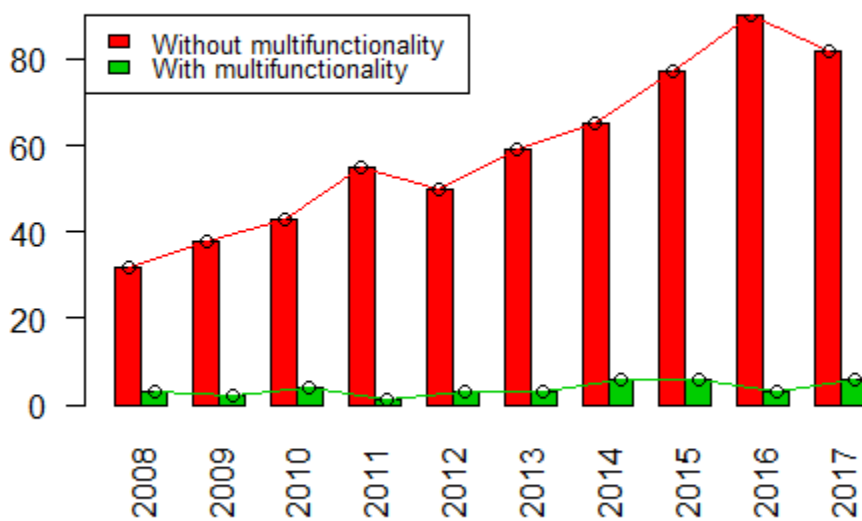
Evolution of the proportion of studies with Climate Change



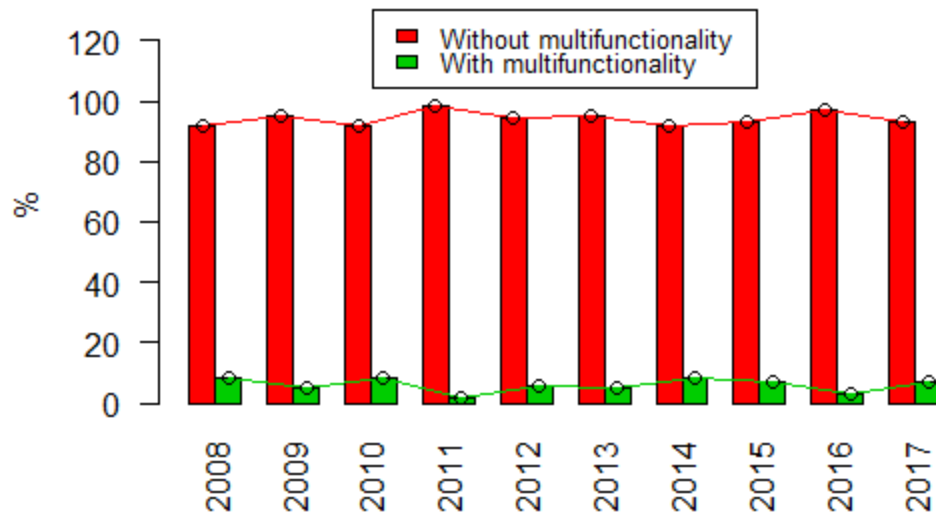
```
##
## Welch Two Sample t-test
##
## data: tbl1.2008_2012[2, ] and tbl1.2013_2017[2, ]
## t = -4.714, df = 7.9696, p-value = 0.001529
```

```
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -8.937012 -3.062988
## sample estimates:
## mean of x mean of y
## 3.4 9.4
##
##
## Welch Two Sample t-test
##
## data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]
## t = -3.2385, df = 5.6361, p-value = 0.01937
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.08410046 -0.01105973
## sample estimates:
## mean of x mean of y
## 0.07043786 0.11801796
```

Evolution of the number of studies with multifunctionality



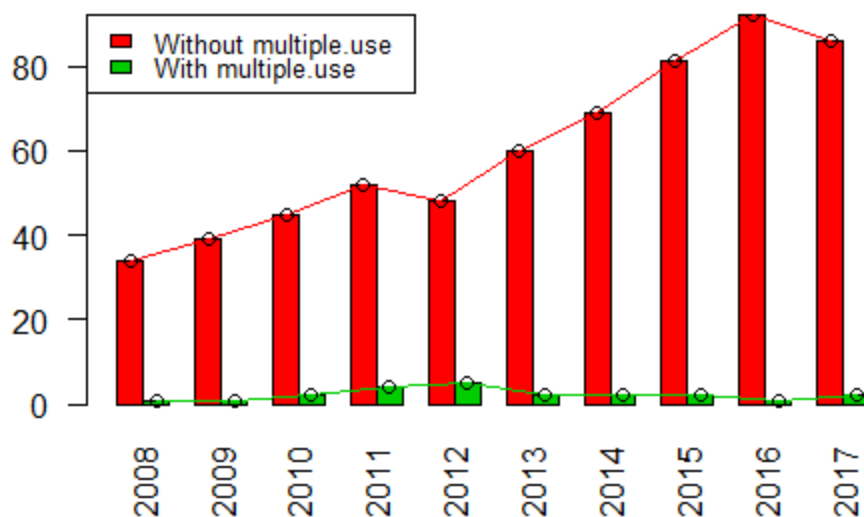
### Evolution of the proportion of studies with multifunctionality



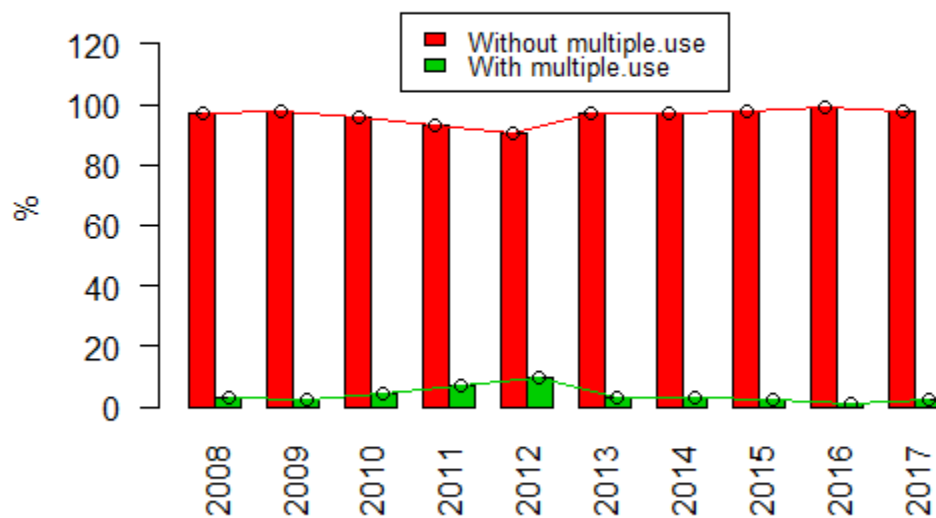
```
##
## Welch Two Sample t-test
##
## data: tbl1.2008_2012[2, ] and tbl1.2013_2017[2, ]
## t = -2.4597, df = 7.1269, p-value = 0.04289
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -4.30737111 -0.09262889
## sample estimates:
## mean of x mean of y
## 2.6 4.8
##
##
## Welch Two Sample t-test
##
## data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]
## t = -0.13226, df = 7.3472, p-value = 0.8983
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.03869656 0.03455992
## sample estimates:
## mean of x mean of y
## 0.05905632 0.06112464
```



Evolution of the number of studies with multiple.use



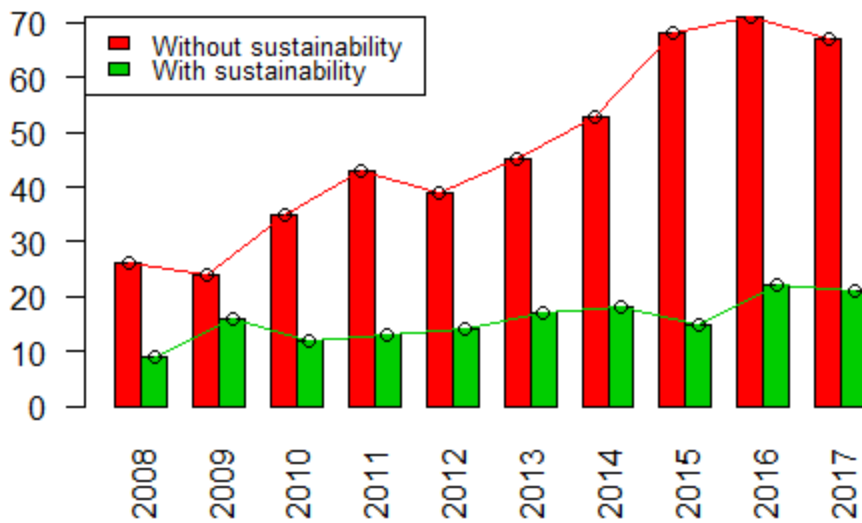
Evolution of the proportion of studies with multiple.use



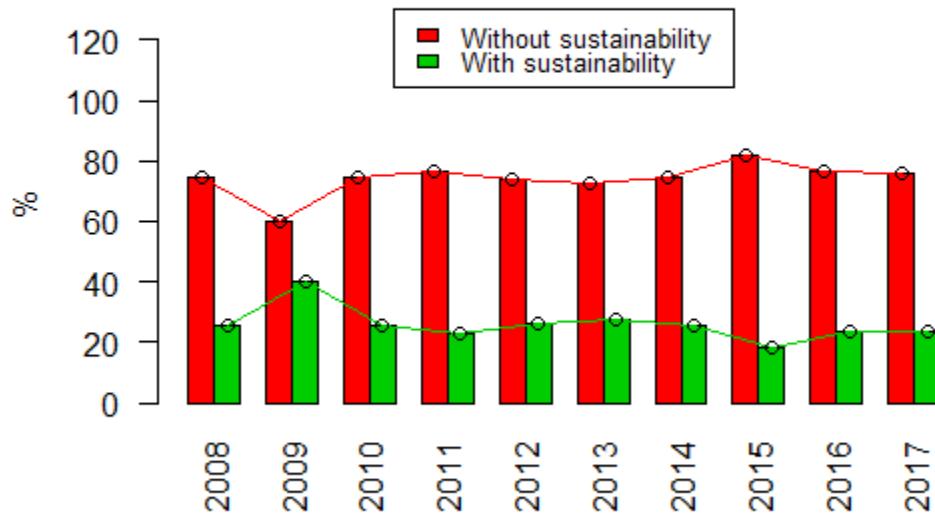
```
##
## Welch Two Sample t-test
##
## data: tb1.2008_2012[2, ] and tb1.2013_2017[2, ]
## t = 0.95618, df = 4.4831, p-value = 0.3877
```

```
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -1.427477 3.027477
## sample estimates:
## mean of x mean of y
## 2.6 1.8
##
##
## Welch Two Sample t-test
##
## data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]
## t = 2.0882, df = 4.5892, p-value = 0.09611
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.00762321 0.06517897
## sample estimates:
## mean of x mean of y
## 0.05237856 0.02360069
```

Evolution of the number of studies with sustainability



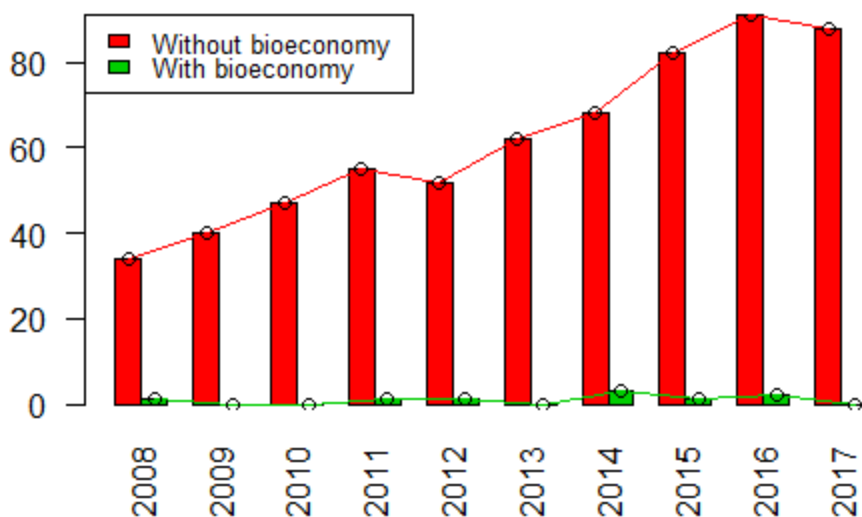
Evolution of the proportion of studies with sustainability



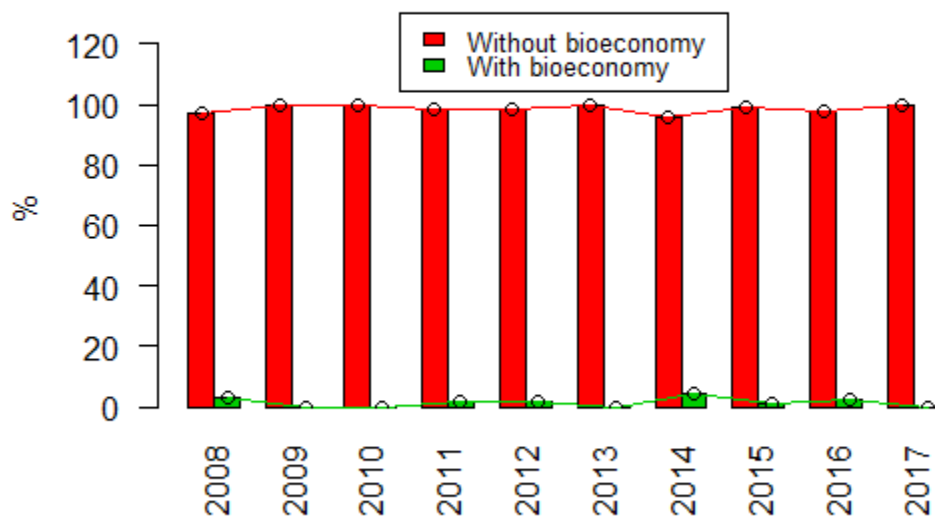
```
##
## Welch Two Sample t-test
##
## data: tb1.2008_2012[2, ] and tb1.2013_2017[2, ]
## t = -3.3486, df = 7.91, p-value = 0.01027
```

```
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -9.80204 -1.79796
## sample estimates:
## mean of x mean of y
## 12.8 18.6
##
##
## Welch Two Sample t-test
##
## data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]
## t = 1.3311, df = 5.9955, p-value = 0.2315
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.03775933 0.12780842
## sample estimates:
## mean of x mean of y
## 0.2817512 0.2367266
```

Evolution of the number of studies with bioeconomy



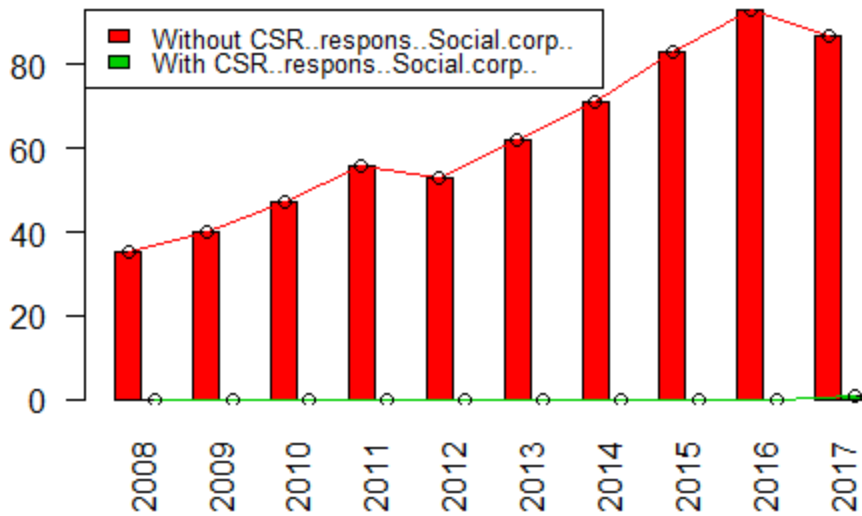
Evolution of the proportion of studies with bioeconomy



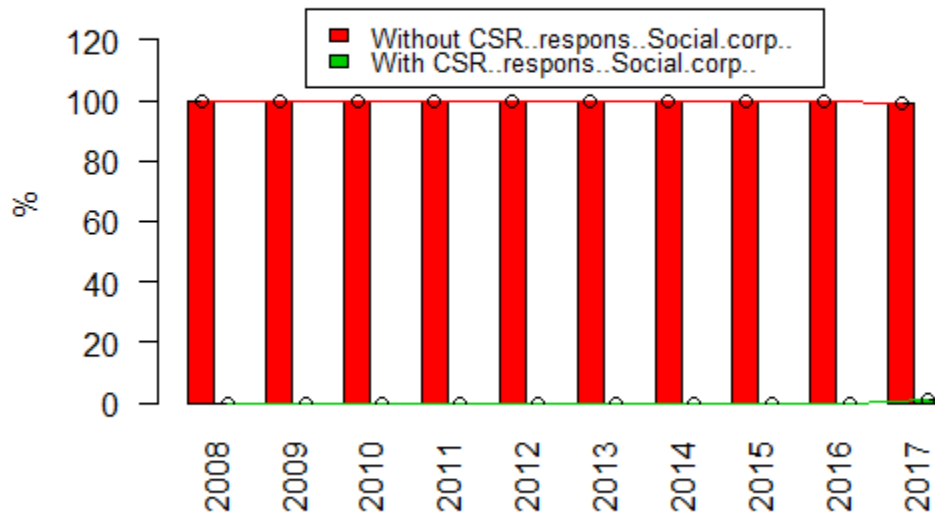
```
##
## Welch Two Sample t-test
##
## data: tbl1.2008_2012[2, ] and tbl1.2013_2017[2, ]
## t = -0.94868, df = 5.3691, p-value = 0.3835
```

```
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -2.1927285 0.9927285
## sample estimates:
## mean of x mean of y
## 0.6 1.2
##
##
## Welch Two Sample t-test
##
## data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]
## t = -0.21669, df = 7.2504, p-value = 0.8344
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.02488138 0.02067715
## sample estimates:
## mean of x mean of y
## 0.01305930 0.01516142
```

Evolution of the number of studies with CSR..respons..Social.corp..



Evolution of the proportion of studies with CSR..respons..Social.corp..

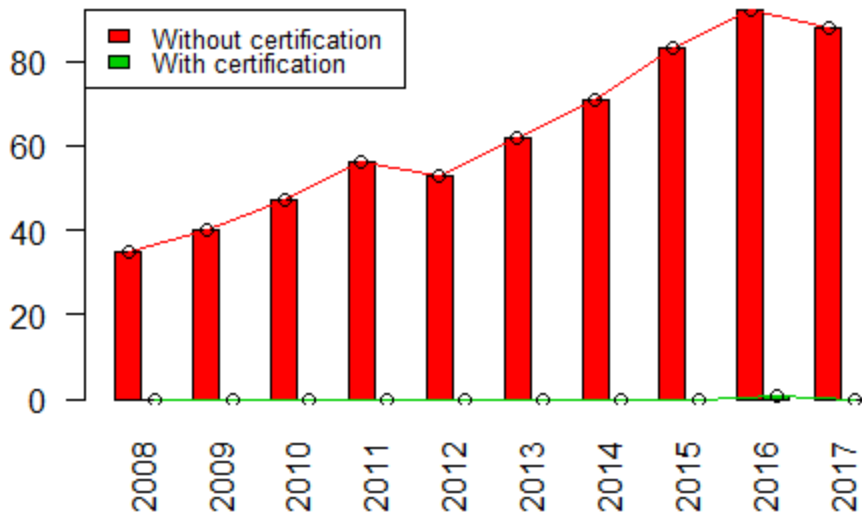


```
##
## Welch Two Sample t-test
##
## data: tb1.2008_2012[2, ] and tb1.2013_2017[2, ]
## t = -1, df = 4, p-value = 0.3739
```

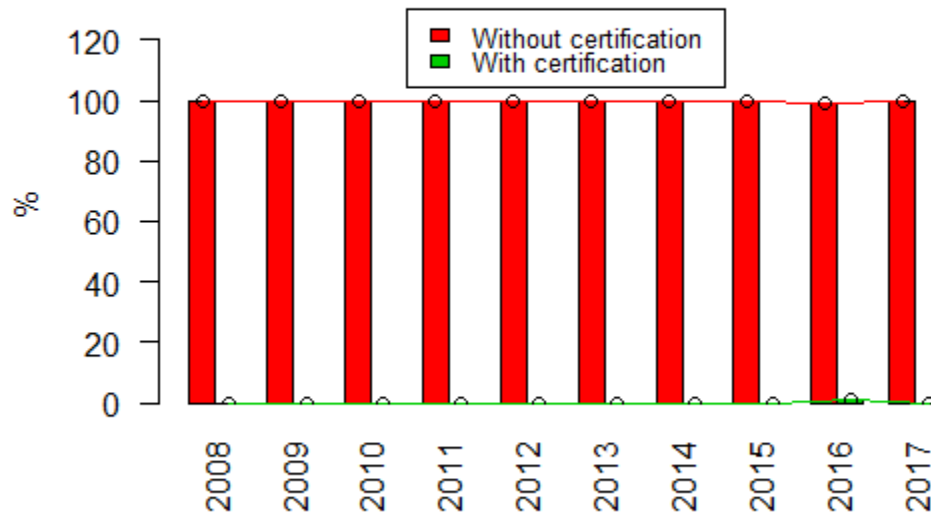
```
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.755289 0.355289
## sample estimates:
## mean of x mean of y
## 0.0 0.2
##
##
## Welch Two Sample t-test
##
## data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]
## t = -1, df = 4, p-value = 0.3739
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.008582830 0.004037375
## sample estimates:
## mean of x mean of y
## 0.000000000 0.002272727
```



Evolution of the number of studies with certification



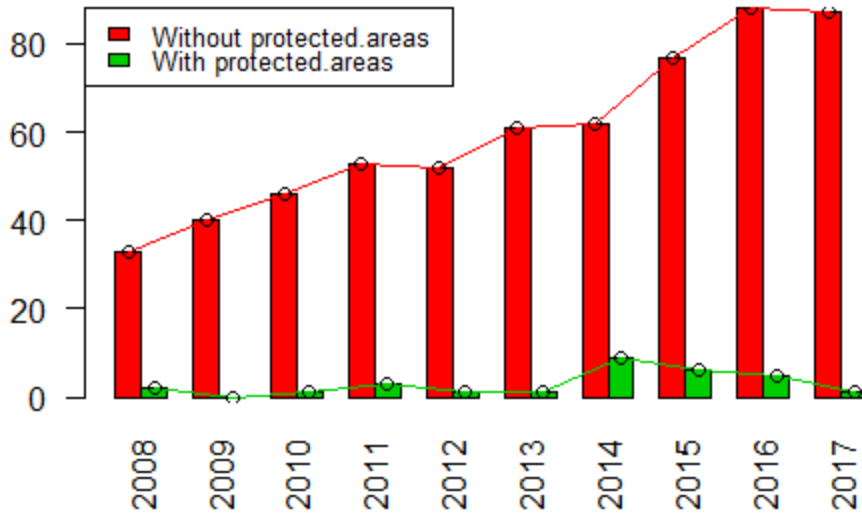
Evolution of the proportion of studies with certification



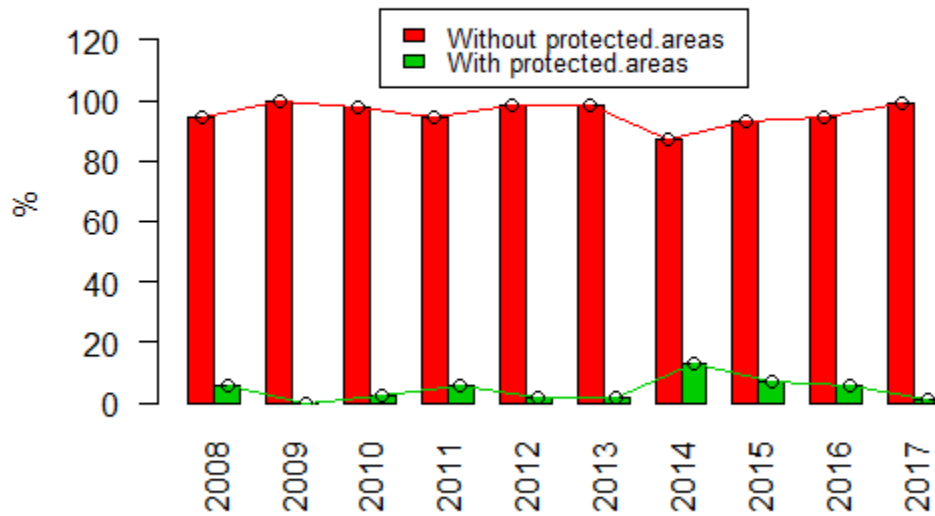
```
##  
## Welch Two Sample t-test  
##  
## data: tbl1.2008_2012[2, ] and tbl1.2013_2017[2, ]  
## t = -1, df = 4, p-value = 0.3739
```

```
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.755289 0.355289
## sample estimates:
## mean of x mean of y
## 0.0 0.2
##
##
## Welch Two Sample t-test
##
## data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]
## t = -1, df = 4, p-value = 0.3739
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.008121387 0.003820312
## sample estimates:
## mean of x mean of y
## 0.000000000 0.002150538
```

Evolution of the number of studies with protected areas



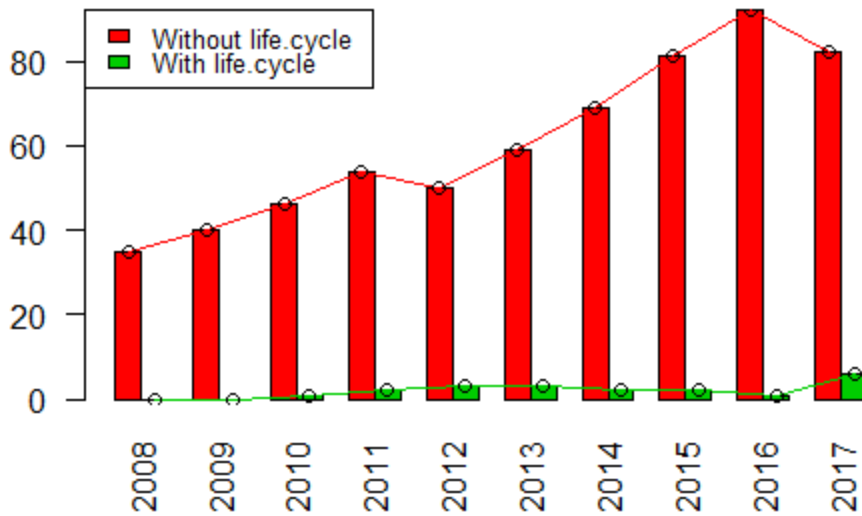
Evolution of the proportion of studies with protected areas



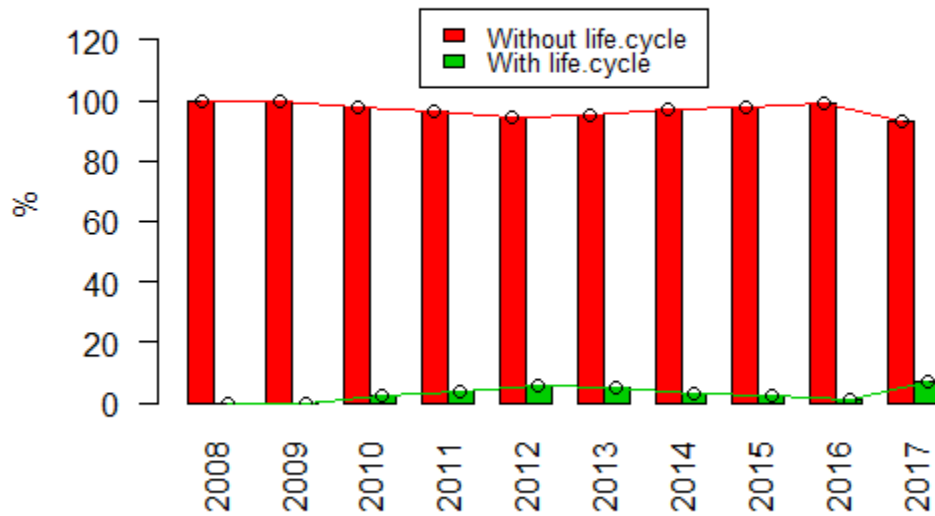
```
##
## Welch Two Sample t-test
##
## data: tbl1.2008_2012[2, ] and tbl1.2013_2017[2, ]
## t = -1.8534, df = 4.8708, p-value = 0.1246
```

```
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -7.19427 1.19427
## sample estimates:
## mean of x mean of y
## 1.4 4.4
##
##
## Welch Two Sample t-test
##
## data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]
## t = -1.0915, df = 6.0134, p-value = 0.3168
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.08389617 0.03211736
## sample estimates:
## mean of x mean of y
## 0.03017176 0.05606117
```

Evolution of the number of studies with life.cycle



Evolution of the proportion of studies with life.cycle



```
##
## Welch Two Sample t-test
##
## data: tbl1.2008_2012[2, ] and tbl1.2013_2017[2, ]
## t = -1.5396, df = 7.035, p-value = 0.1673
```

```
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -4.0549147 0.8549147
## sample estimates:
## mean of x mean of y
## 1.2 2.8
##
##
## Welch Two Sample t-test
##
## data: tbl2.2008_2012[2, ] and tbl2.2013_2017[2, ]
## t = -0.89177, df = 7.957, p-value = 0.3987
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.04736007 0.02096314
## sample estimates:
## mean of x mean of y
## 0.02271893 0.03591740
```

## Hypothesis 16

16. There is a higher probability of using GIS hybridized with some MCDM technique in the forestry area than in other areas.

==> We accept the Hypothesis.

==> Those studies within the forestry field are more likely (statistically significant) to use GIS with respect to fishery and with respect to agriculture

```
table(db$FOR, db$GIS)

##
##   0 1
## 0 266 88
## 1 168 106

chisq.test(table(db$FOR, db$GIS))

##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: table(db$FOR, db$GIS)
## X-squared = 13.192, df = 1, p-value = 0.0002811

summary(glm(db$GIS ~ db$FOR, family="binomial"))

##
## Call:
## glm(formula = db$GIS ~ db$FOR, family = "binomial")
##
## Deviance Residuals:
##   Min     1Q   Median     3Q    Max
## -0.9891 -0.9891 -0.7560  1.3782  1.6685
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.1062    0.1230  -8.995 < 2e-16 ***
## db$FOR      0.6456    0.1747   3.696 0.000219 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##   Null deviance: 776.50  on 627  degrees of freedom
## Residual deviance: 762.72  on 626  degrees of freedom
## AIC: 766.72
##
## Number of Fisher Scoring iterations: 4
```

```

db$Campo_f <- relevel(as.factor(db$Campo), "FOR")
summary(glm(db$GIS ~ db$Campo_f, family="binomial"))

##
## Call:
## glm(formula = db$GIS ~ db$Campo_f, family = "binomial")
##
## Deviance Residuals:
##   Min     1Q   Median     3Q      Max
## -0.9415 -0.9415 -0.7479  1.4333  1.7194
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -0.5839   0.1338  -4.364 1.28e-05 ***
## db$Campo_fAGR -0.5470   0.1911  -2.862 0.00421 **
## db$Campo_fFISH -0.6353   0.3429  -1.853 0.06390 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##   Null deviance: 711.04  on 590  degrees of freedom
## Residual deviance: 701.59  on 588  degrees of freedom
## (37 observations deleted due to missingness)
## AIC: 707.59
##
## Number of Fisher Scoring iterations: 4

```