

R code for the statistical analysis of the paper: ‘Taking advantage of sampling designs in Bayesian spatial small area survey studies’

Preparations

Libraries loading

```
if (!"pbugs" %in% installed.packages() | packageVersion("pbugs") < "0.1.4") {  
  remotes::install_github(repo = "fisabio/pbugs")  
}  
library(pbugs)  
  
library(spdep)  
  
## Loading required package: sp  
## Loading required package: spData  
## To access larger datasets in this package, install the spDataLarge  
## package with: `install.packages('spDataLarge',  
## repos='https://nowosad.github.io/drat/', type='source')`  
## Loading required package: sf  
## Linking to GEOS 3.9.1, GDAL 3.2.1, PROJ 7.2.1  
library(RColorBrewer)
```

Data loading

```
load("data/survey.Rdata")  
head(survey, n = 3)  
  
##   neigh NHIC   age ownhouse  
## 1    60    1 20-39        No  
## 2    31    0 20-39        No  
## 3    18    0 20-39        No  
  
# data.frame not supplied due to confidentiality restrictions  
  
# data corresponding to the Barcelona health survey questionnaires  
  
# Columns:  
  
# neigh: neighborhood  
  
# NHIC: Non-high income origin country  
  
# age: age group
```

```

# ownhouse: owner of his/her living house

load("data/neighs.Rdata")
head(neighs,n = 3)

##   gold.standard   n      N   p.0.19   p.20.39   p.40.59   p.60.79   p.80...
## 1    0.4586879 194 47359 0.1679301 0.3849743 0.2876116 0.1160920 0.04339196
## 2    0.2637934  46 15569 0.1109256 0.4253324 0.2786306 0.1318646 0.05324684
## 3    0.2023961  72 14941 0.1169266 0.3748745 0.2738103 0.1591594 0.07522923
##   p.NotOwn      p.Own
## 1 0.8043601 0.1956399
## 2 0.8180046 0.1819954
## 3 0.7428956 0.2571044

# (exhaustive) data on Barcelona neighborhoods corresponding to the city census

# Columns:

# gold.standard: percentage of people from non-high income origin country

# n: number of people in the neighborhood interviewed in the Barcelona health survey

# N: number of people in the neighborhood

# p.XX.YY: percentage of people in the neighborhood with age from XX to YY

# p.(Not)Own: Percentage of people in teh neighborhood owning their current home

load("data/geo.Rdata")
# Cartography (carto) and adjacencies list (adjacencies)

```

Simple Random Sampling (Case study 1)

Frequentist SRS estimate

```

# frequentist SRS estimate
pi.SRS<-tapply(survey$NHIC,survey$neigh,mean)

# Correlation with the gold standard
cor.SRS<-cor(pi.SRS,neighs$gold.standard)
cor.SRS

## [1] 0.6123257

# SQRT.MSE
MSE.SRS<-sqrt(mean((pi.SRS-neighs$gold.standard)^2))
MSE.SRS

## [1] 0.08322505

# CIs
sd.pi.SRS<-sqrt(pi.SRS*(1-pi.SRS)/neighs$n*(1-neighs$n/neighs$N))
CIl.SRS<-pi.SRS-qt(0.975,neighs$n)*sd.pi.SRS
CIu.SRS<-pi.SRS+qt(0.975,neighs$n)*sd.pi.SRS
# mean CI length

```

```
L.interval.SRS<-mean(2*qt(0.975,neighs$n)*sd.pi.SRS)
L.interval.SRS
```

```
## [1] 0.2707645
```

```
# Empirical coverage of the CIs
```

```
Cover.SRS<-mean(CI1.SRS<neighs$gold.standard & neighs$gold.standard<CIu.SRS)
Cover.SRS
```

```
## [1] 0.8082192
```

```
# Moran's I
```

```
nbNeigh<-nb2listw(poly2nb(carto),style="W")
# Moran's I for the gold.standard
I.true<-moran(neighs$gold.standard,nbNeigh,n=length(carto),Szero(nbNeigh))
# Moran's I for the SRS estimator
I.SRS<-moran(pi.SRS,nbNeigh,n=length(carto),Szero(nbNeigh))
c(I.true$I,I.SRS$I)
```

```
## [1] 0.2021324 0.1294423
```

```
# The gold standard shows larger spatial dependence than SRS
```

```
# Moran's I distribution under SRS
```

```
I.SRS.sample<-vector()
set.seed(1)
for(i in 1:1000){
  I.SRS.sample[i]<-moran(pi.SRS+rt(73,df=neighs$n)*sd.pi.SRS,nbNeigh,n=length(carto),Szero(nbNeigh))$I
}
# Probability of Moran's I distribution under SRS to be lower than that of the gold standard
mean(I.SRS.sample<I.true)
```

```
## [1] 0.986
```

Bayesian SRS estimate

```
obs<-tapply(survey$NHIC,survey$neigh,sum)
# Pis' posterior distribution: dbeta(obs+1,n-obs+1)
# Posterior Mode: (alpha-1)/(alpha+beta-2)=obs/n coincides with pi.SRS.
```

```
# CIs
```

```
CI1.SRSbayes<-qbeta(0.025,obs+1,neighs$n-obs+1)
CIu.SRSbayes<-qbeta(0.975,obs+1,neighs$n-obs+1)
L.interval.SRSbayes<-mean(CIu.SRSbayes-CI1.SRSbayes)
c(L.interval.SRS,L.interval.SRSbayes)
```

```
## [1] 0.2707645 0.2516170
```

```
Cover.SRSbayes<-mean(CI1.SRSbayes<neighs$gold.standard & neighs$gold.standard<CIu.SRSbayes)
c(Cover.SRS,Cover.SRSbayes)
```

```
## [1] 0.8082192 0.9589041
```

```
# Avoiding the normality assumption improves the empirical coverage problems of the previous CIs
```

```
# Moran's I
```

```
I.SRSbayes<-I.SRS
# Moran's I for the bayes estimator coincides with the SRS since their estimates also coincide
```

```

I.SRSbayes.sample<-vector()
set.seed(1)
for(i in 1:1000){
I.SRSbayes.sample[i]<-moran(rbeta(73,obs+1,neighs$n-obs+1)
      ,nbNeigh,n=length(carto),Szero(nbNeigh))$I
}
mean(I.SRSbayes.sample<I.true)

```

```
## [1] 0.969
```

```
# The lack of spatial dependence of pi.SRS is not solved by the bayes estimates.
```

Spatially dependent Bayesian estimates

```

# WinBUGS Model
modSRSsp <- function() {
  for (i in 1:nIndiv) {
    Y[i] ~ dbern(p[i])
    logit(p[i]) <- mu+sd.theta*theta[neigh[i]]
  }

  mu~dflat()
  sd.theta~dunif(0,10)

  # lCAR distribution for theta
  for (j in 1:nNeigh) {
    theta[j] ~ dnorm(media.theta[j], prec.theta[j])
    prec.theta[j] <- (1 - lambda + lambda * nadj[j])
    media.theta[j] <- (lambda / (1 - lambda + lambda * nadj[j])) *
      sum(theta.map[(index[j] + 1):index[j + 1]])

    logit(pi[j])<-mu+sd.theta*theta[j]
  }
  for (k in 1:nadj.tot) {
    theta.map[k] <- theta[map[k]]
  }
  lambda~dunif(0,1)
}

# Data
dataSRSsp <- list(
  Y = survey$NHIC,
  neigh = survey$neigh,
  nNeigh = nrow(neighs),
  nIndiv = sum(neighs$n),
  nadj.tot = length(adjacencies$adj),
  nadj = adjacencies$num,
  map = adjacencies$adj,
  index = c(0, cumsum(adjacencies$num))
)

# Parameters to save
parametersSRSsp <- c("mu", "lambda", "sd.theta", "pi")

```

```

# Inits
initsSRSSp <- function() {
  list(
    theta= rnorm(dataSRSSp$nNeigh, 0,1), mu=rnorm(1),sd.theta=runif(1)
  )
}

set.seed(1)
resultSRSSp <- pbugs(
  data          = dataSRSSp,
  inits          = initsSRSSp,
  parameters.to.save = parametersSRSSp,
  model.file     = modSRSSp,
  n.chains       = 3,
  n.iter         = 6000,
  n.burnin       = 1000,
  n.thin = 1,
  bugs.seed = 1
)

#  $\hat{\pi}^{\text{spatial}}$  estimates
Y.miss<-matrix(nrow=dim(resultSRSSp$sims.list$pi)[1],ncol=dim(resultSRSSp$sims.list$pi)[2])
for(i in 1:(dim(resultSRSSp$sims.list$p)[1])){
  Y.miss[i,]<-rbinom(dim(neighs)[1],neighs$N-neighs$n,resultSRSSp$sims.list$pi[i,])
}
Y.sample<-tapply(survey$NHIC,survey$neigh,sum)
p.spatial<-matrix(nrow=dim(resultSRSSp$sims.list$pi)[1],ncol=dim(resultSRSSp$sims.list$pi)[2])
for(i in 1:(dim(resultSRSSp$sims.list$p)[1])){
  p.spatial[i,]<-(neighs$n/neighs$N)*(Y.sample/neighs$n)+((neighs$N-neighs$n)/neighs$N)*Y.miss[i,]/(nei
}

# Correlation with the gold standard
cor(neighs$gold.standard,apply(p.spatial,2,mean))

## [1] 0.7468466
cor(neighs$gold.standard,obs/neighs$n)

## [1] 0.6123257

# SQRT.MSE
sqrt(mean((neighs$gold.standard-apply(p.spatial,2,mean))^2))

## [1] 0.04904433
sqrt(mean((neighs$gold.standard-(obs/neighs$n))^2))

## [1] 0.08322505

# CIs
CIl.SRSSp<-apply(p.spatial,2,quantile,0.025)
CIu.SRSSp<-apply(p.spatial,2,quantile,0.975)
L.interval.SRSSp<-mean(CIu.SRSSp-CIl.SRSSp)
c(L.interval.SRSbayes,L.interval.SRSSp)

## [1] 0.2516170 0.1721787

```

```

Cover.SRSsp<-mean(CI1.SRSsp<neighs$gold.standard & neighs$gold.standard<CIu.SRSsp)
c(Cover.SRS, Cover.SRSbayes, Cover.SRSsp)

## [1] 0.8082192 0.9589041 0.9452055

# Moran's I
I.SRSsp.sample<-apply(p.spatial,1,function(x){moran(x,nbNeigh,n=length(carto),Szero(nbNeigh))$I})
mean(I.SRSsp.sample)

## [1] 0.2328906
mean(I.SRSsp.sample<I.true)

## [1] 0.6742667
# The spatial dependence of this estimator matches that observed at population level.

# \hat{\pi}^star estimates
# Correlation with the gold standard
cor(neighs$gold.standard,resultSRSsp$mean$pi)

## [1] 0.7469775
cor(neighs$gold.standard,obs/neighs$n)

## [1] 0.6123257
# SQRT.MSE
sqrt(mean((neighs$gold.standard-resultSRSsp$mean$pi)^2))

## [1] 0.04901569
sqrt(mean((neighs$gold.standard-(obs/neighs$n))^2))

## [1] 0.08322505
# CIs
CI1.SRSsp<-resultSRSsp$summary[grepl("pi",rownames(resultSRSsp$summary)),3]
CIu.SRSsp<-resultSRSsp$summary[grepl("pi",rownames(resultSRSsp$summary)),7]
L.interval.SRSsp<-mean(CIu.SRSsp-CI1.SRSsp)
c(L.interval.SRSbayes,L.interval.SRSsp)

## [1] 0.2516170 0.1719827
Cover.SRSsp<-mean(CI1.SRSsp<neighs$gold.standard & neighs$gold.standard<CIu.SRSsp)
c(Cover.SRS, Cover.SRSbayes, Cover.SRSsp)

## [1] 0.8082192 0.9589041 0.9452055

# Moran's I
I.SRSsp.sample<-apply(resultSRSsp$sims.list$pi,1,function(x){moran(x,nbNeigh,n=length(carto),Szero(nbNeigh))$I})
mean(I.SRSsp.sample)

## [1] 0.2333322
mean(I.SRSsp.sample<I.true)

## [1] 0.6712
# The spatial dependence of this estimator matches that observed at population level.

```

Stratified sampling estimator (Age)

7

```

L.interval.str<-mean(2*qt(0.975,neighs$n)*sd.pi.str,na.rm=T)
c(mean(CIu.SRS[!is.na(pi.str)]-CIl.SRS[!is.na(pi.str)]),L.interval.str)

## [1] 0.2055585 0.1009214
# The stratified CIs are substantially narrower than the SRSs

Cover.str<-mean((pi.str-qt(0.975,neighs$n)*sd.pi.str)<neighs$gold.standard & neighs$gold.standard<(pi.s
Cover.str

## [1] 0.6
# ICs Coverage for SRS wherever pi.str is not missing
mean(CI1.SRS[!is.na(pi.str)]<neighs$gold.standard[!is.na(pi.str)] & neighs$gold.standard[!is.na(pi.str)]

## [1] 0.8666667
# The coverage of the ICs is also far lower than for SRS

# Moran's I
#####
I.str<-moran(pi.str,nbNeigh,n=length(carto),Szero(nbNeigh),NAOK = TRUE)$I
I.str

## [1] 0.04356376

aux<-neighs$gold.standard
aux[is.na(pi.str)]<-NA
I.true.NA<-moran(aux,nbNeigh,n=length(carto),Szero(nbNeigh),NAOK = TRUE)$I
I.true.NA

## [1] 0.1409699
# Moran's I for the stratified estimator is lower than for the gold standard

# Randomization
I.str.sample<-vector()
for(i in 1:1000){I.str.sample[i]<-moran(rnorm(length(carto),pi.str,sd.pi.str),nbNeigh,n=length(carto),S
mean(I.str.sample<I.true.NA)

## [1] 1
# Moran's I of the stratified estimator does not fit that of the gold.standard

# Moran's I for SRS excluding those outcomes where pi.str is missing
aux<-pi.SRS
aux[is.na(pi.str)]<-NA
I.SRS.NA<-moran(aux,nbNeigh,n=length(carto),Szero(nbNeigh),NAOK = TRUE)$I
I.SRS.NA

## [1] 0.04798209

I.SRS.NA.sample<-vector()
for(i in 1:1000){I.SRS.NA.sample[i]<-moran(rnorm(length(carto),aux,sd.pi.SRS),nbNeigh,n=length(carto),S
mean(I.SRS.NA.sample<I.true.NA)

## [1] 1

```


Ratio estimator (House owning)

```
pi.own<-cbind(
  tapply(survey$NHIC[survey$ownhouse=="No"],factor(survey$neigh[survey$ownhouse=="No"],levels = 1:73),mean),
  tapply(survey$NHIC[survey$ownhouse=="Yes"],factor(survey$neigh[survey$ownhouse=="Yes"],levels = 1:73),mean),
  c(mean(is.na(pi.own)),mean(apply(is.na(pi.own),1,any)))

## [1] 0.02739726 0.05479452

# 2.7% of the owning-neighborhood combinations are not represented in the sample, what will make a 5.5%

# Proportion of owning-groups-neighborhoods combinations with sample sizes between 1 and 5
(sum(0<table(factor(survey$neigh[survey$ownhouse=="No"],levels = 1:73)) & table(factor(survey$neigh[survey$ownhouse=="No"],levels = 1:73))) & table(factor(survey$neigh[survey$ownhouse=="No"],levels = 1:73))) +
sum(0<table(factor(survey$neigh[survey$ownhouse=="Yes"],levels = 1:73)) & table(factor(survey$neigh[survey$ownhouse=="Yes"],levels = 1:73))))

## [1] 0.0890411

# Ratio estimator including house owning information
pi.ratio<-rowSums(pi.own*cbind(neighs$p.NotOwn,neighs$p.Own))

# Correlation with the gold standard
cor.ratio<-cor(pi.ratio[!is.na(pi.ratio)],neighs$gold.standard[!is.na(pi.ratio)])
cor.ratio

## [1] 0.7272673

cor(pi.SRS[!is.na(pi.ratio)],neighs$gold.standard[!is.na(pi.ratio)])

## [1] 0.6990941

# The ratio estimator shows an evident improvement over the SRS estimator

# SQRT.MSE
sqrt(mean((pi.ratio[!is.na(pi.ratio)]-neighs$gold.standard[!is.na(pi.ratio)])^2))

## [1] 0.06227874

sqrt(mean((pi.SRS[!is.na(pi.ratio)]-neighs$gold.standard[!is.na(pi.ratio)])^2))

## [1] 0.06888978

# The improvement is also evident in terms of MSE

# CIs
sd.pi.ratio<-sqrt(rowSums(pop.age*pop.age*pi.ratio*(1-pi.ratio)/(neighs$n-1)*(neighs$N-neighs$n)/neighs$n))
L.interval.ratio<-mean(2*qt(0.975,neighs$n)*sd.pi.ratio,na.rm=T)
c(mean(CIu.SRS[!is.na(pi.ratio)]-CIl.SRS[!is.na(pi.ratio)]),L.interval.ratio)

## [1] 0.2316223 0.1151147

# The length of the ratio estimator is substantially lower for the ratio estimator

Cover.ratio<-mean((pi.ratio-qt(0.975,neighs$n)*sd.pi.ratio)<neighs$gold.standard & neighs$gold.standard<pi.ratio+qt(0.975,neighs$n)*sd.pi.ratio)

Cover.ratio

## [1] 0.6666667

mean(CIu.SRS[!is.na(pi.ratio)]<neighs$gold.standard[!is.na(pi.ratio)] & neighs$gold.standard[!is.na(pi.ratio)]<CIl.SRS[!is.na(pi.ratio)])

## [1] 0.826087
```

```
# Anyway, the coverage of the CIs is lower than for SRSs, and lower than expected
```

```
# Moran, s I
```

```
I.ratio<-moran(pi.ratio,nbNeigh,n=length(carto),Szero(nbNeigh),NAOK = TRUE)$I  
I.ratio
```

```
## [1] 0.1497097
```

```
aux<-neighs$gold.standard
```

```
aux[is.na(pi.ratio)]<-NA
```

```
I.true.NA<-moran(aux,nbNeigh,n=length(carto),Szero(nbNeigh),NAOK = TRUE)$I  
c(I.ratio,I.true.NA)
```

```
## [1] 0.1497097 0.1788096
```

```
I.ratio.sample<-vector()
```

```
set.seed(1)
```

```
for(i in 1:1000){I.ratio.sample[i]<-moran(rnorm(length(carto),pi.ratio,sd.pi.ratio),nbNeigh,n=length(ca)  
mean(I.ratio.sample<I.true.NA)
```

```
## [1] 0.806
```

```
# I.SRS considering as missing values those values with missing pi.str
```

```
aux<-pi.SRS
```

```
aux[is.na(pi.ratio)]<-NA
```

```
I.ratio.NA<-moran(aux,nbNeigh,n=length(carto),Szero(nbNeigh),NAOK = TRUE)$I  
I.ratio.NA
```

```
## [1] 0.1583015
```

```
I.ratio.NA.sample<-vector()
```

```
set.seed(1)
```

```
for(i in 1:1000){I.ratio.NA.sample[i]<-moran(rnorm(length(carto),aux,sd.pi.SRS),nbNeigh,n=length(carto)  
mean(I.ratio.NA.sample<I.true.NA)
```

```
## [1] 0.838
```

Full (stratified-ratio-spatial) model

```
# Model
```

```
modFull <- function() {  
  for (i in 1:nIndiv) {  
    Y[i] ~ dbern(p[i])  
    logit(p[i]) <- mu +  
      beta.age[age[i]] +  
      beta.own[own[i]] +  
      sd.theta*theta[neigh[i]]  
  }  
}
```

```
mu ~ dflat()
```

```
sd.theta ~ dunif(0,10)
```

```
beta.age[3] <- 0
```

```
for (k in 1:2) {  
  beta.age[k] ~ dflat()  
}
```

```
for (k in 4:nAge) {
```

```

    beta.age[k] ~ dflat()
  }
  beta.own[1] <- 0
  beta.own[2] ~ dflat()

  lambda ~ dunif(0, 1)

  # lCAR distribution for theta
  for (k in 1:nNeigh) {
    theta[k] ~ dnorm(media.theta[k], prec.theta[k])
    prec.theta[k] <- (1 - lambda + lambda * nadj[k])
    media.theta[k] <- (lambda / (1 - lambda + lambda * nadj[k])) *
      sum(theta.map[(index[k] + 1):index[k + 1]])
  }
  for (k in 1:nadj.tot) {
    theta.map[k] <- theta[map[k]]
  }

  for(iNeigh in 1:nNeigh){
    for(iAge in 1:nAge){
      for(iOwn in 1:nOwn){
        logit(pi.pred[iAge,iOwn,iNeigh])<- mu +
          beta.age[iAge] +
          beta.own[iOwn] +
          sd.theta*theta[iNeigh]
      }
      pi.pred2[iAge,iNeigh]<-inprod2(pi.pred[iAge,,iNeigh],p.own[iNeigh,])
    }
    pi[iNeigh]<-inprod2(pi.pred2[,iNeigh],p.age[iNeigh,])
  }
}

# Data
dataFull <- list(
  Y = survey$NHIC,
  age = as.numeric(survey$age),
  own = as.numeric(survey$ownhouse),
  nOwn = length(unique(survey$ownhouse)),
  nAge = length(unique(survey$age)),
  neigh = survey$neigh,
  nNeigh = nrow(neighs),
  nadj.tot = length(adjacencies$adj),
  nadj = adjacencies$num,
  map = adjacencies$adj,
  index = c(0, cumsum(adjacencies$num)),
  nIndiv = sum(neighs$n),
  p.age = as.matrix(pop.age),
  p.own = cbind(neighs$p.NotOwn,neighs$p.Own)
)

# Inits and parameters
parametersFull <- c(
  "mu", "lambda", "sd.theta", "beta.age", "beta.own", "theta", "pi"

```

```

)

initsFull <- function() {
  list(
    mu = rnorm(1, 0, 0.1),
    beta.age = c(rnorm(2), NA, rnorm(dataFull$nAge - 3)),
    beta.own = c(NA, rnorm(1)),
    lambda = runif(1),
    sd.theta = runif(1),
    theta = rnorm(dataFull$nNeigh, 0.1)
  )
}

# Model run
set.seed(1)
resultFull <- pbugs(
  data = dataFull,
  inits = initsFull,
  parameters.to.save = parametersFull,
  model.file = modFull,
  n.chains = 3,
  n.iter = 6000,
  n.burnin = 1000,
  DIC = FALSE,
  bugs.seed = 1
)

# Correlation
cor(neighs$gold.standard, resultFull$mean$pi)

## [1] 0.7962962
cor(neighs$gold.standard, resultSRSsp$mean$pi)

## [1] 0.7469775
cor(obs/neighs$n, neighs$gold.standard)

## [1] 0.6123257
# SQRT.MSE
sqrt(mean((neighs$gold.standard - resultFull$mean$pi)^2))

## [1] 0.04336361
sqrt(mean((neighs$gold.standard - resultSRSsp$mean$pi)^2))

## [1] 0.04901569
sqrt(mean((neighs$gold.standard - (obs/neighs$n))^2))

## [1] 0.08322505
# CIs
CI1.Full <- resultFull$summary[grepl("pi", rownames(resultFull$summary)), 3]
CIu.Full <- resultFull$summary[grepl("pi", rownames(resultFull$summary)), 7]
L.interval.Full <- mean(CIu.Full - CI1.Full)
c(L.interval.SRSbayes, L.interval.SRSsp, L.interval.Full)

```

```
## [1] 0.2516170 0.1719827 0.1412398
Cover.Full<-mean(CI1.Full<neighs$gold.standard & neighs$gold.standard<CIu.Full)
c(Cover.SRS,Cover.SRSbayes,Cover.SRSsp,Cover.Full)

## [1] 0.8082192 0.9589041 0.9452055 0.9589041
# Moran's I
I.full<-moran(resultFull$mean$pi,nbNeigh,n=length(carto),Szero(nbNeigh))$I
I.full

## [1] 0.3050834
I.Full.sample<-apply(resultFull$sims.list$pi,1,function(x){moran(x,nbNeigh,n=length(carto),Szero(nbNeigh))})
mean(I.Full.sample<I.true)

## [1] 0.6437126
# The spatial dependence observed in the gold standard matches that observed in the full estimator.
```

Figure 1

```
dev.new()

par(mfrow=c(2,2), mar=c(0,0,2,0))
pi.SRS.cut<-cut(pi.SRS,c(-0.1,0.05,0.10,0.15,0.20,0.25,1.1))
cols.SRS<-brewer.pal(name = "Blues",n=6)[pi.SRS.cut]
plot(carto,col=cols.SRS)
legend(x="bottomright",legend=c("<5%", "5%-9%", "10%-14%", "15%-19%", "20%-24%", ">=25%"),fill=brewer.pal(name = "Blues",n=6))
title("SRS")

pi.spatial.cut<-cut(resultSRSsp$mean$pi,c(-0.1,0.05,0.10,0.15,0.20,0.25,1.1))
cols.spatial<-brewer.pal(name = "Blues",n=6)[pi.spatial.cut]
plot(carto,col=cols.spatial)
title("Spatial SRS")

pi.Full.cut<-cut(resultFull$mean$pi,c(-0.1,0.05,0.10,0.15,0.20,0.25,1.1))
cols.Full<-brewer.pal(name = "Blues",n=6)[pi.Full.cut]
plot(carto,col=cols.Full)
title("Full estimator")

pi.gold<-cut(neighs$gold.standard,c(-0.1,0.05,0.10,0.15,0.20,0.25,1.1))
cols.gold<-brewer.pal(name = "Blues",n=6)[pi.gold]
plot(carto,col=cols.gold)
title("Gold standard")
```