

ITS Health Intervention

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Install packages required for the analysis (uncomment if needed)

```
install.packages("lmtest") ; install.packages("Epi") install.packages("tsModel"); install.packages("vcd")
```

load the packages

```
library(foreign) library(tsModel) library("lmtest") library("Epi") library("splines") library("vcd")
```

read data from csv file

```
data <- read.csv("sicily.csv") head(data) data
```

1. Descriptive Statistics

compute the standardized rates

```
datarate <- with(data, aces/stdpop*105) # start the plot, excluding the points and the x-axis plot(datarate, type="n", ylim=c(0, 10000),  
ylab="Std rate x 10,000", bty="l", xaxt="n") # shade the post intervention period grey rect(36, 0, 60, 300, col=grey(0.9), border=F)  
# plot the observed rate for pre-intervention period points(datarate[datasmokban==0], cex=0.7) # specify  
the x-axis (i.e. time units) axis(1, at=0:5*12, labels=F) axis(1, at=0:4*12+6, tick=F, labels=2002:2006) # add a  
title title("Sicily, 2002-2006")
```

```
summary(data)
```

```
# tabulate aces before and after the smoking ban summary(dataaces[datasmokban==0]) summary(dataaces[datasmokban==1])  
summary(datarate[datasmokban==0]) summary(datarate[datasmokban==1])
```

2. Regression model

```
# Poisson with the standardised population as an offset model1 <- glm(aces ~ offset(log(stdpop)) + smokban  
+ time, family=poisson, data) summary(model1) summary(model1)$dispersion round(ci.lin(model1, Exp =  
T), 3) datanew <- data.frame(stdpop = mean(datastdpop), smokban=rep(c(0,1), c(360, 240)), time=  
1:600/10, month=rep(1:120/10, 5)) # We generate predicted values based on the model in order to create a plot  
pred1 <- predict(model1, type="response", datanew)/mean(datastdpop)*105 # This can then be plotted along with a scatter graph(s  
rate x 10,000", bty="l", xaxt="n") rect(36, 0, 60, 300, col=grey(0.9), border=F) points(datarate, cex =  
0.7) axis(1, at = 0 : 5 * 12, labels = F) axis(1, at = 0 : 4 * 12 + 6, tick = F, labels = 2002 :  
2006) lines((1 : 600/10), pred1, col = 2) title("Sicily, 2002 - 2006") datanew <- data.frame(stdpop =  
mean(datastdpop), smokban=0, time=1:600/10, month=rep(1:120/10, 5)) # generate predictions under the  
counterfactual scenario and add it to the plot pred1b <- predict(model1, datanew, type="response")/mean(datastdpop)*
```

```
105lines(datanewtime,pred1b,col=2,lty=2) # return the data frame to the scenario including the intervention
datanew <- data.frame(stdpop=mean(data$stdpop),smokban=rep(c(0,1),c(360,240)), time=1:600/10,month=rep(1:120/10,5))
```

#3. methodological issues

```
#Model checking and autocorrelation # Check the residuals by plotting against time res2 <- residuals(model2,type="deviance") plot(data$time,res2,ylim=c(-5,10),pch=19,cex=0.7,col=grey(0.6),
main="Residuals over time",ylab="Deviance residuals",xlab="Date") abline(h=0,lty=2,lwd=2) # Further check for autocorrelation by examining the autocorrelation and # partial autocorrelation functions acf(res2)
pacf(res2)
```

#4. adjusting for seasonality

There are various ways of adjusting for seasonality - here we use harmonic

terms specifying the number of sin and cosine pairs to include (in this

case 2) and the length of the period (12 months)

```
model3 <- glm(aces ~ offset(log(stdpop)) + smokban + time + harmonic(month,2,12), family=quasipoisson, data)
summary(model3) summary(model3)$dispersionround(ci.lin(model3,Exp = T),3)#EFFECTSci.lin(model3,Exp = T)["smokban",5 : 7]#TRENDExp(coef(model3)["time"] * 12)#Weagaincheckthemodelandautocorrelationfunctionsres3 <- residuals(model3,type = "deviance")plot(res3,ylim = c(-5, 10),pch = 19,cex = 0.7,col = grey(0.6),main = "Residualsovertime",ylab = "Devianceresiduals",xlab = "Date")abline(h = 0,lty = 2,lwd = 2)acf(res3)pacf(res3)#predictandplotoftheseasonallyadjustedmodelpred3 <- predict(model3,type = "response",datanew)/mean(data$stdpop)10^5 plot(datarate,type = "n",ylim = c(120,300),xlab = "Year",ylab = "Std rate x 10,000",bty = "l",xaxt = "n")rect(36,120,60,300,col = grey(0.9),border = F)points(datarate,cex=0.7) axis(1,at=0:512,labels=F) axis(1,at=0:412+6,tick=F,labels=2002:2006) lines(1:600/10,pred3,col=2) title("Sicily, 2002-2006") # it is sometimes difficult to clearly see the change graphically in the # seasonally adjusted model, therefore it can be useful to plot a straight # line representing a 'deseasonalised' trend # this can be done by predicting all the observations for the same month, in # this case we use June pred3b <- predict(model3,type="response",transform(datanew,month=6))/ mean(data$stdpop)10^5 #this can then be added to the plot as a dashed line lines(1:600/10,pred3b,col=2,lty=2)
```

5. Additional material

add a change-in-slope

we parameterize it as an interaction between time and the ban indicator

```
model4 <- glm(aces ~ offset(log(stdpop)) + smokbantime + harmonic(month,2,12), family=quasipoisson, data)
summary(model4) round(ci.lin(model4,Exp=T),3) # predict and plot the 'deseasonalised' trend # compare it with the step-change only model pred4b <- predict(model4,type="response",transform(datanew,month=6))/ mean(data$stdpop) * 105plot(datarate,type="n",ylim=c(120,300),xlab="Year",ylab="Std rate x 10,000",bty="l",xaxt="n") rect(36,120,60,300,col=grey(0.9),border=F) points(data$rate,cex=0.7) axis(1,at=0:512,labels=F) axis(1,at=0:412+6,tick=F,labels=2002:2006) lines(1:600/10,pred3b,col=2) lines(1:600/10,pred4b,col=4) title("Sicily, 2002-2006") legend("topleft",c("Step-change only","Step-change + change-in-slope"),lty=1,
```

```
col=c(2,4),inset=0.05,bty="n",cex=0.7) # test if the change-in-slope improve the fit # the selected test
here is an F-test, which accounts for the overdispersion, # while in other cases a likelihood ratio or wald
test can be applied anova(model3,model4,test="F") # not surprisingly, the p-value is similar to that of the
interaction term
```