Environmental Health Impact Assessment using R $_{\rm Mapping Risks}$

In this session we will use R to do some disease mapping. We will work through an example of how to create basic maps in R by creating a map of Mexico City. We will then move on to an example of creating smoothed SMRs and plot them on a map by working with data on hospital admissions for chronic obstructive pulmonary disease (COPD) for England between 2001–2010. All data required for this practical can be found in the folder Data. You will need the following files

- shapefiles and information for Mexico City split by municipalities (cdmx.shp, cdmx.dbf)
- population count and density for Mexico City split by municipalities (cdmx.csv)
- shapefiles and information for England split by local authorities (englandlocalauthority.shp, englandlocalauthority.dbf)
- observed numbers of hospital admissions by local authority (copdmortalityobserved.csv)
- expected numbers of hospital admissions by local authority (copdmortalityobserved.csv).

Preliminaries

For this practical, we need the following packages

- spdep Package to create spatial objects (such as neighbourhood matrix).
- shapefiles Package to read and write shapefiles.
- CARBayes Package to fit spatial GLMMs.
- rgdal Package to handle spatial objects.

As in Practical 1, we use the install.packages() function to download and install the packages that we need.

```
# Installing required packages
install.packages("spdep")
install.packages("shapefiles")
install.packages("CARBayes")
install.packages("rgdal")
```

We use the library() function to load them into the R library.

```
# Loading required packages into the library
library(spdep)
library(shapefiles)
library(CARBayes)
library(rgdal)
```

Before reading in any data for this practical you will need to ensure that you are in the correct folder. As explained in Practical 1, you can use the setwd() function

```
setwd("Chosen_Directory_Path")
```

If you cannot get the setwd() to work, go to Session > Set Working Directory > Choose Directory in the toolbar on the top.

Remember, more information about any of the functions used here can be found by typing help(function_name) or ?function_name into R.

Creating maps of Scotland

To create maps, we use something called 'shapefiles'. Shapefiles contain location, shape, and attributes of geographic features such as country borders. The files Scotland_County.shp, and Scotland_County.dbf contain the location, shape, and attributes of Scotland by county. These were obtained from http://www.gadm.org. The functions read.shp() and read.dbf() will read these shapefiles into R.

```
# Reading in borders
shp_Scotland_C <- read.shp(shp.name = "Scotland_County.shp")
dbf_Scotland_C <- read.dbf(dbf.name = "Scotland_County.dbf")</pre>
```

The file Scotland_County.csv contains the population of Scotland by county and we will use this to create maps. These are in csv format, so we use the read.csv() function.

```
# Read population data for Scotland
pop_Scotland_C <- read.csv('Scotland_County.csv', row.names = 1)</pre>
```

To check that the data has been read into R correctly, we can use the head() and function, which prints the first six rows of a dataset.

```
# Printing first six rows
head(pop_Scotland_C)
```

	Name	Pop2016	PopPercChange2016
1	Aberdeenshire	262190	0.09
2	Aberdeen	229840	-0.22
3	Angus	116520	-0.33
4	Argyll and Bute	87130	0.28
5	Clackmannanshire	51350	-0.02
6	Dumfries and Galloway	149520	-0.10

We can see that this dataset contains the following variables:

- Name County name,
- Pop2016 Population count in 2016,
- PopPercChange2016 Percentage change in population between 2015 and 2016.

Lets create a map of population in 2016 for Scotland. To plot the population data in a map, we need to attach them to the shapefiles. The function combine.data.shapefile() allows us to combine shapefiles to plot later.

We use downloaded population data to create the map here. If you have your own data, we can use that later. To plot the map, we use the **spplot()** function.

```
# Scaling population counts (to 1000s
Scotland_C$Pop2016 <- Scotland_C$Pop2016/1000
# Creating map of population counts in Scotland
spplot(obj = Scotland_C, # Spatial object to be plotted
zcol = c("Pop2016"), # Choice of the column the object you are plotting.
main = "Population (in 1000s)", # Plot title
at = seq(0,650, length.out=20), # Break points for legend
col = 'transparent', # Colour for borders
col.regions = hsv(0.6, seq(0.2, 1, length.out=20), 1)) # Create a set of colours
```

Population (in 1000s)



Activities

- Carefully change the above code to create a map of the percentage change in population between 2015 and 2016 in Scotland.
- In the folder Data there are also shapefiles for Scotland split by administrative boundary (Scotland_Admin.shp, Scotland_Admin.dbf, Scotland_Admin.csv). Add your own data to the Scotland_Admin.csv file and carefully change the above code to create your own map of Scotland by administrative boundary.

COPD in England

We now look at example into the hospital admission rates for chronic obstructive pulmonary disease (COPD) in England between 2001–2010. In England, there are 324 local authority administrative areas each with an observed and expected number of cases. The expected numbers were calculated using indirect standardisation by applying the age–sex specific rates for the whole of England to the age–sex population profile of each of the areas.

Reading in data and shapefiles

To create SMR maps, we need to read in the relevant shapefiles. The files englandlocalauthority.shp and englandlocalauthority.dbf contain the location, shape, and attributes of English local authorities. The functions read.shp() and read.dbf() will read these shapefiles into R.

```
# Reading in borders
shp <- read.shp(shp.name="englandlocalauthority.shp")
dbf <- read.dbf(dbf.name="englandlocalauthority.dbf")</pre>
```

The observed and expected COPD counts in England by local authority need to be read into R. These are in csv format, so we use the read.csv() function.

```
# Reading in observed numbers of hospital admissions in England by local authority
observed <- read.csv(file="copdmortalityobserved.csv", row.names=1)</pre>
```

```
# Reading in expected numbers of hospital admissions in England by local authority
expected <- read.csv(file="copdmortalityexpected.csv", row.names=1)</pre>
```

To check that the data has been read into R correctly, we can use the head() function, which prints the first six rows of a dataset.

5

91

72

53

Printing first six rows of the observed counts head(observed) name Y2001 Y2002 Y2003 Y2004 Y2005 Y2006 Y2007 OOAA City of London LB 2 0 3 1 1 1 00AB Barking and Dagenham LB 100 100 122 93 136 97 OOAC 89 Barnet LB 110 102 106 99 97 OOAD Bexley LB 109 113 113 96 97 94 113 OOAE 70 Brent LB 69 89 59 61 48 00AF Bromley LB 120 129 135 124 128 117 120 Y2008 Y2009 Y2010 OOAA 1 0 1 OOAB 96 101 78 84 78 OOAC 89 OOAD 89 93 93 OOAE 46 55 43 OOAF 106 107 113 # Printing first six rows of the expected counts head(expected) E2001 E2002 E2003 E2004 E2005 E2006 OOAA 2.648915 2.68106 2.727112 2.749562 2.808655 2.915977 OOAB 63.946730 63.41700 62.567863 61.444884 60.677119 59.678672 00AC 121.795213 121.91534 122.451050 123.201898 124.449563 125.982868 00AD 90.201336 91.24645 91.949050 92.754781 93.674540 94.598593 00AE 76.876437 77.18529 78.017980 78.967493 80.422828 81.785325 00AF 131.182934 132.30521 133.257442 134.520920 136.441229 137.382528 E2007 E2008 E2009 E2010 OOAA 3.021586 3.114696 3.237998 3.237998 OOAB 58.487583 57.701932 57.250524 57.250524 00AC 127.088805 128.825149 131.374946 131.374946 00AD 95.447131 96.832061 97.651369 97.651369 00AE 83.651266 85.265264 87.089119 87.089119 00AF 138.634021 139.508507 140.634084 140.634084

To familiarise ourselves with the data, we can summarise it using the summary() function. This will allow us to check for anomalies in our data.

Summarising the observed counts summary(observed)

	name	Y2001	Y200)2	Y2003
Adur CD	: 1	Min. : 2	.00 Min. :	0.00 Mi	n. : 3.00
Allerdale CD	: 1	1st Qu.: 35	.00 1st Qu.:	38.00 1s	t Qu.: 38.00
Amber Valley	CD: 1	Median : 50	.00 Median :	52.00 Me	dian : 52.00
Arun CD	: 1	Mean : 68	.01 Mean :	69.63 Me	an : 73.44
Ashfield CD	: 1	3rd Qu.: 83	. <mark>50</mark> 3rd Qu.:	80.75 3r	d Qu.: 83.25
Ashford CD	: 1	Max. :445	.00 Max. :	438.00 Ma	x. :480.00
(Other) :318					
Y2004		Y2005	Y2006	Y20	07
Min. : 1.0	DO Min.	: 1.00	Min. : 1.0	0 Min.	: 5.00
1st Qu.: 35.0	00 1st	Qu.: 37.00	1st Qu.: 35.0	00 1st Qu.	: 37.00
Median : 49.	50 Medi	an : 51.00	Median : 49.0	0 Median	: 50.00
Mean : 66.6	57 Mean	: 69.37	Mean : 67.0	7 Mean	: 68.17

3rd Qu.: 81.25	3rd Qu.: 80.50	3rd Qu.: 81.00	3rd Qu.: 79.00
Max. :428.00	Max. :395.00	Max. :428.00	Max. :456.00
Y2008	Y2009	Y2010	
Min. : 1.00	Min. : 0.00	Min. : 1.00	
1st Qu.: 37.00	1st Qu.: 36.00	1st Qu.: 38.00	
Median : 51.00	Median : 50.00	Median : 51.00	
Mean : 71.40	Mean : 67.04	Mean : 68.81	
3rd Qu.: 84.25	3rd Qu.: 78.00	3rd Qu.: 81.25	
Max. :463.00	Max. :394.00	Max. :441.00	

Summarising the expected counts

<pre>summary(expected)</pre>			
E2001	E2002	E2003	E2004
Min. : 2.649	Min. : 2.681	Min. : 2.727	Min. : 2.75
1st Qu.: 39.066	1st Qu.: 39.456	1st Qu.: 39.849	1st Qu.: 40.60
Median : 51.766	Median : 52.671	Median : 53.487	Median : 54.29
Mean : 62.944	Mean : 63.589	Mean : 64.139	Mean : 64.72
3rd Qu.: 74.292	3rd Qu.: 74.974	3rd Qu.: 74.701	3rd Qu.: 74.02
Max. :370.913	Max. :371.271	Max. :369.861	Max. :368.87
E2005	E2006	E2007	E2008
Min. : 2.809	Min. : 2.916	Min. : 3.022	Min. : 3.115
1st Qu.: 41.646	1st Qu.: 42.497	1st Qu.: 43.203	1st Qu.: 44.262
Median : 54.765	Median : 55.506	Median : 56.552	Median : 57.522
Mean : 65.440	Mean : 66.180	Mean : 67.022	Mean : 67.950
3rd Qu.: 75.003	3rd Qu.: 75.260	3rd Qu.: 75.790	3rd Qu.: 76.935
Max. :368.565	Max. :367.838	Max. :368.026	Max. :368.291
E2009	E2010		
Min. : 3.238	Min. : 3.238		
1st Qu.: 45.062	1st Qu.: 45.062		
Median : 58.077	Median : 58.077		
Mean : 68.901	Mean : 68.901		
3rd Qu.: 78.166	3rd Qu.: 78.166		
Max. :368.940	Max. :368.940		

We can see that **observed** has the following variables:

- name Name of local authority,
- Y20XX Observed number of hospital admissions for COPD in the year 20XX.

We can see that **expected** has the following variables:

• Y20XX - Expected number of hospital admissions for COPD (calculated using indirect standardisation) in the year 20XX.

Activities

- Does it look like R has read in the data correctly?
- Are there any extreme values in our dataset?
- Can you find which local authorities have the smallest and largest observed counts in England in 2010? HINT: Use subset()

Modelling the Raw SMRs

Now that we have read in the data, we can calculate raw SMRs. Remember that

```
\mathrm{SMR} = \frac{\mathrm{observed}}{\mathrm{expected}}
```

```
# Calculating the raw SMRs
SMR_raw <- observed[ ,-1]/expected</pre>
```

To change attributes of a dataset such as the column names, we use the names() function.

It is important that we check that no errors have occurred at any stages, so we check by summarising the results using the head() and summary() functions.

<i># Printing first</i>	six rows of raw S	MRs	
<pre>head(SMR_raw)</pre>			
SMR2001	SMR2002 SMR2003	SMR2004 SMR20	005 SMR2006 SMR2007
00AA 0.7550261 0	.0000000 1.1000648	0.3636943 0.35604	123 0.3429382 1.6547601
OOAB 1.5638016 1	.5768644 1.9498828	1.5135516 2.24137	721 1.6253713 1.5558858
00AC 0.9031554 0	.8366462 0.8656520	0.7223915 0.79550	030 0.7699460 0.5665330
00AD 1.2084078 1	.2384043 1.2289415	1.0349871 1.20630	043 1.0253852 0.9848384
00AE 0.8975442 1	.1530694 0.8972291	0.7471429 0.75849	0.5869024 0.6335828
00AF 0.9147531 0	.9750183 1.0130766	0.9217897 0.93813	329 0.8516367 0.8655884
SMR2008	SMR2009 SMR2010		
00AA 0.3210586 0	.0000000 0.3088328		
00AB 1.6637225 1	.7641760 1.3624329		
00AC 0.6520466 0	.5937205 0.6774503		
00AD 0.9191171 0	.9523676 0.9523676		
00AE 0.5394928 0	.6315370 0.4937471		
00AF 0.7598103 0	.7608397 0.8035037		
# Summarising rat	w SMRs		
<pre>summary(SMR_raw)</pre>			
SMR2001	SMR2002	SMR2003	SMR2004
Min. :0.3883	Min. :0.0000	Min. :0.3616	Min. :0.2778
1st Qu.:0.7900	1st Qu.:0.8272	1st Qu.:0.8519	1st Qu.:0.7636
Median :0.9496	Median :1.0168	Median :1.0209	Median :0.9266
Mean :1.0349	Mean :1.0508	Mean :1.0895	Mean :0.9812
3rd Qu.:1.2526	3rd Qu.:1.2364	3rd Qu.:1.3071	3rd Qu.:1.1858
Max. :1.9861	Max. :2.2181	Max. :2.2483	Max. :1.9811
SMR2005	SMR2006	SMR2007	SMR2008
Min. :0.3326	Min. :0.3429	Min. :0.3509	Min. :0.3211
1st Qu.:0.7592	1st Qu.:0.7415	1st Qu.:0.7533	1st Qu.:0.7695
Median :0.9573	Median :0.9101	Median :0.9305	Median :0.9404
Mean :1.0126	Mean :0.9726	Mean :0.9743	Mean :1.0069
3rd Qu.:1.2083	3rd Qu.:1.1586	3rd Qu.:1.1679	3rd Qu.:1.1979
Max. :2.2414	Max. :2.0805	Max. :1.8528	Max. :2.0567
SMR2009	SMR2010		
Min. :0.0000	Min. :0.3088		
1st Qu.:0.7452	1st Qu.:0.7682		
Median :0.8777	Median :0.9337		
Mean :0.9328	Mean :0.9639		
3rd Qu.:1.0934	3rd Qu.:1.1335		
Max. :1.8507	Max. :2.3856		

Activities

- Does it look like the SMRs have been estimated correctly?
- Are there any strange values?

Mapping the Raw SMRs

To plot these SMRs to a map, we need to attach them to the shapefiles. The function combine.data.shapefile() allows us to combine shapefiles to plot later.

Now that the estimates are attached to the shapefile, the function **spplot()** allows us to create a map which colours the local authorities by the SMR estimate.



Activities

- Do you notice anything about this plot?
- Are there any extreme values?
- If so, do you believe that these are the truth or perhaps sampling error?

Modelling the smoothed SMRs

To calculate the smoothed SMRs, we first need to create a 'neighbourhood' structure. The functions poly2nb() and nb2mat() can be used to create this.

```
# Creates the neighbourhood
W.nb <- poly2nb(SMRspatial_raw, row.names = rownames(SMRspatial_raw))
# Creates a matrix for following function call
W.mat <- nb2mat(W.nb, style="B")</pre>
```

Here, we use 'first neighbours' to define our structure, so any local authority that shares a border with another are considered neighbours.

The function S.CARleroux() allows us to use this neighbourhood structure and performs a Bayesian analysis, to create a smoothed set of observed values as discussed in the lecture.

```
# Running smoothing model
model <- S.CARleroux(formula=observed$Y2010~offset(log(expected$E2010)), # Model Formula
    family="poisson", # Choosing Poisson Regression
    W=W.mat, # Neighbourhood matrix
    burnin=20000, # Number of burn in samples
    n.sample=100000, # Number of MCMC samples
    thin=10,
    fix.rho=TRUE,
    rho=1)</pre>
```

The new smoothed values can be extracted from the model output and divided by the expected values to allow comparison between the two methods.

```
# Creating a dataset with smoothed SMRs in 2010
SMR2010 <- model$fitted.values / expected$E2010
SMR_smooth <- as.data.frame(SMR2010, row.names = rownames(observed))</pre>
```

Again, we check that no errors have occurred, by summarising the results using the head() and summary() functions.

```
# Printing first six rows of smoothed SMRs
head(SMR_smooth)
       SMR2010
00AA 0.9969077
00AB 1.2598926
00AC 0.6855420
00AD 0.9729904
00AE 0.5986103
00AF 0.8603609
# Summarising smoothed SMRs
summary(SMR_smooth)
   SMR2010
Min.
       :0.5458
 1st Qu.:0.7944
Median :0.9229
Mean :0.9645
3rd Qu.:1.0810
Max. :1.7414
```

Activities

- Does it look like the SMRs have been estimated correctly?
- Are there any extreme values?

Mapping the Smoothed SMRs

Similarly to before, we attach the values of the smoothed SMRs to the shapefile using the combine.data.shapefile() function and create a map using the spplot() function.



Activities

- What do you notice about this new plot?
- Are there any extreme values?
- Are there any differences between the smoothed and raw estimates?

Repeat this analysis for another year to see if the results are similar. Carefully go through the previous sections and change any references from 2010 to any year that you wish between 2001–2009.