

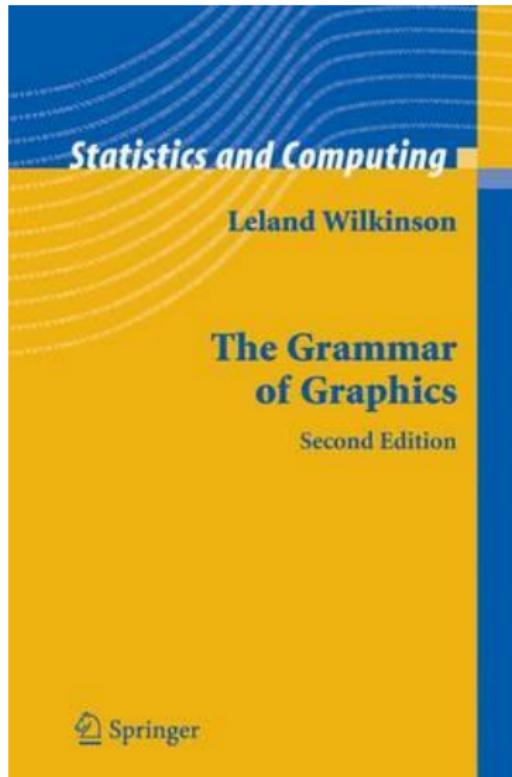
# ggplot2 and maps

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# The grammar of graphics

Hadley Wickham's **ggplot2** package implements the grammar of graphics described in Leland Wilkinson's book by the same title. It offers a very flexible and efficient way of generating plots based on data and is gaining more and more popularity.



# The ggplot() function

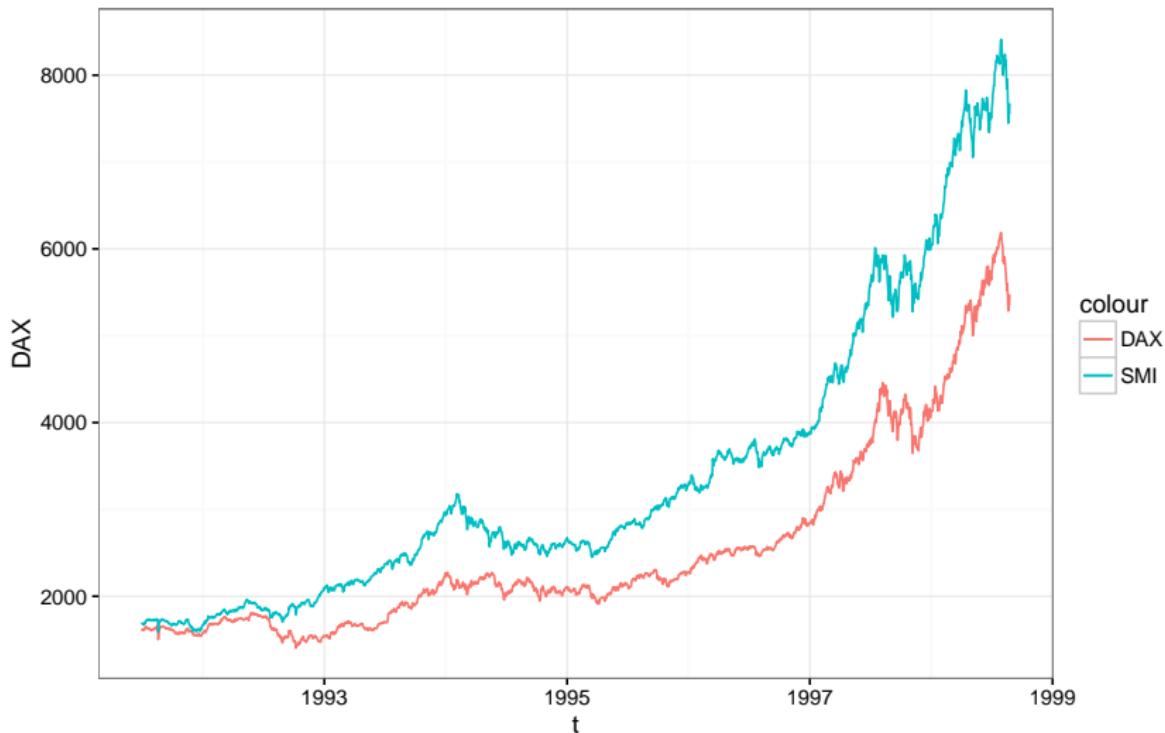
In the ggplot2 package, the default plotting function is called `ggplot()`. It is relatively easy to use. Let us see some examples:

```
library(ggplot2)
data(EuStockMarkets)
data.eu <- as.data.frame(EuStockMarkets)
t <- time(EuStockMarkets)
data.eu <- data.frame(t, data.eu)
stock.plot <- ggplot(data=data.eu, aes(x=t)) +
  geom_line(aes(y=DAX, col='DAX')) +
  geom_line(aes(y=SMI, col='SMI')) +
  theme_bw()
class(stock.plot)
```

```
## [1] "gg"      "ggplot"
```

# ggplot() – example plot 1

```
## Don't know how to automatically pick scale for object of
```



# The ggplot() function – example 1

```
summary(stock.plot)
```

```
## data: t, DAX, SMI, CAC, FTSE
##      [1860x5]
## mapping: x = t
## facetting: facet_null()
## -----
## mapping: y = DAX, colour = DAX
## geom_line: na.rm = FALSE
## stat_identity: na.rm = FALSE
## position_identity
##
## mapping: y = SMI, colour = SMI
## geom_line: na.rm = FALSE
## stat_identity: na.rm = FALSE
## position_identity
```

## ggplot() – another example

```
stock.plot <- ggplot(data=data.eu, aes(x=t, y=DAX)) +  
  geom_boxplot() +  
  geom_line() +  
  theme_bw()  
summary(stock.plot)
```

```
## data: t, DAX, SMI, CAC, FTSE  
## [1860x5]  
## mapping: x = t, y = DAX  
## facetting: facet_null()  
## -----  
## geom_boxplot: outlier.colour = NULL, outlier.shape = 19  
## stat_boxplot: na.rm = FALSE  
## position_dodge  
##  
## geom_line: na.rm = FALSE  
## stat_identity: na.rm = FALSE
```

# ggplot() – another example plot

```
## Don't know how to automatically pick scale for object of
```



# Visualising John Snow cholera data

On the 31 August 1854 a major outbreak of cholera occurred in London's SOHO. A physician, John Snow, put all reported deaths on a map of London and identified the focal point of the epidemics. It turned out, that the area has been supplied in water by a particular pump. Snow ordered the pump being closed and stopped the epidemics. He also provided an indirect proof that cholera is a waterborne disease. Let us try to recreate his work using modern tools.

# The data

First, we need to get Snow's original data in the digital form.  
Luckily, it can be obtained from, e.g.  
[\[http://blog.rtwilson.com/updated-snow-gis-data/\]](http://blog.rtwilson.com/updated-snow-gis-data/).

We will need a couple of packages to work with maps:

```
# an extension of ggplot2 for spatial data
# visualisations
library(ggmap)
# various tools, e.g. to convert between datums
library(maptools)
```

```
## Loading required package: sp
```

```
## Checking rgeos availability: TRUE
```

# The data cted.

```
library(sp)
# a Geospatial Data Abstraction Library,
# also useful for datum conversions etc.
library(rgdal)

## rgdal: version: 1.1-10, (SVN revision 622)
## Geospatial Data Abstraction Library extensions to R suc
## Loaded GDAL runtime: GDAL 1.11.4, released 2016/01/25
## Path to GDAL shared files: /Library/Frameworks/R.framework/Versions/3.3/Resources/library/rgdal/gdal
## Loaded PROJ.4 runtime: Rel. 4.9.1, 04 March 2015, [PJ_VERTCS]
## Path to PROJ.4 shared files: /Library/Frameworks/R.framework/Versions/3.3/Resources/library/rgdal/proj
## Linking to sp version: 1.2-3

# for Voronoi tessellation
library(deldir)

## deldir 0.1-12
```

# Reading the data

```
# download SOHO map from Google Maps  
google.london <- get_map(c(-.137,51.513), zoom=16)
```

```
## Map from URL : http://maps.googleapis.com/maps/api/stat
```

```
# and make it to a ggmap object
```

```
london <- ggmap(google.london)
```

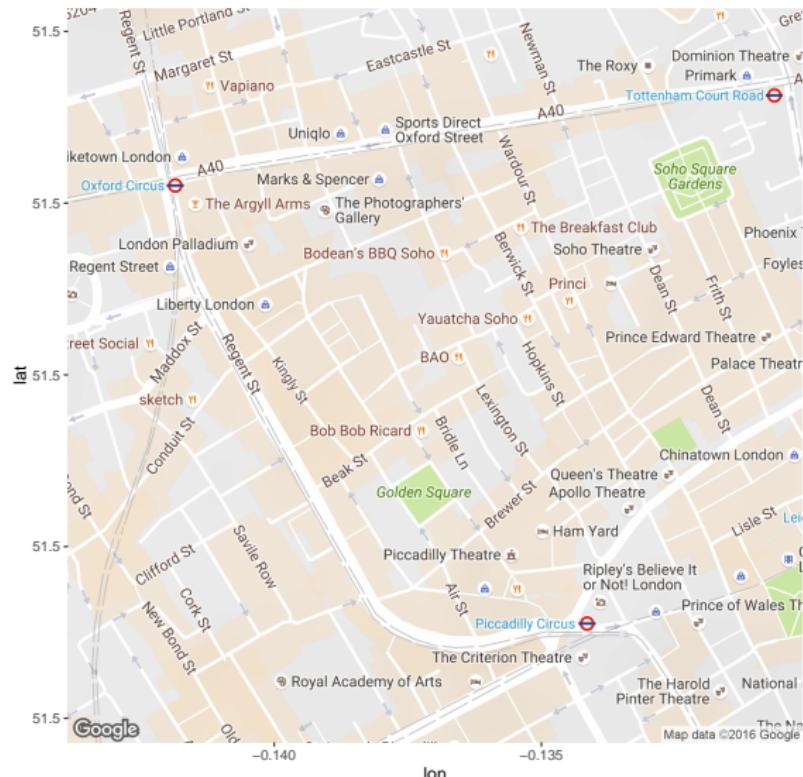
```
# now, read the downloaded Snow data
```

```
deaths <- readShapePoints("~/Dropbox/Rcourse/Labs/Lab - maps")
```

```
pumps <- readShapePoints("~/Dropbox/Rcourse/Labs/Lab - maps")
```

# Plot London

london



# Make a data frame for ggplot2

```
tmp.deaths <- data.frame(deaths@coords)
tmp.pumps <- data.frame(pumps@coords)
tmp <- rbind(tmp.deaths, tmp.pumps)
# we need a column telling ggplot if it
# is a death case or a pump
tmp$type <- c(rep('death', times=dim(tmp.deaths)[1]),
              rep('pump', times=dim(tmp.pumps)[1]))
```

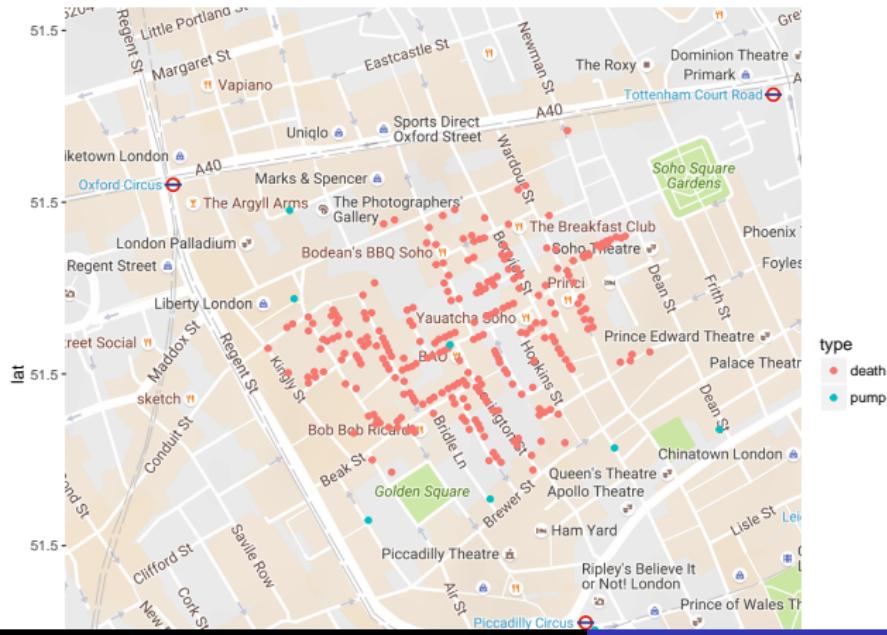
## Transform the datum

```
# Transform coordinates to WGS84 datum used by Google
# Check EPSG codes online

# create object of coordinates class
coordinates(tmp)=~coords.x1+coords.x2
# set the projection in the object
proj4string(tmp)=CRS("+init=epsg:27700")
# transform the projection to WGS84
tmp = spTransform(tmp, CRS("+proj=longlat +datum=WGS84"))
# adjust in the data frame
tmp <- data.frame(tmp@coords, type=tmp@data$type)
```

# Plot Snow's data

```
london +  
  geom_point(mapping=  
    aes(x=coords.x1, y=coords.x2, col=type),  
    data=tmp)
```



## Further analyses

Well, so far, so good, but it still does not give the answer to our question on where the cholera source is...

```
# do Voronoi tessellation
voronoi <- deldir(tmp[tmp$type=='pump',])

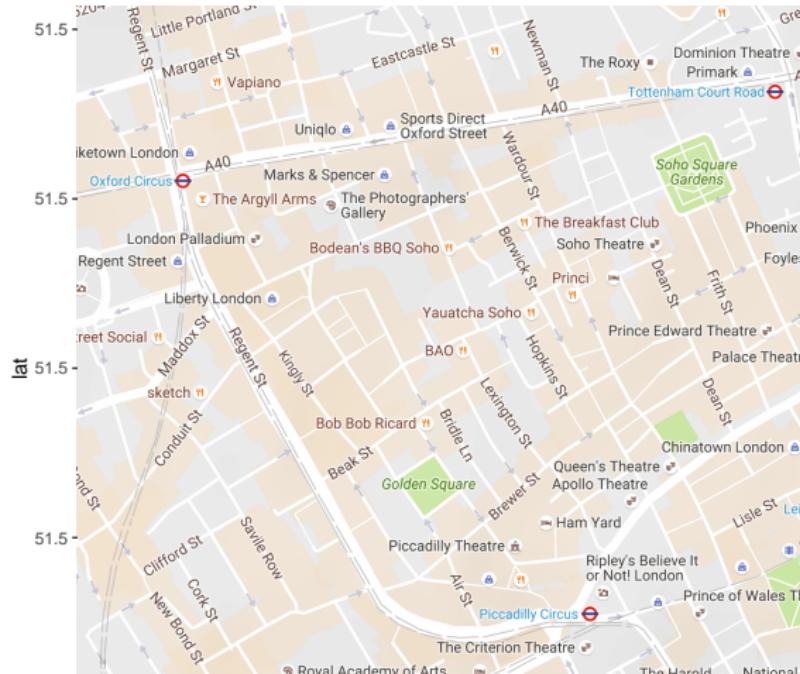
## PLEASE NOTE: The components "delsgs" and "summary"
## object returned by deldir() are now DATA FRAMES rather
## matrices (as they were prior to release 0.0-18).
## See help("deldir").

## PLEASE NOTE: The process that deldir() uses for det-
## duplicated points has changed from that used in version
## 0.0-9 of this package (and previously). See help("deldir")
```

# Step 1

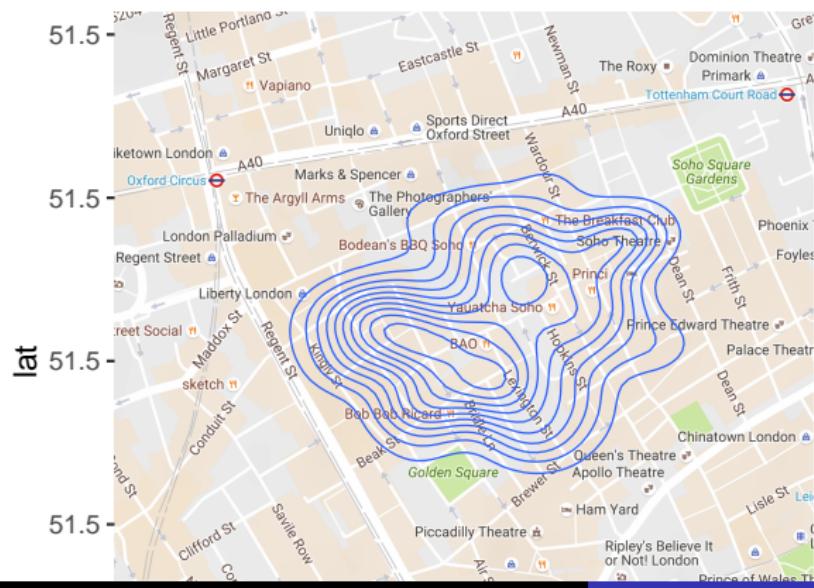
```
# plot SOHO
```

```
snow.plot <- london  
snow.plot
```



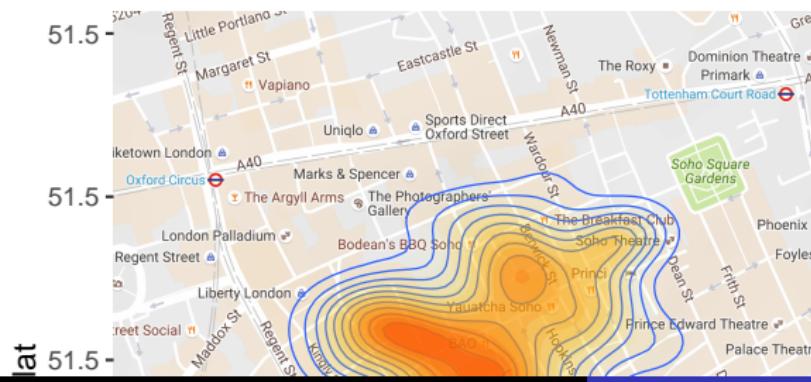
## Step 2

```
# plot death density lines
snow.plot <- snow.plot + geom_density2d(data = tmp[tmp$type == "death"], aes(x = coords.x1, y = coords.x2), size = 0.3)
snow.plot
```



# Step 3

```
# plot death gradient
snow.plot <- snow.plot + stat_density2d(data = tmp[tmp$type == "death"], aes(x = coords.x1, y = coords.x2, fill = ..level.., alpha = ..level..), size = 0.01, bins = 16, geom = "polygon") +
  scale_fill_gradient(low = "yellow",
    high = "red", guide = FALSE) +
  scale_alpha(range = c(0, 0.3), guide = FALSE)
snow.plot
```

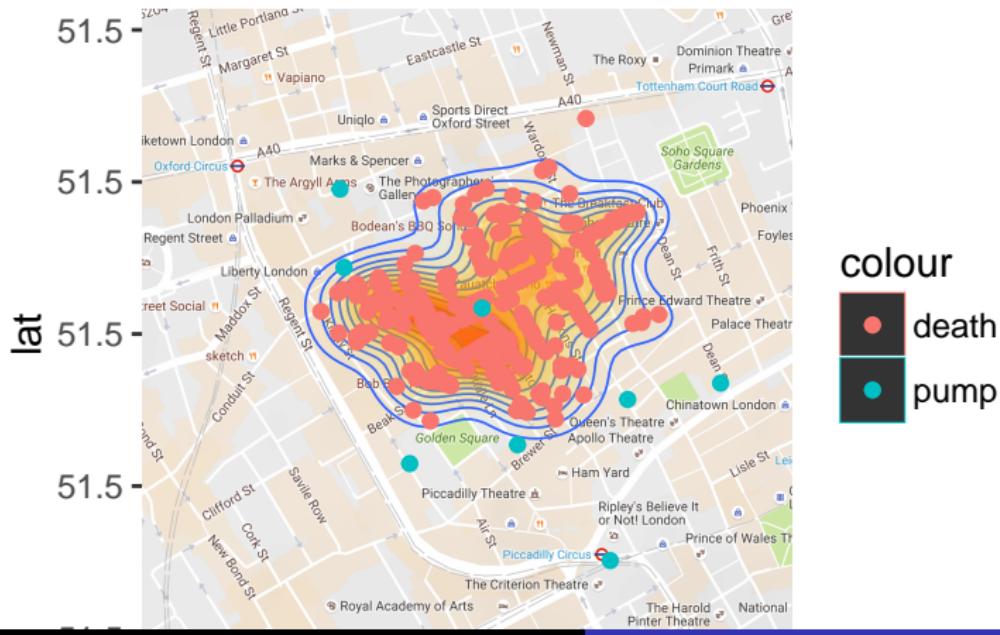


lat

# Step 4

```
# plot pumps and deaths
```

```
snow.plot <- snow.plot + geom_point(mapping = aes(x = coords.x1,  
y = coords.x2, col = type), data = tmp)  
snow.plot
```



# Step 5

```
# plot Voronoi tesselation
```

```
snow.plot <- snow.plot + geom_segment(aes(x = x1,  
y = y1, xend = x2, yend = y2), size = 0.7,  
data = voronoi$dirsgs, color = "grey")  
snow.plot
```

