# **Network Constrained Spatial Point Patterns Analy- sis**

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#### Overview

**Network constrained Spatial Point Patterns Analysis (NetSPAA)** is a collection of spatial point patterns analysis methods special developed for analysing spatial point event occurs on or alongside network. The spatial point event can be locations of traffic accident or childcare centre for example. The network, on the other hand can be a road network or river network.

In this hands-on exercise, you are going to gain hands-on experience on using appropriate functions of **spNetwork** package:

- to derive network constrained kernel density estimation (NetKDE), and
- to perform network G-function and k-function analysis

#### The Data

In this study, we will analyse the spatial distribution of childcare centre in Punggol planning area. For the purpose of this study, two geospatial data sets will be used. They are:

- *Punggol\_St*, a line features geospatial data which store the road network within Punggol Planning Area.
- *Punggol\_CC*, a point feature geospatial data which store the location of childcare centres within Punggol Planning Area.

Both data sets are in ESRI shapefile format.

## Installing and launching the R packages

In this hands-on exercise, four R packages will be used, they are:

- spNetwork, which provides functions to perform Spatial Point Patterns Analysis such as kernel density estimation (KDE) and K-function on network. It also can be used to build spatial matrices ('listw' objects like in 'spdep' package) to conduct any kind of traditional spatial analysis with spatial weights based on reticular distances.
- rgdal, which provides bindings to the 'Geospatial' Data Abstraction Library (GDAL) (>= 1.11.4) and access to projection/transformation operations from the PROJ library. In this exercise, rgdal will be used to import geospatial data in R and store as sp objects.
- sp, which provides classes and methods for dealing with spatial data in R. In this exercise, it will be used to manage SpatialPointsDataFrame and SpatiaLinesDataFrame, and for performing projection transformation.

• **tmap** which provides functions for plotting cartographic quality static point patterns maps or interactive maps by using leaflet API.

Use the code chunk below to install and launch the four R packages.

#### **Data Import and Preparation**

The code chunk below uses st\_read() of **sf** package to important Punggol\_St and Punggol\_CC geospatial data sets into RStudio as sf data frames.

```
Reading layer `Punggol_St' from data source
  `D:\tskam\IS415-GAA\In-class_Ex\In-class_Ex03\data\geospatial'
  using driver `ESRI Shapefile'
Simple feature collection with 2642 features and 2 fields
Geometry type: LINESTRING
Dimension: XY
Bounding box: xmin: 34038.56 ymin: 40941.11 xmax: 38882.85 ymax: 44801.27
Projected CRS: SVY21 / Singapore TM
```

```
Reading layer `Punggol_CC' from data source
  `D:\tskam\IS415-GAA\In-class_Ex\In-class_Ex03\data\geospatial'
  using driver `ESRI Shapefile'
Simple feature collection with 61 features and 1 field
Geometry type: POINT
Dimension: XYZ
Bounding box: xmin: 34423.98 ymin: 41503.6 xmax: 37619.47 ymax: 44685.77
z_range: zmin: 0 zmax: 0
Projected CRS: SVY21 / Singapore TM
```

We can examine the structure of the output SpatialDataFrame in RStudio. Alternative, code chunk below can be used to print the content of network SpatialLineDataFrame and childcare Spatial-PointsDataFrame by using the code chunk below.

```
str(network)
str(childcare)
```

When I exploring spNetwork's functions, it came to my attention that spNetwork is expecting the geospatial data contains complete CRS information.

In the code chunk below, *spTransform()* of **sp** package is used to assign EPSG code to the Spatial-DataFrames. The epsg:3414 is the code for svy21.

#### Visualising the Geospatial Data

Before we jump into the analysis, it is a good practice to visualise the geospatial data. There are at least two ways to visualise the geospatial data. One way is by using *plot()* of Base R as shown in the code chunk below.

```
plot(network)
plot(childcare,add=T,col='red',pch = 19)
```

LINK\_ID

ST\_NAME



To visualise the geospatial data with high cartographic quality and interactive manner, the mapping function of **tmap** package can be used as shown in the code chunk below.

```
tmap_mode('view')
tm_shape(childcare) +
  tm_dots() +
```

```
tm_shape(network) +
tm_lines()
```



tmap\_mode('plot')

## Network Constrained KDE (NetKDE) Analysis

In this section, we will perform NetKDE analysis by using appropriate functions provided in **sp-Network** package.

#### Preparing the lixels objects

Before computing NetKDE, the SpatialLines object need to be cut into lixels with a specified minimal distance. This task can be performed by using with <code>lixelize\_lines()</code> of spNetwork as shown in the code chunk below.

What can we learned from the code chunk above:

- The length of a lixel, lx length is set to 750m, and
- The minimum length of a lixel, *mindist* is set to 375m.

After cut, if the length of the final lixel is shorter than the minimum distance, then it is added to the previous lixel. If NULL, then mindist = maxdist/10. Also note that the segments that are already shorter than the minimum distance are not modified

**Note:** There is another function called *lixelize lines.mc()* which provide multicore support.

#### Generating line centre points

Next, *lines\_center()* of **spNetwork** will be used to generate a SpatialPointsDataFrame (i.e. samples) with line centre points as shown in the code chunk below.

```
samples <- lines_center(lixels)</pre>
```

The points are located at center of the line based on the length of the line.

#### **Performing NetKDE**

We are ready to computer the NetKDE by using the code chunk below.

```
densities <- nkde(network,
                  events = childcare,
                  w = rep(1, nrow(childcare)),
                  samples = samples,
                  kernel name = "quartic",
                  bw = 300,
                  div= "bw",
                  method = "simple",
                  digits = 1,
                  tol = 1,
                  grid\_shape = c(1,1),
                  max depth = 8,
                  agg = 5, #we aggregate events within a 5m radius (faster
calculation)
                  sparse = TRUE,
                  verbose = FALSE)
```

What can we learn from the code chunk above?

- kernel\_name argument indicates that **quartic** kernel is used. Are possible kernel methods supported by spNetwork are: triangle, gaussian, scaled gaussian, tricube, cosine ,triweight, epanechnikov or uniform.
- *method* argument indicates that **simple** method is used to calculate the NKDE. Currently, sp-Network support three popular methods, they are:
  - method="simple". This first method was presented by Xie et al. (2008) and proposes an intuitive solution. The distances between events and sampling points are replaced by network

distances, and the formula of the kernel is adapted to calculate the density over a linear unit instead of an areal unit.

- method="discontinuous". The method is proposed by Okabe et al (2008), which equally "divides" the mass density of an event at intersections of lixels.
- method="continuous". If the discontinuous method is unbiased, it leads to a discontinuous kernel function which is a bit counter-intuitive. Okabe et al (2008) proposed another version of the kernel, that divide the mass of the density at intersection but adjusts the density before the intersection to make the function continuous.

The user guide of **spNetwork** package provide a comprehensive discussion of *nkde()*. You should read them at least once to have a basic understanding of the various parameters that can be used to calibrate the NetKDE model.

#### **Visualising NetKDE**

Before we can visualise the NetKDE values, code chunk below will be used to insert the computed density values (i.e. densities) into *samples* and *lixels* objects as *density* field.

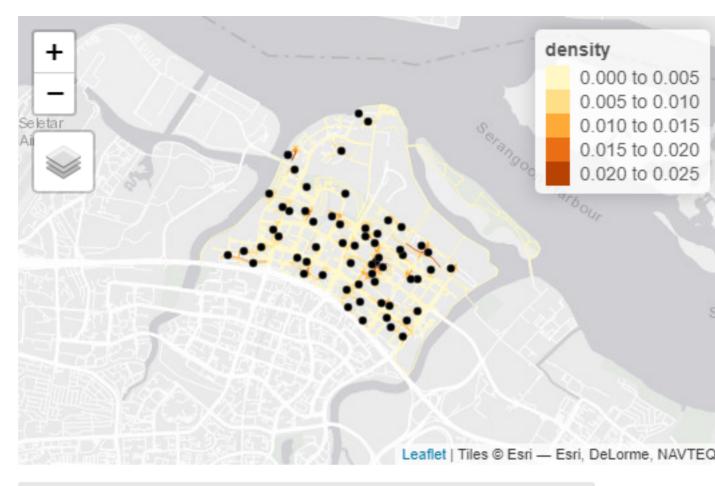
```
samples$density <- densities
lixels$density <- densities</pre>
```

Since svy21 projection system is in meter, the computed density values are very small i.e. 0.0000005. The code chunk below is used to resale the density values from number of events per meter to number of events per kilometer.

```
# rescaling to help the mapping
samples$density <- samples$density*1000
lixels$density <- lixels$density*1000</pre>
```

The code below uses appropriate functions of tmap package to prepare interactive and high cartographic quality map visualisation.

```
tmap_mode('view')
tm_shape(lixels)+
  tm_lines(col="density")+
tm_shape(childcare)+
  tm_dots()
```



```
tmap_mode('plot')
```

The interactive map above effectively reveals road segments (darker color) with relatively higher density of childcare centres than road segments with relatively lower density of childcare centres (lighter color)

### **Temporal Network KDE**

#### Importing the data

```
data(bike_accidents)
```

## **Data Preparation**

#### Converting the Date field to a numeric field (counting days)

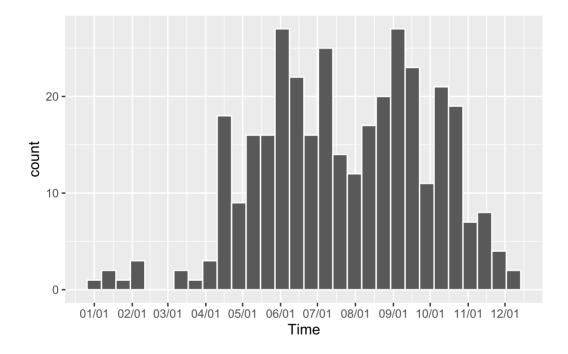
```
bike_accidents$Time <- as.POSIXct(bike_accidents$Date, format = "%Y/%m/%d")
start <- as.POSIXct("2016/01/01", format = "%Y/%m/%d")
bike_accidents$Time <- difftime(bike_accidents$Time, start, units = "days")</pre>
```

```
bike_accidents$Time <- as.numeric(bike_accidents$Time)

months <- as.character(1:12)
months <- ifelse(nchar(months)==1, paste0("0", months), months)
months_starts_labs <- paste("2016/",months,"/01", sep = "")
months_starts_num <- as.POSIXct(months_starts_labs, format = "%Y/%m/%d")
months_starts_num <- difftime(months_starts_num, start, units = "days")
months_starts_num <- as.numeric(months_starts_num)
months_starts_labs <- gsub("2016/", "", months_starts_labs, fixed = TRUE)</pre>
```

#### Visualising the temporal patterns

```
ggplot(bike_accidents) +
  geom_histogram(aes(x = Time), bins = 30, color = "white") +
  scale_x_continuous(breaks = months_starts_num, labels = months_starts_labs)
```



```
bike_accidents <- subset(bike_accidents, bike_accidents$Time >= 90)
```

We can now calculate the kernel density values in time for several bandwidths.

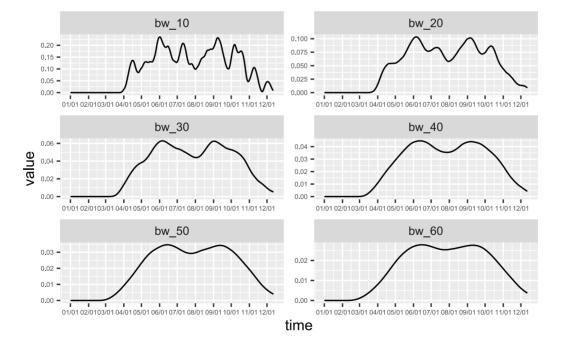
```
w <- rep(1,nrow(bike_accidents))
samples <- seq(0, max(bike_accidents$Time), 0.5)

time_kernel_values <- data.frame(
  bw_10 = tkde(bike_accidents$Time, w = w, samples = samples, bw = 10, kernel_name)</pre>
```

```
= "quartic"),
  bw_20 = tkde(bike_accidents$Time, w = w, samples = samples, bw = 20, kernel_name
= "quartic"),
  bw_30 = tkde(bike_accidents$Time, w = w, samples = samples, bw = 30, kernel_name
= "quartic"),
  bw_40 = tkde(bike_accidents$Time, w = w, samples = samples, bw = 40, kernel_name
= "quartic"),
  bw_50 = tkde(bike_accidents$Time, w = w, samples = samples, bw = 50, kernel_name
= "quartic"),
  bw_60 = tkde(bike_accidents$Time, w = w, samples = samples, bw = 60, kernel_name)
= "quartic"),
  time = samples
)
```

```
df_time <- reshape2::melt(time_kernel_values,id.vars = "time")
df_time$variable <- as.factor(df_time$variable)</pre>
```

```
ggplot(data = df_time) +
  geom_line(aes(x = time, y = value)) +
  scale_x_continuous(breaks = months_starts_num, labels = months_starts_labs) +
  facet_wrap(vars(variable), ncol=2, scales = "free") +
  theme(axis.text = element_text(size = 5))
```



It seems that a bandwidth between 30 and 40 days capture the bimodal shape of the temporal dimension of bike accidents. The lower densities during July and August are likely caused by the lower traffic due to holidays.

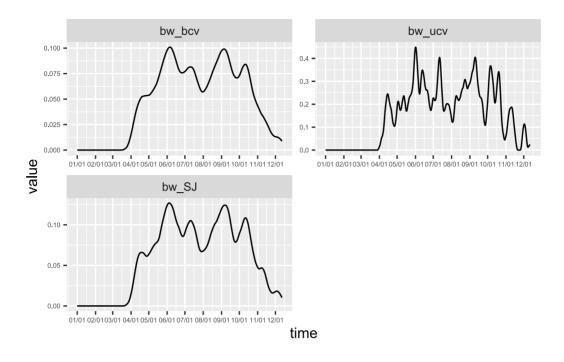
It is also possible to use the classical functions from R to find a bandwidth with a data-driven approach.

```
bw1 <- bw.bcv(bike_accidents$Time, nb = 1000, lower = 1, upper = 80)
bw2 <- bw.ucv(bike_accidents$Time, nb = 1000, lower = 1, upper = 80)
bw3 <- bw.SJ(bike_accidents$Time, nb = 1000, lower = 1, upper = 80)</pre>
```

```
time_kernel_values <- data.frame(
    bw_bcv = tkde(bike_accidents$Time, w = w, samples = samples, bw = bw1,
kernel_name = "quartic"),
    bw_ucv = tkde(bike_accidents$Time, w = w, samples = samples, bw = bw2,
kernel_name = "quartic"),
    bw_SJ = tkde(bike_accidents$Time, w = w, samples = samples, bw = bw3, kernel_name
= "quartic"),
    time = samples
)</pre>
```

```
df_time <- reshape2::melt(time_kernel_values,id.vars = "time")
df_time$variable <- as.factor(df_time$variable)</pre>
```

```
ggplot(data = df_time) +
  geom_line(aes(x = time, y = value)) +
  scale_x_continuous(breaks = months_starts_num, labels = months_starts_labs) +
  facet_wrap(vars(variable), ncol=2, scales = "free") +
  theme(axis.text = element_text(size = 5))
```



## **Preparing and Exploring Spatial Point Data**

```
tm_shape(mtl_network) +
  tm_lines(col = "black") +
  tm_shape(bike_accidents) +
  tm_dots(col = "red", size = 0.1)
```



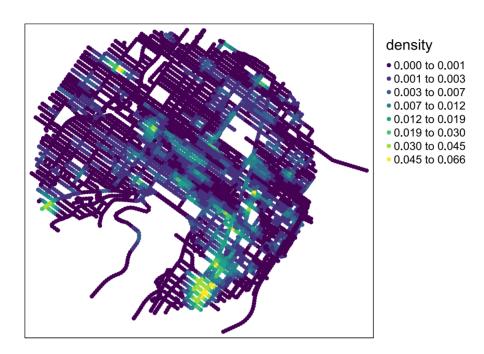
#### Creating sample points

```
lixels <- lixelize_lines(mtl_network, 50)
sample_points <- lines_center(lixels)</pre>
```

#### Calculating the densities

#### Visualising the NKDE

```
tm_shape(sample_points) +
  tm_dots(col = "density", style = "kmeans", n = 8, palette = "viridis", size =
0.05) +
  tm_layout(legend.outside = TRUE)
```



## **Spatio-temporal KDE**

```
cv_scores <- bws_tnkde_cv_likelihood_calc(</pre>
 bw net range = c(200, 1100),
 bw net step = 100,
 bw_time_range = c(10,70),
 bw time step = 10,
 lines = mtl_network,
 events = bike_accidents,
 time_field = "Time",
 w = rep(1, nrow(bike_accidents)),
 kernel_name = "quartic",
 method = "discontinuous",
 diggle correction = FALSE,
 study_area = NULL,
 max_depth = 10,
 digits = 2,
 tol = 0.1,
 agg = 10,
 sparse=TRUE,
 grid_shape=c(1,1),
```

```
sub_sample=1,
verbose = FALSE,
check = TRUE)
```

#### Choosing sample in times (every 10 days)

```
sample_time <- seq(0, max(bike_accidents$Time), 10)</pre>
```

#### Calculating densities

```
tnkde densities <- tnkde(lines = mtl network,</pre>
                   events = bike accidents,
                   time_field = "Time",
                   w = rep(1, nrow(bike accidents)),
                   samples loc = sample points,
                   samples_time = sample_time,
                   kernel name = "quartic",
                   bw_net = 700, bw_time = 60,
                   adaptive = TRUE,
                   trim_bw_net = 900,
                   trim bw time = 80,
                   method = "discontinuous",
                   div = "bw", max depth = 10,
                   digits = 2, tol = 0.01,
                   agg = 15, grid\_shape = c(1,1),
                   verbose = FALSE)
```

#### Creating a color palette for all the densities

```
all_densities <- c(tnkde_densities$k)
color_breaks <- classIntervals(all_densities, n = 10, style = "kmeans")</pre>
```

#### generating a map at each sample time

## Creating a gif with all the maps

```
tmap_animation(all_maps, filename = "images/animated_map.gif",
               width = 1000, height = 1000, dpi = 300, delay = 50)
Creating frames
===
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```

===
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==
<del></del>
==

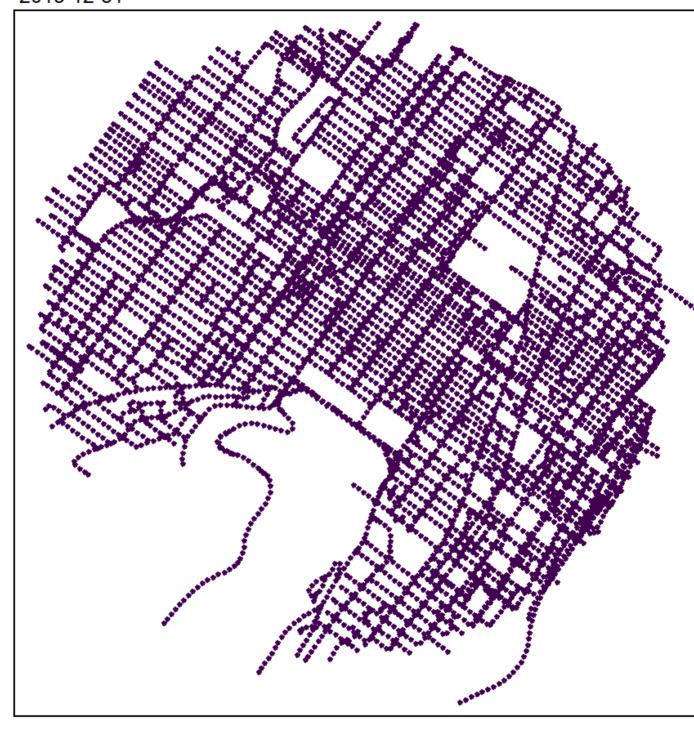
```
==

Creating animation
Animation saved to D:\tskam\IS415-GAA\In-class_Ex\In-class_Ex\In-class_Ex03\images\animated_map.gif
```

## Displaying the animated TNKDE

knitr::include\_graphics("images/animated\_map.gif")

## 2015-12-31



## References

- sp Network: Spatial Analysis on Network
- Network Kernel Density Estimate
- Details about NKDE
- Network k Functions