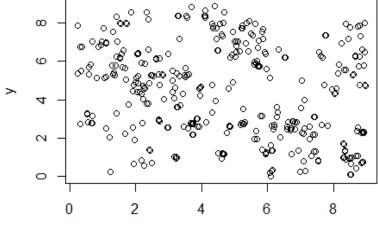
Steve Harms April 22, 2018

```
2)
```

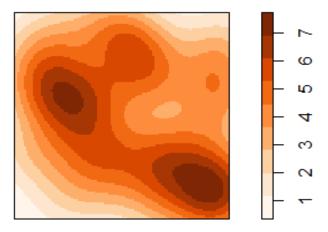
a)

```
canes <- ppp(x, y, window=owin(c(0,9),c(0,9)))
#a plot of the data
plot(x = x, y = y)</pre>
```



#plot of the density using Diggle's Bandwidth
image(density(canes, diggle=T), main = "Estimated Intensity", col = brewer.pal(n = 9, name = "Oranges"))

Estimated Intensity

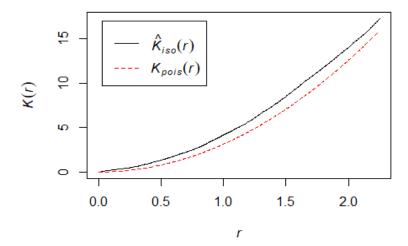


It appears there may be some clustering based on the plot of the intensity, but the plot of the original data is inconclusive.

```
b)
```

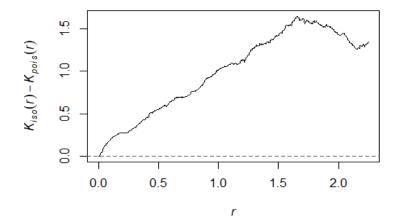
```
#K-function
Kc <- Kest(canes, correction="Ripley")
plot(Kc, main="Estimated K function")</pre>
```

Estimated K function



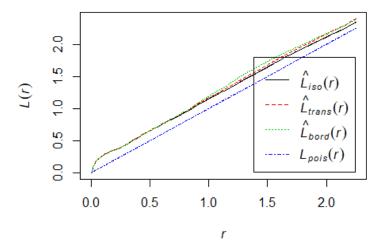
plot(Kc, iso - theo ~ r, main="Deviation from CSR (K-function)")
abline(h=0, lty=2, col="gray40")

Deviation from CSR (K-function)



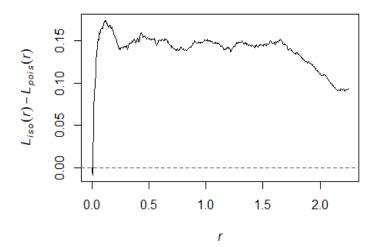
#L-function
Lc <- Lest(canes)
plot(Lc, main = "Estimated L Function")</pre>

Estimated L Function



```
plot(Lc, iso - theo ~ r, main="Deviation from CSR (L-Function)")
abline(h=0, lty=2, col="gray40")
```

Deviation from CSR (L-Function)

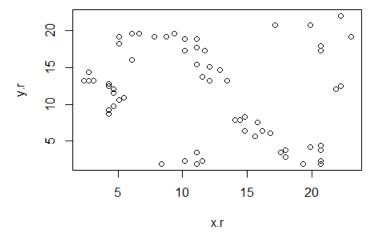


Both the K- and L- functions have positive values near 0, indicating that the plot is clustered. However, note that the deviation from CSR for the L- function is actually less than 0 for very small distance from origin, which indicates that this process may not actually be clustered.

3)

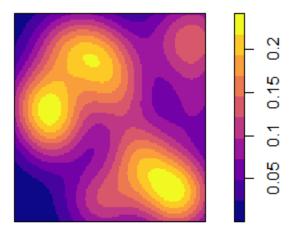
a)

```
trees <- ppp(x.r, y.r, window=owin(c(0,23),c(0,23)))
#a plot of the data
plot(x = x.r, y = y.r)</pre>
```



```
#plot of the density
plot(density(trees, diggle=T), main = "Estimated Intensity", col = plasma(10))
```

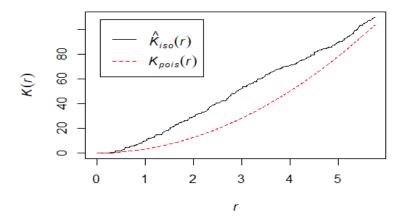
Estimated Intensity



b)

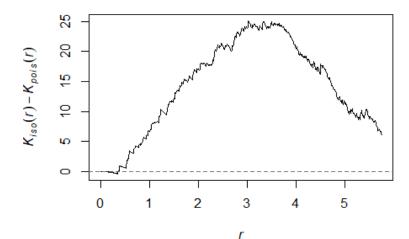
#K-function
Ks <- Kest(trees, correction="Ripley")
plot(Ks, main="Estimated K function")</pre>

Estimated K function



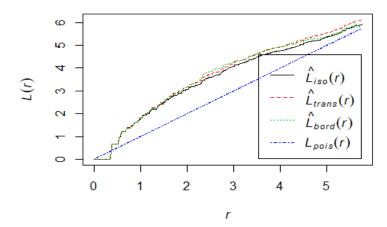
plot(Ks, iso - theo ~ r, main="Deviation from CSR (K)")
abline(h=0, lty=2, col="gray40")

Deviation from CSR (K)



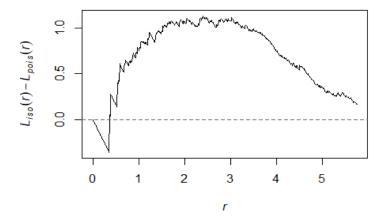
```
#L-function
Ls <- Lest(trees)
plot(Ls, main = "Estimated L Function")</pre>
```

Estimated L Function



plot(Ls, iso - theo ~ r, main="Deviation from CSR (L)")
abline(h=0, lty=2, col="gray40")

Deviation from CSR (L)



While our intuition suggests that this process should be clustered around trees, the K- and L-functions and actually negative near 0, indicating that this process is actually somewhat regular/unclustered.