Geolocation of mobile devices

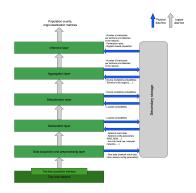
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Context

The European project ESSnet on Big Data II

- A modular statistical process was designed
- It goes from the raw telco data to the final target population estimates
- A set of R packages were developed to carry out the computations that this process requires
- destim package was developed for the geolocation layer



destim is just a Hidden Markov Model estimation package. While some existing HMM estimation could have been used . . .

- The transition matrix is expected to be highly sparse. However, many HMM estimation packages are designed for dense transition matrices.

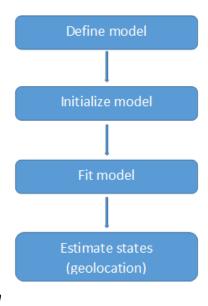
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- *destim* is optimized for one specific kind of constraints: the equalities between two transition probabilities (*isotropy*). This is too specific and we don't know any other package with this characteristic.
- In general, it is an unusual HMM estimator, because it is designed for a high number of states but a small number of parameters.

How destim works

- The initialization finds transition probabilities that satisfy the constraints, and also computes a QR decomposition to find a minimal set of parameters.
- Rcpp and RcppEigen are used to improve performance, specially for sparse linear algebra computations.
- For likelihood optimization, constrained optimization packages are used: solnp and donlp2. It is also possible to use constrOptim from package stats.



Basic usage (I)

```
model <- HMM(5)
model <- addtransition(model,c(1,2))</pre>
model <- addtransition(model,c(2,3))</pre>
model <- addtransition(model,c(3,4))</pre>
model <- addtransition(model,c(4,5))</pre>
model <- addtransition(model,c(5,1))</pre>
model <- addconstraint(model,c(2,4))</pre>
emissions(model)<-matrix(c(0.3, 0.3, 0.7, 0.9, 0.9,
                              0.7, 0.7, 0.3, 0.1, 0.1),
                            nrow = 5, ncol = 2)
model <- initparams(model)</pre>
model <- minparams(model)</pre>
logLik(model,c(1L,2L))
## [1] 2.056399
model <- fit(model, c(1L, 2L), init = T, method = "constrOptim")</pre>
logLik(model,c(1L,2L))
## [1] 1.213004
```

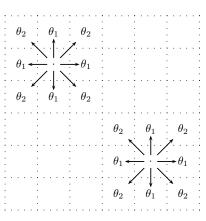
Basic usage (II)

```
sstates(model, c(1L, 2L))
## 5 x 2 sparse Matrix of class "dgCMatrix"
##
## [1,] 0.07884862 0.46667993
## [2,] 0.09784651 0.07884862
## [3,] 0.22716072 0.26821705
## [4,] 0.06279566 0.08818847
## [5,] 0.53334849 0.09806593
scpstates(model, c(1L,2L))
## 5 x 5 sparse Matrix of class "dgCMatrix"
##
## [1,] 9.685594e-10 7.884862e-02 .
## [2,] .
                    2.804493e-09 0.09784651 .
## [3,] .
                          0.17037054 0.05679018 .
## [4.] .
                                             0.03139828 0.03139737
## [5,] 4.666799e-01.
                                                        0.06666856
```

The *rectangle* model

- An isotropic model for a square grid.
- No matter how big the map is, it only has two parameters.
- It is an easy model, good for testing.

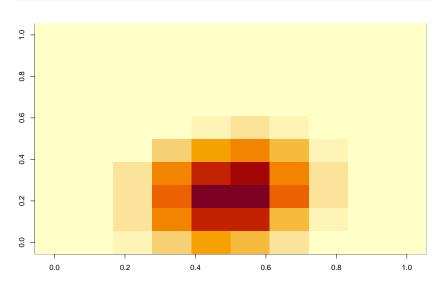
```
model <- HMMrectangle(100,100)
nstates(model)
## [1] 10000
ntransitions(model)
## [1] 88804
nconstraints(model)
## [1] 88802</pre>
```



Example with the rectangle model (I)

Example with the rectangle model (II)

image(matrix(sstates(model, obs)[,4], ncol = 10))



Some remarks about computacional efficiency

- The package is expected to scale well for map size, because sparsity. Very long gaps between detection events might be problematic.
- A desktop computer can carry out the computations for a few devices in a small country. RAM size would still be a limitation though.
- As the number of devices to estimate is expected to be huge, it is an embarrassingly parallel problem.
- Keeping a small number of parameters makes also easier the task of the optimizer.

Thank you!