© SoDA Supercomputing with R part 2 Agent-based model of all neighbourhoods in the Netherlands

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About me



Background

• PhD in **Statistics** (UU)

Structural Equation Models, high-dimensional data, regularization & penalization, algorithms & optimization

- Assistant professor at Methodology & Statistics, UU Human Data Science group, teaching Data Science master courses
- Team lead for the **ODISSEI Social Data Science (SoDa) team** Advancing data- & computation-intensive research in social science

https://odissei-soda.nl/



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We help social scientists with data intensive & computational research

Our goal is to enhance the evidence base and impact of social science by bringing the added value of new data sources and new data analysis techniques into social research in the Netherlands



About me



Relevant experience

• Experience with **parallel programming, supercomputing, large simulations**

statistics, social sciences, a bit of neuroscience (structural MRI), and a bit of bioinformatics (microarrays, epigenetics)

- Native in **R**, capable in **Python** dabbled in C++, C#, web languages, Julia, and more
- Many research **consultations**
- Strongly advocating for **open science** Make everything available all the time!
- Almost no experience with agent-based models!!!

About you

Write down in one short sentence why you are here / what you hope to learn

The remainder of today How to structure R projects for running analyses on a SURF supercomputer

The remainder of today

Time	Title
11:00	Lecture: computational limits in social science
11:45	Hands-on: a parallel agent-based model in R
12:30	Break
13:30	Lecture: supercomputing with R
14:15	Hands-on: submitting an R array job
15:15	Break
15:30	Lecture: combining & analysing the results
16:15	Conclusion & Q&A

Hands-on 1

Hands-on 2

Lecture 3





Computational limits in social science

Experimental research in soc. sci.

- Come up with research question
- Design experiment
- Run experiment
- Analyze results (perform statistical test)
- Make inferences about found effect

Observational research in soc. sci.

- Come up with research question
- Collect data
- Create statistical model
- Make inferences about model (pay attention to assumptions)

Computational research in soc. sci.?

- Come up with research question
- Create generative / computational model
- Generate data from computational model
- (compare computational model data with real data)
- Make conclusions about computational model (pay attention to assumptions)

Psych trend: theory construction

A *formal model* captures the principles of the explanatory theory in a set of equations or rules (as *implemented in a computer program or simulation*).

The theorist can then examine whether the theory, as implemented in the formal model, does in fact **generate the phenomena** as a matter of course, either **in a simulation** study or through analytic derivations.

Borsboom et al. (2021), Theory Construction Methodology: A Practical Framework for Building Theories in Psychology <u>doi.org/10.1177/1745691620969647</u>

A COMPUTATIONAL MODEL OF PANIC



Figure 4. Individual Differences in Vulnerability to Panic Attacks. We simulated perturbations to arousal of varying strength (inducing arousal of .25, .50, and .75) in three conditions: low, medium, and high arousal schema (*S*=.25, .50, and .75, respectively). To

Robinaugh et al. (2019), A Computational Model of Panic Disorder <u>10.31234/osf.io/km37w</u>

71

With more parameter settings?



(this is not real, just for illustration)

- Used in economics, sociology, ecology, finance, spatial planning, social psychology, and more
- Create agents who interact in an environment
- Each agent has **rules** based on theory
- Simulating the system means applying these rules repeatedly
- Then you can investigate the system



```
child_a ← function(sand) sand + runif(1, 0, 0.1)
child_b ← function(sand) if (sand > 0.95) runif(1, 0.05, 0.2) else sand
sand_vec ← numeric(100)
sand_vec[1] ← 0.5
for (i in 2:100) {
   sand_vec[i] ← child_a(sand_vec[i-1])
   sand_vec[i] ← child_b(sand_vec[i])
}
plot(sand_vec, type = "l")
```



Time

Interim conclusion

- Computational methods used by social scientists to formalize & investigate theories
- 2. Simulation from computational models to inspect phenomena following from model
- 3. Do this for different parameter settings
- 4. (2) and (3) may take a long time -> computational limits reached!

abm • • • •

Schelling's model of segregation

• Famous example of ABM in social behaviour

- What are the causes of de facto segregation in society?
- Theoretical / formal model of population dynamics
- Implemented as an agent-based model
- Conclusions drawn based on phenomena resulting from this model

- Environment: two-dimensional grid
- Agents belong to one of two groups
- Agents want to live close to others like themselves
 - Agents have preference (B_a) for the proportion of neighbours like them (B)
- If $B < B_a$, then move to random free location on grid
- Else stay

in-group preference
Ba ← 0.5

```
# initialize population
pop ← init_population(c(0.5, 0.5))
```

```
# occupation matrix
M ← matrix(data = pop, nrow = N)
```

```
# happiness matrix
H ← matrix(data = FALSE, nrow = N, ncol = N)
```

```
# run for 50 iterations
for (i in 1:50) {
    H ← compute_happiness(M, Ba)
    M ← move_agents(M, H)
```











 Micro behaviour: happy or unhappy with current location -> move or stay

• Macro phenomenon: how does the distribution of agents over the grid look?

- Schelling's finding: for groups of equal size with $B_a \gtrsim 0.33$, the system is likely to end in a **segregated** state
- Below that, the system will stay in a **mixed** / random state

Conclusion

Even if there is only a mild in-group preference, the world might still end up very segregated!

(keep assumptions in mind ⁽ⁱⁱⁱ⁾)

- Note: **randomness** in initialization & in movement to different locations
- At which *B_a* will the system segregate?
- Need to run this model many times for different *B_a* and compute expectation (average over the iterations)
- Monte carlo simulation

Some more parameters you may want to vary:

- Number of distinct populations
- Relative population sizes
- Number of free spots in the grid
- Neighbour preference
- Radius for looking at neighbours
- Other extensions...



This will take a long time

Speeding up the ABM

Two options

faster

Ē

Write faster, optimized code parallel

Run multiple ABMs at the same time









Speeding up slow code

- There isn't one solution for all types of code
- Speeding up slow code takes time
- Investigate smarter algorithms for your problem!
- If you are rewriting your R code, use vectorized & matrix operations where possible (faster than loops!)
- Use benchmarking to check the speed & memory usage of your functions (I like bench::mark())

Speeding up slow code

- Another step: rewrite in C++
- Depending on problem, this may yield great speedup

• In R: **Rcpp** package helps with this

sourceCpp("my_cpp_function.cpp")

An example of Rcpp speedup is in the hands-on later

# A tibble: 2 x 13												
	expression	min	median	`itr/sec`	mem_alloc	`gc/sec`	n_itr	n_gc	total_time	result		
	<bch:expr></bch:expr>	<bch:tm></bch:tm>	<bch:t></bch:t>	<dbl></dbl>	<bch:byt></bch:byt>	<dbl></dbl>	<int></int>	<dbl></dbl>	<bch:tm></bch:tm>	<list></list>		
1	r	724.2 ms	724.2ms	1.38	4.76MB	5.52	1	4	724 ms	<null></null>		
2	срр	65.6ms	71.2ms	13.8	41.42MB	11.9	7	6	506ms	<null></null>		
#	with 3	more var	iables: n	nemory <lis< td=""><td>st>, time <</td><td><list>, g</list></td><td>c <list< td=""><td>:></td><td></td><td></td></list<></td></lis<>	st>, time <	<list>, g</list>	c <list< td=""><td>:></td><td></td><td></td></list<>	:>				

Parallel programming

• Many problems are of the "embarrassingly parallel" type Little to no effort required to separate problem into number of parallel tasks

- ABM itself is **not** embarrassingly parallel: time step 3 requires results from time step 2!
- Running the whole ABM several times to average over uncertainty **is** embarrassingly parallel

Parallel programming

- Computers nowadays can do more than one task at a time: threads
 - Often: 4 or 8 threads
 - Bigger computers have 12, 16 or 32 threads
 - Depending on computer, potential speedup of 32 times! (remember that our Rcpp effort gave ~10 times)
- Parallel programming is built into R (package parallel)

library(parallel)

```
# create a function to run in parallel
my_func ← function(i) sprintf("this is iteration %i", i)
```

```
# instantiate 8 workers
clus ← makeCluster(8)
```

```
# run the function for iteration 1:100
parSapply(clus, 1:100, my_func)
```

```
#> [1] "this is iteration 1" "this is iteration 2" "this is iteration 3"
#> [4] "this is iteration 4" "this is iteration 5" "this is iteration 6"
#> [7] "this is iteration 7" "this is iteration 8" "this is iteration 9"
#> [10] "this is iteration 10" "this is iteration 11" "this is iteration 12"
#> [13] "this is iteration 13" "this is iteration 14" "this is iteration 15"
#> [16] "this is iteration 16" "this is iteration 17" "this is iteration 18"
#> [19] "this is iteration 19" "this is iteration 20" "this is iteration 21"
```

stop the workers (we're done with them now)
stopCluster(clus)

An example of parallel programming is in the hands-on later

Interim conclusion

- 1. Today we are working with the Schelling agent-based model
- 2. Running the abm with different settings takes a long time
- 3. We can program the abm itself more efficiently
- 4. We can perform the abm in parallel

Let's try it out!

Hands-on session 1