

A Thread-Based Parallel Implementation of the Open Source Framework MATSim Episim

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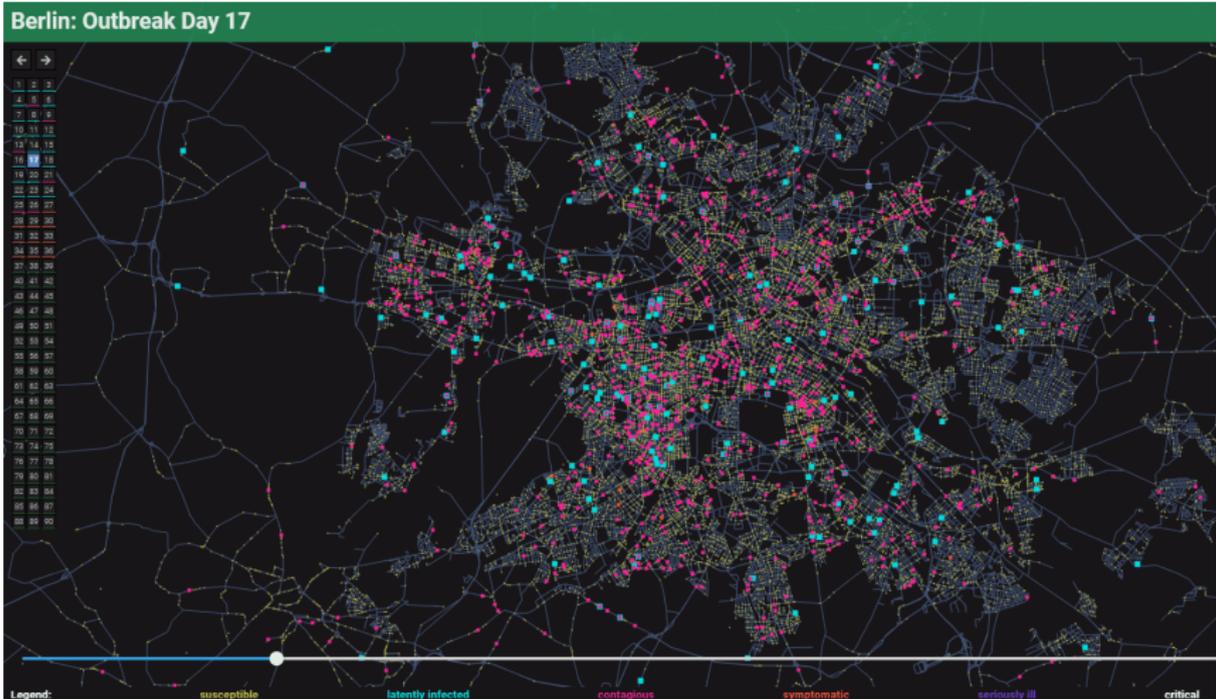
June 21, 2021

MATSim is an open-source framework for implementing large-scale agent-based transport simulations.



Source: [1]

MATSim Episim is an epidemic simulation based on MATSim.



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Where based means:

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But apart from that, Episim is independent single-threaded Java code.

The following elements are in nearly every agent-based epidemiological simulation:

Synthetic population

id	age	gender	household-id
1324019	42	f	382741
1324020	29	m	395832

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Locations

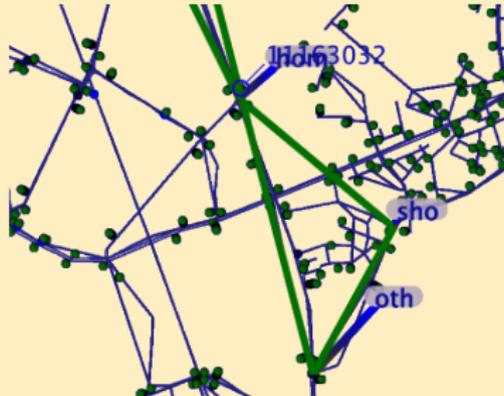
id	type
342813	shopping
9856325	public transport

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Locations

Movement profile



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Movement profile

person-id	time	location-id
1324020	10:00 - 10:20	9856325
1324020	10:20 - 11:40	342813

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Synthetic population

Locations

Movement profile

Infection model

$$p(\text{infect}|\text{contact}) \approx \theta \cdot \frac{\text{sheddingRate} \cdot \text{intake}}{\text{roomSize} \cdot \text{airExchange}} \cdot \text{duration}$$

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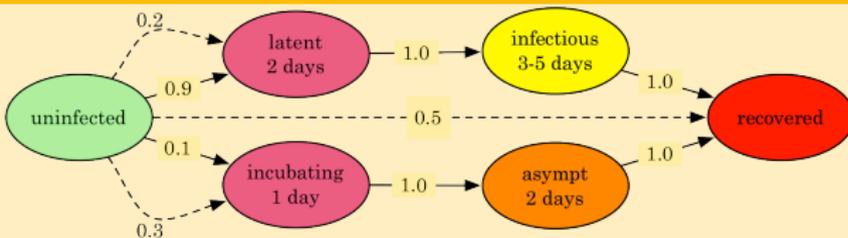
Synthetic population

Locations

Movement profile

Infection model

Disease progression model



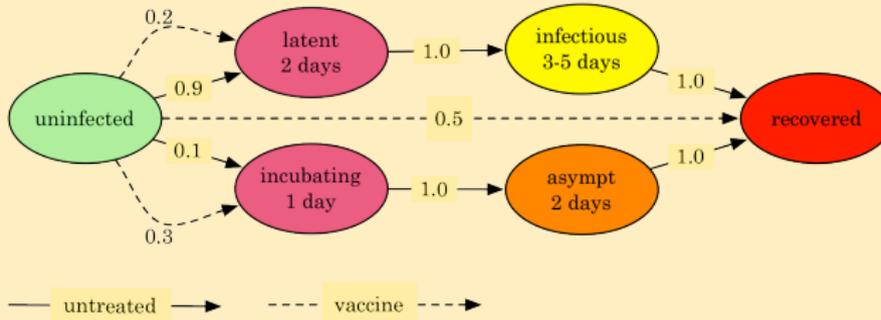
Source: [3]

A promising parallelization approach builds on two properties:

Parallelization Strategy

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Health state changes are not immediately



Source: [3]

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Health state changes are not immediately

The movement profiles can be divided among the different locations

time	action	person-id	location-id
10:00	arrive	1324020	9856325
10:05	leave	493592	9856325
10:07	leave	948539	9856325

⇒ This allows independent processing of events for each location, with a barrier at the end of a simulated day.

About 80% of the runtime is used to process the movement profiles.

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For parallelizing the movement profiles:

- ▶ In the initialization each location n is assigned to a task-id.
- ▶ The fact that locations are visited by different numbers of people is taken into account.
- ▶ To handle the movement profile events an array of `CompletableFutures` is run asynchronously.

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In the remaining part of the code, two time-demanding loops could be parallelized by converting them to a Java8 `ParallelStream`.

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Performance loss: $\approx 20\%$

The most used simulation configuration is for the Berlin metro area (ca. 5 mio people), with one agent for every four persons.

- ▶ 10000-20000 simulations per report (every 3 weeks)
- ▶ Runtime (single-threaded, one simulation): ≈ 2 h
- ▶ Memory usage: ≈ 11 GB
- ▶ File I/O: 500 / 34 MB

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Standard compute node of HLRN's Lise cluster:

- ▶ 2 Intel Cascade 9242 (= 96 cores)
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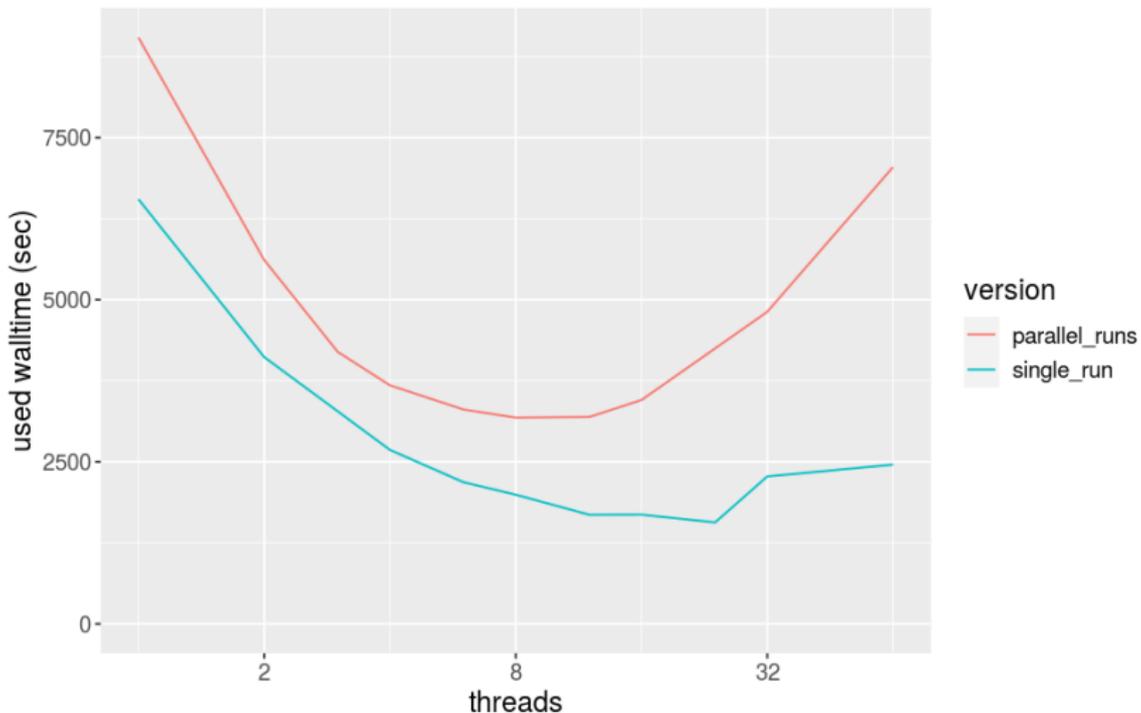
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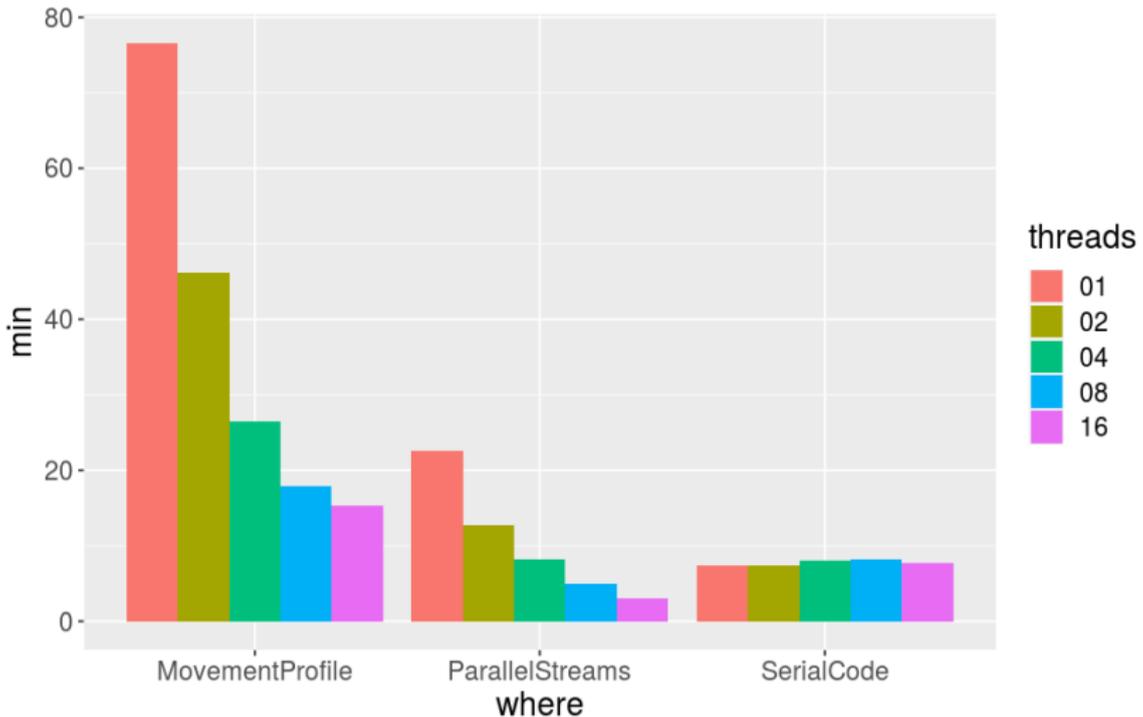
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Because of the memory constraint only 32 simulation can be started in parallel.

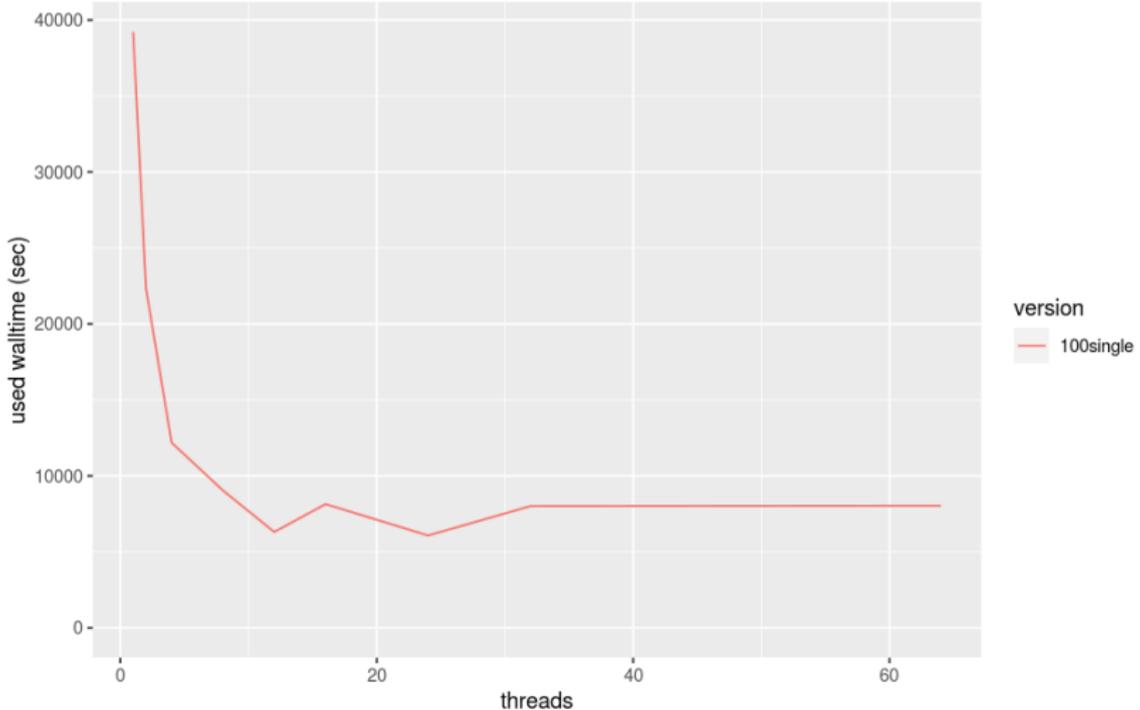
Walltime of Berlin scenario with 25% population



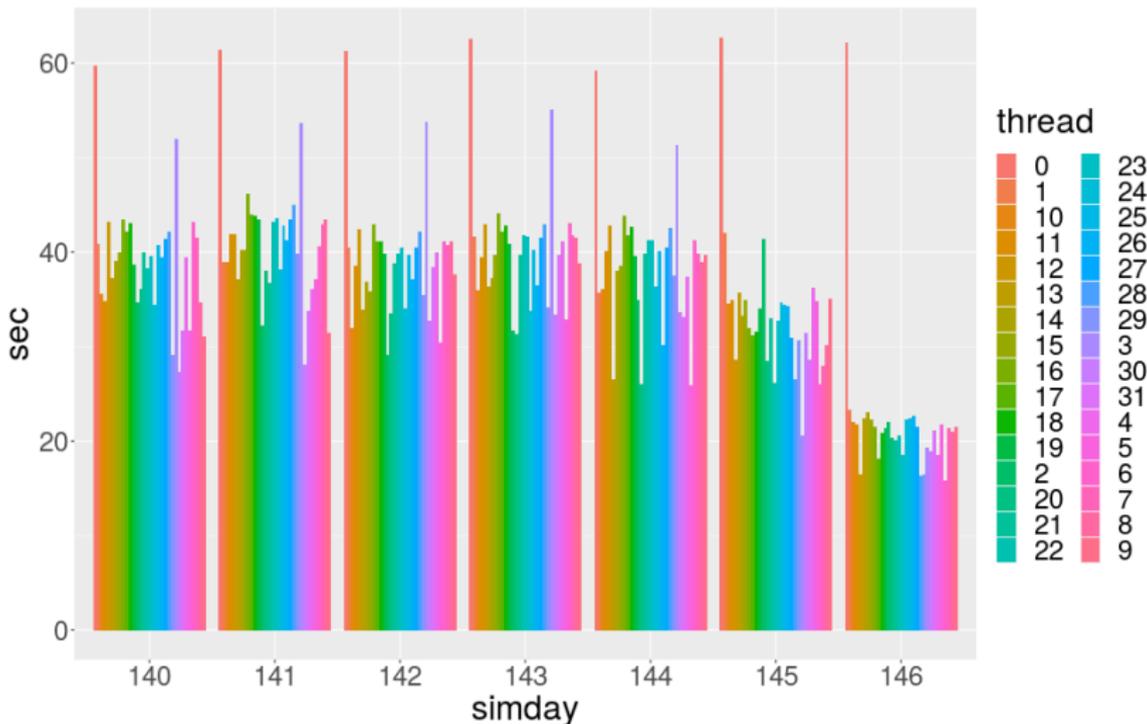
Walltime of different parts (version: single_run)



Walltime of Berlin scenario with 100% population

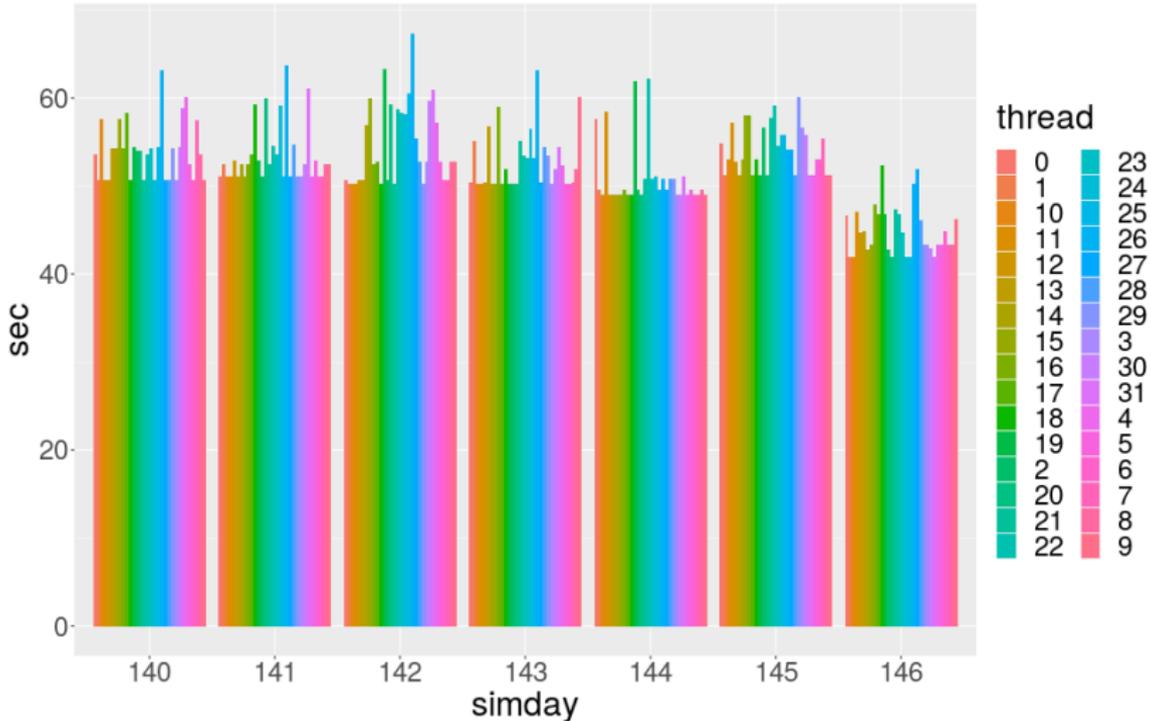


Time used for handling the movement profiles for separate days

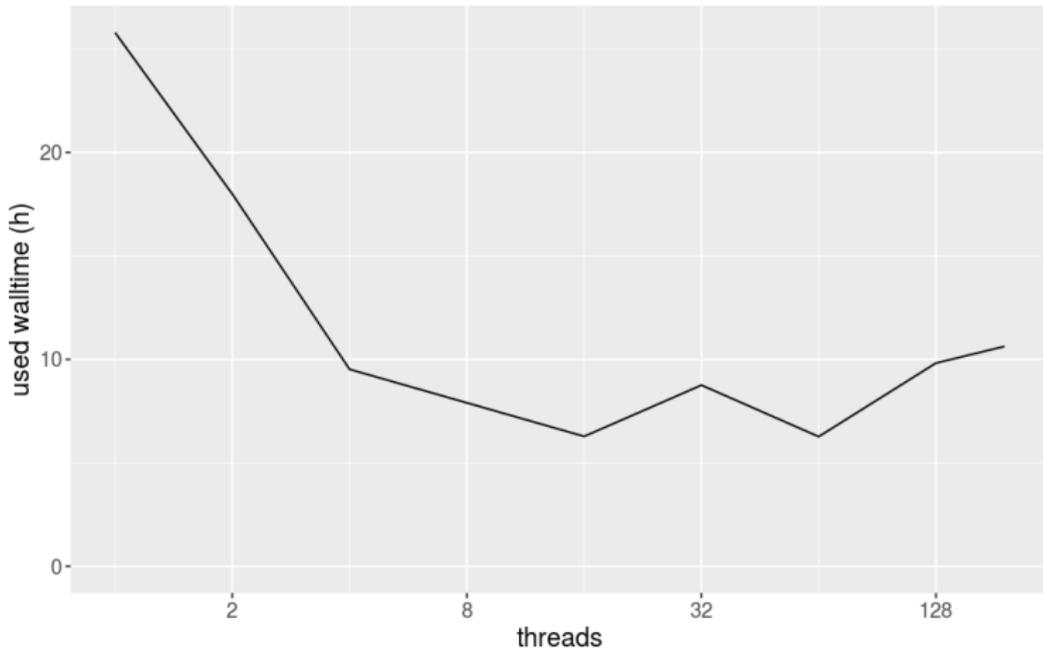


Germany 25%: Runtime per Thread

Time used for handling the movement profiles for separate days



Walltime of Germany scenario with 25% population



Links:

- ▶ <https://covid-sim.info/>: MATSim Episim portal with model results, published reports ...
- ▶ <https://github.com/matsim-org/matsim-episim-libs>: MATSim Episim source code (the work presented here is in the branch `multithreaded`).

Figure Sources:

[1] <https://www.matsim.org/gallery/poznan>

[2] <https://covid-sim.info/v3?day=17>

[3] C. L. Barrett, K. R. Bisset, S. G. Eubank, Xizhou Feng and M. V. Marathe, "EpiSimdemics: An efficient algorithm for simulating the spread of infectious disease over large realistic social networks," *SC '08: Proceedings of the 2008 ACM/IEEE Conference on Supercomputing*, 2008, pp. 1-12

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Thank you for your attention!