

Four compartment epidemic random walk model

Téo Granger^(a), Thomas Michelitsch^(a), Bernard Collet^(a) Michael Bestehorn^(b), Alejandro P. Riascos^(c)

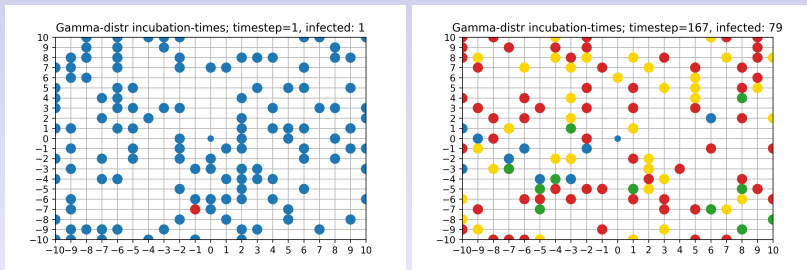


Figure: Colors of indicate health status S C I R of the walkers.

- $Z \gg 1$ random walkers navigate independently on a $N \times N$ square-lattice, jumping with probability $1/4$ to any of four neighbor lattice points
- Each walker is in one of the states ('compartments')

S : susceptible (for infection)

C : incubated, infected but not infectious

I : infected and infectious

R : recovered and immune

(a) Sorbonne Univ., Institute ∂' Alembert, Campus Jussieu, Paris

(b) Institute of statistical Physics, Technical Univ. of Cottbus-Senftenberg, Germany

(c) Institute of Physics, Complex Systems Department, UNAM, Mexico City

Infection rule :

If: **S** meets **I** on the same lattice point – collision of I and S walkers

Then: the S walker gets infected with probability P_{inf}

performing transition

S \rightarrow **C** with probability P_{inf}

followed by delayed transition **C** \rightarrow **I** \rightarrow **R** \rightarrow **S**

with random sojourn times t_C, t_I, t_R in the compartments

drawn from probability density functions

$$\mathbb{P}(t_{C,I,R} \in [\tau, \tau + d\tau]) = K_{C,I,R}(\tau)d\tau$$

Evolution equations $s(t) + c(t) + j(t) + r(t) = 1$ (constant population without deaths)

$$\frac{d}{dt}s(t) = -\mathcal{A}(t) + (\mathcal{A} \star K_C \star K_I \star K_R)(t)$$

$$\frac{d}{dt}c(t) = \mathcal{A}(t) - (\mathcal{A} \star K_C)(t)$$

$$\frac{d}{dt}j(t) = (\mathcal{A} \star K_C)(t) - (\mathcal{A} \star K_C \star K_I)(t)$$

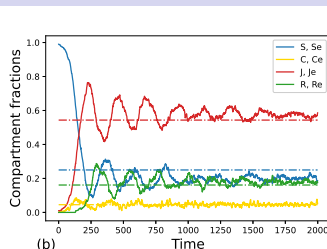
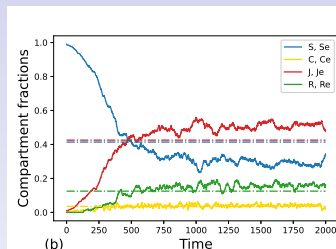
$$\frac{d}{dt}r(t) = (\mathcal{A} \star K_C \star K_I)(t) - (\mathcal{A} \star K_C \star K_I \star K_R)(t)$$

$(a \star b)(t) = \int_0^t a(\tau)b(t - \tau)d\tau$ stands for convolution

$\mathcal{A}(t)$ infection rate, assumption $\mathcal{A}(t) = \beta j(t)s(t)$ nonlinear function of $j(t)$ and $s(t)$ describing probability of collision of I and S walkers.

Four compartment SCIRS model with superspreaders (long-range jumpers)

δ -kernels: $K_{C,I,R}(\tau) = \delta(\tau - t_{C,I,R})$, i.e. constant t_C, t_I, t_R for all walkers



Left plot: Epidemic evolution without superspreaders, all walkers perform local jumps.

Right plot: 30% superspreaders = fraction of walkers performing long-range jumps

Simulation with $Z = 100$ walkers, N^2 lattice points ($N = 41$), density of walkers $Z/N^2 \approx 0.06$, infection probability $P_{inf} = 0.9$ with $t_C = 10$, $t_R = 40$ and large time of illness $t_I = 130$ sojourn times. We count at each time increment the fractions $s(t) = \frac{Z_s(t)}{Z}$, $c(t) = \frac{Z_c(t)}{Z}$, $j(t) = \frac{Z_I(t)}{Z}$, $r(t) = \frac{Z_R(t)}{Z}$.

References:

- [1] W.O. Kermack, A.G. McKendrick, A contribution to the mathematical theory of epidemics, Proc. Roy. Soc. A 115, 700–721 (1927)
 - [2] T. Granger, T.M. Michelitsch, M. Bestehorn, A. P. Riascos, B. A. Collet, *In Press*, Phys. Rev. E, Preprint: arXiv:2210.09912 (2022).
 - [3] M. Bestehorn, T. M Michelitsch, B. A. Collet, A. P. Riascos, A. F. Nowakowski, Phys. Rev. E, 105, 024205, (2022).
- For animated simulations and details of the model consult:



<https://sites.google.com/view/scirs-model-supplementaries/accueil>