

# Benchmarking road network growth models

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models

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[Cats and Birch, 2021]

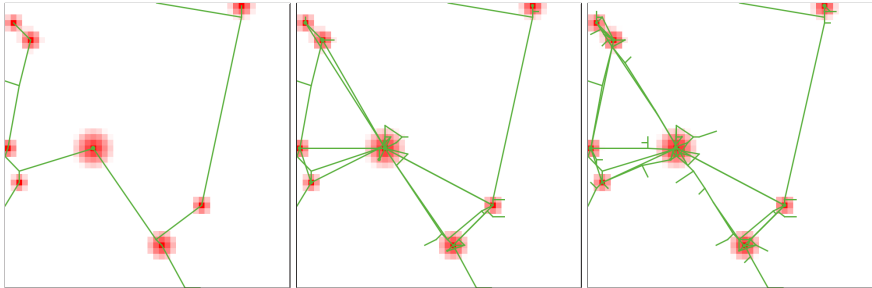
[Szell et al., 2021]

[Cats et al., 2020]

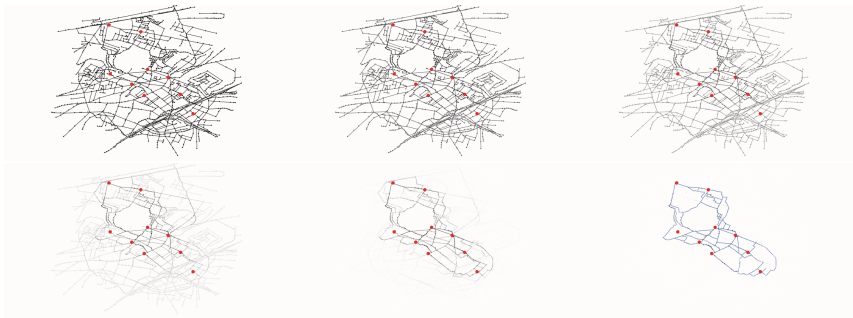


At each time step, with a fixed population density:

- 1** Add new nodes preferentially to population and connect them
- 2** Variable heuristic for new links, among: nothing, random, gravity-based deterministic breakdown, gravity-based random breakdown (from [Schmitt, 2014]), cost-benefits (from [Louf et al., 2013]), biological network generation (based on [Tero et al., 2010])



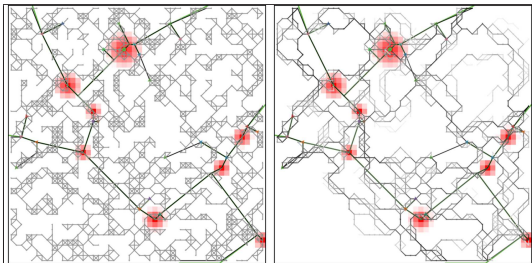
Model introduced by [Tero et al., 2010]: exploration and reinforcement by a slime mould searching for resources



*Application to the design of optimal bus routes in [Raimbault, 2018]*

Adding new links with biological heuristic:

- 1 Create network of potential new links, with existing network and randomly sampled diagonal lattice
- 2 Iterate for  $k$  increasing ( $k \in \{1, 2, 4\}$  in practice) :
  - Using population distribution, iterate  $k \cdot n_b$  times the slime mould model to compute new link capacities
  - Delete links with capacity under  $\theta_d$
  - Keep the largest connected component
- 3 Planarize and simplify final network

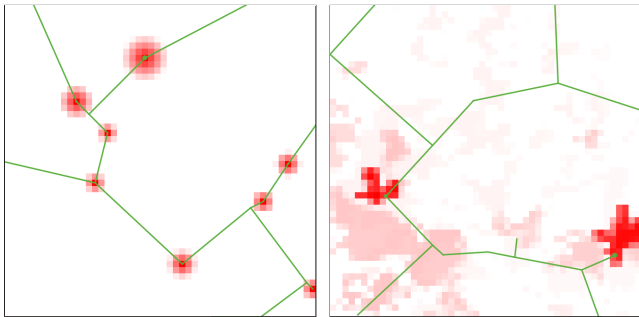


Intermediate steps for biological network generation

Heuristic	Param.	Name	Process	Domain	Default
Base	$l_m$	added links	growth	0; 100	10
	$d_G$	gravity distance	potential	0; 5000	500
	$d_0$	gravity shape	potential	0; 10	2
	$k_h$	gravity weight	potential	0; 1	0.5
	$\gamma_G$	gravity hierarchy	potential	0.1; 4	1.5
Random	$\gamma_R$	random selection	hierarchy	0.1; 4	1.5
	$\theta_R$	random threshold	breakdown	1; 5	2
Cost-benefits	$\lambda$	compromise	compromise	0; 0.1	0.05
Biological	$n_b$	iterations	convergence	40; 100	50
	$\theta_b$	biological th.	threshold	0.1; 1.0	0.5

**Synthetic setup:** rank-sized monocentric cities, simple connection with border nodes to avoid border effects

**Real setup:** Population density raster at 500m resolution (European Union, from Eurostat)



**Stopping conditions:** fixed final time; fixed total population; fixed network size.

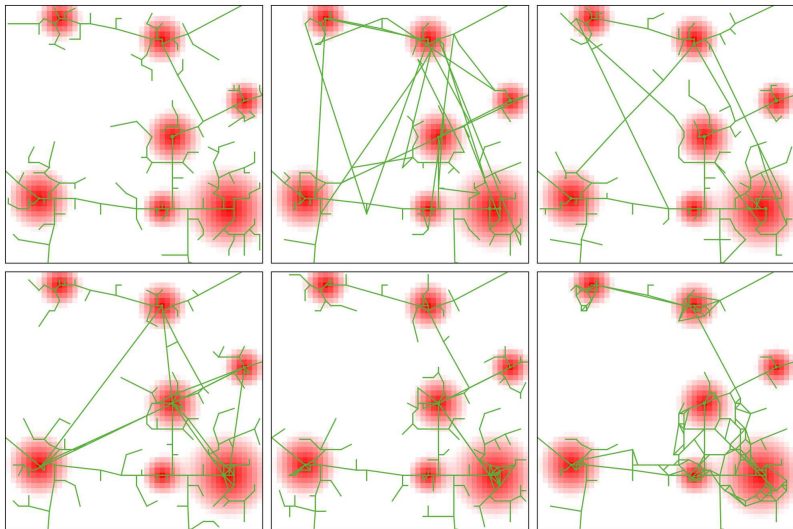


Network Topology measured by:

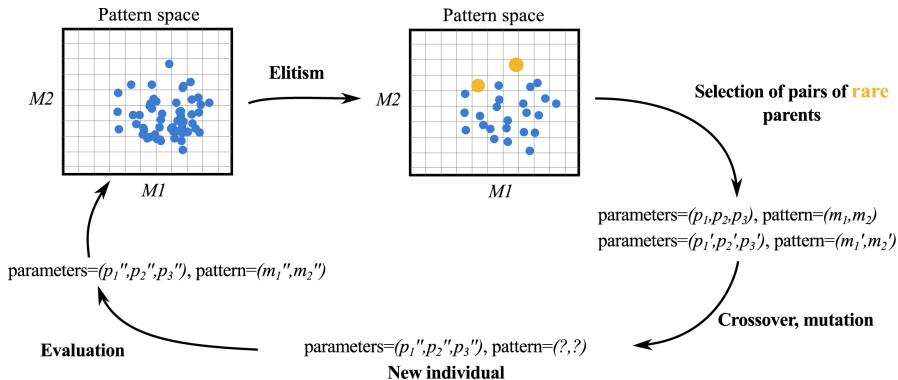
- Average betweenness and closeness centralities
- Efficiency (network pace relative to euclidian distance)
- Mean path length, diameter



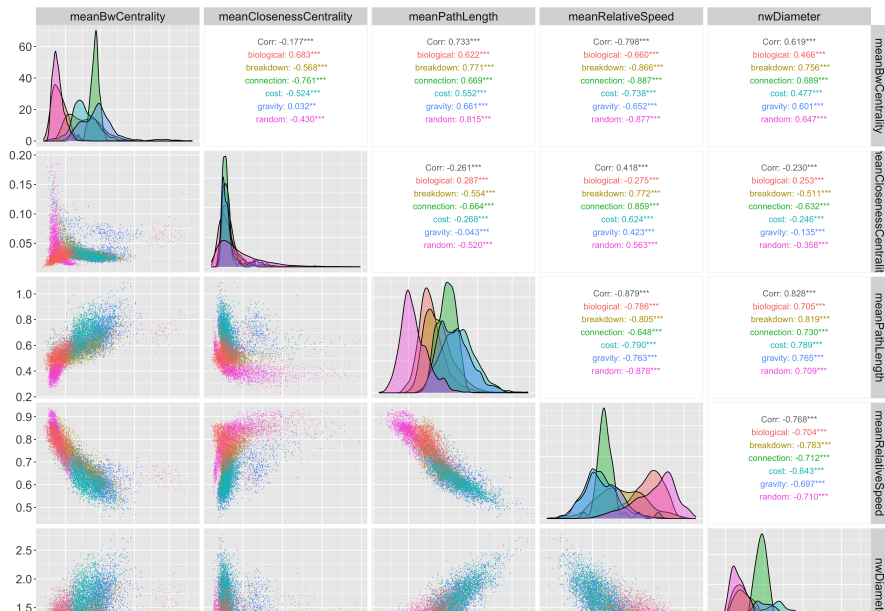
# Example of generated networks

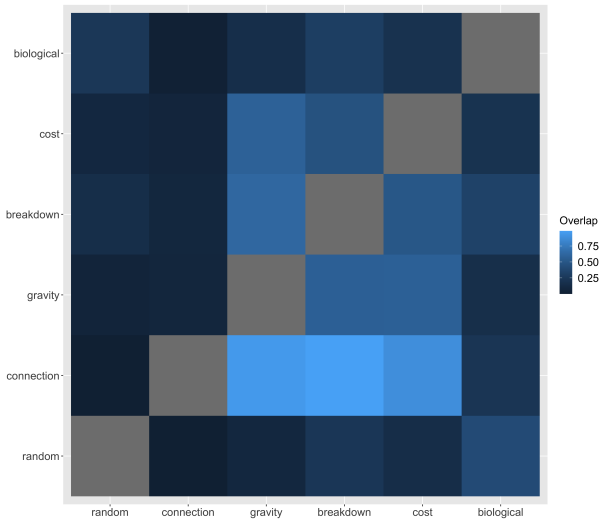


*In order: connection; random; deterministic breakdown; random breakdown; cost-driven; biological.*



# Indicators feasible space










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## Open repositories

<https://github.com/JusteRaimbault/NetworkGrowth>



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110(22):8824–8829.



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Tero, A., Takagi, S., Saigusa, T., Ito, K., Bebbber, D. P., Fricker, M. D., Yumiki, K., Kobayashi, R., and Nakagaki, T. (2010).  
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