

Impact of Spatial Seeding on Metapopulation Models for Disease Dynamics



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Introduction

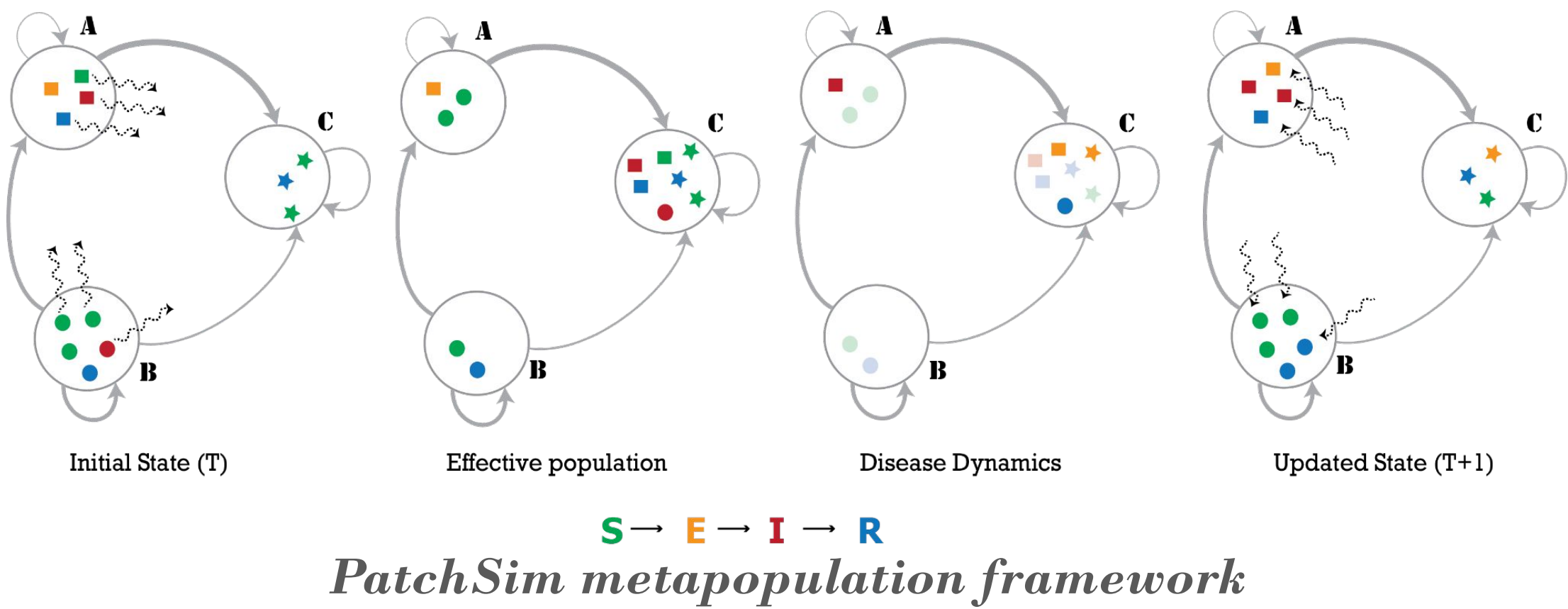
- Metapopulation models combine SEIR models with network models
- Metapopulation models provide insights into individual regions without significant computational cost
- Distinct features arise from accounting for mobility within the model
- PatchSim [1] is a metapopulation model developed at the BII NSSAC
- PatchFlow [2] is a repository of datasets containing mobility data from radiation modeling that can be used in PatchSim
- The goal is for the results to not only provide insights into PatchSim/PatchFlow itself but also for better pandemic preparedness

Purpose

- The purpose of this experiment is to deepen the understanding on mobility and spatial heterogeneity, esp. metapopulation models
- Understanding should be brought on how disease spread is altered by different networks and seeding locations
- Contrasting with a homogeneous model for entire country

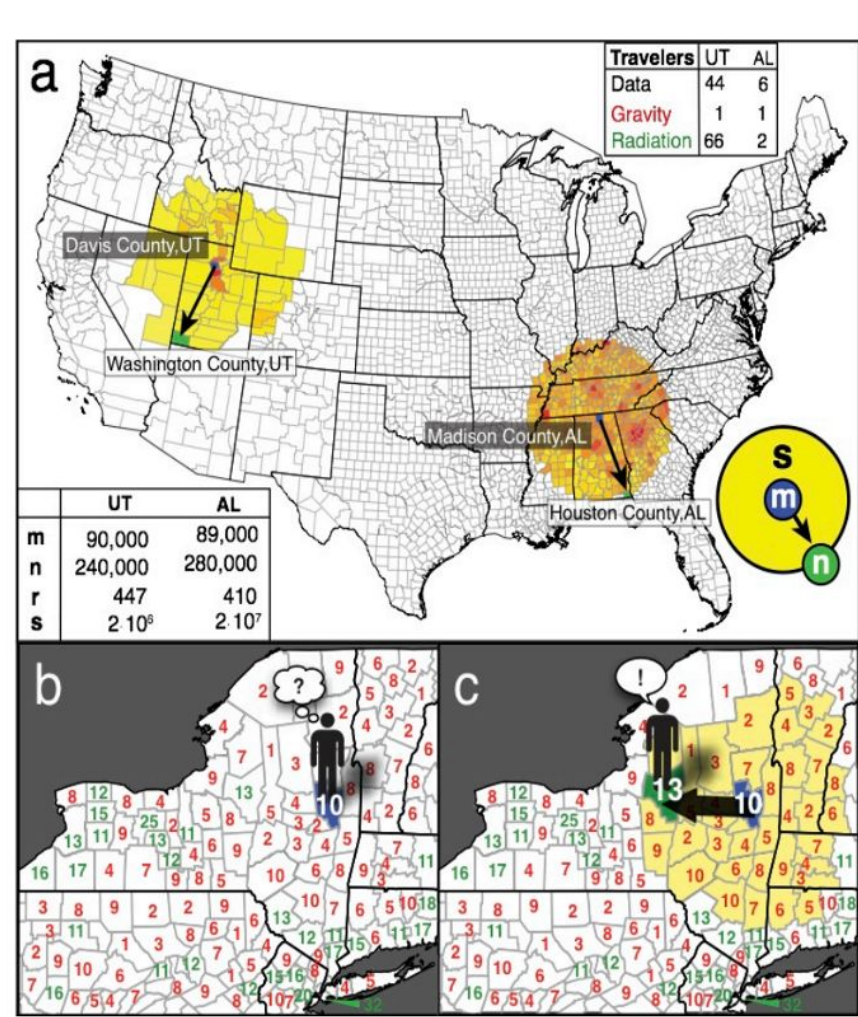
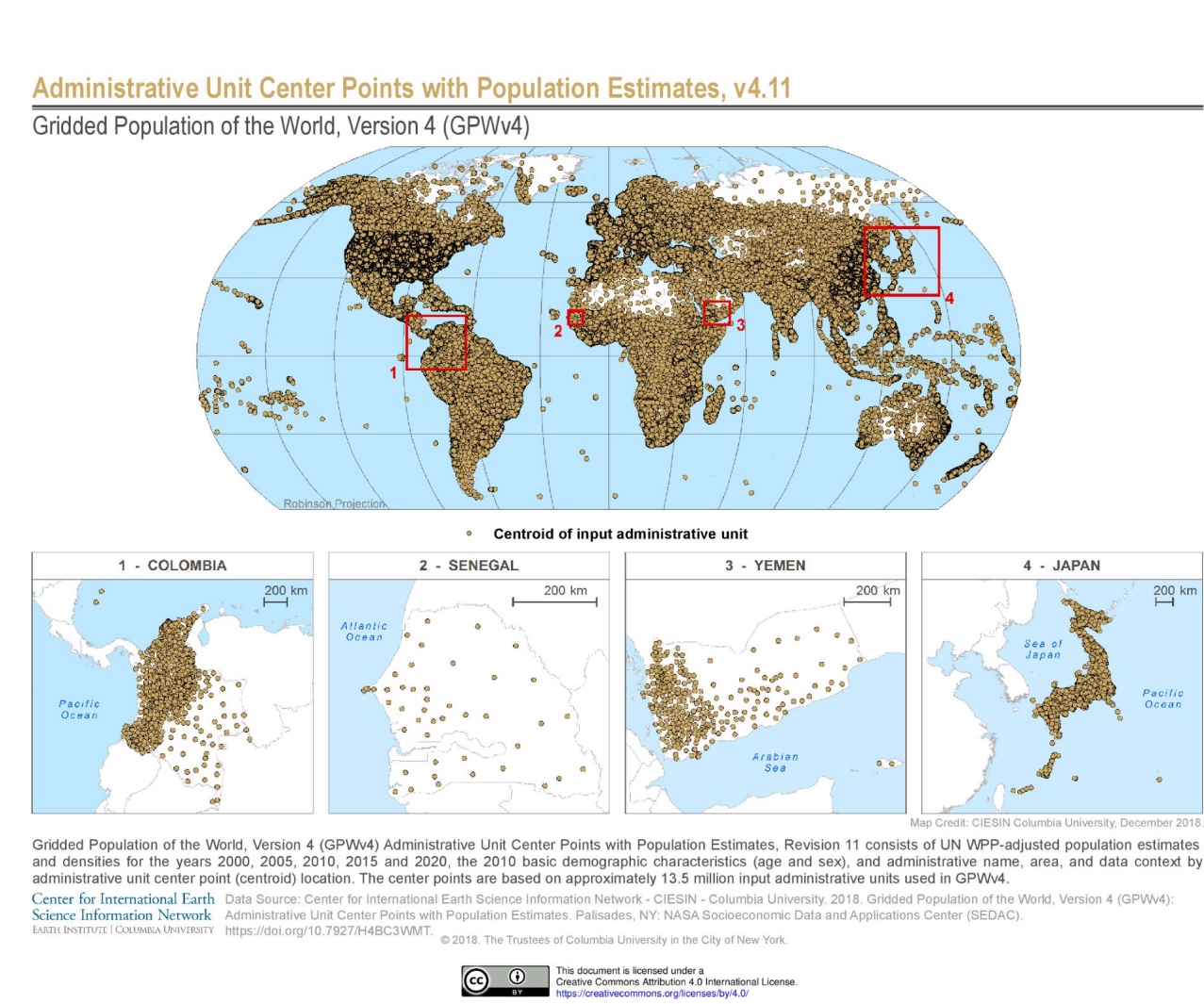
Patchsim Description

- Metapopulation based disease model both SEIR models and network models
 - SEIR difference equations for dynamics within region
 - Network models account for mobility among subregions
- Has been used in various contexts including resource allocation [3], forecasting [4], and scenario-based projections [5]



Patchflow Description

- PatchFlow is a set of precalculated networks for simulating the metapopulation SEIR model at a global level
- Country specific mobility networks are based on a radiational model
- Population distributions are obtained from Gridded Populations of the World (GPWv4), and administrative regions were obtained from GADM
- Admin level is the resolution of the network - for instance admin 1 in USA is states and 2 is counties
- The radiation parameter (rad) determines what percent of the population travels in the network [6]
- Network and Population files are designed to be compatible with PatchSim

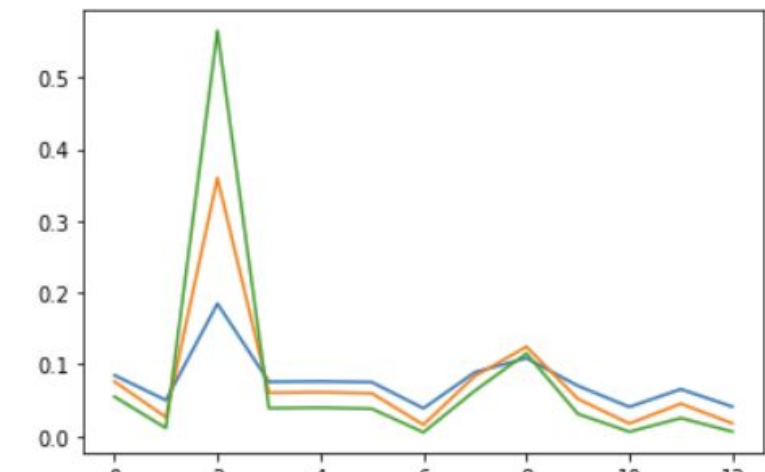


$$\langle T_{ij} \rangle = T_i \frac{m_i n_j}{(m_i + s_{ij})(n_i + n_j + s_{ij})},$$

- m_i - population of source region i
- n_j - population of destination region j
- T_i - number of commuters from source region i
- d_{ij} = distance from region i to region j
- s_{ij} - population residing within d_{ij} from region i

Study Design

- Independent variables are admin level (admin), radiation parameter (rad), seeding scaler (skew) and exposure rate (β)
- Only countries with a population over one million are included
- admin0 is a homogeneous SEIR model with one patch, and compared against admin1 and admin2 simulations
- $\text{rad} \in [1\%, 5\%, 10\%]$, $\text{skew} \in [1, 2, 3]$, $\beta \in [0.3, 0.5, 0.7]$
- Total number of initial cases are a proportion of the national population (20 per 100k)
- Initial cases within patches are taken as a proportion relative to the patch's population exponentiated by the skew parameter: $P^{\text{skew}} / \sum P^{\text{skew}} \times \text{Total}$
- A higher skew leads to more initial cases concentrated in higher populated patches as shown below (skew = 1, 2, 3)
- Such skewing allows the network to play a larger role
- Note that due to full connectivity, total attack rate remains unchanged for a given β



Exposure Rate	Cumulative Attack Rate (%)
0.45	21.45
0.5	37.47
0.55	49.63
0.6	59.08
0.65	66.52
0.7	72.47

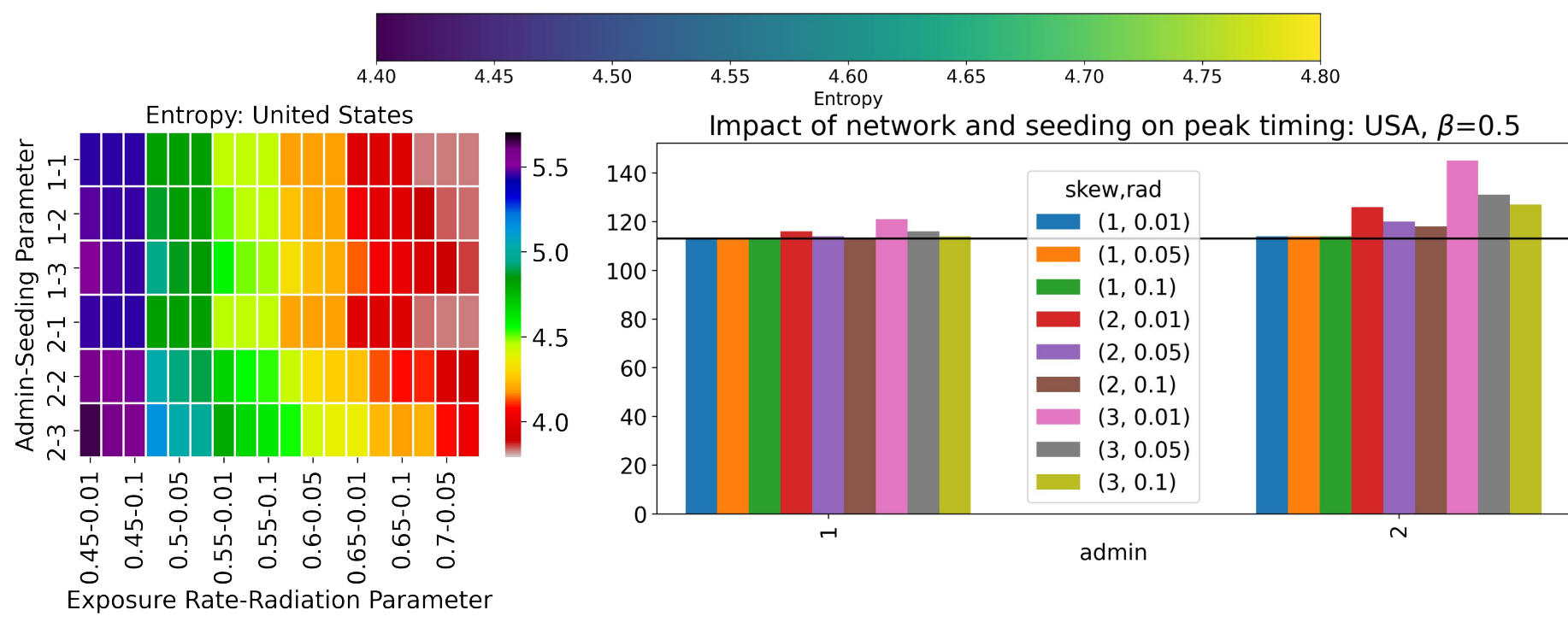
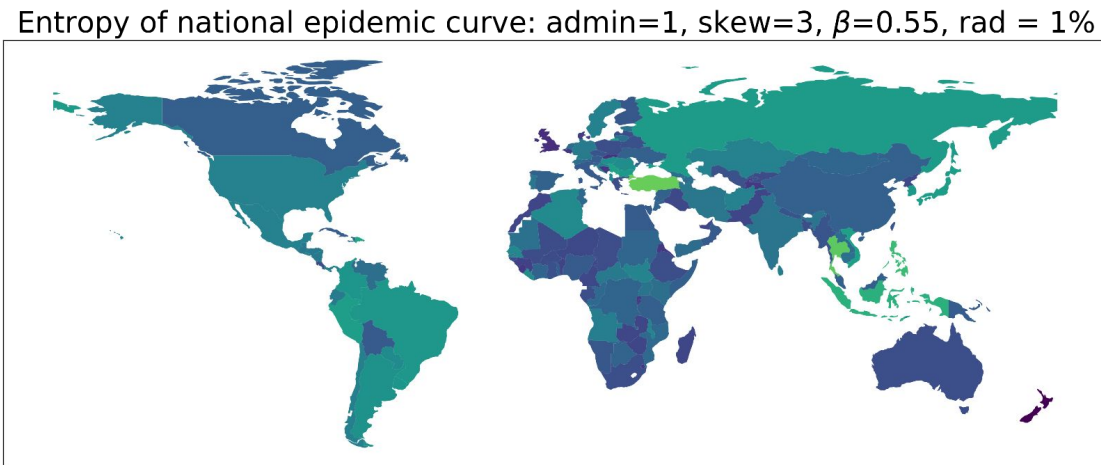
Metrics

- Number of days to cross 1%, 50% and 99% of total cases
- Shannon Entropy measurements of the normalized epidemic curve [7]

$$H(X) = - \sum_{i=1}^n P(x_i) \log_b P(x_i)$$

Initial Results

- Impact of exposure rate on entropy is non-monotonic
 - Higher β lead to sharper epidemics, while lower β leads to faster extinction
- With fixed β , as admin level increases, entropy increases due to localized mixing within the regions
 - This is further accentuated by the skew in seeding and the rad parameter of the network
- Uniform seeding (skew=1) does not significantly change peak times, whereas for higher skew, the effect of rad parameter is more pronounced



Current and Future Work

- Impact of core and peripheral seeding using the network [8]
- Impact of heterogeneity in prior immunity & vaccination [9]
- Role of adaptive interventions at local/global levels [10]

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