

## Generation intervals in space

Jonathan Dushoff<sup>1</sup>, Sang Woo Park<sup>1</sup>, and David Champredon<sup>2</sup>

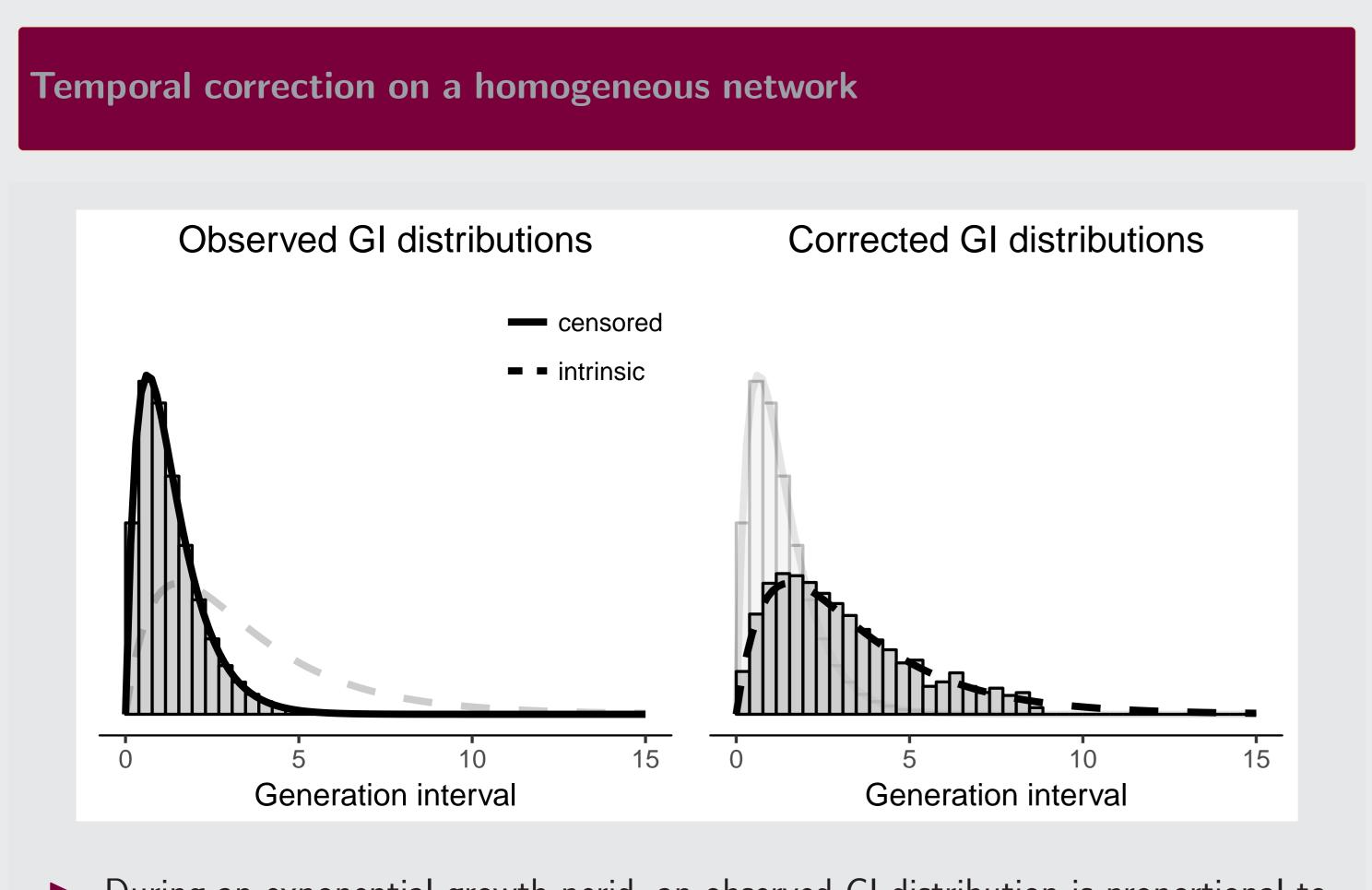
<sup>1</sup>McMaster University, Hamilton, Ontario, Canada; <sup>2</sup>York University, Toronto, Ontario

## Introduction

- ► Generation interval (GI) measures time between when a person is infected and when that person infects another person
- ▶ GI distributions,  $g(\tau)$ , link speed, r, and strength,  $\mathcal{R}$ , of an epidemic [1]

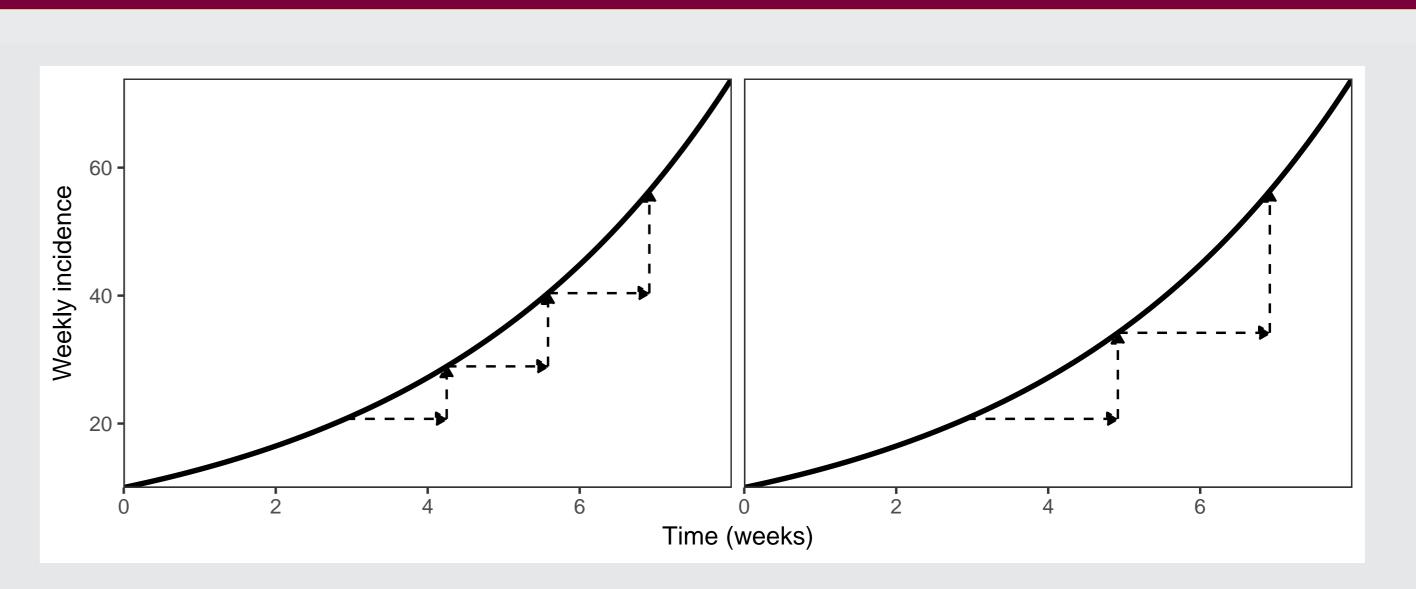
$$1/\mathcal{R} = \int g( au) \exp(-r au) d au$$

- ► Previous work showed that measuring GI through contact tracing data can introduce bias [2]
- Trapman et al. [3] demonstrated that network structure can affect  $\mathcal{R}$  but it also has effect on GI distributions



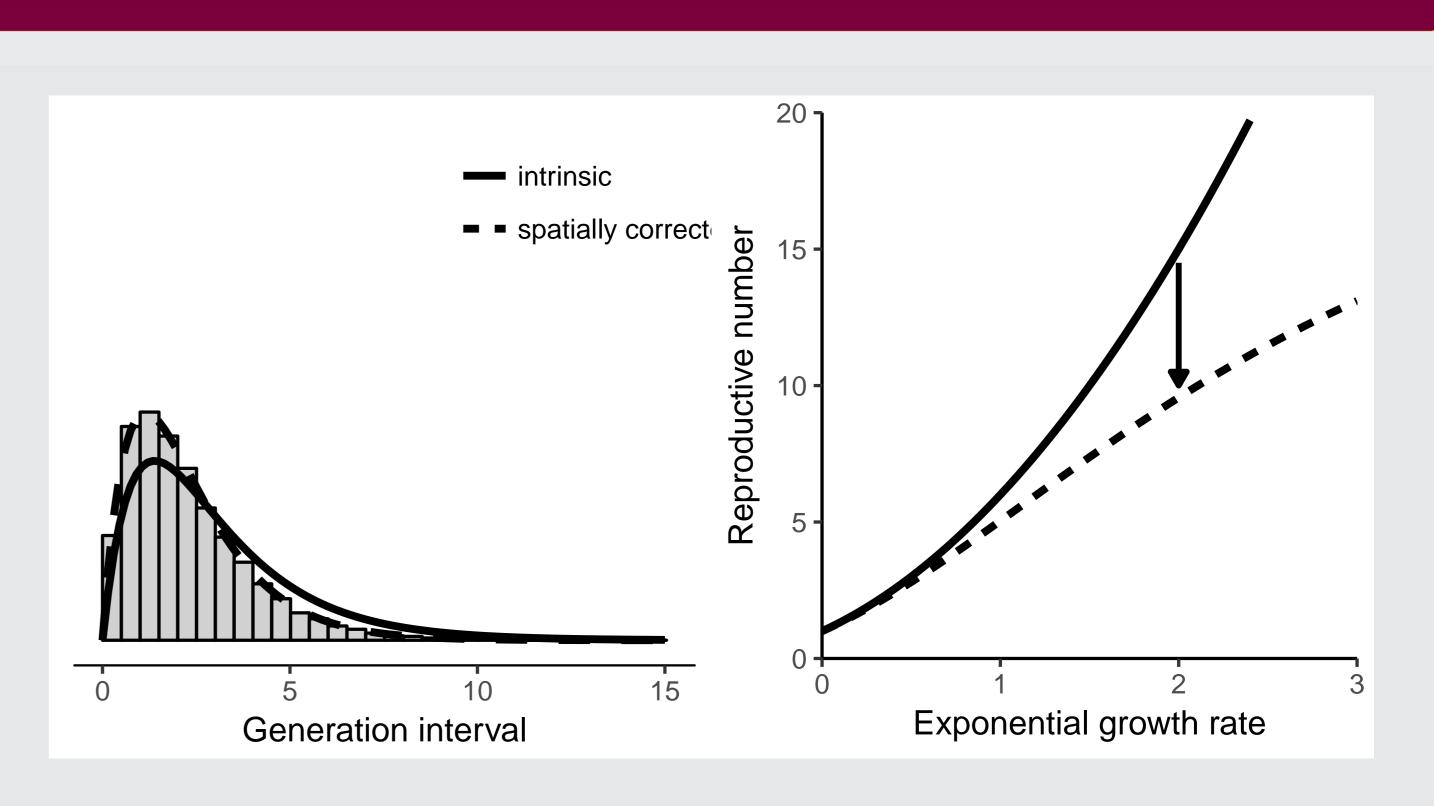
- During an exponential growth perid, an observed GI distribution is proportional to  $g(\tau) \exp(-r\tau)$ .
- ▶ By weighting an observed distribution by  $exp(r\tau)$ , the intrinsic GI distribution can be recovered





Longer generation interval requires higher  $\mathcal{R}$  given fixed exponential growth rate r.

## Temporal correction on an empirical network

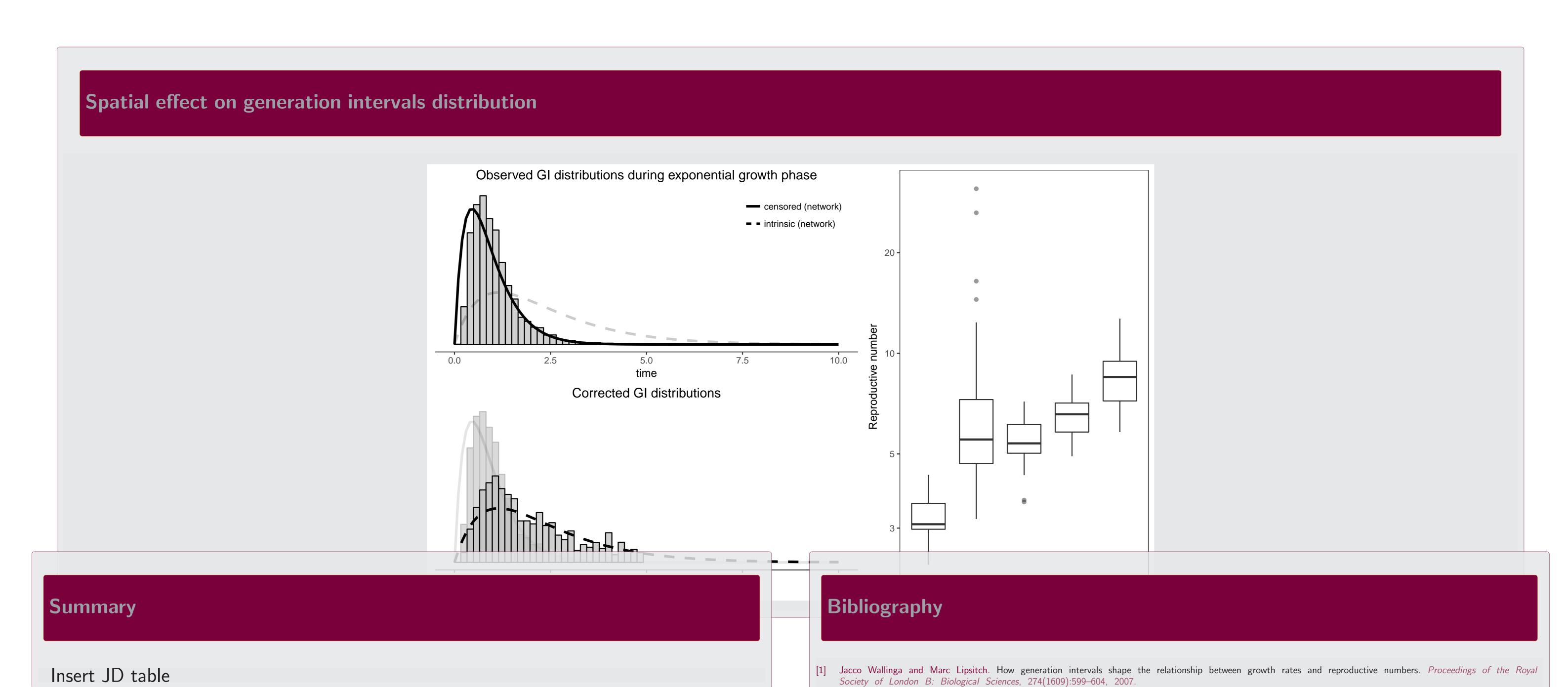


- Something about network/spatial effect
- $ightharpoonup \mathcal{R}$  estimate based on corrected GI distributions matches empirical  $\mathcal{R}$ . Meanwhile, using intrinsic GI distributions and observed GI distributions can over/underestimate  $\mathcal{R}$ .

[2] David Champredon and Jonathan Dushoff. Intrinsic and realized generation intervals in infectious-disease transmission. In Proc. R. Soc. B, volume 282, page

[3] Pieter Trapman, Frank Ball, Jean-Stéphane Dhersin, Viet Chi Tran, Jacco Wallinga, and Tom Britton. Inferring r0 in emerging epidemicsthe effect of

common population structure is small. Journal of the Royal Society Interface, 13(121):20160288, 2016.



20152026. The Royal Society, 2015.