



## UvA-DARE (Digital Academic Repository)

### Modeling the Impact of White-Plague Coral Disease in Climate Change Scenarios

Zvuloni, A.; Artzy-Randrup, Y.; Katriel, G.; Loya, Y.; Stone, L.

**DOI**

[10.1371/journal.pcbi.1004151](https://doi.org/10.1371/journal.pcbi.1004151)

**Publication date**

2015

**Document Version**

Other version

**Published in**

PLoS Computational Biology

**License**

CC BY

[Link to publication](#)

**Citation for published version (APA):**

Zvuloni, A., Artzy-Randrup, Y., Katriel, G., Loya, Y., & Stone, L. (2015). Modeling the Impact of White-Plague Coral Disease in Climate Change Scenarios. *PLoS Computational Biology*, 11(6), e1004151. <https://doi.org/10.1371/journal.pcbi.1004151>

**General rights**

It is not permitted to download or to forward/distribute the text or part of it without the consent of the author(s) and/or copyright holder(s), other than for strictly personal, individual use, unless the work is under an open content license (like Creative Commons).

**Disclaimer/Complaints regulations**

If you believe that digital publication of certain material infringes any of your rights or (privacy) interests, please let the Library know, stating your reasons. In case of a legitimate complaint, the Library will make the material inaccessible and/or remove it from the website. Please Ask the Library: <https://uba.uva.nl/en/contact>, or a letter to: Library of the University of Amsterdam, Secretariat, Singel 425, 1012 WP Amsterdam, The Netherlands. You will be contacted as soon as possible.

*UvA-DARE is a service provided by the library of the University of Amsterdam (<https://dare.uva.nl>)*

Figure S2

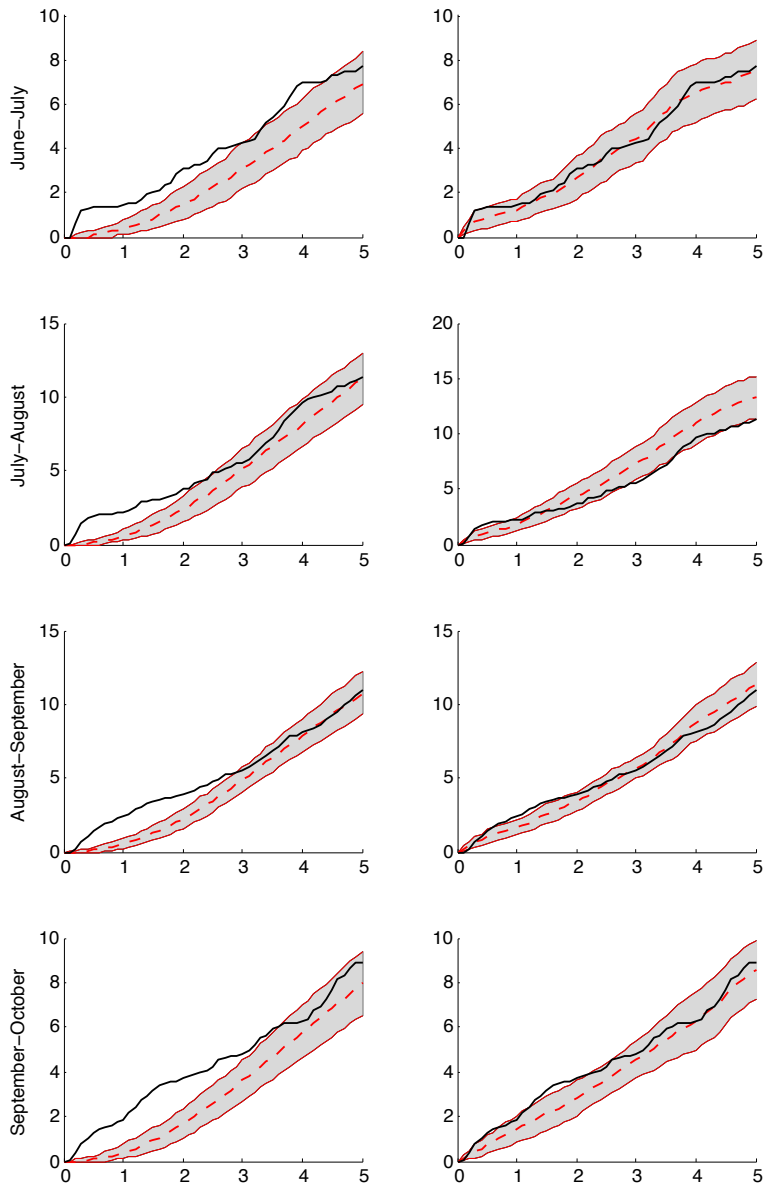
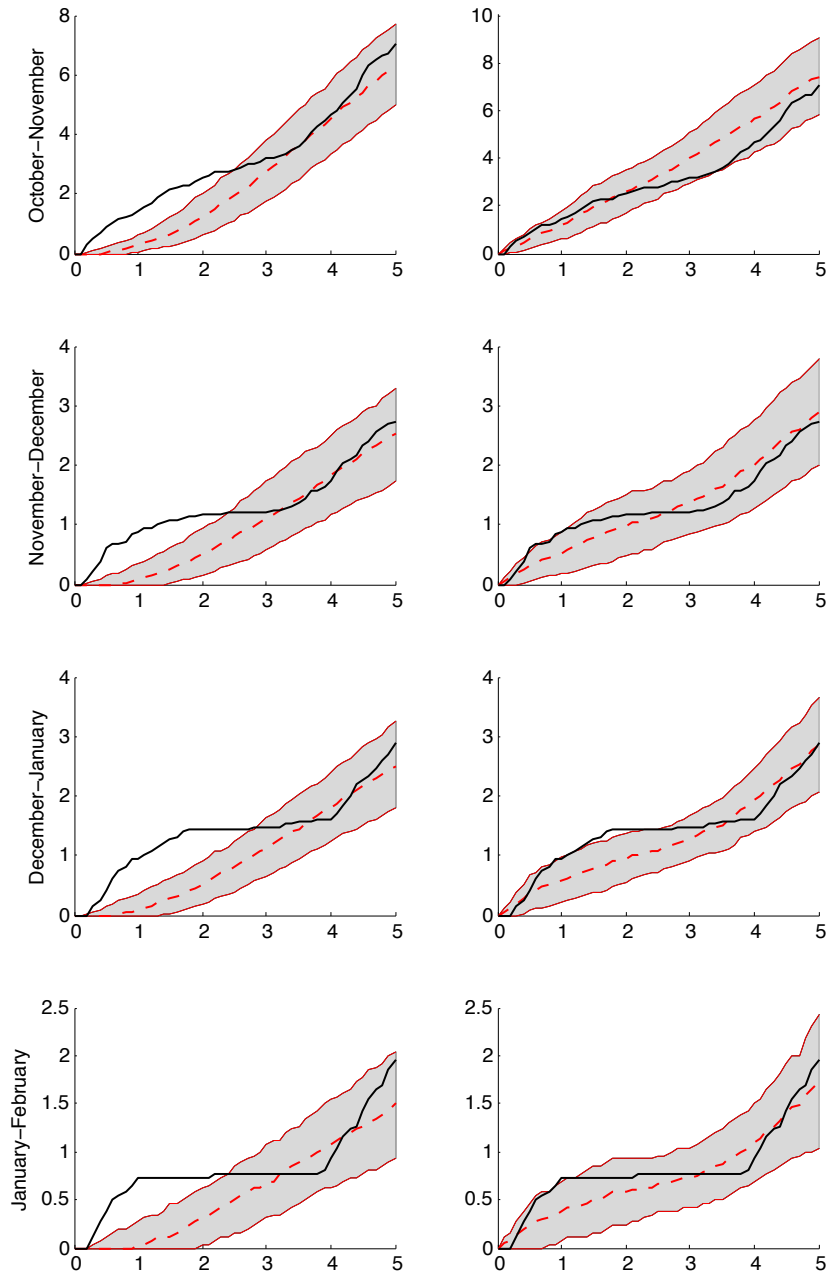
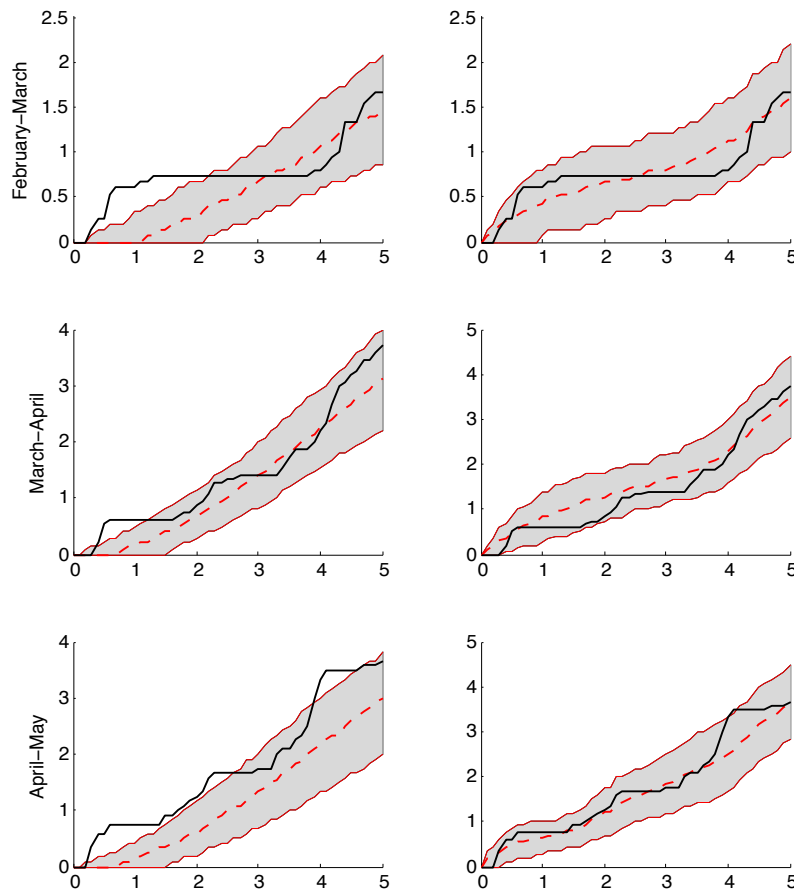


Figure S2 continued



**Figure S2 continued**



**Figure S2. Plots of the spatiotemporal index  $n(r)$  calculated for pairs of sequential sampling dates (see text).** The black line represents the observed  $n(r)$  values (eqn 1) for corals infected with white-plague disease (WPD). The shaded areas are the Monte Carlo 95% confidence interval (CI) envelopes, representing two different null expectations: **(a)** new infections develop randomly within the studied plot, independent of the spatial location of infected corals from the previous month; and **(b)** new infections develop according to the spatiotemporal model (eqn 2). For distance scales ( $r$ ) where  $n(r)$  values fall within the envelope, the spatial distribution of infected corals does not differ significantly from the null distribution. Infected corals are significantly more aggregated (/over-dispersed) where the observed  $n(r)$  values fall above (/below) the CI envelope.

In all cases the NICs observed in the field appeared to form aggregations around PICs over distance scales of up to 4.5 m. That is, in all cases the hypothesis that the NICs were infected by a random process of disease transmission independent of the

spatial location of the PICs was rejected. Almost in all cases, the observed  $n(r)$  was purely within the expectation of the spatiotemporal model (eqn 2) for all distance scales  $r$ . However, in a few cases the observed  $n(r)$  was found to be greater than the upper bound of the 95% CI envelope generated by the model realizations for certain distance scales (see, for example, August-September 2006).