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Daniella Lowenberg@danilowenbergDryad Product ManagerMake Data Count LeadCalifornia Digital Library



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Principles Governing Establishment versus Collapse of HIV-1 Cellular Spread

Hataye, Jason, National Institute of Allergy and Infectious Diseases,

https://orcid.org/0000-0003-1986-5752

Casazza, Joseph, National Institute of Allergy and Infectious Diseases

Best, Katharine, Los Alamos National Laboratory

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Liang, C. Jason, National Institute of Allergy and Infectious Diseases

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Henry, Amy, National Institute of Allergy and Infectious Diseases

Laboune, Farida, National Institute of Allergy and Infectious Diseases

Maldarelli, Frank, Frederick National Laboratory for Cancer Research

Douek, Daniel, National Institute of Allergy and Infectious Diseases

Hengartner, Nicolas, Los Alamos National Laboratory

Yamamoto, Takuya, National Institute of Biomedical Innovation, Health and Nutrition

Keele, Brandon, Frederick National Laboratory for Cancer Research, (1) https://orcid.org/0000-0002-2381-1151

Perelson, Alan, Los Alamos National Laboratory, https://orcid.org/0000-0002-2455-0002





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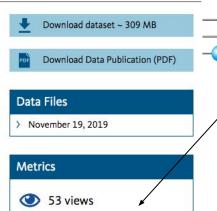
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Citation

Hataye, Jason et al. (2019), Principles Governing Establishment versus Collapse of HIV-1 Cellular Spread, v7, Dryad, Dataset, https://doi.org/10.5061/dryad.wdbrv15j3

Abstract

A population at low census might go extinct, or instead transition into exponential growth to become firmly established. Whether this pivotal event occurs for a within-host pathogen can be the difference between health and illness. Here we define the principles governing whether HIV-1 spread among cells fails or becomes established, by coupling stochastic modeling with laboratory experiments. Following ex vivo activation of latently-infected CD4 T cells without de novo infection, stochastic cell division and death contributes to high variability in the magnitude of initial virus release.

Transition to exponential HIV-1 spread often fails due to release of an insufficient amount of replication-competent virus. Establishment of exponential growth occurs when virus produced from multiple infected cells exceeds a critical population size. We quantitatively define the crucial transition to exponential viral spread. Thwarting this process would prevent HIV transmission or rebound from the latent reservoir.

Methods

The file "code-data-HatayeJ.zip" contains two experimental data tables and script code in R, details are in the file "README.txt".

These files were generated for this research publication:

Principles Governing Establishment versus Collapse of HIV-1 Cellular Spread

Hataye JM et al. Cell Host & Microbe, 2019

https://doi.org/10.1016/j.chom.2019.10.006

See this publication for details. It has an extensive methods section. The HIV env sequencing for this study was deposited at Genbank (https://www.ncbi.nlm.nih.gov/genbank/) with accession numbers MN515491-MN516420.

LEDOULIU

critical threshold

viral dynamics

exponential growth

tipping point

License

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Usage Notes

To verify the integrity (verify intact download) of the "code-data-HatayeJ.zip" file, one can check the SHA-256 hash of this file. On Mac OS X, this can be done by opening a terminal, typing "cd Desktop" to change to the Desktop directory (if you put the file there), and typing "shasum -a 256 code-data-HatayeJ.zip". Note that you may need to first unzip the direct download from Dryad to do the SHA-256 for "code-data-HatayeJ.zip".

The SHA-256 for "code-data-Hataye].zip" is

92fff454a014518690deb0c2f29592b17993cdc1566db6ef5cf019d3552c99e9

Funding

Division of Intramural Research, National Institute of Allergy and Infectious Diseases (USA),

National Institutes of Health (USA), Award: R01-Al028433

Department of Energy (USA), Award: Contract 89233218CNA000001

National Cancer Institute (USA), Award: Contract HHSN261200800001E

National Institutes of Health (USA), Award: R01-OD011095

National Institutes of Health (USA), Award: R01-OD011095

National Institutes of Health (USA), Award: P01-Al131365



Community Building

"Data Preservation, Sharing, and Discovery: Challenges for Small Science in the Digital Era"

A report of a workshop held May 16 -17th 2007 at NESCent, Durham, NC

Sponsors: The National Evolutionary Synthesis Center (NESCent) and the University of North Carolina at Chapel Hill, School of Information and Library Science, Metadata Research Center (<MRC>)

Organizers: Jane Greenberg (MRC), Hilmar Lapp (NESCent) and Todd Vision (NESCent).

Participants: Participants included project participants from the NESCent-MRC digital data sharing initiative, representatives from major evolutionary biology journals and societies, and experts from the information, library and computer science communities.

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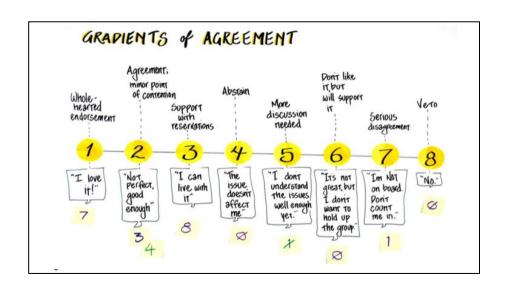


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31

Community-led Data Publishing

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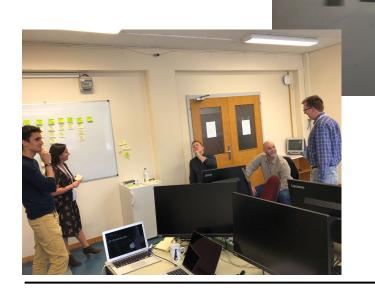
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