



May 22nd, 11:30 AM - 12:00 PM

Determinants of the efficacy of HIV latency reversing agents and implications for drug and treatment design

Ruian Ke

North Carolina State University at Raleigh, rke2@ncsu.edu

Follow this and additional works at: <http://scholarscompass.vcu.edu/bamm>



Part of the [Disease Modeling Commons](#), [Non-linear Dynamics Commons](#), and the [Virus Diseases Commons](#)

<http://scholarscompass.vcu.edu/bamm/2016/May22/12>

This Event is brought to you for free and open access by the Dept. of Mathematics and Applied Mathematics at VCU Scholars Compass. It has been accepted for inclusion in Biology and Medicine Through Mathematics Conference by an authorized administrator of VCU Scholars Compass. For more information, please contact libcompass@vcu.edu.

DETERMINANTS OF THE EFFICACY OF HIV LATENCY REVERSING AGENTS AND IMPLICATIONS FOR DRUG AND TREATMENT DESIGN

Ruian Ke^{1*}, Jessica M. Conway², Alan S. Perelson³

¹ Department of Mathematics, North Carolina State University

² Department of Mathematics, Pennsylvania State University

³Theoretical Biology and Biophysics (T-6), Los Alamos National Laboratory, USA

HIV eradication studies have focused on developing latency reversing agents (LRAs) to activate ('shock') HIV expression in latently infected cells so that these cells are purged through virus- or immune-mediated killing ('kill'). However, even in the presence of LRAs, HIV transcription may not be permanently turned on, and it is not clear which steps in the latency reversing process determine the rate of latent reservoir reduction. This makes evaluating the efficacy of candidate LRAs and predicting long-term treatment outcomes difficult. Furthermore, since almost all LRAs studied thus far have certain toxicities, it is likely that LRA treatment has to be structured/pulsed in a long-term treatment regimen to avoid severe side effects. How drug properties affect efficacy in structured treatment regimens is not clear, and this has to be taken into consideration when evaluating candidate LRAs.

We constructed a mathematical model that describes the dynamics of latently infected cells under both continuous and structured/pulsed LRA treatment. Using the model, we find that in addition to 'shock' and 'kill' rates, a previously understudied parameter, the rate at which HIV expression is deactivated both during and after LRA treatment, plays an important role in determining the efficacy of LRAs. This parameter determines the duration of HIV activation and the fate of activated cells. We further identified conditions/properties that allow LRAs to work better in structured treatment regimens than in a continuous treatment regimen. With the same amount of total dosing, pulsing treatment regimens can be more effective than a continuous treatment regimen when the 'shock' rate is high and the HIV deactivation rate after treatment stops is low. Therefore, in addition to increasing 'shock' and 'kill' rates, drug development should focus on drugs that minimize the HIV deactivation rate (i.e., maximize the duration of HIV expression) both during and after treatment.

Presenter correspondence address:

406E Cox Hall,

North Carolina State University

Raleigh, NC27695, U.S.A.