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Basic reproduction number for in-host viral infections IRMA THALIA RODRIGUEZ, HANA DOBROVOLNY, Texas Christian Univ — In epidemiology, the basic reproduction number  $(R_0)$  is used to measure the spread potential of a disease. It is defined as the number of secondary infections produced by a first infection in a homogeneous susceptible population. To calculate  $R_0$  for within host models, modelers use the methods commonly used for epidemiological models. However, the infection dynamics of epidemiology models differ from the simple in-host viral model in that epidemiological models assume direct contact between infected and susceptible individuals while in-host infectious use the virus as an intermediary agent. We examine the meaning of the basic reproduction number in the context of a simple in-host viral infection model.

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