Within-host dynamics

Understanding within-host host-parasite interactions (dynamically)

Lots of molecular biology, genetics (*recognition* mechanisms and *effector* mechanisms), won’t deal with that now.

Interaction between different components of the immune system (modeled at different levels of detail/realism), parasite populations (maybe in multiple compartments?)

Longitudinal data (rare), distributional data.

Bonhoeffer, Coffin, and Nowak (1997)

300-fold decline in viral RNA in some patients, but not clearance

“virus load paradox”: if is initially 50, we would have to reduce it to slightly above 1 but never **below** 1 to see these results.

* add a drug-resistant type
* add mutation
* add immune responses ()
* homeostasis of infectible cells (logistic growth)
* virus-induced killing of uninfected cells (e.g. gp120 shedding)
* differential effects of drug on different types
* distribution of infectibility

Brown et al. (2006)

## References

Bonhoeffer, S, J M Coffin, and M A Nowak. 1997. “Human Immunodeficiency Virus Drug Therapy and Virus Load.” *Journal of Virology* 71 (4): 3275–8. <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC191463/>.

Brown, Sam P, Stephen J Cornell, Mark Sheppard, Andrew J Grant, Duncan J Maskell, Bryan T Grenfell, and Pietro Mastroeni. 2006. “Intracellular Demography and the Dynamics of Salmonella Enterica Infections.” *PLoS Biol* 4 (11): e349. <https://doi.org/10.1371/journal.pbio.0040349>.

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