

*Read:* Chapter 2.3–2.7: Bistability in the lactose operon on *Escherichia coli*: A comparison of differential equation and Boolean network models. By R. Robeva and N. Yildirim, pages 47–73.

1. Recall another model of the *lac* operon that we saw last time:

$$\begin{aligned}f_M &= A, \\f_B &= M, \\f_A &= (B \wedge L_m) \vee L \vee (A \wedge \overline{B}).\end{aligned}$$

Does this model exhibit bistability? Why or why not?

2. Recall our original 3-variable Boolean model of the *lac* operon:

$$\begin{aligned}f_M &= \overline{G_e} \wedge (L \vee L_e), \\f_E &= M, \\f_L &= \overline{G_e} \wedge ((E \wedge L_e) \vee (L \wedge \overline{E})).\end{aligned}$$

Since this model cannot distinguish between basal, medium, and high levels of lactose, it cannot exhibit bistability. Add a new parameter  $L_{em}$  that stands for “at least medium levels of extracellular lactose”. Modify the Boolean functions so the new model exhibits bistable behavior for medium lactose concentrations.

### Summary of relevant literature.

A review of bistability and a discussion of the necessity of positive feedback loops can be found in [Fer02]. Bistable switches govern standard cell processes such as cell fate determination [FM98], cell cycle regulation [PSF03, SMC<sup>+</sup>03], and maintenance of epigenetic traits [HBL03].

A study of bistability in the *lac* operon was published in a 2004 *Nature* paper [OTL<sup>+</sup>04]. The ODE models of the *lac* operon that exhibit bistability were published in [YM03], in [YSHM04]. Bistability was observed in the original Boolean network model of the *lac* operon [VCS11]. The arabinose (*ara*) operon in *E. coli* was recently modeled with a Boolean network and was shown to exhibit bistability [JM17], which was observed experimentally in [SH97].

Another well-studied bistable switch governs the lytic and lysogenic cell cycles in the phage lambda virus. A logical model was published in 1976 [TGL76]. In 1995, a 4-node Boolean network model was proposed in [TT95]. A Boolean model was proposed in [HL11] where extra variables were used to model the time-delays, and this Boolean model exhibits bistability. This paper also contains a Boolean model of the *lac* operon that incorporates time-delays and exhibits bistability.

## References

- [Fer02] J.E. Ferrell. Self-perpetuating states in signal transduction: positive feedback, double-negative feedback and bistability. *Curr. Opin. Cell Biol.*, 14(2):140–148, 2002.
- [FM98] J.E. Ferrell and E.M. Machleder. The biochemical basis of an all-or-none cell fate switch in *xenopus* oocytes. *Science*, 280(5365):895–898, 1998.
- [HBL03] A.D. Hernday, B.A. Braaten, and D.A. Low. The mechanism by which DNA adenine methylase and PapI activate the pap epigenetic switch. *Mol. Cell*, 12(4):947–957, 2003.
- [HL11] F. Hinkelmann and R. Laubenbacher. Boolean models of bistable biological systems. *Discrete Cont. Dyn. Sys. Ser. S*, 4(6):1443–1456, 2011.
- [JM17] A. Jenkins and M. Macauley. Bistability and asynchrony in a Boolean network model of the L-arabinose operon in *escherichia coli*. *Bull. Math. Biol.*, Under review (minor revisions):q-bio.MN/1611.02656, 2017.
- [OTL<sup>+</sup>04] E.M. Ozbudak, M. Thattai, H.N. Lim, B.I. Shraiman, and A. Van Oudenaarden. Multistability in the lactose utilization network of *escherichia coli*. *Nature*, 427(6976):737–740, 2004.
- [PSF03] J.R. Pomerening, E.D. Sontag, and J.E. Ferrell. Building a cell cycle oscillator: hysteresis and bistability in the activation of Cdc2. *Nature Cell Biol.*, 5(4):346–351, 2003.
- [SH97] D.A. Siegele and J.C. Hu. Gene expression from plasmids containing the *arabid* promoter at subsaturating inducer concentrations represents mixed populations. *Proc. Natl. Acad. Sci.*, 94(15):8168–8172, 1997.
- [SMC<sup>+</sup>03] W. Sha, J. Moore, K. Chen, A.D. Lassaletta, C.-S. Yi, J.J. Tyson, and J.C. Sible. Hysteresis drives cell-cycle transitions in *Xenopus laevis* egg extracts. *Proc. Natl. Acad. Sci.*, 100(3):975–980, 2003.
- [TGL76] R. Thomas, A.-M. Gathoye, and L. Lambert. A complex control circuit. Regulation of immunity in temperate bacteriophages. *Eur. J. Biochem.*, 71(1):211–227, 1976.
- [TT95] D. Thieffry and R. Thomas. Dynamical behaviour of biological regulatory networks—ii. immunity control in bacteriophage lambda. *Bull. Math. Biol.*, 57(2):277–297, 1995.
- [VCS11] A. Veliz-Cuba and B. Stigler. Boolean models can explain bistability in the lac operon. *Journal of Computational Biology*, 18(6):783–794, 2011.
- [YM03] N. Yildirim and M.C. Mackey. Feedback regulation in the lactose operon: a mathematical modeling study and comparison with experimental data. *Biophysical J.*, 84(5):2841–2851, 2003.
- [YSHM04] N. Yildirim, M. Santillan, D. Horike, and M.C. Mackey. Dynamics and bistability in a reduced model of the lac operon. *Chaos*, 14(2):279–292, 2004.