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Delay Differential Equations in Epidemiology

Nsoki Mamie Mavinga

Swarthmore College

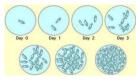
SIMIODE EXPO 2021 Conference February 12, 2021

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"Qu' est-ce que le passé, sinon du présent qui est en retard?" (What is the past, if not the present, which is late?) (Pierre Dac, a French humorist (1893-1975))

Motivation

1. A scientist studying the growth of a population, p(t), may make a very simple assumption that a population grows at a rate directly proportional to its size.



Malthus model:

$$\frac{dp(t)}{dt} = r p(t), \quad t \ge 0$$
$$p(0) = p_0$$

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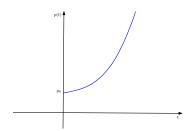
► The solution of the DE is

 $p(t) = p_0 e^{rt}$



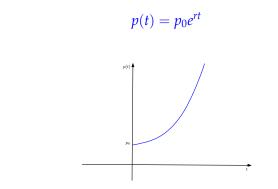
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► The solution of the DE is



► $\lim_{t\to\infty} p(t) = \infty$ whenever r > 0 ("Unlimited growth")

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- 2. Limited population growth (Logistic equation)
- ► In 1838, the Belgian mathematician Pierre Verhulst introduced a model where the population has some self-limitation.

- 2. Limited population growth (Logistic equation)
- ► In 1838, the Belgian mathematician Pierre Verhulst introduced a model where the population has some self-limitation.
- Assume that the per capita growth rate decreases linearly as a function of population.
- The growth equation is given by

$$\frac{dp}{dt} = r\left(1 - \frac{p}{K}\right)p = R(p)p; \qquad p(0) = p_0, \tag{1}$$

where r(> 0) is the intrinsic growth rate; and K(> 0) is the carrying capacity; $R(p) = r(1 - \frac{p}{K})$.

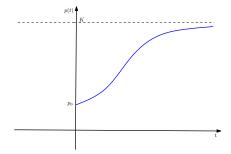
► The Logistic equation (1) assumes that population density negatively affects the per capita growth rate according to $\frac{1}{p}\frac{dp}{dt} = r\left(1 - \frac{p}{K}\right)$ due to environmental degradation.

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The solution is
$$p(t) = \frac{p_0 K}{p_0 - (p_0 - K)e^{-rt}}$$
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3. In 1948, G. E. Hutchinson pointed out that negative effects that high population densities have on the environment influence birth rates at later times due to developmental and maturation delays.

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- 3. In 1948, G. E. Hutchinson pointed out that negative effects that high population densities have on the environment influence birth rates at later times due to developmental and maturation delays.
- Hutchinson modified the logistic equation to incorporate a delay into the growth rate, so R(p) becomes $R(p(t \tau))$:

 $\frac{dp}{dt} = r\left(1 - \frac{p(t-\tau)}{K}\right)p(t) \quad (\text{Hutchinson's eq or logistic DDE}),$ (2)

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where the constant $\tau > 0$ is the time delay.

Example

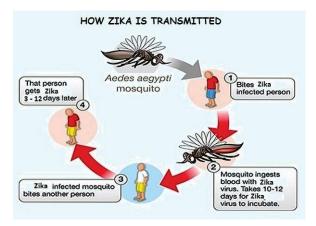
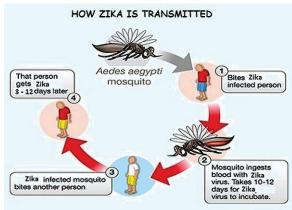


Figure: Transmission Cycle of the Zika virus

http://health.gov.bz/www/images/stories/zika

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Example



Incubation period is the time it takes for the disease to develop inside of a newly infected being (this is the

delay time).

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Figure: Transmission Cycle of the Zika virus

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Definition: Delay differential equation (DDE) is a differential equations in which the current rate of change of the system depends not only on the current state but also on the history of the system. Vector Borne Diseases Model
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- Definition: Delay differential equation (DDE) is a differential equations in which the current rate of change of the system depends not only on the current state but also on the history of the system.
- Consider a simple linear delay-differential equation:

$$y'(t) = -ay(t - \tau), \qquad t > 0,$$
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where $a \in \mathbb{R}$, and $\tau > 0$ is the delay or time lag.

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- A single DDE is capable of producing oscillatory motion, in contrast to a first-order ODE.

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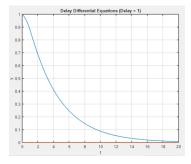


Figure: $y' = -ay(t - \tau)$ with small τ , and a > 0

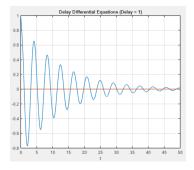


Figure: $y' = -ay(t - \tau)$ with larger τ , and a > 0

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Setting
$$y(\tau t) = u(t)$$
, we get

$$u'(t) = -\beta u(t-1), \tag{4}$$

where $\beta = a\tau$.



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- The only equilibrium solution is $u^*(t) = 0$ for all *t*.
- We look for solutions of the form: $u(t) = ce^{\lambda t}$, where *c* is a constant and the eigenvalues λ are solutions of the transcendental equation:

$$\lambda + \beta e^{-\lambda} = 0$$
 (Characteristics equation) (5)

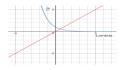
 $\iff \lambda = -\beta e^{-\lambda}$

 Solving and understanding the roots of (5) would be helpful in studying the stability of the equilibrium and the oscillatory behavior of the solution.

STABILITY OF THE ZERO EQUILIBRUIUM

• **Proposition:** Suppose that $\lambda \in \mathbb{R}$.

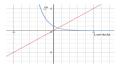
(a) If $\beta < 0$



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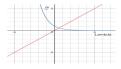


Then (5) has exactly one positive real root λ_0 . $\Rightarrow u(t) = ce^{\lambda_0 t} \to \infty$ as $t \to \infty$, and $u^* = 0$ is unstable.

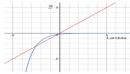
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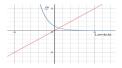
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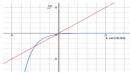
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STABILITY OF THE ZERO EQUILIBRUIUM

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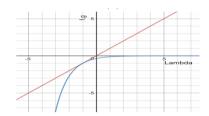


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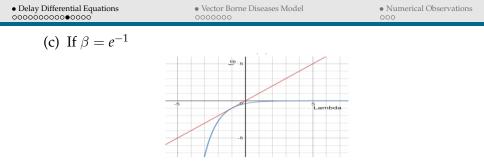


Then it has exactly two negative real roots where $\lambda_1 < -1$ and $-1 < \lambda_2 < 0 \Rightarrow u(t) \rightarrow 0$ as $t \rightarrow \infty$, and $u^* = 0$ is asymptotically stable. • Vector Borne Diseases Model 0000000 Numerical Observations
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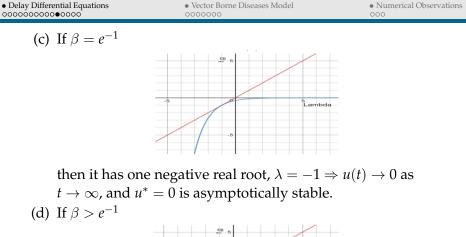
(c) If
$$\beta = e^{-1}$$



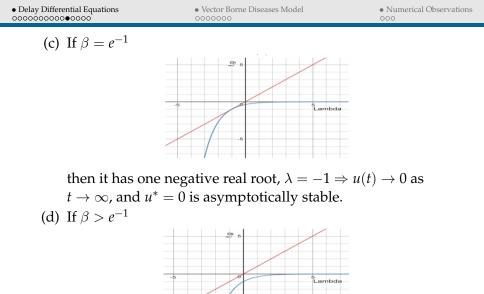
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then it has one negative real root, $\lambda = -1 \Rightarrow u(t) \rightarrow 0$ as $t \rightarrow \infty$, and $u^* = 0$ is asymptotically stable.







then there are no real roots.

• Suppose that $\lambda \in \mathbb{C}$. Set $\lambda = x + iy$.

Separating the real part and imaginary parts of the characteristic equation $\lambda + \beta e^{-\lambda} = 0$, we obtain:

$$\begin{cases} x = -\beta e^{-x} \cos y \\ y = \beta e^{-x} \sin y \end{cases}$$
(6)

$$\Rightarrow \frac{x}{y} = -\cot(y) \Longrightarrow x = -y\cot(y)$$

• Suppose that $\lambda \in \mathbb{C}$. Set $\lambda = x + iy$.

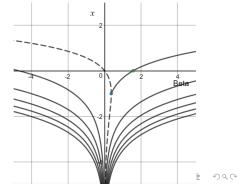
Separating the real part and imaginary parts of the characteristic equation $\lambda + \beta e^{-\lambda} = 0$, we obtain:

$$\begin{cases} x = -\beta e^{-x} \cos y \\ y = \beta e^{-x} \sin y \end{cases}$$
(6)

$$\Rightarrow \frac{x}{y} = -\cot(y) \Longrightarrow x = -y\cot(y)$$

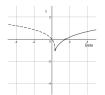
We get the parametric equations:

$$\begin{cases} x = -y \cot(y) \\ \beta = \frac{y}{e^{y \cot(y)} \sin y} \end{cases}$$
(7)



Definition:

The leading roots $\{\lambda_L\} = \{x_L + iy_L\}$ of an equation are those that are such that $x_L > x = Re(\lambda)$ for all $\lambda = x + iy$. **Proposition:**



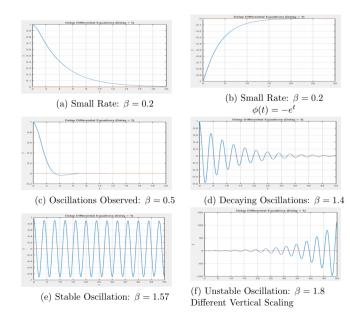
- 1. If $\beta < 0$ then there is only one leading real root that is positive. Therefore, $u^* = 0$ is unstable.
- 2. If $0 < \beta < e^{-1}$ then there is only one leading real root and it is negative. Therefore, $u^* = 0$ is asymptotically stable.
- If e⁻¹ < β < π/2 then there is only one pair of complex conjugate leading roots with negative real part. Therefore, u* = 0 is asymptotically stable.
- 4. If $\beta = \pi/2$ then there is only one pair of complex conjugate leading roots $\pm \frac{\pi}{2}i$. Therefore, $u^* = 0$ is unstable.

Oscillatory behavior: We observe that

- 1. For β small positive then the solution decays exponentially towards the zero equilibrium without any oscillatory behavior.
- 2. When β hits a value round 0.37 ($\approx e^{-1}$), the solution becomes oscillatory but it would still decay to the zero equilibrium.
- 3. When β hits a value around 1.5 ($\approx \pi/2$), oscillations would still take place but the zero equilibrium would no more be stable; the amplitude of the oscillations grows indefinitely as time progress

Theorem: Every Solution of the DDE (4) is oscillatory if and only if $\beta > e^{-1}$.

• Vector Borne Diseases Model 0000000



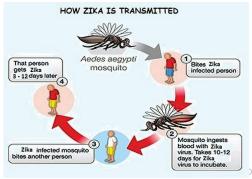
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VECTOR BORNE DISEASES

- Definition: A vector borne disease is a disease transmitted to humans through the bites of an infected arthropod vector (e.g. mosquitoes).
- Malaria and the Zika virus are two well-known examples.
- Understanding the spread of such diseases is vital to their eventual containment and eradication.

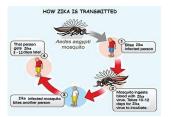
Definition:

Incubation period is the time it takes for the disease to develop inside of a newly infected being (this is the delay time).



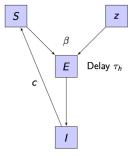
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• Numerical Observations 000



- S = Number of Susceptible Individuals z = Number of Infected Mosquitoes E = Number of Exposed Individuals
- *I* = Number of Infected Individuals
- β = Biting Rate
- c =Disease Recovery Rate

We are interested in the dynamics of infected humans.



 Vector Borne Diseases Model
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Assumptions

- 1. Upon biting an infectious human *I*, with a biting rate β , a susceptible vector becomes infected. And upon biting a susceptible human *S*, an infectious vector *z* infects the bitten human. Infected humans recover from the disease at a rate *c* and they confer no immunity after recovery.
- 2. The size of the human population N is fixed and each human can either be susceptible, exposed, or infected (i.e. S + I + E = N).
- 3. There is an incubation period τ_h in humans, that is a delay between an individual receiving infection and becoming fully infected.

Vector Borne Diseases Model
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- 4. There is an incubation period τ_v in vectors, that is a delay between the vector receiving infection and becoming fully infected.
- 5. The infected vector population is proportional to the infected human population, that is $z(t) = pI(t \tau_v)$.
- 6. The exposed human population (population developing the disease) is proportional to the infected human population, that is E(t) = qI(t).

Vector Borne Diseases Model
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• The Model

From the assumptions, we have the equation:

$$I'(t) = \beta \frac{S(t - \tau_h)}{N} z(t - \tau_h) - cI(t)$$

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Using assumptions 1, 4, and 5 and normalizing, we get a two-lag DDE:

$$I'(t) = [b(1 - eI(t - \tau_h))I(t - \tau_h - \tau_v)] - cI(t),$$
(8)

where $b = \beta p$, e = q + 1, and *I* is the proportion of infected individuals in the population.

When setting $\tau_h = 0$, q = 0, and $\tau_v \neq 0$, we get a previously studied model by Kenneth Cooke (1979): $I'(t) = b[(1 - I(t))I(t - \tau_v)] - cI(t).$

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The equilibria of the model:

•
$$I^* = 0$$
 (the disease-free equilibrium)

►
$$I^* = \frac{1}{e} \left(1 - \frac{c}{b} \right)$$
 (the endemic equilibrium which exists when $R_0 = \frac{b}{c} \ge 1$)

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STABILITY ANALYSIS: APPROACH

• Linearizing around the disease-free zero equilibrium, we derive the following transcendental characteristic equation:

$$\lambda = b e^{(-\tau_v - \tau_h)\lambda} - c \tag{9}$$

Setting $z = (\tau_v + \tau_h)\lambda$, then Eq. (9) becomes:

$$z + a_1 + a_2 e^{-z} = 0, (10)$$

where $a_1 = (\tau_v + \tau_h)c$ and $a_2 = -b(\tau_v + \tau_h)$.

STABILITY ANALYSIS: APPROACH

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where $a_1 = (\tau_v + \tau_h)c$ and $a_2 = -b(\tau_v + \tau_h)$.

• Linearizing around the endemic equilibrium, we derive the equation:

$$\lambda + c = c e^{(-\tau_l - \tau_v)\lambda} + (c - b) e^{-\tau_l \lambda}$$
(11)

Assuming $\tau_v = 0$ and setting $z = \tau_h \lambda$, then Eq. (11) becomes:

$$z + a_1 + a_2 e^{-z} = 0, (12)$$

where $a_1 = \tau_h c$ and $a_2 = -(2c - b)\tau_h$ The stability results follow from the study of the real parts of the roots λ .

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STABILITY ANALYSIS: RESULTS

► The disease-free equilibrium is stable if $R_0 = \frac{b}{c} \le 1$ and unstable if $R_0 > 1$.

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STABILITY ANALYSIS: RESULTS

▶ The disease-free equilibrium is stable if $R_0 = \frac{b}{c} \le 1$ and unstable if $R_0 > 1$.

► The endemic equilibrium is unstable if $0 \le R_0 < 1$. Moreover, if $\tau_v = 0$ and $R_0 > 1$, then there exists a specific b_0 such that $3c < b_0 < \frac{1}{\tau_h} \left[(\pi^2 + \tau_h^2 c^2)^{\frac{1}{2}} + 2\tau_h c \right]$ and a change in stability occurs when $R_0 = \frac{b_0}{c}$. • Vector Borne Diseases Model 0000000 Numerical Observations
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NUMERICAL OBSERVATIONS

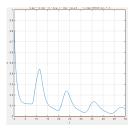


Figure: Stable disease free equilibrium for small values of, transmission rate, *b* Figure: Stable endemic equilibrium for realistic parameters. (From Zika paper by Agusto et al.)

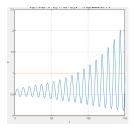


Figure: Unstable Equilibria and Unbounded Solution for even larger values of *b*

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Thank you!

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