# Brilliant: Differential Equations II

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Note: Latex reference: http://tug.ctan.org/info/undergradmath/undergradmath.pdf

# 1 Chapter 1: Basics

# 1.1 Chapter 1: Nonlinear Equations

The two types of problems in this course are:

- Nonlinear equations (several equations on one independent variable)
- Partial differential equations (single equation with several independent variables)

**Linear** equations have solutions like  $y_1, y_2$  that can be combined using any  $c \in \mathbb{R}$  like  $y_1 + cy_2$ .

**Example:** Bacteria in a dish with a lot of food, no deaths

- $b'(t) = r_b b(t), r_b > 0.r_b$  would be the rate of growth.
- This is linear. Reason 1:  $\frac{d}{dt}(y_1+cy_2)=y_1'+cy_2'=r_b(y_1+c_y2)$  since  $y'=r_by(t)$ , and same for y2.
- Also, this works because the solution is  $b(t) = b(0)e^{r_b t}$ , so  $b_1(t) + cb_2(t) = b_1(0)e^{r_b t} + cb_2(0)e^{r_b t} = (b_1(0) + cb_2(0))e^{r_b t}$

**Example:** Logistic equation: Bacteria in a dish with a lot of food, limited by carrying capacity M.

- $b'(t) = r_b b(t) [M b(t)].$
- This is nonlinear. Reason:  $\frac{d}{dt}(y_1'+cy_2')=y_1'+cy_2'=r_b[y_1+cy_2][M-y_1-cy_2]=My_1+Mcy_2-y_1^2-2cy_1y_2-cy_1^2y_2^2$
- $\neq My_1 y_1^2 + Mcy_2 c^2y_2^2$  because of the extra  $-2cy_1y_2$  term.

Sidebar: Note that this equation  $b' = r_b b[M - b]$  is separable, so it can be solved.

- $\frac{db}{dt} = rb[M-b]$
- $\bullet \ \frac{db}{b(M-b)} = rdt$
- $\frac{1}{M}(\frac{1}{b} + \frac{1}{M-b})db = rdt$  after partial fractions work
- $(\ln(b) \ln(M b)) = Mrt + C \Rightarrow \ln(\frac{b}{M b}) = Mrt + C$
- $\frac{b}{M-b} = e^{Mrt}e^C$
- Initial conditions  $b=b(0), t=0 \Rightarrow \frac{b}{M-b} = \frac{b(0)}{M-b(0)} e^{Mrt}$
- $b(1 + \frac{b(0)}{M b(0)}e^{Mrt}) = M\frac{b(0)}{M b(0)}e^{Mrt}$
- $b(M b(0) + b(0)e^{Mrt}) = Mb(0)e^{Mrt}$
- $b = \frac{Mb(0)e^{Mrt}}{M+b(0)[e^{Mrt}-1]}$

This logistic solution will taper off to M at some point. Note that  $\lim_{t\to\infty} b(t) = M$  since the non-exponential terms stop mattering. Also b(t) = M sticks as a constant solution or equilibrium immediately. These equilibria tell us what matters - the long-term behavior of solutions!

Another **Example**: Lotka-Volterra equation pairs: Bacteria (b) and bacteria-killing phages (p), with kill rate k.

- The "product" kb(t)p(t) measures the interactions and kills resulting from this.
- $b'(t) = r_b b(t) kp(t)b(t)$ , or the normal growth rate minus kill rate
- p'(t) = kp(t)b(t) since its population grows as it kills bacteria.
- Equilibria include b = 0, p = 0 and b = 0, p > 0, since these are *constant* solutions, or places where b'(t) = 0, p'(t) = 0.

**Direction fields**, with vector pointing towards  $\langle b'(t), p'(t) \rangle$  (TODO - I think) let us follow the arrows to determine the curve over time. In this case, the bacteria will always go extinct.

However, if we add a new death rate term  $-d_p p(t)$  so  $p'(t) = -d_p p(t) + k p(t) b(t)$ :

- We get an equilibrium at  $b = \frac{d_p}{k}$ ,  $p = \frac{r_b}{k}$ . (Since 0 = b'(t) = rb kpb,  $(\Rightarrow pk = r)$ , 0 = p'(t) dp + kpb,  $(\Rightarrow bk = d)$ )
- But otherwise the solutions swirl around this point. This is called a **cycle**. TODO What is a **limit cycle**?

Note that there are systems where the "solution particle" neither reaches an equilibrium or cycles around one point. The **Lorenz system** famously has this owl-eye shaped double attractor (an example of **strange sets**) where initially close particles diverge unpredictably if the constants  $\rho$ ,  $\sigma$ , b are chosen right:

- $x'(t) = \sigma(y x)$
- $y'(t) = x(\rho z) y$
- z'(t) = xy bz
- TODO

# 1.2 Chapter 1.2: PDEs

Many methods of attack for PDEs

- Separation of variables
- Power series (Note: did we actually touch on this?)
- Fourier Transform

**Example**: Standing wave, where one end of a rope is fixed.

- Vertical displacement from a line of rope: u(x,t) depends on where (x) and when (tt).
- Rope's wave equation is  $u_{tt} = v^2 u_{xx}$ , where v is the "constant wave speed", and the others are the space, time partials.
- Note that  $u = \cos(vt)\sin(x)$  and  $u = \sin(vt)\cos(x)$  both work.
- If you guess the solution has split variables like u = X(x)Y(y)T(t), then, upon substitution and division by X(x)Y(y)T(t),  $\frac{\delta^2 u}{\delta t^2} = v^2 \left[\frac{\delta^2 u}{\delta x^2} + \frac{\delta^2 u}{\delta y^2}\right]$  yields  $\frac{T''(t)}{T(t)} = v^2 \left[\frac{X''(x)}{X(x)} + \frac{Y''(y)}{Y(y)}\right]$
- This method may or may not work. But if it does, it means that since x, y, and t are independent variables, each individual piece must be constant.
- So, for example, if we know  $\frac{X''(x)}{X(x)} = -4\pi^2$ , we can get to  $X(x) = \sin(2\pi x)$
- The wave equation is similar in 3D:  $u_{tt} = v^2[u_{xx} + u_{yy} + u_{zz}]$ , or using the Laplacian,  $u_{tt} = v^2 \nabla^2 u$ . Here, u measures not displacement but expansion/compression of air at (x, y, z), time t.

Using Fourier transforms helps turn difficult PDEs into an easier problem like an ODE. Fourier transforms work best when

- The domain is all of  $\mathbb{R}^n$
- The function *u* vanishes at infinity.

The Fourier transform changes the domain of x to that of  $\omega$ . It comes with the (highly simplified) rule (see Vector Calculus course):  $F\left[\frac{\delta f}{\delta x}\right] = i\omega F[f]$ . **Example**: Drunkard's walk.

- One dimensional: moves left or right in a random way. Starts at x = 0, t = 0.
- u(x,t) is probability of being at point x at time t. Naturally,  $\int_{x=-\infty}^{x=\infty} u(x,t) dx = 1$ .
- Also, it obeys the 1-dD diffusion equation  $\frac{\delta u}{\delta t} = \frac{\delta^2 u}{\delta x^2}$
- The Fourier transform doesn't affect t at all.
- So by taking Fourier transform of both sides of diffusion equation we get

$$-F(u_t) = \frac{\delta}{\delta t}F(u)$$
 since F doesn't care about t.

$$-\frac{\delta^2 u}{\delta x^2} = i\omega F(\frac{\delta u}{\delta x}) = -\omega^2 F(u)$$

- So 
$$\frac{\delta}{\delta t}F(u) = -\omega^2 F(u)$$

– This is solvable as 
$$F(u) = ce^{-\omega^2 t}$$
. Take it on faith that  $c = \frac{1}{2\pi}$  for now. TODO

– Known fact: 
$$F[Ae^{-\frac{ax^2}{2}}] = \sqrt{\frac{1}{2\pi a}}Ae^{\frac{-\omega^2}{2a}}$$

– This means 
$$t = \frac{1}{2a}$$
 and  $a = \frac{1}{2t}$ 

$$-F(u) = \frac{1}{2\pi}e^{-\omega^2t}, F[Ae^{-\frac{ax^2}{2}}] = \sqrt{\frac{1}{2\pi a}}Ae^{\frac{-\omega^2}{2a}} \text{ so } u = Ae^{\frac{-ax^2}{2}}$$

– Solving, you get 
$$A=\sqrt{\frac{1}{4\pi t}}, a=\frac{1}{2t},$$
 so  $u(x,t)=\sqrt{\frac{1}{4\pi t}}e^{-\frac{x^2}{4t}}$ 

# 2 Chapter 2: Nonlinear Equations

#### 2.1 2.1: Lotka-Volterra I

Major ideas:

• phase plane: TODO

• nullcline: TODO

• direction field: TODO

• equilibria: TODO

**Example:** Bacteria vs. phages (again)

- Bacteria unrestrained grow in proportion to their population, so  $\frac{db}{dt} = r_b b(t)$  (solved:  $b(t) = b(0)e^{r_b t}$ )
- Phages unfed decrease in proportion to current size, so  $\frac{dp}{dt} = -d_p p(t)$  (solved:  $p(t) = p(0)e^{-d_p t}$ )
- Bacteria die with likelihood of meeting a phage, and phages increase with likelihood of meeting a bacterium. So the set of equations, for constant k, becomes:
  - $-b'(t) = r_b b(t) k b(t) p(t)$
  - $-p'(t) = -d_n p(t) + kb(t)p(t)$
  - The product of p and b makes our equations nonlinear (WHY?)
  - I guess, very generally,  $b_1p_1 = k$ ,  $b_2p_2 = k$ , but  $(b_1 + b_2)(p_1 + p_2) = b_1p_1 + b_2p_2 + b_1p_2 + b_2p_1 = 2k + b_1p_2 + b_2p_1 \neq 2k$ , so the last two "mixed" terms mean you can't just add solutions  $(b_1, p_1)$  and  $(b_2, p_2)$ .

## General thoughts on this solution:

- So a solution (b(t), p(t)), traces out a curve on the bp-phase plane (b is x-axis, p is y-axis) as time (unrepresented in the plane) continues.
- If we add a unit tangent vector at every point (B, P) aligned with  $(b'(t), p'(t)) = (r_b B k B P, -d_p P + k B P)$ , we can follow the arrows to see the solution over time.
- The above is called a direction field
- This is sometimes hard to sketch analytically, so we can look to the **nullclines**: places where one of the components of the direction field is zero.
- In this case,  $r_bB kBP = (r_b kP)B = 0$  when P = 0 or  $P = \frac{r_b}{k}$ , and  $-d_pP + kBP = (kB d_p)P = 0$  when P = 0 or  $B = \frac{d_p}{k}$ .
- The **upshot of nullclines** (since we don't care about  $P, B \leq 0$ ): The lines  $B = \frac{d_p}{k}, P = \frac{r_b}{k}$  divide the plane into pieces where the components of this (continuous) function pair can't change sign.
- For instance,  $B > \frac{d_p}{k}$ ,  $P < \frac{r_b}{k}$  means  $r_b b k b p > 0$ ,  $-d_p p + k d p > 0$ , so both populations are growing here. This helps to sketch the curve.
- The curve looks like a counterclockwise whirlpool around the  $(B, P) = (\frac{d_p}{k}, \frac{r_b}{k})$ . (bacteria grow with low but growing phages; bacteria decrease as phages overwhelm; both decrease as phages starve; bacteria start coming back)
- The center point is a (constant **equilibrium**) solution, and other solutions swirl around it but don't get attracted or repelled.

There are a few types of equilibria:

- This one is a **center** around which solutions circle.
- A stable equilibrum would see small upsets come back to an unchanging state.
- An unstable equilibrum would see small upsets create wildly divergent paths.

## 2.2 2.2: Lotka-Volterra II

In the Bacteria-Phage system, we can't yet prove everything rotates around the **center**. Let's do that.

Developing a **conserved quantity** will help to do that. **Example**: Block on a horizontal spring with mass m, spring constant  $k_s$ :

- x(t): Displacement from rest position.
- $v(t) = \frac{dx}{dt}$ : Horizontal velocity
- $\frac{dv}{dt} = -\frac{k_s}{m}x(t)$  by Hooke's law, I think.
- Suppose there's some Energy function E(x,v). By chain rule  $\frac{d}{dt}E(x(t),v(t)) = \frac{dE}{dx}\frac{dx}{dt} + \frac{dE}{dv}\frac{dv}{dt}$
- =  $\frac{dE}{dx}v \frac{k_s}{m}\frac{dE}{dv}x$ . If we set E as conserved, as in E'(t) = 0, then  $\frac{dE}{dx}v = \frac{k_s}{m}\frac{dE}{dv}x$
- We can eyeball and see that  $E = \frac{1}{2}k_sx^2 + \frac{1}{2}mv^2$  solves this equation, or we can assume  $E(x,v) = F(x) + G(v) \Rightarrow 0 = E'(t) = F'(x)v \frac{k_s}{m}G'(v)x = 0$  from the above equations and guess from there.
- This means in the xv phase space, that there's a fixed E such that the particle follows the ellipse  $E = \frac{1}{2}k_sx^2 + \frac{1}{2}mv^2$  in phase space around the solution point (0,0).

Extended Example: Continuing on finding a conserved quantity for Bacteria / Phage:

- We need to find U(b(t), p(t)) such that U'(t) = 0, or by chain rule  $\frac{\delta U}{\delta b} \frac{\delta b}{\delta t} + \frac{\delta U}{\delta p} \frac{\delta p}{\delta t} = 0$
- Subbing in,  $\frac{\delta U}{\delta b}[r_b b kbp] + \frac{\delta U}{\delta p}[-d_p p + kbp] = 0$
- A hint suggests finding U such that  $\frac{\delta U}{\delta b} = -\frac{d_p}{b} + k$ ,  $\frac{\delta U}{\delta p} = -\frac{r_b}{p} + k$  to make terms cancel.
- Integrating these gives us U as both  $-d_p \ln(b) + kb + Q(p)$  and  $-r_b \ln(p) + kp + R(b)$  so  $U = -d_p \ln(b) r_b \ln(p) + kb + kp$ . This weird curve consistutes a level set in pb-space upon which a solution sits.
- The spring example has an elliptic paraboloid solution. There's an absolute minimum (E = 0 at (0,0)) but level sets become closed loops away from it.

- For the Lotka example, there is a critical point  $(\nabla U = \vec{0})$  when  $\nabla U(b, p) = (\frac{\delta U}{\delta b}, \frac{\delta U}{\delta p}) = (k \frac{d_p}{b}, k \frac{r_b}{p})$ , which is (0, 0) at our known center  $(\frac{d_p}{k}, \frac{r_b}{k})$
- Showing that we always increase gong away from the point  $(\frac{d_p}{k}, \frac{r_b}{k})$  should guarantee us closed level sets.
- One method: Assume we're picking a unit vector  $\vec{v} = \langle \hat{v}_b, \hat{v}_p \rangle$  so that our line from our center is  $\vec{v} = \langle \frac{d_p}{k} + tv_b, \frac{r_b}{k} + tv_b \rangle$ . U = F(b) + G(p) in this case, so sub the b part into F to get  $F(\frac{d_p}{k} + t\hat{v}_b) = d_p[1 \ln(\frac{d_p}{k} + t\hat{v}_b)] + kt\vec{v}_b$ . Taking derivative of that w.r.t t shows it is always positive. Same goes for the G(p) portion of U.
- Another (DF) method: Note that  $\nabla U = (k \frac{d_p}{b}, k \frac{r_b}{p})$ 's grad (second derivative) is always positive. So derivative always has positive curvature (maybe using that term wrong), and we'll always increase around this point.
- Also, we know that the particle travels around the level set (loop) and doesn't reverse course, because then, b'(t) = p'(t) = 0, and we only have that at the center point (nullcline intersection).

#### 2.3 2.3: Linearization

**Extended Example**: Suppose there's a limit to bacterial growth, so we cap our population at  $M_b$ .

- If  $b(t) << M_b$ , things should be similar. If b(t) is nearly  $M_b$ , then growth should approach 0. So, this implies  $\frac{db}{dt} = r_b b(t) \to \frac{db}{dt} = r_b b(t) (1 \frac{b(t)}{M_b})$ . Note: This isn't the only possibility but we'll use it.
- This updates our Lotka-Volterra model to something more complicated:

$$-b'(t) = r_b b(t) (1 - \frac{b(t)}{M_b}) - kb(t)p(t)$$

$$-p'(t) = -d_p p(t) + kb(t)p(t)$$

- Other than b = 0, p = 0, the meaningful nullclines are solved by setting b'(t) = 0 (yielding  $r_b(1 \frac{b}{M_b}) kp = 0$ ) and p'(t) = 0 (yielding  $b = \frac{d_p}{k}$ )
- Note: We'll clean up through some MAGIC non-dimensionalization (how to derive?) to simplify:

$$-x(t) = \frac{1}{M_b}b(\frac{t}{r_b}), y(t) = \frac{k}{r_b}(\frac{t}{r_b}), \alpha = \frac{d_p}{r_b}, \beta = \frac{kM_b}{r_b}$$

- Gives us new equations:  $\frac{dx}{dt} = x(t)[1-x(t)] x(t)y(t), \frac{dy}{dt} = -\alpha y(t) + \beta x(t)y(t)$
- And new nullclines:  $x + y = 1, x = \frac{a}{b}$

- So there's an equilibrium point in the positive xy quadrant if:  $y = 1 x = 1 \frac{\alpha}{\beta}$  and y > 0 implies  $1 \frac{\alpha}{\beta} > 0 \Rightarrow \frac{\alpha}{\beta} < 1$
- Looking at the direction field, it appears solutions swirl around and are attracted into the center point  $(\frac{\alpha}{\beta}, 1 \frac{\alpha}{\beta})$ , making it a **stable equilibrum**

This is similar to the block-spring example, if a damping term  $-\frac{\gamma}{m}v$  is added.

- $\frac{dx}{dt} = v, \frac{dv}{dt} = -\frac{k_s}{m}x \frac{\gamma}{m}v$
- This can be thought of in matrix terms:  $\frac{d}{dt} \begin{pmatrix} x(t) \\ v(t) \end{pmatrix} = \begin{pmatrix} 0 & 1 \\ -\frac{k_s}{m} & -\frac{\gamma}{m} \end{pmatrix} \begin{pmatrix} x(t) \\ v(t) \end{pmatrix}$  Call the matrix A.
- From Diff Eq I, the solution is exp(tA) (matrix exponential), making x(t) a linear combination of  $e^{\lambda t}$  or possibly  $te^{\lambda t}$  terms, with the eigenvalues as  $\lambda s$ .
- The eigenvalues in this case, using the quadratic formula, could be:
  - Two real, distinct, negative roots. So, these  $e^{\lambda t}$  terms decay, and x(t) levels off.
  - Two distinct complex roots with real part  $-\frac{\gamma}{2m} < 0$ . This ends up being some sines and cosines multiplied by  $e^{-\frac{\gamma t}{2m}}$ , which decays too.
  - Finally, if we have a repeated negative real eigenvalue, we have solution  $x(t) = Ae^{-\frac{\gamma t}{2m}} + Bte^{-\frac{\gamma t}{2m}}$ , also decaying.
  - So any disturbance in the spring will oscillate and come to rest at x(t) = v(t) = 0 quickly.

So with linear systems  $\vec{x}'(t) = A\vec{x}(t)$ , the eigenvalues determine what happens around the equilibrium point. However, the **bacteria-phage model is non-linear**. Here is **how we linearize** for nearby solutions in a nonlinear system:

- Set small disturbance  $\delta x(t) << 1, \delta y(t) <<$  so  $x(t) = \frac{\alpha}{\beta} + \delta x(t), y(t) = 1 \frac{\alpha}{\beta} + \delta y(t)$
- Since they're small, all powers like  $\delta x(t)^2$  and  $\delta x(t)\delta y(t)$  are considered basically zero.
- So substitute  $x(t) \to \frac{\alpha}{\beta} + \delta x(t), y(t) \to 1 \frac{\alpha}{\beta} + \delta y(t)$  into our  $\frac{dx}{dt}$  and  $\frac{dy}{dt}$  equations.
- This gives us the A solving  $\frac{d}{dt} \begin{pmatrix} \delta x(t) \\ \delta y(t) \end{pmatrix} = A \begin{pmatrix} \delta x(t) \\ \delta y(t) \end{pmatrix}$ , which is  $A = \begin{pmatrix} -\frac{\alpha}{\beta} & -\frac{\alpha}{\beta} \\ \beta \alpha & 0 \end{pmatrix}$  after working through the substitution.

 Finding the eigenvalues here yields the same situation as the block-spring example: decays in all situations.

It turns out through the **Hartman-Grobman Theorem** that  $\vec{x}'(t) = \vec{F}(\vec{x}(t))$ , for some continuously differential vector field F, if we linearize near equilibrium  $x_0$ , then what falls out of this A approach works if the eigenbalues aren't all purely imaginary.

It turns out the uncapped bacteria system from before looks like  $\frac{d}{dt} \begin{pmatrix} \delta x(t) \\ \delta y(t) \end{pmatrix} =$ 

 $\begin{pmatrix} 0 & -1 \\ \alpha & 0 \end{pmatrix} \begin{pmatrix} \delta x(t) \\ \delta y(t) \end{pmatrix}$ , with characteristic equation  $\lambda^2 + \alpha = 0, \alpha > 0$ . This means both values are imaginary, and we had to use the conserved quantity appraoach!

#### 2.4 2.4: Hartman-Grobman Theorem

**Extended Example**: Consider a phage that dies off quicky:

- $\frac{db}{dt} = r_b b(t) k_b b(t) p(t)$ ,  $\frac{dp}{dt} = -r_p p(t) = 0 \cdot b(t) p(t)$ , where  $k_p$  is the zero (phages don't increase), and  $k_b$  is still the kill factor for the bacteria.
- In this base, b(t) = p(t) = 0 is the only equilibrium
- Non-dimensionalize as  $x(t) = b(\frac{t}{r_b}), y(t) = \frac{k_b}{r_b} p(\frac{t}{r_b}), \alpha = \frac{r_p}{r_b}$
- This makes the equations  $x'(t) = x(t) x(t)y(t), y'(t) = -\alpha y(t)$ , and the nullclines therefore x(t) = 0, y(t) = 1, y(t) = 0
- Looking at this six-section dierection field, we see that solutions exactly on the y-axis are attracted to equilibrium (0,0), and other are repelled.
- This makes sense since if the bacteria is 0, the phage die and approach (0,0), otherwise the bacteria multiply and win (so it's a *saddle point*)
- The way to tell: linearize the equations.  $\frac{d}{dt} \begin{pmatrix} x(t) \\ y(t) \end{pmatrix} \approx \begin{pmatrix} 1 & 0 \\ 0 & -\alpha \end{pmatrix} \begin{pmatrix} x(t) \\ y(t) \end{pmatrix}$  since, if x(t), y(t) << 1, x(y)t(t) = 0.
- Then the eigenvalues are  $\lambda = 1, -\alpha$  so the solution is  $Ae^t, Be^{-\alpha t}$  for x(t), y(t) (TODO respectively?) Hartman-Grobman ensures this is the general solution.

However, let's solve directly and see if we come to the same result.

- $y'(t) = -\alpha y(t), y(0) = y_0 \Rightarrow y(t) = y_0 e^{-\alpha t}$
- With this in hand,  $\frac{dx}{dt} = x(t) x(y)y(t) = x(t)[1 y_0e^{-\alpha t}], x(0) = x_0$  separates out to

$$-\frac{dx}{x} = [1 - y_0 e^{-\alpha t}]dt$$

$$-\ln(x) = \left[t + \frac{y_0}{\alpha}e^{-\alpha t}\right] + C$$

$$-x = e^C e^t \exp\left(\frac{y_0}{\alpha}e^{-\alpha t}\right)$$

$$-x(0) = x_0 \Rightarrow e^C = x_0 e^{-\frac{y_0}{\alpha}}$$

$$-\Rightarrow x(t) = x_0 e^t \exp\left(\frac{y_0}{\alpha}(e^{-\alpha t} - 1)\right)$$

But how do we deform the phase plane so this looks linear? We need some mapping  $\vec{h}(x,y) = \langle u(x,y), v(x,y) \rangle$  that is continuous and invertible (so we don't "damage" the phase plane). This is called a **homeomorphism**.

- So near the equilibrium (0,0), the equations  $y'(t) = -\alpha y(t), y(0) = y_0 \Rightarrow y(t) = y_0 e^{-\alpha t}$  linearized for  $\delta x, \delta y$  must be similar to those for u(x(t), y(t), v(x(t), y(t)))
- This means we need  $\frac{du}{dt} = u, \frac{dv}{dt} = -\alpha v$
- After doing the substitution, we see that  $v = v_0 e^{-\alpha t}$  exactly mimics  $y(t) = y_0 e^{-\alpha t}$  for the phage solution. So we take v = y.
- Therefore, we know that since  $u = u_0 e^t$  and  $x(ty) = x_0 \exp(t + \frac{y_0}{\alpha}(e^{-\alpha t} 1))$ , that we need  $u(x(t), y(t) = u(x_0, y_0)e^t$
- And this is satisfied if we guess  $u(x,y) = xe^{-y}\alpha$  and work it out.
- This function  $\vec{h}(x,y) = (u,v) = \langle xe^{-\frac{y}{\alpha}}, y \rangle$  is invertible by  $(x,y) = \langle ue^{\frac{v}{\alpha}}, v \rangle$ , which is continuous.

#### 2.5 2.5: Application - Lasers

Lasers create excited atoms, which then emit photos while transitioning to an unexcited state. This system has a close analogue with the previous phages (like photons) and bacteria (like atoms) model.

- n(t): number of photons in the laser;  $r_g$ : rate of photons gained (created by excited atoms transitioning to unexcited state);  $r_l$ : rate of photons lost (emitted)
- $\Rightarrow \frac{dn}{dt} = r_g r_l$  by definition.
- We can assume we're losing a constant k (kill?) portion of photons per unit time, so  $\frac{dn}{dt} = r_q kn(t)$
- e(t): number of excited atoms (that will maybe create photons). Atoms are excited by external energy pump.
- Excited atoms radiate when meeting a photon (which survives the meeting)
- So we can use the same setup from the bacteria: with I the constant of meeting (intersect?),  $r_g = Ie(t)n(t) \Rightarrow n'(t) = Ie(t)n(t) kn(t)$

Mini example: Assume no photons leave (cap the end of the laser)

- k = 0 in this scenario.
- So every meeting creates one more photon  $(n \to n+1)$  while enervating one excited atom  $(e \to e-1)$ . This implies, equivalently:
  - -e+n is a conserved quantity,

$$-e(t) + n(t) = e(0) + n(0),$$

$$- [e(t) + n(t)]' = 0$$

– Then, if 
$$k = 0$$
,  $n'(t) = Ie(t)n(t) - kn(t)$ , and coupled with  $e'(t) + n'(t) = 0$  above, we have  $e'(t) = -Ie(t)n(t)$ 

Extended example: Atoms spontaneously lose energy. This is actually what happens

- $\bullet$  From quantum physics, we have a rate s of atoms just (s)pontaneously losing energy.
- We also have an energy (p)ump that energizes atoms with quantity p.
- Then, our change in (e)xcited atoms is e'(t) = p s Ie(n)(t)
- So our final laser equations are e'(t) = p s Ie(n)(t), n'(t) = Ie(t)n(t) kn(t)
- If we want to find the smallest p guaranteeing  $n \ge 1$  (there's at least one photo) at equilibrium (e'(t) = n'(t) = 0):

$$-n'(t) = 0 \Rightarrow Ien = kn \Rightarrow n(Ie - k) = 0.$$
 If  $n \neq 0, \Rightarrow e = \frac{k}{I}$ 

$$-e'(t) = 0 \Rightarrow Ien = p - se$$

– Together, 
$$p - se = Ien = kn \Rightarrow kn + se = p \Rightarrow kn + s\frac{k}{I} = p$$

$$-n \ge 1 \Rightarrow p \le k + \frac{ks}{I}$$

- **Another tactic**: We could also assume we *start out at equilibrium*, so  $n_0, e_0$  are constant solutions.

- Solving 
$$n' = 0 = Ie_0n_0 - kn_0, e' = 0 = Ie_0n_0 - se_0 + p$$
, we find equilibria  $n_0 = \frac{p}{k} - \frac{s}{I}, e_0 = \frac{k}{I}$ 

- Then, 
$$n_0 \ge 1 \Rightarrow \frac{p}{k} - \frac{s}{I} \ge 1 \Rightarrow p \ge k + \frac{ks}{I}$$

Non-dimensionalization time:

• Scale against  $e_0(=\frac{k}{I}), n_0(=\frac{p}{k}-\frac{s}{I})$  like this:  $x(t)=\frac{n(\alpha t)}{n_0}, y(t)=\frac{e(\alpha t)}{e_0}$ 

- NOTE: What does this do? This makes (1,1) the equilibrium, as  $x(t) = \frac{n_0}{n_0} = 1, y(t) = \frac{e_0}{e_0} = 1$ !
- What  $\alpha$  lets us the n' = Ien kn, e' = -Ien se + p and write

$$-\frac{dx}{dt} = x(t)y(t) - x(t)$$

$$-\frac{dy}{dt} = \frac{1}{k}(\frac{pI}{k} - s)[1 - x(t)y(t)] + \frac{s}{k}[1 - y(t)]$$

$$-x' = \frac{\alpha n'(\alpha t)}{n_0} = xy - x = \frac{Ie(\alpha t)n(\alpha t)}{kn_0} - \frac{n(\alpha t)}{n_0}$$

$$-\frac{\alpha Ien - \alpha kn(\alpha t)}{n_0} = \frac{Ie(\alpha t)n(\alpha t)}{kn_0} - \frac{n(\alpha t)}{n_0}$$

$$-\alpha Ie - \alpha k = \frac{Ie(\alpha t)}{k} - 1 \Rightarrow \alpha (Ie - k) = \frac{Ie - k}{k} \Rightarrow \alpha = \frac{1}{k}$$

- This solves the x equation, and I suppose it can be validated in the y equation (tediously).
- If we chunk up our (somehow positive?) constants as  $c = \frac{1}{k}(\frac{pI}{k} s), d = \frac{s}{k}$ , we end up with y' = c[1 xy] + d[1 y]
- We only care about x, y > 0, so x' = 0 = xy x = x(y 1) implies y = 1 is a nullcline
- $-y'=0=c[1-xy]+d[1-y]=c-cxy+d-dy\Rightarrow c+d=y(d+cx)\Rightarrow y=\frac{c+d}{d+cx}$  a scaled and shifted hyperbola.

#### Look at the solutions:

- It appears we have a counterclockwise swirl around (1,1), and nearby solutions tend toward this equilibrium.
- Hartman-Grobman: rewrite our linearized solution in neighborhood of (1,1) as  $x(t) = 1 + \delta x(t), y(t) = 1 + \delta y(t)$

- Using 
$$x' = xy - x$$
,  $y' = c[1 - xy] + d[1 - y]$  and  $\frac{d}{dt} \begin{pmatrix} \delta x(t) \\ \delta y(t) \end{pmatrix} = A \begin{pmatrix} \delta x(t) \\ \delta y(t) \end{pmatrix}$ , we can solve and write  $A = \begin{pmatrix} 0 & 1 \\ -c & -c - d \end{pmatrix}$ 

- Eigenvalues:  $\lambda = \frac{1}{2}(-c d \pm \sqrt{(c+d)^2 4c})$ 
  - \* If square root term is zero, we have repeated eigenvalue, so  $\delta x(t), \delta y(t)$  are combos of  $e^{-\frac{c+d}{2}}, te^{-\frac{c+d}{2}}$ , which decays
  - \* If square root term is greater than zero, we have two distinct real, negative eigenvalues (since c, d are positive), so this decays.

- \* If square root term is less than zero, we have distinct complex eigenvalues, but combos of  $e^{-\frac{c+d}{2}}\cos(\frac{1}{2}\sqrt{-(c+d)^2+4c}), e^{-\frac{c+d}{2}}\sin(\frac{1}{2}\sqrt{-(c+d)^2+4c})$  decay too
- \* Note: I suppose Hartman-Grobman can't work in purely imaginary eigenvalue scenario, because these kinds of functions don't converge or diverge without a term outside the sin or cos
- \* And in any case, since these lambdas aren't strictly imaginary, Hartman-Grobman works.

# 2.6 2.6: Liapunov Equations

We had some intuition that "nearby" solutions would fall into an equilibrium, but what does "nearby" mean? **Liapunov Equations** help us here. What is the "basin of attraction"?

- Suppose we turn the pump off (p = 0), and set spontaneous enervation equal to photon leak s = k.
- (TODO?) Somehow we can rescale to  $\frac{dx}{dt} = Ie(t)n(t) kn(t), \frac{dy}{dt} = -Ie(t)n(t) kn(t)$  which (TODO??) gives us  $\frac{dx}{dt} = xy x, \frac{dy}{dt} = xy y$
- This means equilibria (x' = y' = 0) exist at (0,0),(1,1)
- If we're turning the pump off, we're looking at equilibrium (0,0). Linearizing, we get  $x' = -\delta x, y' = -\delta y$ , so a matrix of  $\begin{pmatrix} -1 & 0 \\ 0 & -1 \end{pmatrix}$
- With repeated non-imaginary (H-G applies!) eigenvalues -1, -1, we can see that both  $e^{-t}$ ,  $te^{-t}$  decay, and we get sucked into the origin.

But how do we prove this? Let's find a conserved quantity U'(x(t), y(t)) = 0

- $U'(x(t), y(t)) = \frac{\delta U}{\delta x} \frac{\delta x}{\delta t} + \frac{\delta U}{\delta y} \frac{\delta y}{\delta t} = \frac{\delta U}{\delta x} x(y-1) + \frac{\delta U}{\delta y} y(x-1)$
- Setting  $\frac{\delta U}{\delta x}x = -x + 1$ ,  $\frac{\delta U}{\delta y}y = y 1$  makes this zero
- Solving those two by separating variables and combining, we get  $U = -x + y + \ln(|\frac{x}{y}|)$
- So if we're stabiling f = (x-y) (why?), we see  $(x-y)' = x'-y' = (xy-x)-(xy-y) = x-y = f \Rightarrow f = e^{-t}$
- With  $x(0) = x_0, y(0) = y_0 \Rightarrow f(0) = x_0 y_0, f = x(t) y(t) = (x_0 y_0)e^{-t}$
- How to express y(t) while eliminating x(t), knowing  $x(y) y(t) = (x_0 y_0)e^{-t}$  and  $U(x,y) = y x + \ln(|\frac{x}{y}|)$  is conserved? The trick:  $U(x_0,y_0) = U(x,y)$  since it doesn't change!

$$-y_0 - x_0 + \ln(\frac{x}{y}) = y - x + \ln(\frac{x}{y}) = -(x_0 - y_0)e^{-t} + \ln(\frac{x}{y})$$

$$- (1 - e^{-t})(y_0 - x_0) = \ln(\frac{x/y}{x_0/y_0})$$

– Defining for convenience, 
$$f = \exp((1 - e^{-t})(y_0 - x_0))$$
, then  $f \frac{y}{y_0} = \frac{x}{x_0}$ 

– Sub in to 
$$x - y = (x_0 - y_0)e^{-t}$$
:  $y[\frac{x_0}{y_0}f - 1] = (x_0 - y_0)e^{-t}$ 

- Solve for 
$$y : y = \frac{y_0(x_0 - y_0)e^{-t}}{x_0f(t) - y_0}$$

- Combine with above to get 
$$x = \frac{x_0(x_0 - y_0)e^{-t}f(t)}{x_0f(t) - y_0}$$

- So with equilibria (0,0),(1,1), the direction field computer plot shows us attracted to (0,0) (no laser action) pretty much anywhere left and down from (1,1) in the x,y phase plane.
- Apparently the linearized solutions near 0,0 are  $x_{lin} = x_0 e^{-t}$ ,  $y_{lin} = y_0 e^{-t}$  (WHY?)
- Looking above, if  $(x_0 y_0) \approx 0$ , then  $f(t) \approx 1$ , and  $x, y \to x_{lin}, y_{lin}$

On to **Liapunov** functions, which will tell us perhaps the size of the "basin of convergence", unlike Hartman-Grobman, which just says there is a neighborhood.

# A **Liapunov** function U(x,y) is

- Continuously differentiable
- With a unique minimum  $(x_0, y_0)$ , usually aligned to be U's only zero.
- $U'(x(t), y(t)) \le 0$ . Everything "flows downhill";
- Tailor made for the problem, hard to find.

## Back to the rescaled laser example

$$\bullet \ x'(t) = x(t)y(t) - x(t)$$

• 
$$y'(t) = c[1 - x(t)y(t)] + d[1 - y(t)], c, d > 0$$

- Analogy: The damped-block spring system 
$$\frac{d}{dt} \begin{pmatrix} x(t) \\ v(t) \end{pmatrix} = \begin{pmatrix} 0 & 1 \\ -\frac{k_s}{m} & -\frac{\gamma}{m} \end{pmatrix} \begin{pmatrix} x(t) \\ v(t) \end{pmatrix}$$

- When 
$$\gamma = 0$$
, we know  $E(x, v) = \frac{1}{2}k_sx^2 + \frac{1}{2}mv^2$  is conserved when looking at  $E'$ 

$$-\gamma = 0 \Rightarrow x' = v, v' = -\frac{k_s}{m}x$$

$$-\frac{dE}{dt} = (\frac{1}{2}k_sx^2 + \frac{1}{2}mv^2)' = 0 \text{ since } \frac{1}{2}(k_sxx' + mvv') = \frac{1}{2}(kxv + mv\frac{-k}{m}x) = 0$$

- But if 
$$\gamma \neq 0$$
,  $\frac{d}{dt}E(x(t), v(t)) = \frac{d}{dt}[\frac{1}{2}k_sx^2 + \frac{1}{2}mv^2] = kxx' + mvv'$ 

- $= kx(v) + mv(\frac{-k_s}{m}x \frac{\gamma}{m}v) = -\gamma v(t)^2 = \frac{dE}{dt}$
- Total spring energy is then decreasing in the fluid.
- Brilliant has Cool visualization of spiraling down into the "bowl" of x, y with E as the z dimension, equilibrium (0,0,0)
- We need to choose a  $\gamma$ -fied E-like function that decreases for pairs  $\delta x(t), \delta y(t)$ . We can choose, like E, some  $u(\delta x, \delta y) = \frac{1}{2}C_1[\delta x]^2 + C_2[\delta y]^2$ .
- Choosing  $C_1=c, C_2=1$  gives us  $\frac{d}{dt}u(\delta x(t), \delta y(t))=\frac{d}{dt}(\frac{c}{2}[\delta x(t)]^2+\frac{1}{2}[\delta y(t)]^2)$
- $= c(dx)(dx)' + dy(dy)' = c(dx)(dy) + dy(-c(dx) (c+d)(dy)) = -(c+d)[\delta y(t)]^{2}$
- So  $u = \frac{d}{dt}(\frac{c}{2}[\delta x(t)]^2 + \frac{1}{2}[\delta y(t)]^2)$  is an energy function that could work for the laser

Finally, we want to construct a function that

- Doesn't increase (derivative negative) on any pairs x, y > 0 (pulls down)
- Is near equal to  $u = \frac{c}{2}(x-1)^2 + \frac{1}{2}(y-1)^2$  near (1,1). (the energy function for block-spring above)
- With x'-=xy-x, y'=c-cxy+d-dy, plus the identity near  $z\approx 1$  of  $\ln(z)\approx (z-1)-\frac{1}{2}(z-1)^2...$
- You can find  $U(x,y) = c(x-1) + (y-1) c \ln(x) \ln(y)$  that satisfies all of these
- It therefore shows that pumped laser solutions tend to equilibrium (1,1) in the long term.

TODO: So this is enough to establish a convergence to an equilibrium?

- Find an equilibrium  $(x_0, y_0)$
- Find an energy function u that decreases for all pairs  $(\delta x(t)\delta y(t))$  near the minimum.
- Find a Liapunov function U function that decreases EVERYWHERE along x(t), y(t) (in our domain, like x, y > 0)
- Ensure that U = u in the neighborhood of the equilibrium.
- Then Liapunov's theorem somehow makes this work (TODO)?

# 2.7 2.7: Dog chasing a duck (Limit Cycles)

This is a pair of nonlinear equations to determine if a dog in the pond's interior catches a duck who skates along the border.

#### • Variables:

- $r_p$ : Radius of pond.
- $-\vec{r_H}$ : Distance of duck to center (always the radius of the pond)
- $-\vec{l}$ : Displacement of dog from duck, which is of some length R at any point.
- $-\theta$ : Duck's position in the lake (think polar coordinates)
- $-\phi$ : Angle between  $\vec{r_H}$  and  $\vec{l}$ .
- Duck always swims at speed  $r_p\theta'(t)$ , and dog swims at k>0 times this, or  $kr_p\theta'(t)$ .
- Therefore  $\vec{r_H} = \langle r_p \cos(\theta), r_p \sin(\theta) \rangle$ . It's just the polar coordinates.
- Doing some geometry gets you  $\vec{l} = R \langle \cos(\theta + \phi, \sin(\theta + \phi)) \rangle$
- We can establish  $\vec{T} = \vec{r_H} \vec{l}$  and dog's speed squared  $||T'(t)||^2 = (\vec{r_H}' \vec{l'}) \cdot (\vec{r_H}' \vec{l'}) = ||\vec{r_H}'||^2 + ||\vec{l'}||^2 2\vec{r_H}'\vec{l'}|^2$
- Naturally, this  $||T'||^2$  is also equal to the constant  $(kr_p\theta')^2$ . Our diff equations will fall out of these.
- $\vec{r_H}^2 = r_p^2 [\theta'(t)]^2$  since duck's speed is constant.  $\vec{l} = (R')^2 + R^2 [\theta' + \phi']^2$  after working it out.
- Finally, after using identities  $\cos(\theta + \phi) = \cos(\theta)\cos(\phi) \sin(\theta)\sin(\phi), \sin(\theta + \phi) = \cos(\theta)\sin(\phi) + \sin(\theta)\cos(\phi)$ , we can work out  $-2\vec{r_H}'\vec{l'} = -2r_p\theta'[R\cos(\phi)(\theta' + \phi') + \sin(\phi)R']$
- After rescaling R to  $\rho$  such that  $\frac{R}{r_p} = \rho$  and diving our speed equation by constant  $r_p\theta'$ , we end up with speed equation  $k^2 = [\rho(1 + \frac{d\phi}{d\theta} \cos(\phi))]^2 + (\frac{d\rho}{d\theta} \sin(\phi))^2$
- We propose that there are some solutions here for the **pursuit equations**. We'll ignore the generalized form an focus on one set
  - $-\rho(1+\frac{d\phi}{d\theta})-\cos(\phi)=0, \frac{d\rho}{d\theta}-\sin(\phi)=-k$  do work in the above. (Doesn't prove others don't work)
  - This leaves our equations as  $\frac{d\phi}{d\theta} = \frac{\cos(\phi)}{\rho} 1, \frac{d\rho}{d\theta} = -k + \sin(\phi)$
  - However, there aren't simple equilibria here. In no world with  $k \neq 0$  does the dog sit still (or the duck).

- Supposing k < 1 and  $R, \phi$  are fixed (dog never gets closer and just loops), this means he's going in a circle, since the two legs of a triangle  $(\vec{l}, \vec{r_p})$  and the interior angle  $(\phi)$  are fixed, so this fixes length of the third leg, which is a radius
- You can also use dog's position vectors  $x(t) = r_p \cos(\theta) R \cos(\theta + \phi), y(t) = r_p \sin(\theta) R \sin(\theta + \phi)$  and trig identities to prove  $x(t)^2 + y(t)^2 = r_p^2 + R^2 2r_pR\cos(\phi)$
- If k < 1, then solving  $\frac{d\rho}{d\theta} = 0 = -k + \sin(\phi) \Rightarrow \sin(\phi) = k \Rightarrow \phi = \sin^{-1}(k)$  and  $\rho = \cos(\phi) = \cos(\sin^{-1}(k)) = \sqrt{1 k^2}$ 
  - \* Quick proof of  $\cos(\sin^{-1}(k)) = \sqrt{1 k^2}$ :
  - \*  $\cos^2(\sin^{-1}(k)) + \sin^2(\sin^{-1}(k)) = 1 \Rightarrow \cos(\sin^{-1}(k)) = 1 \sin^2(\sin^{-1}(k))$
  - $* = 1 k^2 \Rightarrow \cos(\sin^{-1}(k)) = \sqrt{1 k^2}$
- When k < 1, the direction field seems to have attractive equilibriua but **GOTCHA**: there are  $\phi$  angles that differ by  $2\pi$  units, so they're the same. The direction field is a cylinder with circumference  $2\pi$ , and there are other solutions tracking toward  $(\sin^{-1}(k), \sqrt{1-k^2})$
- Linearizing, assume we are near our equilibrium point and  $\phi = \sin^{-1} k + \delta \phi$ ,  $\rho = \sqrt{1 k^2} + \delta \rho$ .
- We can also remember that  $f(x + \delta x) \approx f(x) + f'(x)\delta x$  from calculus.
- $\frac{d}{d\theta}[\delta\rho] = \frac{d}{d\theta}[\rho \sqrt{1 k^2}] = \frac{d\rho}{d\theta} \frac{d}{d\theta}\sqrt{1 k^2} = -k + \sin(\phi)$
- $-=-k+\sin(\sin^{-1}(k)+\delta\phi)$  and by the calculus rule  $\frac{d}{d\theta}[\delta\rho]=-k+\sin(\sin^{-1}(k))+\cos(\sin^{-1}(k))\delta\phi=\sqrt{1-k^2}\delta\phi$
- And for  $\frac{d}{d\theta}[\delta\phi] = \frac{d}{d\theta}\phi \frac{d}{d\theta}(\sin^{-1}(k)) = \frac{\cos(\phi)}{\rho} 1$
- Using multivariable hint  $f(x + \delta x, y + \delta y) \approx f(x, y) + \frac{\delta f}{\delta x} \delta x + \frac{\delta f}{\delta y} \delta y$ ,
- $f = \frac{\cos(\sin^{-1}(k) + \delta\phi)}{\sqrt{1 k^2} + \delta\rho} 1 \approx \frac{\sqrt{1 k^2}}{\sqrt{1 k^2}} 1 + \frac{-\sin(\sin^{-1}(k))}{\sqrt{1 k^2}} \delta\phi \frac{\cos(\sin^{-1}(k))}{1 k^2} \delta\rho$
- $-=-\frac{k\delta\phi+\delta\rho}{\sqrt{1-k^2}}$
- So  $\frac{d}{d\theta} \begin{pmatrix} \delta \phi \\ \delta \rho \end{pmatrix} \begin{pmatrix} -\frac{k}{\sqrt{1-k^2}} & -\frac{1}{\sqrt{1-k^2}} \\ \sqrt{1-k^2} & 0 \end{pmatrix} \begin{pmatrix} \delta \phi \\ \delta \rho \end{pmatrix}$ , and the eigenvalues aren't purely imegi-

nary, and the real part is negative, so all decay. Therefore, the equilibrium at  $(\sin^{-1}(k), \sqrt{1-k^2})$  attracts nearby solutions.

- There aren't solutions (for K < 1?), but numerically solved, the dog catches at k > 1, and for  $k \le 1$ , swims out to a path approaching a circle. This is a **limit cycle**, an isolated trajectory that closes on itself.

#### 2.8 Poincare-Bendixson Theorem

Limit cycles in the real world: a chemical reaction in perpetual oscialiation!

Key concept - **trapping region**: a region in phase plane on some region D, with differential solutions touching every point, where the direction field sees every boundary arrow point IN. This means:

- The solution has to stay in D.
- Any solution that self-intersects forms a cycle in the phase plane.
- The three conceivable ways a solution can "snake" around forever (the **Poincare-Bendixson theorem** says it):
  - Approaches a closed loop in D.
  - Approaches a fixed point in D (possibly a special case of the last bullet)
  - Cycle: Snake eats its own tail
- A non-cycling solution is the only other possibility just a point equilibirum.

**Example:** Chemical oscillatory reaction.

- x is concentration of  $I^-$ , y is concentration of  $ClO_2^-$  ions in some reaction.
- a is positive, and clearly  $x, y \ge 0$  in the physical world.
- Otherwise meaningless equations:  $\frac{dx}{dt} = 5a x \frac{4xy}{1+x^2}, \frac{dy}{dt} = x(\frac{4y}{1+x^2})$
- Solve for equilibria by setting  $\frac{dx}{dt} = \frac{dy}{dt} = 0$ 
  - Denote  $Q = \frac{y}{1+x^2}$
  - First equation implies x(1+4Q) = 5a
  - Second equation, plus knowing  $x \neq 0, \Rightarrow x(1-Q) = 0 \Rightarrow Q = 1$
  - $-Q = 1 \Rightarrow 5x = 5a \Rightarrow x = a$
  - $\Rightarrow 1 = \frac{y}{1+x^2} \Rightarrow y = 1 + a^2$
  - Only solution pair is  $(a, 1 + a^2)$

Linearizing the solution around  $(a, 1 + a^2)$ 

• 
$$x = a + \delta x, y = 1 + a^2 + \delta y \Rightarrow \frac{dx}{dt} = \frac{d[\delta x]}{dt}, \frac{dy}{dt} = \frac{d[\delta y]}{dt}$$

• Call 
$$f = \frac{d[\delta x]}{dt} = 5a - x - \frac{4xy}{1+x^2}$$
,

• Approximate 
$$f(x + \delta x, y + \delta y) \approx f(x, y) + \frac{\delta f}{\delta x} \delta x + \frac{\delta f}{\delta y} \delta y$$

• 
$$f(x,y)(a,1+a^2) = 5a - x - \frac{4xy}{1+x^2}(a,1+a^2) = 5a - a - 4(a\frac{1+a^2}{1+a^2}) = 0$$

$$\bullet \ \ \frac{\delta f}{\delta x}\delta x(a,1+a^2) = (-1 - \frac{(1+x^2)(4y-2x4xy}{(1+x^2)^2})\delta x(a,1+a^2)) = (-1 - 4 - \frac{8a^2}{1+a^2})\delta x = \frac{-5+3a^2}{1+a^2}\delta x$$

• 
$$\frac{\delta f}{\delta y}\delta y(a,1+a^2) = \frac{-4x}{1+x^2}\delta y(a,1+a^2)) = \frac{-4a}{1+a^2}\delta y(a,1+a^2)$$

• Call 
$$g = \frac{d[\delta y]}{dt} = x - \frac{xy}{1+x^2}$$

• Approximate 
$$g(x + \delta x, y + \delta y) \approx g(x, y) + \frac{\delta g}{\delta x} \delta x + \frac{\delta g}{\delta y} \delta y$$

• 
$$g(x,y)(a,1+a^2) = x - \frac{xy}{1+x^2}(a,1+a^2) = a - a\frac{1+a^2}{1+a^2} = 0$$

• 
$$\frac{\delta g}{\delta x}\delta x(a,1+a^2) = (1-\frac{(1+x^2)y-xy^2x}{(1+x^2)^2})\delta x = (1-\frac{(1+a^2)^2-2a^2(1+a^2)}{(1+a^2)^2})\delta x = 2a^2\delta x$$

• 
$$\frac{\delta g}{\delta y}\delta y(a, 1+a^2) = \frac{-x}{1+x^2}\delta y = \frac{-a}{1+a^2}\delta y$$

$$\bullet \ \Rightarrow \frac{d}{dt} \begin{pmatrix} \delta x \\ \delta y \end{pmatrix} = \frac{1}{1+a^2} \begin{pmatrix} 3a^2 - 5 & -4a \\ 2a^2 & -a \end{pmatrix} \begin{pmatrix} \delta x \\ \delta y \end{pmatrix}$$

• Let's arbitrarily choose  $a = 2 \Rightarrow (a, 1 + a^2) = (2, 5)$ . The coefficient matrix ends up being  $\frac{1}{5} \begin{pmatrix} 7 & -8 \\ 8 & -2 \end{pmatrix}$ , which has eigenvalues with a positive real  $\pm$  some i component. So, Hartman-Grobman applies and we don't decay into our point but push away.

### We want to build the trapping region.

- Remember,  $\frac{dx}{dt} = 10 x \frac{4xy}{1+x^2}$ ,  $\frac{dy}{dt} = x(1 \frac{y}{1+x^2})$  subbing in 2 for a)
- On the left, if x = 0 we see  $\frac{dx}{dt} = 10$ ,  $\frac{dy}{dt} = 0$ . So we're pointing right (into the first quadrant region)
- On the bottom, if y = 0, we're pointing at  $\langle 10 x, x \rangle$  (into the region).
- On the right, for some x = b,  $10 b \frac{4b}{1+b^2}y$  will make sure we point left.
- On the top, for some y = c,  $x(1 \frac{c}{1+x^2} < 0$  makes sure we point down.
- Assume, since we're encircling (2,5), that  $b \geq 3, c \geq 6$  for comfort.
- To satisfy all of these, note  $x(1-\frac{c}{1+x^2})<0\Rightarrow 1-\frac{c}{1+x^2}<0\Rightarrow c>1+x^2,0< x< b\Rightarrow c>1+b^2\Rightarrow \sqrt{c-1}>b$

- And for 0 < y < c, note that  $10 x \frac{4xy}{1+x^2} < 10 b < 0$ .
- Pick b = 11, say, implying  $11 < \sqrt{c-1}$ , so then 123 < c. So (b, c) = (11, 124) ensures oscillation around (2, 5) without leaving that region.

Tricky: How to reduce this region? No real way except simulation or some tricks. If we PRESUME a cycle, we can prove the cycle extens to the left of x = 3 or  $x_{min} < 3$ 

- META trick: Don't worry if you have unsolvable integrals maybe you can cancel them out. Run with what you have.
- Trick: Assume x(t+T) = x(t), y(t+T) = y(t) for some T > 0, or that there's a PERIOD T.
- $\int_0^T \frac{dx}{dt} dt = x(T) x(0) = 0$ ,  $\int_0^T \frac{dy}{dt} dt = y(T) y(0) = 0$  by fundamental theorem.
- Our equations again:  $\frac{dx}{dt} = 10 x \frac{4xy}{1+x^2}, \frac{dy}{dt} = x(1 \frac{y}{1+x^2})$
- So  $0 = \int_0^T [10 \int x(t) 4 \int \frac{x(t)y(t)}{1 + x(t)^2}] dt$  by the first equation
- $0 = \int x(t) \int \frac{x(t)y(t)}{1+x(t)^2} dt$  by the second.
- Subtract four times the second from the first to get  $0 = 10T 5 \int_0^T x(t) \Rightarrow 2T = \int_0^T x(t)dt \ge int_0^T x_{min}dt = Tx_{min}$
- So  $2 \ge x_{min}$

#### 2.9 Chaos and the Lorenz Equation

What enabled mathematical **chaos** (unpredictability in nonlinear differential equations) was really computers and seeing simulated solutions.

The (simplified) **Lorenz system** are these equations

- $\bullet \ \frac{dx}{dt} = \sigma(y x)$
- $\frac{dy}{dt} = x(\rho z) y$
- $\frac{dx}{dt} = xy bz$
- All with  $\sigma, \rho, b > 0$

Solving the equations, we see equilibria for these are:

- (0,0) always
- The two solutions  $(\pm \sqrt{b(\rho-1)}, \pm \sqrt{b(\rho-1)}, \rho-1)$  when  $\rho > 1$ .

Looking at  $0 < \rho < 1$  specifically:

- Linearizing is simple, with  $x(t) = \delta x(t), y(t) = \delta y(t), z(t) = \delta z(t)$  and linearized system:
- $\frac{d[\delta x]}{dt} = \sigma(\delta y \delta x)$
- $\frac{d[\delta y]}{dt} = \rho \delta x \delta y$
- $\frac{d[\delta z]}{dt} = -b\delta z$
- $\bullet \ \, \frac{d}{dt} \begin{pmatrix} \delta x \\ \delta y \\ \delta z \end{pmatrix} \approx \begin{pmatrix} -\sigma & \sigma & 0 \\ \rho & -1 & 0 \\ 0 & 0 & -b \end{pmatrix} \begin{pmatrix} \delta x \\ \delta y \\ \delta z \end{pmatrix}$
- Characteristic equation is  $(-b \lambda)[(1 + \lambda)(\sigma + \lambda) \sigma\rho] = 0$
- Eigenvalues are -b < 0 and  $\lambda = \frac{1}{2}[-(\sigma+1) \pm \sqrt{(\sigma+1)^2 4\sigma(1-\rho)}]$
- If  $\rho < 1$ , we have distinct, real, negative eigenvalues, and a locally attractive equilibrium by Hartman-Grobman.

But if  $\rho < 1$  globally attactive? Find a Liapunov function.

- Requirement is that the function U(x(t), y(t), z(t)) is minimized at the equilibrium, and that as time progresses, U decreases (so we're sucked into the bowl)
- We suppose that  $U(x,y,z)=ax^2+y^2+z^2$  and using  $xy\leq \frac{1}{2}x^2+\frac{1}{2}y^2$ :
  - Identity derivation:  $0 \le (x-y)^2 \Rightarrow 0 \le x^2 2xy + y^2 \Rightarrow xy \le \frac{1}{2}(x^2 + y^2)$
- $\bullet \ \ \tfrac{\delta U}{\delta x} x'(t) + \tfrac{\delta U}{\delta y} y'(t) + \tfrac{\delta U}{\delta z} z'(t) = 2ax\sigma(y-x) + 2yx(\rho-z) 2y^2x + 2zxy 2bz^2$
- $\bullet = 2(a\sigma + \rho)xy 2a\sigma x^2 2y^2 2bz^2$
- **GOTHCA**: We can't choose  $a = -\frac{\rho}{\sigma}$  since then  $U = -\frac{\rho}{\sigma}x^2 + y^2 + z^2$  isn't minimized at (0,0,0)! So a needs to be positive.
- Choosing  $a = \frac{1}{\sigma} \Rightarrow a\sigma = 1$ , with  $\rho < 1 \Rightarrow 2(a\sigma + \rho)xy 2a\sigma x^2 2y^2 2bz^2 < 2a\sigma(2xy x^2 y^2) 2bz^2 \le -2bz^2$  by the identity above.
- Then  $U = \frac{1}{\sigma}x^2 + y^2 + z^2$  decreases as  $t \to \infty$  and is minimized at the globally attractive (0,0,0)

If  $\rho > 1$  things get chaotic. Instead of one equilibrium, we have two new ones at  $(\pm \sqrt{b(\rho - 1)}, \pm \sqrt{b(\rho - 1)}, \rho - 1)$ . Everything **bifurcates**, or qualitatively shifts when inching past  $\rho = 1$ :

- We have three equilibria.
- The origin turns into a saddle equilibrium.

- Linearizing around  $(\alpha, \alpha, \rho 1)$  with  $\alpha$  denoting  $\sqrt{b(\rho 1)}$  (pretty straightforward), we get characteristic equation for A of  $-\lambda^3 (\sigma + b + 1)\lambda^2 b(\sigma + \rho)\lambda 2\sigma b(\rho 1) = 0$
- Problem is, setting  $\rho = 1$  drops the  $(\rho 1)$  term and we have  $-\lambda(\lambda^2 + (\sigma + 1 + b)\lambda + b(\sigma + 1)) = 0$ , with solutions  $\lambda = 0, -b, -\sigma 1$ .
- The last two solutions are attractive, but the zero doesn't work for Hartman-Grobman.
- If we set  $\lambda = (\rho 1)\Delta r$  when nudging  $\rho$  just over 1, we ignore all  $\lambda^2, \lambda^3$ ... as negligible and get  $-b(\sigma + \rho)(\rho 1)\Delta r 2\rho b(\rho 1) \approx 0$
- This means  $\Delta r \approx -\frac{2\rho}{\rho+\sigma}$ , or that this nudged root has to be negative when  $\rho$  is near 1.
- More rigorously, we could have proven the roots of the equation are negative for small  $\rho 1 > 0$
- In any case, this means that the near-zero root is negative, so  $(\alpha, \alpha, \rho 1)$  attracts locally.
- We can show that this applies the same for  $(-\alpha, -\alpha, \rho 1)$

How do equilibria change as we change  $\rho$ ?

- We saw the What about as we dial past  $\rho = 1$ , our origin equilibrium changes from globally attractive to saddle point.
- In going from a stable equilibrium with negative real-part eigenvalues (attractors) to (0,0,0) as a saddle (mix of negative and positive real parts), we necessarily have a point where the eigenvalues' real parts are zero.
- In other words,  $\lambda = ia$  for some real a.
- Subbing ia into our  $-\lambda^3 (\sigma + b + 1)\lambda^2 b(\sigma + \rho)\lambda 2\sigma b(\rho 1) = 0$ , we end up getting  $[(\sigma + b + 1)a^2 2\sigma b(\rho 1)] + i[a^3 (b(\sigma + \rho)a)] = 0$
- Then we need  $a^3 b(\sigma + \rho)a = 0 \Rightarrow a = 0, a = \pm, \sqrt{b(\sigma + \rho)}$
- If a=0, the real part isn't zero. But subbing  $a=\pm\sqrt{b(\sigma+\rho)}$  gives us solutions for a set of  $\rho=\frac{\sigma(\sigma+b+3)}{\sigma-b-1}$
- So, when moving past this value, our two new equilibria change from locally attractive to saddles too.

Can we create a trapping region?

• The hint: The solutions have to pass through every ellipsoid of form  $R^2 = \rho x^2 + ]sigmay^2 + \sigma(z-2p)^2$ 

- What we need to prove: At every point on the boundary, the direction field points "in", or more specifically, the angle between inward normal and direction field is acute.
- This also means that the gradient  $\nabla g$  of the level set  $R^2 = \rho x^2 + \sigma y^2 + \sigma (z 2r\rho)^2$  is the normal. This is  $\langle 2\rho, 2\sigma, 2(z-2\rho) \rangle$
- So  $-\nabla g \cdot \langle \frac{dx}{dt}, \frac{dy}{dt}, \frac{dz}{dt} \rangle > 0 \Rightarrow \dots \Rightarrow 2\rho\sigma x^2 + 2\sigma y^2 + 2\sigma z^2 4\rho\sigma z > 0$
- Use  $R^2 \rho x^2 \sigma (z 2p)^2 = \sigma y^2 \Rightarrow \rho \sigma x^2 + 2\sigma y^2 + 2\sigma z^2 4\rho \sigma z > 0$
- This simplifies the dot product to  $2R^2 8\rho^2\sigma + 4\sigma\rho z + 2\rho(\sigma 1)x^2 > 0$
- Since the  $x^2$  term is always positive, we just need to set R to clear zero when z is its most negative (x = 0, y = 0, z =  $2\rho \frac{R}{\sqrt{\sigma}}$ ). If we churn a little more we can see that setting  $R > 2\sqrt{\sigma}\rho$  will provide a trapping region.

Question: Do the confined solutions fill up the whole (ellipsoid) container?

- Looking at the divergence (volume change of a cube over time) of the solution will tell us.
- For unspecified reasons,  $\frac{1}{v(t)}v'(t) = \nabla \cdot \langle \sigma(y-x), x(\rho-z) y, xy bz \rangle = -\sigma 1 b$
- Solving,  $v(t) = v(0)e^{-t(\sigma+1+b)}$
- This means the volume decays to 0, so therefore, our line is confined to a smaller and smaller space (but not just a point, I guess?)
- TODO