

> #OK to post  
#Julian Herman, 8th November, 2021, Assignment 18

> #1)

> C := proc(a, b, c, d, e) local rate :

rate :=  $\frac{b}{a}$  : #rate = number of eggs one chicken lays per day

print(d·rate·e) :

end proc:

> C( $\frac{3}{2}, \frac{3}{2}, \frac{3}{2}, 3, 3$ )

6

(1)

> #2)

> W := proc(a, b, k) local C :

C :=  $\left(\frac{1}{a} - \frac{1}{b}\right) \cdot \left(\frac{1}{k-1}\right)$  : #equation simplifies the same every time:  $\left\{A + B = \frac{1}{a}, A + C = \frac{1}{b}, B = k \cdot C\right\} \Rightarrow \left\{A + k \cdot C = \frac{1}{a}, A + C = \frac{1}{b}\right\} \Rightarrow$

$\Rightarrow$  subtracting eq2 from eq1  $\Rightarrow k \cdot C - C = \frac{1}{a} - \frac{1}{b} \Rightarrow C(k-1) = \frac{1}{a} - \frac{1}{b} \Rightarrow C = \left(\frac{1}{a} - \frac{1}{b}\right) \cdot \left(\frac{1}{k-1}\right)$

#C represents the speed of work done by C in the units "cisterns per hour," therefore, the reciprocal of C = number of hours for C alone to fill one cistern

print $\left(\frac{1}{C}\right)$  :

end proc:

> W(4, 5, 2)

20

(2)

> #3)iii)

> solve({x·(1 - x - y) = 0, x·(3 - 2·x - y) = 0}, {x, y})  
 $\{x=0, y=y\}, \{x=2, y=-1\}$

(3)

> with(LinearAlgebra) :

> #For equilibrium point (2,-1):

> evalf(Eigenvalues([[ -2, -2], [-4, -2]]))

$\begin{bmatrix} 0.828427124 \\ -4.828427124 \end{bmatrix}$

(4)

> #(2,-1) is UNSTABLE because 0.828427124 is > 0... The real parts of ALL of the eigenvalues must be NEGATIVE for the equilibrium point to be stable.

> #For equilibrium point (0,y):  
 > Eigenvalues ([[1 - y, 0], [3 - y, 0]])

$$\begin{bmatrix} 0 \\ 1 - y \end{bmatrix} \quad (5)$$

> #(0, y) is UNSTABLE because the first eigenvalue is always 0, **and** 0 is NOT < 0. (borderline case when y = 1 ... could be called semi - stable because **then** the eigenvalues are [[0], [0]]? not sure though)

> #4)  
 > read `Users/julianherman/Documents/Rutgers/Fall 2021/Dynamical Models In Biology/HW/M18.txt`  
 > Help18( )

*Dis2(F,x,y,pt,h,A), SIRS(s,i,beta,gamma,nu,N)* (6)

> Dis2([x\*(1 - x - y), x\*(3 - 2\*x - y)], x, y, [2.01, -1.01], 0.01, 10)[990..1000]  
 [[9.90, [Float(-∞), Float(-∞)], [9.91, [Float(-∞), Float(-∞)], [9.92, [Float(-∞), Float(-∞)], [9.93, [Float(-∞), Float(-∞)], [9.94, [Float(-∞), Float(-∞)], [9.95, [Float(-∞), Float(-∞)], [9.96, [Float(-∞), Float(-∞)], [9.97, [Float(-∞), Float(-∞)], [9.98, [Float(-∞), Float(-∞)], [9.99, [Float(-∞), Float(-∞)], [10.00, [Float(-∞), Float(-∞)]]]]]]]]]] (7)

> #This confirms that equilibrium point (2,-1) is unstable because a small change (0.01) away from the point, and it tends to (-infinity,-infinity).

> Dis2([x\*(1 - x - y), x\*(3 - 2\*x - y)], x, y, [0.01, 10.01], 0.01, 10)[999..1000]  
 [[9.99, [1.293672666 × 10<sup>-43</sup>, 10.00221360]], [10.00, [1.177213489 × 10<sup>-43</sup>, 10.00221360]]] (8)

> Dis2([x\*(1 - x - y), x\*(3 - 2\*x - y)], x, y, [-0.01, -1.01], 0.01, 10)[999..1000]  
 [[9.99, [Float(-∞), Float(-∞)], [10.00, [Float(-∞), Float(-∞)]]] (9)

> #For the family of equilibrium points (0, y) where y is any real number, it sometimes tends to the same neighborhood as the point (0, y) but **not** exactly = (0, y) as **in** the first case above **for** (0, 10) ⇒ (0.01, 10.01) tends **to** about the same (1.293672666 × 10<sup>-43</sup>, 10.00221360) ... However, **in** the second case (0, 1) ⇒ (-0.01, -1.01) tends **to** (-infinity, -infinity) ... therefore, (0, y) is UNSTABLE

> #5)  
 > F := SIRS(s, i, 0.01, 1, 1, 50) :  
 > L := Dis2(F, s, i, [20, 30], 0.01, 10) :  
 > L[-5...-1]  
 [[9.97, [49.73062516, 0.08865256516]], [9.98, [49.73199151, 0.08820691426]], [9.99, [49.73335085, 0.08776351567]], [10.00, [49.73470322, 0.08732235788]], [10.01, [49.73604867, 0.08688342946]]] (10)

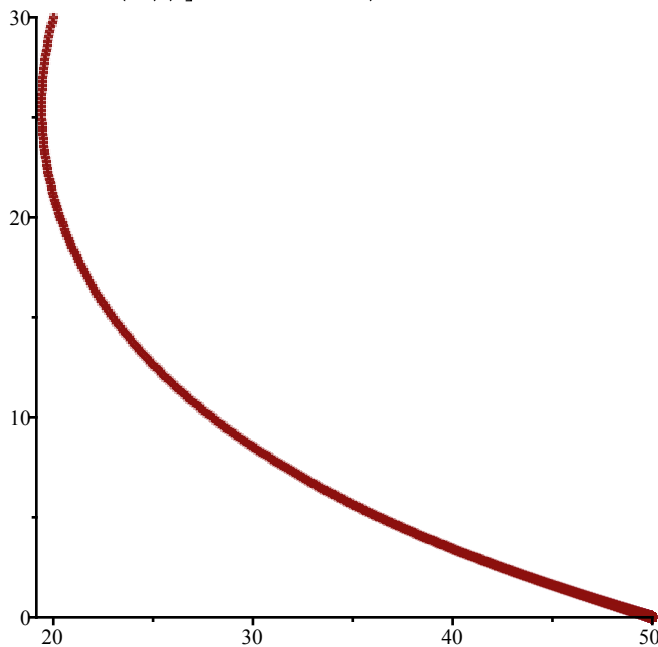
> #N=50. N-S-I=R... 50-49.73604867-0.08688342946=0.1770679005... not enough iterations yet, but removed and infected go to 0!

```
> L := Dis2(F, s, i, [20, 30], 0.01, 50) :
> L[-5 ... -1]
[[49.97, [49.99999950, 1.728549414 × 10-10]], [49.98, [49.99999950, 1.719906667
× 10-10]], [49.99, [49.99999950, 1.711307133 × 10-10]], [50.00, [49.99999950,
1.702750598 × 10-10]], [50.01, [49.99999950, 1.694236845 × 10-10]]] (11)
```

```
> #With more iterations: Changed 'A' variable of Dis2() to 50 so there are more iterations... N=50.
N-S-I=R... 50-49.99999950-1.694236845 × 10-10 = 4.998305763 × 10-7.
..It is apparent that removed and infected go to 0!
```

```
> #Plot for this below:
```

```
> plot([seq(L[i][2], i = 1 ..nops(L))], style = point)
```



```
> #With Beta=0.01, gamma=1, nu=1, N=50: infected (vertical axis) goes to 0 and susceptible
(horizontal axis) goes to N=50. the epidemic is eradicated! This is because  $N=50 < \frac{\nu}{\text{beta}}$ 
= 100
```

```
> F := SIRS(s, i, 0.01, 1, 1, 80) :
```

```
> L := Dis2(F, s, i, [50, 30], 0.01, 50) :
```

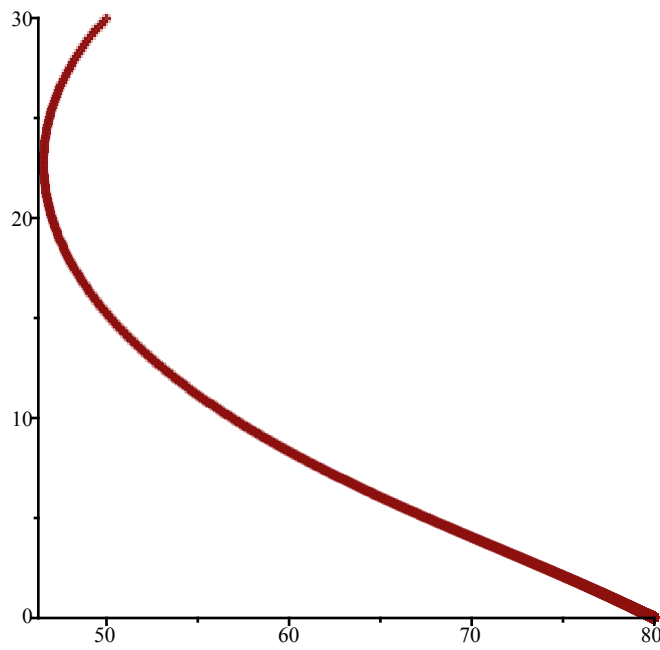
```
> L[-5 ... -1]
```

```
[[49.97, [79.99928055, 0.0003197488413]], [49.98, [79.99928198, 0.0003191093206]], (12)
[49.99, [79.99928342, 0.0003184710791]], [50.00, [79.99928486, 0.0003178341141]],
[50.01, [79.99928629, 0.0003171984232]]]
```

```
> #Changed 'A' variable of Dis2() to 50 so there are more iterations... N=80. N-S-I=R... 80
-79.99928629-0.0003171984232=0.0003965115768... It is apparent that the removed and
infected go to 0!
```

```
> #Plot for this below:
```

```
> plot([seq(L[i][2], i = 1 ..nops(L))], style = point)
```



> #With Beta=0.01, gamma=1, nu=1, N=80: infected (vertical axis) goes to 0 and susceptible (horizontal axis) goes to N=80. The epidemic is eradicated! This is because  $N=80 < \frac{\nu}{\text{beta}}$   
= 100

> F := SIRS(s, i, 0.01, 1, 1, 120) :

> L := Dis2(F, s, i, [90, 30], 0.01, 50) :

> L[-5 ... -1]

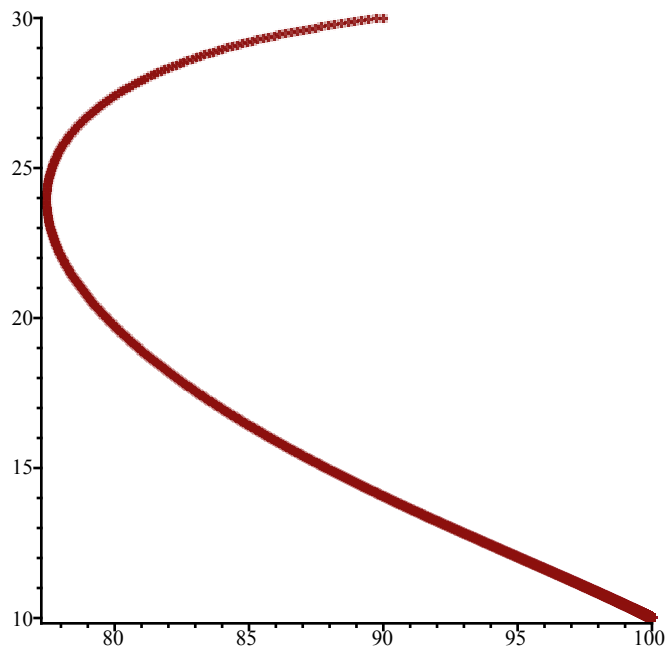
[[49.97, [99.99983227, 10.00007241]], [49.98, [99.99983268, 10.00007224]], [49.99, [99.99983308, 10.00007207]], [50.00, [99.99983348, 10.00007190]], [50.01, [99.99983388, 10.00007173]]]

(13)

> #Changed 'A' variable of Dis2() to 50 so there are more iterations... N=120. N-S-I=R... 120 -99.99983388-10.00007173=10.00009439. Susceptible=100, Infected=10, Removed=10. The epidemic was not eradicated!

> #Plot for this below:

> plot([seq(L[i][2], i = 1 ..nops(L))], style = point)



- > #With  $\text{Beta}=0.01$ ,  $\text{gamma}=1$ ,  $\text{nu}=1$ ,  $N=120$ : infected (vertical axis) goes to 10 and susceptible (horizontal axis) goes to  $N=100$ . Therefore,  $\text{removed} = N-S-I=120-100-10=10$ . The epidemic is NOT eradicated! This is because  $N=120 > \frac{\nu}{\text{beta}} = 100$ .
- > #In the long run, with  $N=120$  as in the example above, 10 individuals will be infected!

OK to post

Julian Herman, 11/8/21, Assignment 18

$$3) i) \quad x'(t) = x(t)(1 - x(t) - y(t))$$
$$y'(t) = x(t)(3 - 2 \cdot x(t) - y(t))$$

$$\text{Let } F(x(t), y(t)) = x(t)(1 - x(t) - y(t))$$

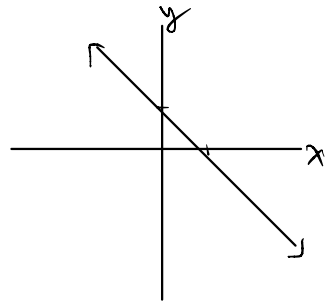
Then at equilibrium:

$$F(x, y) = x(1 - x - y) = 0$$

$$\boxed{x=0}$$

$$1 - x - y = 0$$

$$(eq. 1) \quad \boxed{y = 1 - x}$$



$$\text{Let } G(x(t), y(t)) = x(t)(3 - 2 \cdot x(t) - y(t))$$

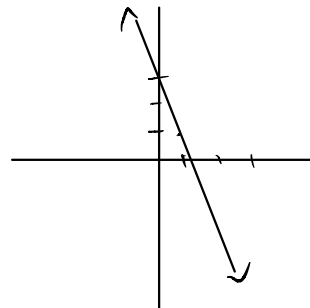
then at equilibrium:

$$G(x, y) = x(3 - 2x - y) = 0$$

$$\boxed{x=0}$$

$$3 - 2x - y = 0$$

$$(eq. 2) \quad \boxed{y = 3 - 2x}$$



$$(eq. 1) = (eq. 2)$$

$$1-x = 3-2x$$

$$x = 2$$

$$y = 1-x = 1-2 = -1$$

E.Q. points:

$$(0, y)$$

$$(2, -1)$$

when  $x=0$ ,  $y$   
can be any  
real number.

$(0, y)$  represents  
an infinite family  
of E.Q. pt's

$$ii) \text{ Jacobian} = \begin{bmatrix} F_x & F_y \\ b_x & b_y \end{bmatrix} = \begin{bmatrix} 1-2x-y & -x \\ 3-4x-y & -x \end{bmatrix}$$

more generally: let  $x = x_1$   
 $y = x_2$

$$\begin{bmatrix} 1-2x_1-x_2 & -x_1 \\ 3-4x_1-x_2 & -x_1 \end{bmatrix}$$

For E.Q. point:  $(2, -1)$ :  $J = \begin{bmatrix} -2 & -2 \\ -4 & -2 \end{bmatrix}$

infinite  
set of  
equilibrium  
points  $x=0$ ,  $y = \text{anything}$

$$\rightarrow (0, y): J = \begin{bmatrix} 1-y & 0 \\ 3-y & 0 \end{bmatrix}$$

REFER TO MAPLE SECTION