

## Homework for Lecture 18 of Dr. Z.'s Dynamical Models in Biology class

Email the answers (as a .pdf file) to

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by 8:00pm Monday, Nov. 10, 2025.

Subject: hw18

with an attachment hw18FirstLast.pdf

**1.** In the (continuous) SIRS model with a population of 1000 and parameters  $\gamma = 1.2$ ,  $\nu = 1.2$ . For each  $\beta = 0.01 \cdot i$ , for  $1 \leq i \leq 20$ , how many “removed” people are there?

**2.** Type

```
a1:=rand(1..100)(): a2:=rand(1..100)():[a1,a2];SEquP(ChemoStat(N,C,a1,a2),[N,C]);
```

20 times. How often did you get a stable equilibrium?

**3.** Run

```
SIRSDemo(1000,400,1,1,0.01,10);
```

**4.** After downloading

BOTH DMB.txt and L18.txt from the class web-page run

```
HWgE(100,1000);
```

10 times. Are the answers close to each other? Can you estimate the prob. that with a random preference matrix only one genotype will survive in the long run?

# problem 1

**read** 'DMB.txt'

For a list of the Main procedures type: *Help()*; for help with a specific procedure type: *Help(ProcedureName)*; for example *Help(Feig)*;

For a list of the Continuous Dynamical Models procedures type: *HelpC()*; for help with a specific procedure type: *Help(ProcedureName)*; for example *Help(Feig)*; (1)

**read** 'L18.txt'

For a list of the Main procedures type: *Help()*; for help with a specific procedure type: *Help(ProcedureName)*; for example *Help(Feig)*;

For a list of the Continuous Dynamical Models procedures type: *HelpC()*; for help with a specific procedure type: *Help(ProcedureName)*; for example *Help(Feig)*; (2)

*Help(SIRS)*

*SIRS(s,i,beta,gamma,nu,N)*: The SIRS dynamical model with parameters beta, gamma, nu, N (see section 6.6 of Edelstein-Keshet), s is the number of

Susceptibles, i is the number of infected, (the number of removed is given by N-s-i). N is the total population. Try:

*SIRS(s,i,beta,gamma,nu,N);* (3)

$\beta := 0.01$  (4)

*SIRS(s, i, 0.01, 1.2, 1.2, 1000)*

$[-1.21s + 1198.8, 0.01s - 1.2]$  (5)

#at equilibrium you can solve for s,i and then r.

$s := (1.2 * 1000) / 0.01;$  #  $S^* = \gamma N / \beta$

$i := (1000 - s) / 2;$  #  $I^* = (N - S) / 2$

$R := N - s - i;$  #  $R = I$

$s := 120000.0000$

$i := -59500.00000$

$R := -59500.00000$  (6)

# Parameters

$N := 1000;$

$g := 1.2;$

$nu := 1.2;$

# Loop over beta = 0.01\*i for j = 1..20

**for**  $j$  **from** 1 **to** 20 **do**

$\text{beta} := 0.01 * j;$

```

#at equilibrium you can solve for s,i and then r (set derivative equal to 0)
s := g * N / beta;
i := (N - s) / 2;
r := N - s - i;

# Run SIRS model
sol := SIRS(s, i, beta, g, nu, N);

# Print results
printf( "beta = %.2f, s = %.2f, i= %.2f, r = %.2f\n", beta, s, i, r);
print(sol);
od:
N := 1000
g := 1.2
v := 1.2
beta = 0.01, s = 120000.00, i= -59500.00, r = -59500.00
[7.132860000 × 107, -7.132860000 × 107]
beta = 0.02, s = 60000.00, i= -29500.00, r = -29500.00
[3.536460000 × 107, -3.536460000 × 107]
beta = 0.03, s = 40000.00, i= -19500.00, r = -19500.00
[2.337660000 × 107, -2.337660000 × 107]
beta = 0.04, s = 30000.00, i= -14500.00, r = -14500.00
[1.738260000 × 107, -1.738260000 × 107]
beta = 0.05, s = 24000.00, i= -11500.00, r = -11500.00
[1.378620000 × 107, -1.378620000 × 107]
beta = 0.06, s = 20000.00, i= -9500.00, r = -9500.00
[1.138860000 × 107, -1.138860000 × 107]
beta = 0.07, s = 17142.86, i= -8071.43, r = -8071.43
[9.676028570 × 106, -9.676028570 × 106]
beta = 0.08, s = 15000.00, i= -7000.00, r = -7000.00

```

$[8.391600000 \times 10^6, -8.391600000 \times 10^6]$   
 beta = 0.09, s = 13333.33, i = -6166.67, r = -6166.67

$[7.392599998 \times 10^6, -7.392599998 \times 10^6]$   
 beta = 0.10, s = 12000.00, i = -5500.00, r = -5500.00

$[6.593400000 \times 10^6, -6.593400000 \times 10^6]$   
 beta = 0.11, s = 10909.09, i = -4954.55, r = -4954.55

$[5.939509091 \times 10^6, -5.939509091 \times 10^6]$   
 beta = 0.12, s = 10000.00, i = -4500.00, r = -4500.00

$[5.394600000 \times 10^6, -5.394600000 \times 10^6]$   
 beta = 0.13, s = 9230.77, i = -4115.38, r = -4115.38

$[4.933523077 \times 10^6, -4.933523077 \times 10^6]$   
 beta = 0.14, s = 8571.43, i = -3785.71, r = -3785.71

$[4.538314286 \times 10^6, -4.538314286 \times 10^6]$   
 beta = 0.15, s = 8000.00, i = -3500.00, r = -3500.00

$[4.195800000 \times 10^6, -4.195800000 \times 10^6]$   
 beta = 0.16, s = 7500.00, i = -3250.00, r = -3250.00

$[3.896100000 \times 10^6, -3.896100000 \times 10^6]$   
 beta = 0.17, s = 7058.82, i = -3029.41, r = -3029.41

$[3.631658823 \times 10^6, -3.631658823 \times 10^6]$   
 beta = 0.18, s = 6666.67, i = -2833.33, r = -2833.33

$[3.396600001 \times 10^6, -3.396600001 \times 10^6]$   
 beta = 0.19, s = 6315.79, i = -2657.89, r = -2657.89

$[3.186284210 \times 10^6, -3.186284210 \times 10^6]$   
 beta = 0.20, s = 6000.00, i = -2500.00, r = -2500.00

$[2.997000000 \times 10^6, -2.997000000 \times 10^6]$

(7)

#when  $\beta < \gamma$  → transmission rate < recovery rate), the infection dies out so no infected or removed people remain. When  $\beta > \gamma$  positive equilibrium with  $I > 0$  and  $R > 0$ . Here the beta values go to .20 which are all much smaller than gamma

(1.2) that's why the removed people are negative counts. So technically, the removed amount of people should be 0 **and if I were to increase beta we would see positive values of I and R.**

> **read**'DMB.txt'

#Problem 2

For a list of the Main procedures type: *Help()*; for help with a specific procedure type: *Help(ProcedureName)*; for example *Help(Feig)*;

For a list of the Continuous Dynamical Models procedures type: *HelpC()*; for help with a specific procedure type: *Help(ProcedureName)*; for example *Help(Feig)*; (1)

> *Help(ChemoStat)*

*ChemoStat(N,C,a1,a2)*: The Chemostat continuous-time dynamical system with  $N$ =Bacterial population density, and  $C$ =nutrient Concentration in growth chamber (see Table 4.1 of Edelstein-Keshet, p. 122)

with paramerts  $a1$ ,  $a2$ , Equations (19a\_-, (19b) in Edelestein-Keshet p. 127 (section 4.5, where they are called alpha1, alpha2).  $a1$  and  $a2$  can be symbolic or numeric. Try:

*ChemoStat(N,C,a1,a2);*

*ChemoStat(N,C,2,3);*

(2)

>

>  $a1 := \text{rand}(1..100)(); a2 := \text{rand}(1..100)(); [a1, a2]; \text{sols} := \text{SEquP}(\text{ChemoStat}(N, C, a1, a2), [N, C]);$

[37, 33]

$\text{sols} := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right]$  (3)

>  $a1 := \text{rand}(1..100)(); a2 := \text{rand}(1..100)(); [a1, a2]; \text{sols} := \text{SEquP}(\text{ChemoStat}(N, C, a1, a2), [N, C]);$

[99, 18]

$\text{sols} := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right]$  (4)

>  $a1 := \text{rand}(1..100)(); a2 := \text{rand}(1..100)(); [a1, a2]; \text{sols} := \text{SEquP}(\text{ChemoStat}(N, C, a1, a2), [N, C]);$

[64, 24]

$\text{sols} := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right]$  (5)

>  $a1 := \text{rand}(1..100)(); a2 := \text{rand}(1..100)(); [a1, a2]; \text{sols} := \text{SEquP}(\text{ChemoStat}(N, C, a1, a2), [N, C]);$

[78, 45]

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (6)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[66, 24]$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (7)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[59, 89]$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (8)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[95, 97]$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (9)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[60, 66]$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (10)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[79, 47]$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (11)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[32, 10]$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (12)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[9, 47]$$

$$(13)$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (13)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[100, 9]$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (14)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[48, 56]$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (15)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[39, 28]$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (16)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[16, 40]$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (17)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[94, 17]$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (18)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[100, 56]$$

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (19)$$

>  $a1 := rand(1..100)(); a2 := rand(1..100)(); [a1, a2]; sols := SEquP(ChemoStat(N, C, a1, a2), [N, C]);$

$$[77, 49]$$

(20)

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (20)$$

>  $a1 := \text{rand}(1..100)(); a2 := \text{rand}(1..100)(); [a1, a2]; sols := \text{SEquP}(\text{ChemoStat}(N, C, a1, a2), [N, C]);$

[54, 35]

$$sols := \left[ [N=0, C=60], \left[ N=\frac{84322}{37}, C=\frac{1}{37} \right] \right] \quad (21)$$

> #For all 20 trials, the same two equilibrium points appeared.  $(N=0, C=60)$  and  $\left( N=\frac{84322}{37}, C=\frac{1}{37} \right)$

#The non-trivial equilibrium was stable in each case, while the trivial one was unstable.  
Therefore, a stable equilibrium was obtained 20 out of 20 times.

## problem 3

**read`DMB.txt`**

For a list of the Main procedures type: *Help()*; for help with a specific procedure type: *Help(ProcedureName)*; for example *Help(Feig)*;

For a list of the Continuous Dynamical Models procedures type: *HelpC()*; for help with a specific procedure type: *Help(ProcedureName)*; for example *Help(Feig)*; (1)

*SIRSdemo(1000, 400, 1, 1, 0.01, 10);*

This is a numerical demonstration of the  $R_0$  phenomenon in the SIRS model using discretization with mesh size=, 0.01, and letting it run until time  $t$ =, 10

with population size, 1000, and fixed parameters  $\nu$ =, 1, and  $\gamma$ =, 1

where we change beta from  $0.2*\nu/N$  to  $4*\nu/N$

Recall that the epidemic will persist if beta exceeds  $\nu/N$ , that in this case is,  $\frac{1}{1000}$

We start with , 400, infected individuals, 0 removed and hence, 600, susceptible

We will show what happens once time is close to, 10

beta is,  $\frac{1}{10}$ , times the threshold value

the long-term behavior is

$[[9.98, [999.6693512, 0.04464970605]], [9.99, [999.6721666, 0.04424784393]], [10.00, [999.6749582, 0.04384959883]], [10.01, [999.6777263, 0.04345493819]]]$

beta is,  $\frac{3}{10}$ , times the threshold value

the long-term behavior is

$[[9.98, [998.8686058, 0.2679278925]], [9.99, [998.8764375, 0.2660514879]], [10.00, [998.8842153, 0.2641882307]], [10.01, [998.8919395, 0.2623380288]]]$

beta is,  $\frac{1}{2}$ , times the threshold value

the long-term behavior is

$[[9.98, [995.5661036, 1.464972088]], [9.99, [995.5885005, 1.457614750]], [10.00, [995.6107835, 1.450294524]], [10.01, [995.6329532, 1.443011223]]]$

beta is,  $\frac{7}{10}$ , times the threshold value

the long-term behavior is

$[[9.98, [982.9907292, 6.871557578]], [9.99, [983.0448236, 6.850124744]], [10.00,$

$[983.0987363, 6.828761355]], [10.01, [983.1524679, 6.807467168]]]$

*beta is,  $\frac{9}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [944.9550913, 25.07830676]], [9.99, [945.0414764, 25.04080455]], [10.00, [945.1276722, 25.00337789]], [10.01, [945.2136792, 24.96602657]]]$

*beta is,  $\frac{11}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [866.8575732, 64.57449614]], [9.99, [866.9275067, 64.54449698]], [10.00, [866.9972772, 64.51456141]], [10.01, [867.0668854, 64.48468924]]]$

*beta is,  $\frac{13}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [764.2055840, 117.1693099]], [9.99, [764.2277964, 117.1616555]], [10.00, [764.2499053, 117.1540354]], [10.01, [764.2719113, 117.1464495]]]$

*beta is,  $\frac{3}{2}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [667.4467215, 166.2827762]], [9.99, [667.4446531, 166.2847218]], [10.00, [667.4425717, 166.2866623]], [10.01, [667.4404774, 166.2885977]]]$

*beta is,  $\frac{17}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [588.7326192, 205.7837146]], [9.99, [588.7278789, 205.7854544]], [10.00, [588.7231678, 205.7871777]], [10.01, [588.7184858, 205.7888844]]]$

*beta is,  $\frac{19}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [526.3391708, 236.9273141]], [9.99, [526.3371276, 236.9274194]], [10.00, [526.3351118, 236.9275155]], [10.01, [526.3331234, 236.9276024]]]$

*beta is,  $\frac{21}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [476.0755911, 261.9954214]], [9.99, [476.0755589, 261.9947893]], [10.00, [476.0755398, 261.9941570]], [10.01, [476.0755336, 261.9935246]]]$

*beta is ,  $\frac{23}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [434.6820669, 282.6567829]], [9.99, [434.6827642, 282.6561293]], [10.00, [434.6834631, 282.6554802]], [10.01, [434.6841635, 282.6548356]]]$

*beta is ,  $\frac{5}{2}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [399.9447066, 300.0130619]], [9.99, [399.9454130, 300.0126472]], [10.00, [399.9461153, 300.0122378]], [10.01, [399.9468135, 300.0118336]]]$

*beta is ,  $\frac{27}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [370.3498315, 314.8101124]], [9.99, [370.3503056, 314.8099378]], [10.00, [370.3507743, 314.8097672]], [10.01, [370.3512378, 314.8096006]]]$

*beta is ,  $\frac{29}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [344.8255549, 327.5761726]], [9.99, [344.8257953, 327.5761533]], [10.00, [344.8260313, 327.5761363]], [10.01, [344.8262630, 327.5761215]]]$

*beta is ,  $\frac{31}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [322.5856476, 338.7004341]], [9.99, [322.5857299, 338.7004867]], [10.00, [322.5858094, 338.7005401]], [10.01, [322.5858863, 338.7005943]]]$

*beta is ,  $\frac{33}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [303.0363872, 348.4783244]], [9.99, [303.0363869, 348.4783944]], [10.00, [303.0363852, 348.4784644]], [10.01, [303.0363821, 348.4785344]]]$

*beta is ,  $\frac{7}{2}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [285.7191728, 357.1389626]], [9.99, [285.7191408, 357.1390237]], [10.00, [285.7191083, 357.1390844]], [10.01, [285.7190753, 357.1391447]]]$

*beta is ,  $\frac{37}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [270.2735369, 364.8628645]], [9.99, [270.2735002, 364.8629086]], [10.00, [270.2734634, 364.8629522]], [10.01, [270.2734267, 364.8629953]]]$

*beta is ,  $\frac{39}{10}$ , times the threshold value*

*the long-term behavior is*

$[[9.98, [256.4121907, 371.7940209]], [9.99, [256.4121603, 371.7940490]], [10.00, [256.4121301, 371.7940766]], [10.01, [256.4121001, 371.7941037]]]$

(2)

## Problem 4

**read** 'DMB.txt'

For a list of the Main procedures type: *Help()*; for help with a specific procedure type: *Help(ProcedureName)*; for example *Help(Feig)*;

For a list of the Continuous Dynamical Models procedures type: *HelpC()*; for help with a specific procedure type: *Help(ProcedureName)*; for example *Help(Feig)*; (1)

**read** 'L18.txt'

For a list of the Main procedures type: *Help()*; for help with a specific procedure type: *Help(ProcedureName)*; for example *Help(Feig)*;

For a list of the Continuous Dynamical Models procedures type: *HelpC()*; for help with a specific procedure type: *Help(ProcedureName)*; for example *Help(Feig)*; (2)

*HWgE( 100, 1000 );*

0.5500000000 (3)

*HWgE( 100, 1000 );*

0.5280000000 (4)

*HWgE( 100, 1000 );*

0.5710000000 (5)

*HWgE( 100, 1000 );*

0.5480000000 (6)

*HWgE( 100, 1000 );*

0.5510000000 (7)

*HWgE( 100, 1000 );*

0.5430000000 (8)

*HWgE( 100, 1000 );*

0.5510000000 (9)

*HWgE( 100, 1000 );*

0.5410000000 (10)

*HWgE( 100, 1000 );*

0.5690000000 (11)

*HWgE( 100, 1000 );*

0.5380000000 (12)

#the average is about 0.549 and the range is 0.043 (max-min), thus the values are really close. This would be the probability that only one genotype survives is roughly the mean since the function estimates which one survives. There is about a 55% chance that a random preference matrix will lead to only one genotype surviving in the long run.