

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)

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

*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

```

$$\left[8.391600000 \times 10^6, -8.391600000 \times 10^6\right]$$

$$\text{beta} = 0.09, \text{ s} = 13333.33, \text{ i} = -6166.67, \text{ r} = -6166.67$$

$$\left[7.392599998 \times 10^6, -7.392599998 \times 10^6\right]$$

$$\text{beta} = 0.10, \text{ s} = 12000.00, \text{ i} = -5500.00, \text{ r} = -5500.00$$

$$\left[6.593400000 \times 10^6, -6.593400000 \times 10^6\right]$$

$$\text{beta} = 0.11, \text{ s} = 10909.09, \text{ i} = -4954.55, \text{ r} = -4954.55$$

$$\left[5.939509091 \times 10^6, -5.939509091 \times 10^6\right]$$

$$\text{beta} = 0.12, \text{ s} = 10000.00, \text{ i} = -4500.00, \text{ r} = -4500.00$$

$$\left[5.394600000 \times 10^6, -5.394600000 \times 10^6\right]$$

$$\text{beta} = 0.13, \text{ s} = 9230.77, \text{ i} = -4115.38, \text{ r} = -4115.38$$

$$\left[4.933523077 \times 10^6, -4.933523077 \times 10^6\right]$$

$$\text{beta} = 0.14, \text{ s} = 8571.43, \text{ i} = -3785.71, \text{ r} = -3785.71$$

$$\left[4.538314286 \times 10^6, -4.538314286 \times 10^6\right]$$

$$\text{beta} = 0.15, \text{ s} = 8000.00, \text{ i} = -3500.00, \text{ r} = -3500.00$$

$$\left[4.195800000 \times 10^6, -4.195800000 \times 10^6\right]$$

$$\text{beta} = 0.16, \text{ s} = 7500.00, \text{ i} = -3250.00, \text{ r} = -3250.00$$

$$\left[3.896100000 \times 10^6, -3.896100000 \times 10^6\right]$$

$$\text{beta} = 0.17, \text{ s} = 7058.82, \text{ i} = -3029.41, \text{ r} = -3029.41$$

$$\left[3.631658823 \times 10^6, -3.631658823 \times 10^6\right]$$

$$\text{beta} = 0.18, \text{ s} = 6666.67, \text{ i} = -2833.33, \text{ r} = -2833.33$$

$$\left[3.396600001 \times 10^6, -3.396600001 \times 10^6\right]$$

$$\text{beta} = 0.19, \text{ s} = 6315.79, \text{ i} = -2657.89, \text{ r} = -2657.89$$

$$\left[3.186284210 \times 10^6, -3.186284210 \times 10^6\right]$$

$$\text{beta} = 0.20, \text{ s} = 6000.00, \text{ i} = -2500.00, \text{ r} = -2500.00$$

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

#when $\beta < \gamma \rightarrow$ 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 β values go to .20 which are all much smaller than γ (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 β 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 := rand(1 ..100) ( ) : a2 := rand(1 ..100) ( ) : [a1, a2]; sols := SEquP( ChemoStat(N, C, a1, a2), [N, C]);
```

[37, 33]

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

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

[99, 18]

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

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

[64, 24]

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

```
> a1 := rand(1 ..100) ( ) : a2 := rand(1 ..100) ( ) : [a1, a2]; sols := SEquP( 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)

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

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

[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 R0 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.*