

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?

1. $i=1$; $R = 10$
2. $i=2$; $R = 40$
3. $i=3$; $R = 90$
4. $i=4$; $R = 160$
5. $i=5$; $R = 249$
6. $i=6$; $R = 358$
7. $i=7$; $R = 487$
8. $i=8$; $R = 635$
9. $i=9$; $R = 803$
10. $i=10$; $R = 990$
11. $i=11$; $R = 1000$
12. $i=12$; $R = 1000$
13. $i=13$; $R = 1000$
14. $i=14$; $R = 1000$
15. $i=15$; $R = 1000$
16. $i=16$; $R = 1000$
17. $i=17$; $R = 1000$
18. $i=18$; $R = 1000$
19. $i = 19$; $R = 1000$
20. $i= 20$; $R = 1000$

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? **I got a stable equilibrium 20/20 times.**

3. Run `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 β from $0.2 \cdot \nu/N$ to $4 \cdot \nu/N$

Recall that the epidemic will persist if β exceeds ν/N , that in this case is $1/1000$

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

We will show what happens once time is close to 10, β is , $1/10$, 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]]]

3

beta is , --, times the threshold value

10

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]]]

1

beta is , -, times the threshold value

2

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]]]

7

beta is , --, times the threshold value

10

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]]]

9

beta is , --, times the threshold value

10

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]]]

11

beta is , --, times the threshold value

10

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]]]

13

beta is , --, times the threshold value

10

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]]]

3
beta is , -, times the threshold value
2

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]]]

17
beta is , --, times the threshold value
10

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]]]

19
beta is , --, times the threshold value
10

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]]]

21
beta is , --, times the threshold value
10

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]]]

23
beta is , --, times the threshold value
10

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]]]

5
beta is , -, times the threshold value
2

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]]]

27

beta is , --, times the threshold value
10

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]]]

29

beta is , --, times the threshold value
10

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]]]

31

beta is , --, times the threshold value
10

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]]]

33
beta is , --, times the threshold value
10

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]]]

7
beta is , -, times the threshold value
2

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]]]

37
beta is , --, times the threshold value
10

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]]]

39

beta is , --, times the threshold value
10

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]]]

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.

1. .548
2. .529
3. .57
4. No sol.
5. No sol.
6. .548
7. No sol.
8. .551
9. .581
10. No sol.

It appears like there is approximately a 40% chance only one genotype will survive for a random matrix.