

Modeling a Fungal Competition on a Plant.

Steven Glasford

November 29, 2020

Content

1 Introduction

2 Modeling

3 Execution and Code

4 Graphs and Analysis

5 Conclusion

Pathogens are costly.

Pathogens make living things sick.

COVID-19 is a significant pathogen.

Pathogens can end up costing gigantically.

2016 alone saw \$540 billion in agricultural damages from plant pathogens.



What is fungal competition?

We will be looking at parasitic pathogenic fungi, such as leaf rust or maple tar spots.

Limited resources, fighting for resources, not each other.

The host has finite resources and fungus does not want to kill plant (loses all of its food).



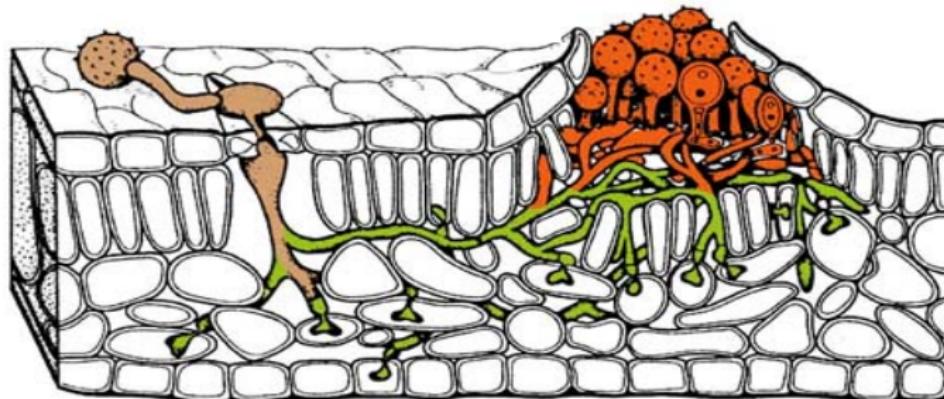
Closer inspection of Fungi

The pathogens we will be investigating exploit leaf tissue.

Mycelia expand into the current leaf.

Spore production gets to more leaves.

Penetration Mycelial growth Sporulation



Matthias Hahn, 2000

Content

1 Introduction

2 Modeling

3 Execution and Code

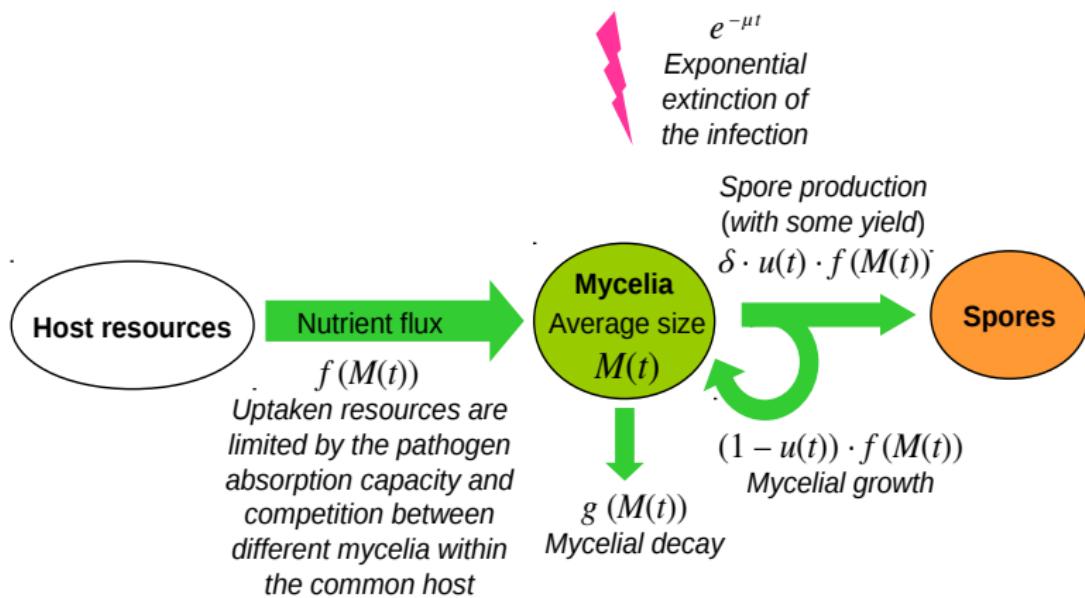
4 Graphs and Analysis

5 Conclusion

Description of fungal equations.

We focus on a single season, on a single plant, without evolution.

Unrealistic in the wild (wild is complicated), we can still get useful results, especially in laboratory conditions.



Convert to equations

Using the information from the previous slide, we can get:

$$\begin{cases} \frac{dM_1(t)}{dt} = (1 - u_1(t)) f_1(M_1(t), M_2(t)) - g(M_1(t)), \\ \frac{dM_2(t)}{dt} = (1 - u_2(t)) f_2(M_1(t), M_2(t)) - g(M_2(t)), \\ f_i(M_1, M_2) \stackrel{\text{def}}{=} \nu(n_1 M_1 + n_2 M_2) \rho(M_i), \quad i = 1, 2, \\ 0 \leq u_i(t) \leq 1, \quad i = 1, 2, \quad t \in I_T, \\ M_1(0) = M_1^0, \quad M_2(0) = M_2^0. \end{cases}$$

Assumptions

n is the lesion density and is constant.

$$f(M) \stackrel{\text{def}}{=} \nu(nM) \rho(M),$$

is the nutrient flux. Where $\rho(M)$ is the amount of resources that flow through a single mycelium. $\nu(nM)$ describes the negative influence of competing mycelia.

We will assume the resident is cohort 1, and the mutant is cohort 2.

Uninvadable strategies

The cohorts are not fighting each other. They are fighting for limited resources.

The competition forms a zero-sum feedback game: the resident defends, the mutant is offensive.

We assume that the cohorts use an uninvadable strategy. Each cohort tries its best and tries to ensure the other cohort does not fully invade.

An uninvadable strategy is also known as an evolutionarily stable strategy.

Uninvadable equations

Let J_i be the marginal fitness (the amount of success of cohort i)

$$J_i(u_1, u_2) = \int_{I_T} u_i(t) f_i(M_1(t), M_2(t)) \delta e^{-\mu t} dt$$

of cohort i .

We can then say that the resident is not invaded if:

$$J(u_1, u_2) \stackrel{\text{def}}{=} J_2(u_1, u_2) - J_1(u_1, u_2) \leq 0.$$

We are most interested in modeling when J forms a saddle point (i.e., $J = 0$).

Differential Game

The previous slides give us the following:

$$\begin{aligned} J(u_1(\cdot), u_2(\cdot)) &= J_2(u_1(\cdot), u_2(\cdot)) - J_1(u_1(\cdot), u_2(\cdot)) = \\ &= \int_{I_T} (u_2(t) f_2(M_1(t), M_2(t)) - u_1(t) f_1(M_1(t), M_2(t))) \delta e^{-\mu t} dt \\ \longrightarrow &\inf_{u_1(\cdot)} \sup_{u_2(\cdot)} \text{ or } \sup_{u_2(\cdot)} \inf_{u_1(\cdot)}, \end{aligned}$$

this describes how the first cohort tries to maximize its resistance to the second, and vice versa.

\inf is the infimum (greatest) \sup is the supremum (least).

$\inf \sup = \sup \inf$ describes a saddle point.

Content

1 Introduction

2 Modeling

3 Execution and Code

4 Graphs and Analysis

5 Conclusion

Numerical Analysis

The equations we are working with are nonsmooth, making them difficult to solve exactly.

We use computers to simulate this system.

We can convert our equations into Hamilton–Jacobi–Isaac equation (HJI).

We can then solve the HJI with ROC-HJ (Reachability, Optimal Control, and Hamilton-Jacobi equations).

Hamilton–Jacobi–Isaac equation

For brevity, we exclude the reasoning behind converting to a Hamiltonian. [2] contain further information

$$\begin{cases} \frac{\partial V(t, M_1, M_2)}{\partial t} + \mathcal{H}\left(t, M_1, M_2, \frac{\partial V(t, M_1, M_2)}{\partial M_1}, \frac{\partial V(t, M_1, M_2)}{\partial M_2}\right) = 0, \\ V(T, M_1, M_2) = 0, \\ t \in [0, T], \quad (M_1, M_2) \in G. \end{cases}$$

Resource Control Strategy

The Hamiltonian minimax (maximin) condition reduces to the following:

$$u_1(t, M_1, M_2) = \begin{cases} 0, & e^{-\mu t} + \frac{\partial V(t, M_1, M_2)}{\partial M_1} < 0, \\ 1, & e^{-\mu t} + \frac{\partial V(t, M_1, M_2)}{\partial M_1} > 0, \\ \text{arbitrary from } [0, 1], & e^{-\mu t} + \frac{\partial V(t, M_1, M_2)}{\partial M_1} = 0, \end{cases}$$
$$u_2(t, M_1, M_2) = \begin{cases} 0, & e^{-\mu t} - \frac{\partial V(t, M_1, M_2)}{\partial M_2} < 0, \\ 1, & e^{-\mu t} - \frac{\partial V(t, M_1, M_2)}{\partial M_2} > 0, \\ \text{arbitrary from } [0, 1], & e^{-\mu t} - \frac{\partial V(t, M_1, M_2)}{\partial M_2} = 0. \end{cases}$$

Where u_1 corresponds to the resource allocation strategy for cohort 1 (resident), and u_2 fits cohort 2 (mutant).

Reverse Time

ROC-HJ works backward (enables the user to start with an outcome and see how it started).

We must rewrite our equations in reverse.

$$\begin{cases} \frac{\partial V(T - \tau, M_1, M_2)}{\partial \tau} + \max_{u_1 \in [0,1]} \min_{u_2 \in [0,1]} (-H(T - \tau, M_1, M_2, u_1, \\ u_2, \frac{\partial V(T - \tau, M_1, M_2)}{\partial M_1}, \frac{\partial V(T - \tau, M_1, M_2)}{\partial M_2})) = 0, \\ V(T - \tau, M_1, M_2) |_{\tau=0} = 0, \\ \tau \in [0, T], \quad (M_1, M_2) \in G, \end{cases}$$

Example and Basic Parameters

We use the Finite Difference Method,
Second-order time discretization,
Find the saddle strategies.

ROC-HJ [1].

```
const int      OPTIM          = MAXMIN;
const int      METHOD         = MFD;
const int      TYPE_SCHEME   = EN02;
```

Figure: Configurations of descriptive variables

Output

ROC-HJ produces a large amount of data in .dat files.

0	0	0.000000000000
1	0	-0.012507240463
2	0	-0.017873483512
3	0	-0.021359387034
4	0	-0.023993289170

Figure: The first five lines from when $\tau = 20$ produced from ROC-HJ.

describes a point on a plane and the direction of the vector.
It needs to be graphed to produce meaningful results.

Content

1 Introduction

2 Modeling

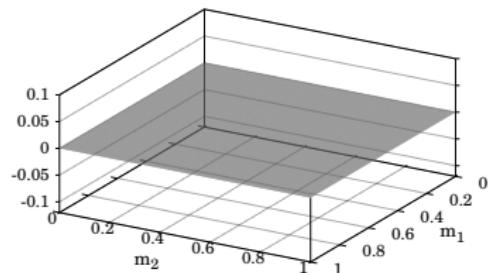
3 Execution and Code

4 Graphs and Analysis

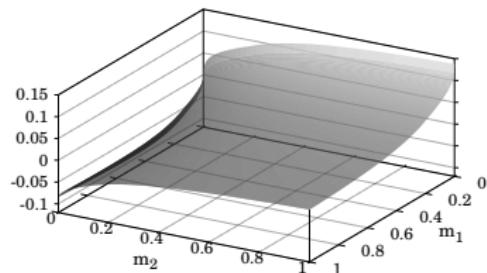
5 Conclusion

Graphs 1/3

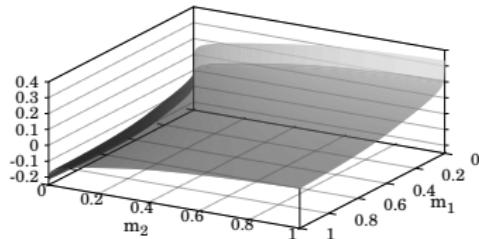
$V \cdot 10^{-4}$, (a) $\tau = 0$ ($t = T = 60$)



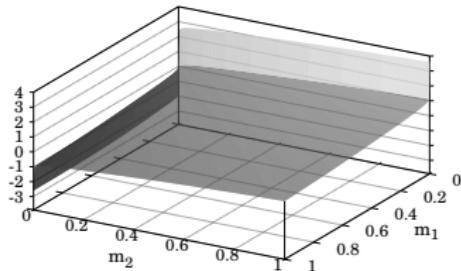
$V \cdot 10^{-4}$, (b) $\tau = 5$ ($t = 55$)



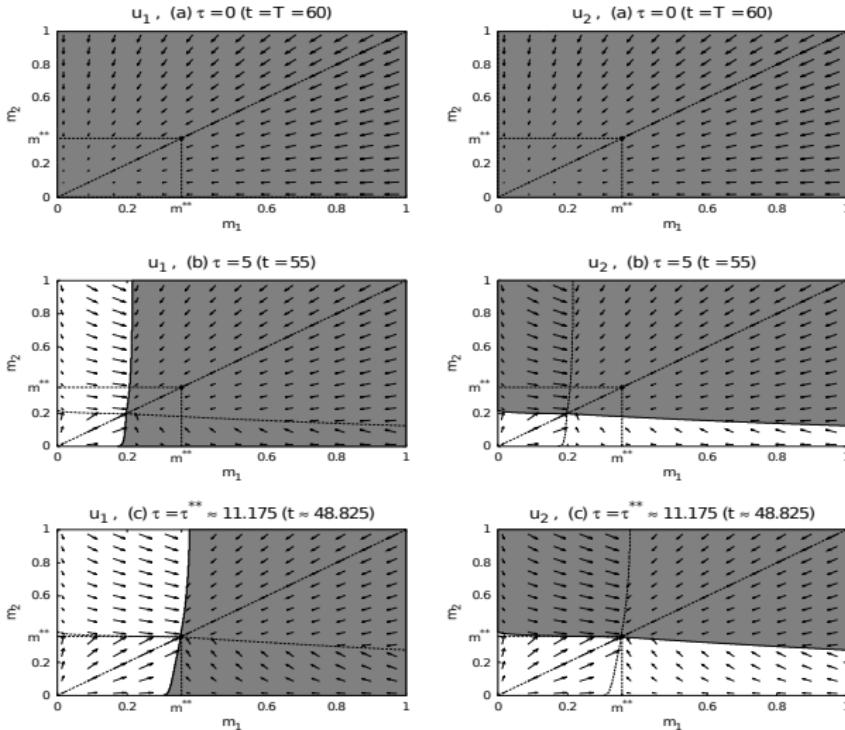
$V \cdot 10^{-4}$, (c) $\tau = \tau^{**} \approx 11.175$ ($t \approx 48.825$)



$V \cdot 10^{-4}$, (d) $\tau = T = 60$ ($t = 0$)



Graphs 2/3



White is mycelial growth, gray is spore production.

Graphs 3/3

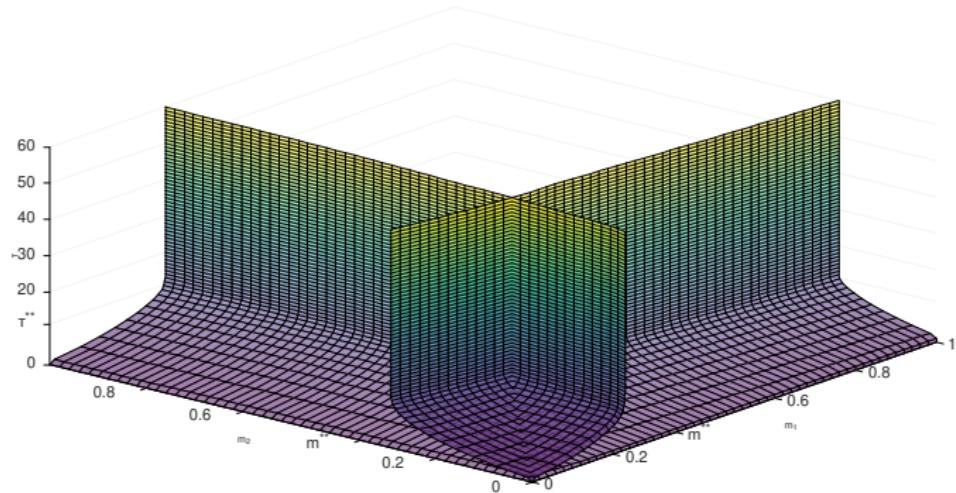


Figure: The four control turnpike switching surfaces in the three-dimensional space (m_1, m_2, τ) . A turnpike is the most efficient route in order to reach the steady-state equilibrium.

Content

1 Introduction

2 Modeling

3 Execution and Code

4 Graphs and Analysis

5 Conclusion

Model Limitations

- Our model is not super general.
- One-seasonal
- No evolution (plant or fungal)
- Single plant
- Ignores plant diversity
- Ignores plant regrowth and other tolerances
- Ignores actual fighting between fungi (some fungi produce toxins to kill opponents)

More research is needed to investigate more dynamic systems.

Still very useful.

End statements

Future analysis will include further analysis and inclusion of previously stated limitations.

We looked at strategies for nearly equivalent fungi competing with each other, fighting for resources.

Thank you, Dr. Ivan Yegorov!

References

-  Desilles A, Zidani H, Bokanowski, O. and J. Zhao.
User's guide for the roc-hj solver: Finite differences and semi-lagrangian methods.
<https://uma.ensta-paristech.fr/soft/ROC-HJ>,
2019-01-21.
-  Ivan Yegorov, Frédéric Grogna, Ludovic Mailleret, Fabien Halkett, and Pierre Bernhard.
A dynamic game approach to unininvadable strategies for biotrophic pathogens.
Dynamic Games and Applications, 10(1):257–296, Mar 2020.