MSc Computational Science joint programme UvA/VU







A DIFFUSION-BASED MODEL OF SPATIAL INTERACTIONS IN ASPERGILLUS SPP. GERMINATION

FIRST EXPERIMENTS

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2 General Assumptions Simulation space

3 Experiment 1: Immediate release
Assumptions
Analytical verification
Results

4 Experiment 2: Slow release
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5 Discussion

INTRODUCTION

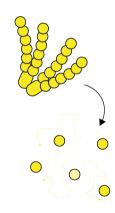








- » Upon inoculation in an aqueous medium, a germination inhibitor diffuses away from the conidium.
 - Experiments have highlighted 1-octen-3-ol as a likely candidate [2, 3, 4]
- » Once its concentration at the spore falls below a certain threshold, the conidium breaks dormancy and enters a swelling phase.
 - It has been observed that in a externally non-inhibited spore swelling begins around 4 hours from inoculation
- » An increased density of spores drives the local inhibitor concentration high, reducing overall germination.
 - Densities above 10⁵ sporesl/mL exhibit germination inhibition[4, 6]









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GENERAL ASSUMPTIONS

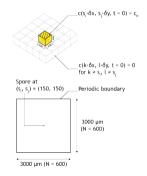








- » 2D lattice \equiv thin layer of medium clamped between 2 impermeable sheets
 - L = $3000 \, \mu \text{m}$
 - Discrete blocks of size $\delta x = \delta y = \delta z = 5 \,\mu \text{m}$
 - -N = 600 subdivisions
 - Lattice volume $V_L=4.5 imes10^7~\mu\mathrm{m}^3=4.5 imes10^{-5}~\mathrm{mL}$
 - Assuming water as extracellular medium
- » Spore \equiv single block in the center of the lattice, inhibitor source
 - Volume $V_s = 125 \, \mu \text{m}^3$
 - Top and bottom side of spore are sealed
- » Inhibitor initial concentrations c(x, y, t = 0):
 - At spore: $c(x_s, y_s, t = 0) = c_0$
 - Elsewhere: c(x, y, t = 0) = 0
 - Assumed non-replenishable during process



GENERAL ASSUMPTIONS









- » Assumed to have known molecular properties of 1-octen-3-ol:
 - Molecular weight: $m_M = 128.21 \,\mathrm{g/mol}$
 - Density: $ho=0.837\,\mathrm{g/mL}$

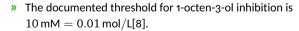


- » Assumed to have similar diffusion properties as glucose:
 - Diffusion constant in water at temperature T = 293.15K:

 $D=600\,\mu\mathrm{m}^2/\mathrm{s}$ [5] (in agar only 5% lower[12])

(i.e. a molecule wanders on average $\langle x \rangle = \sqrt{6 \cdot D} = 60 \ \mu \mathrm{m}$ per second)

- Permeation constant in an artificial lipid bilayer: $P_s=1.9 \times 10^{-6} \ \mu \text{m/s}$ (average from [7] and [1], close to [11])
- Octanol-water partition coefficient: $K pprox 10^{-3}$



» Once below this level, it is assumed swelling begins immediately



Figure: 1-octen-3-ol



Figure: D-Glucose









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- » At t = 0, all the inhibitor is contained in a cell wall at the interface between the spore and the medium, having the same diffusion coefficient as water.
 - Cell wall thickness: $d \approx 130 \, \text{nm}$ (A. fumigatus)[10]
 - Total cell wall volume:

$$V_{\rm cw} \approx 6 \cdot 25 \,\mu{\rm m}^2 \cdot 0.13 \,\mu{\rm m} = 19.5 \,\mu{\rm m}^3$$

- The maximum possible concentration is:

$$c_{
m cw} = rac{
ho V_{
m cw}}{m_{
m M} V_{
m s}} = 6.5\,{
m mol/mL} \cdot 0.156 = 1.014\,{
m mol/mL}$$
 (1)

- Equivalently, $N = N_A c_{\rm cw} V_{\rm cw} = 7.04 \times 10^{10}$ molecules.
- The experiments use an initial concentration $c_0=c_{
 m cw}$.







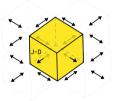




» Diffusion is simulated using the diffusion equation (Fick's Law):

$$\frac{\partial c}{\partial t} = D\nabla^2 c \tag{2}$$

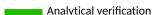
- » Simulation run for $t_{\rm max} = 4 \, {\rm h} = 14\,400 \, {\rm s}$
- » The times for reaching threshold concentrations of $c_t=kc_0$ ($k=\{10^{-1},10^{-2},...,10^{-6}\}$) are recorded.
- » To prevent numerical error, the concentration in the simulations is normalised by a factor $\lambda = \frac{1}{c_0}$.









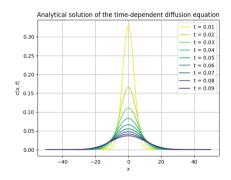


- » Assuming the initial condition is a delta pulse of concentration at coordinates (x_0, y_0)
- » The concentration at an arbitrary position (x, y) at time t can also be computed through the analytical formula[9]:

$$c(x, y, t) = \frac{N^{2/3}}{4\pi Dt} \exp\left(-\frac{(x - x_0)^2 + (y - y_0)^2}{4Dt}\right)$$
 (3)

» At the spore site, this equation becomes:

$$c(x_0, y_0, t) = \frac{(V_s c_0)^{2/3}}{4\pi Dt}$$
 (4)



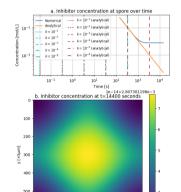








- » Results after 4 hours
- » Numerical vs. analytical
 - Alignment of numerical and analytical solution up to $t \approx 1000$ s, then finite size starts to take effect
 - Under the assumed c₀, the concentration in this period is low enough to make this deviation insignificant (few thousands of molecules).
- » Threshold times follow a power law
- » Under the given c₀ (maximum that the cell wall can fit), threshold of 10 mM reached within a few minutes
- » Diffusion is not a limiting factor









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- » All the inhibitor is contained within the cell wall volume, separated from the medium by a membrane having the properties of an artificial lipid bilayer.
- » Maximum concentration limit
 - Under homogeneous distribution, standard temperature and pressure:

$$c_{
m max} = rac{
ho}{m_{
m M}} = rac{0.837\,{
m g/mL}}{128.21\,{
m g/mol}} pprox 6.5\,{
m mol/mL}$$
 (5

- Equivalently, $N = N_A c_{\rm max} V_s = 4.89 \times 10^{14}$ molecules.











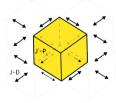
Diffusion assumptions

- » Diffusion through the medium as before
- » Diffusion at the interface follows the flux equation

$$J' = -P_s \Delta c \tag{6}$$

where J' is the flux through the cell wall and Δc is the concentration difference across the membrane.

- » Simulation run for $t_{\text{max}} = 4 \text{ h} = 14400 \text{ s}$
- » The times for reaching threshold concentrations of $c_{\rm t}=kc_0$ ($k=\{10^{-1},10^{-2},...,10^{-6}\}$) are recorded.











Analytical verification

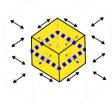
- » Since the concentration outside (c_{out}) the spore with a volume V and a surface A diffuses much faster than inside $(c_{\text{in}}), c_{\text{out}} \approx \text{const.}$
- » Then, an analytical solution is:

$$c_{\rm in}(t) = c_{\rm out} - \Delta c(0)e^{-t/\tau}, \tag{7}$$

where τ is a decay constant:

$$\tau = \frac{V}{AP_s} \tag{8}$$

» Only 2D communication in the numerical simulation \to A is actually the **circumference** of the cross section $\times 1~\mu m!$



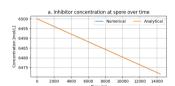


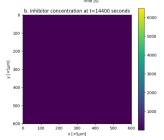






- » Results after 4 hours
- » Numerical vs. analytical
 - Good alignment of numerical and analytical solution (because medium does not saturate in the given time)
- » Almost linear decrease in concentration
- » Threshold of 10 mM never reached, neither is one tenth of the initial concentration













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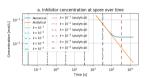


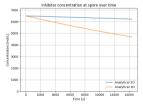




Validity of results

- » Known threshold for 1-octen-3-ol inhibition (10 mM)
 - Experiment 1: threshold reached within minutes
 - Experiment 2: threshold not reached within 4 h
- » The resulting concentrations can be rescaled to different c_0 :
 - Experiment 1: If the starting concentration is based on the entire spore volume V_s, then 1-octen-3-ol threshold reached within 30 minutes. More reasonable result, but unrealistic c₀ and permeability assumptions.
 - Experiment 2: If c₀ is only slightly above threshold, it can be reached within 4 h.
- » The truth lies in between:
 - Permeation of cell wall is between that of membrane and water
 - If permeating through the membrane, facilitated by proteins (e.g. glucose transporters) → increased P_s?
 - Cell wall permeability modulated during transition from dormancy.
 - In three dimensions: more surface area for diffusion





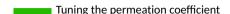


DISCUSSION









» The analytical solution can be rewritten for P_s :

$$P_s = rac{V_{
m s}}{tA} \ln \left(rac{\Delta c(0)}{\Delta c(t)}
ight)$$
 (9)

- » Assuming the inhibitor stops being effective below $c_{
 m t}=10\,{
 m mM}$
- » Keeping the assumptions about c_0 , d, K, V_s and A_s
- » A hypothetical permeation constant P_s can be computed for a diffusion of 4 hours
- » This results in $P_s=0.001\,174\,\mu\mathrm{m/s}$
- » Remains to be interpreted in terms of physical properties

DISCUSSION









- » Focus on 3D systems
- » Investigating the relationship between inhibitor concentration and spore density
- » Investigating effects on the outward flux when the medium is locally saturated

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