MSc Computational Science joint programme UvA/VU







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

FIRST EXPERIMETNS

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November 28, 2024

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1 General Assumptions

Simulation space Inhibitor

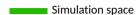
Concentration check

GENERAL ASSUMPTIONS

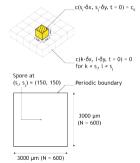








- » 2D lattice \equiv thin layer of medium clamped between 2 impermeable sheets
 - L = 3000 μ m
 - Discrete blocks of size $\delta x = \delta y = \delta z = 5 \,\mu \text{m}$
 - N = 600 subdivisions
 - Lattice volume $V_L = 4.5 imes 10^7 \, \mu \mathrm{m}^3 = 4.5 imes 10^{-5} \, \mathrm{mL}$
 - Assuming water as 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



GENERAL ASSUMPTIONS









- » Initial concentration is assumed to be contained within the cell wall
 - Assumed cell wall thickness: d = 10 nm [6]
 - Total cell wall volume: $V_{cw} \approx 6 \cdot 25 \, \mu \text{m}^2 \cdot 0.01 \, \mu \text{m} = 1.5 \, \mu \text{m}^3$
 - This concentration is non-replenishable.



- Molecular weight: $m_M=128.21\,\mathrm{g/mol}$
- Density: $\rho = 0.837 \, \mathrm{g/mL}$
- Assumed to have similar diffusion properties as glucose:
 - Diffusion constant in water at temperature T = 293.15K:
 - $D=600\,\mu\mathrm{m}^2/\mathrm{s}$ [2] (in agar only 5% lower[8])
 - (i.e. a molecule wanders on average $\langle x \rangle = \sqrt{6 \cdot D} = 60 \, \mu \mathrm{m}$ per second)
 - Permeability constant in an artificial lipid bilayer:
 - $P_s=1.9 imes10^{-6}~\mu\mathrm{m/s}$ (average from [3] and [1], close to [7])
 - Octanol-water partition coefficient: $K \approx 10^{-3}$



Figure: 1-octen-3-ol



Figure: D-Glucose

GENERAL ASSUMPTIONS





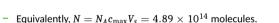


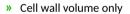


Initial concentration check

- » Full spore volume
 - Homogeneous distribution, standard temperature and pressure:

$$c_{
m max}=rac{
ho}{m_M}=rac{0.837\,{
m g/mL}}{128.21\,{
m g/mol}}pprox 6.5\,{
m mol/mL}$$
 (1)





- The maximum possible concentration is:

$$c_{
m mcw} = rac{
ho V_{
m cw}}{m_{
m W} V_{
m s}} = 6.5 \, {
m mol/mL} \cdot 0.012 = 0.078 \, {
m mol/mL}$$
 (2)

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













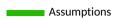
- 1 General Assumptions
 - Simulation space
 - Inhibitor
 - Concentration check
- 2 Experiment 1: Diffusion through water Assumptions Analytical verification
- 3 Experiment 2: Slow release Assumptions Analytical verification
- 4 Results
 Experiment 1
 Experiment 2
- 5 Discussion

EXPERIMENT 1 SETUP





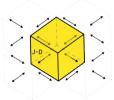




- » At t = 0, all the inhibitor is contained in a cell wall at the interface between the spore and the medium, having the same diffusion coefficient as water.
- » To prevent numerical error, the concentration in the simulations is normalised by a factor $\lambda = \frac{1}{c_0}$.
- » Diffusion is simulated using the diffusion equation (Fick's Law):

$$\frac{\partial c}{\partial t} = D\nabla^2 c \tag{3}$$

- » Simulation run for $t_{\text{max}} = 3600 \,\text{s}$
- » The times for reaching threshold concentrations of $c_t=kc_0$ ($k=\{10^{-1},10^{-2},...,10^{-6}\}$) are recorded.



ANALYTICAL VERIFICATION





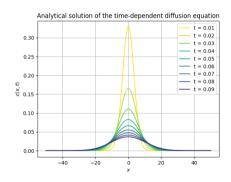


- » Assuming the initial condition is a delta pulse of concentration at coordinates (x_0, y_0)
- » The concentration at an arbitrary position (x, y) at time t can also be computed through the analytical formula[5]:

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

» At the spore site, this equation becomes:

$$c(x_0, y_0, t) = \frac{(V_s c_0)^{2/3}}{4\pi Dt}$$
 (5)









1 General Assumptions

Simulation space

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Concentration check

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EXPERIMENT 2 SETUP









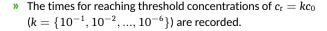
- » All the inhibitor is contained within in a cell wall with the properties of a lipid bilayer at the spore-medium interface.
- » Diffusion through the medium as before
- » Diffusion at the interface follows the flux equation

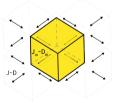
$$J_m = -D_m \nabla c \tag{6}$$

» where J_m is the flux through the cell wall such that

$$D_m = \frac{d}{K} P_s = 1.9 \times 10^{-5} \,\mu\text{m}^2/\text{s}$$
 (7)







ANALYTICAL VERIFICATION







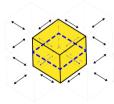
- » Since the concentration outside $(c_{ ext{out}})$ the spore with a volume V and a surface A diffuses much faster than inside $(c_{ ext{in}}), c_{ ext{out}} pprox ext{const.}$
- » Then, an analytical solution is:

$$c_{\rm in}(t) = c_{\rm out} - \Delta c(0)e^{-t/\tau}, \tag{8}$$

where τ is a decay constant:

$$\tau = \frac{V}{AD_m} \tag{9}$$

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









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Simulation space

Concentration shock

2 Experiment 1: Diffusion through water

Assumptions
Analytical verification

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RESULTS

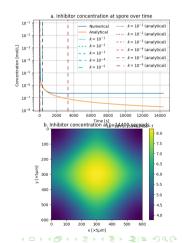








- » Results after 4 hours
- » Numerical vs. analytical
 - Alignment of numerical and analytical solution up to $t \approx 1000$ s, then finite size starts to take effect.
 - Under the assumed c₀, the concentration in this period is low enough to make this deviation insignificant (few thousands of molecules).
- » Threshold times follow a power law
- » Under the given c_0 (maximum that the cell wall can fit), the results seem unrealistic \rightarrow diffusion is too fast



RESULTS

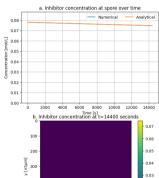


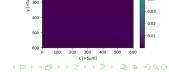






- » Results after 4 hours
- » Numerical vs. analytical
 - Good alignment of numerical and analytical solution (because medium does not saturate in the given time)
- » Almost linear decrease in concentration
- » Diffusion appears too slow, threshold of one tenth never reached











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Simulation space

Inhibitor

Concentration check

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Assumptions

Analytical verification

3 Experiment 2: Slow release

Assumptions

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Experiment '

Experiment 2

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DISCUSSION

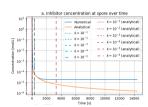


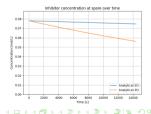






- The documented threshold for 1-octen-3-ol inhibition is $10 \, \text{mM} = 1 \times 10^{-5} \, \text{mol/mL[4]}.$
 - Diffusion through water: threshold immediately reached
 - Diffusion through membrane: threshold not reached
- The resulting concentrations can be rescaled to different c_0 :
 - If the starting concentration is based on the entire spore volume V_s , then Experiment 1 would reach the 1-octen-3-ol threshold within 30 minutes.
 - More reasonable result, but unrealistic initial concentration.
 - Perhaps the initial concentration is only slightly above threshold?
- The truth lies in between:
 - Permeation through the membrane is facilitated by proteins (e.g. glucose transporters) \rightarrow increased P_s ?
 - Cell wall permeability modulated during transition from dormancy.
 - In three dimensions: more surface area for diffusion



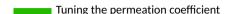


DISCUSSION









» The analytical solution can be rewritten for P_s :

$$P_s = rac{V_{
m cw}}{t K A} \ln \left(rac{\Delta c(0)}{\Delta c(t')}
ight)$$
 (10)

- » Assuming the inhibitor stops being effective below $c=10\,\mathrm{mM}$
- » 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 8 hours
- » This results in $P_s = 0.0016 \,\mathrm{m/s}$
- » Remains to be interpreted in terms of physical properties

DISCUSSION







Future outlook

- » 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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