

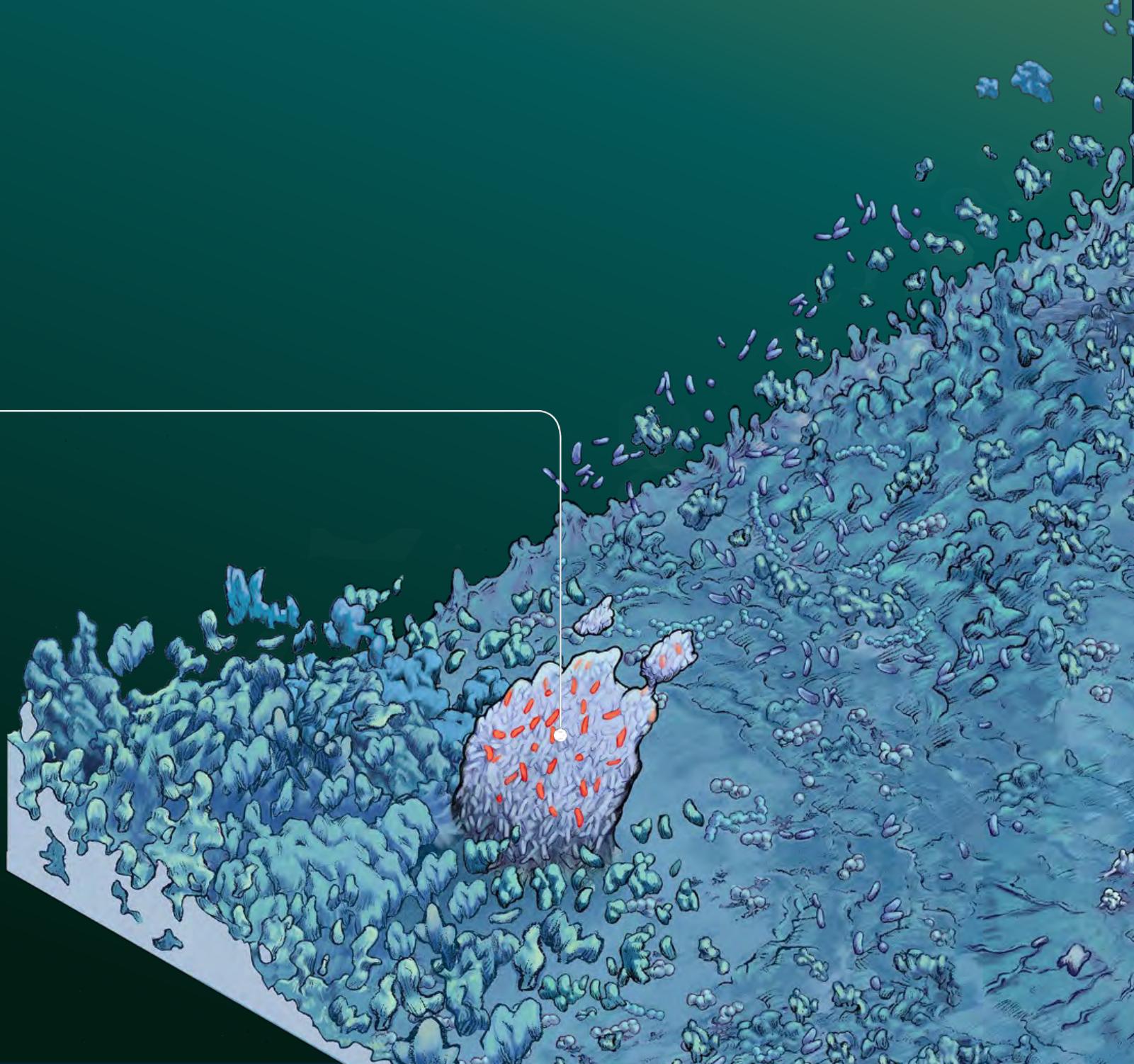
A1 RB-3 : « Biofilms bénéfiques : des communautés aux applications »

Beneficial biofilms

Unlocking the potential of biofilm properties in beneficial
microbes for one health advancements

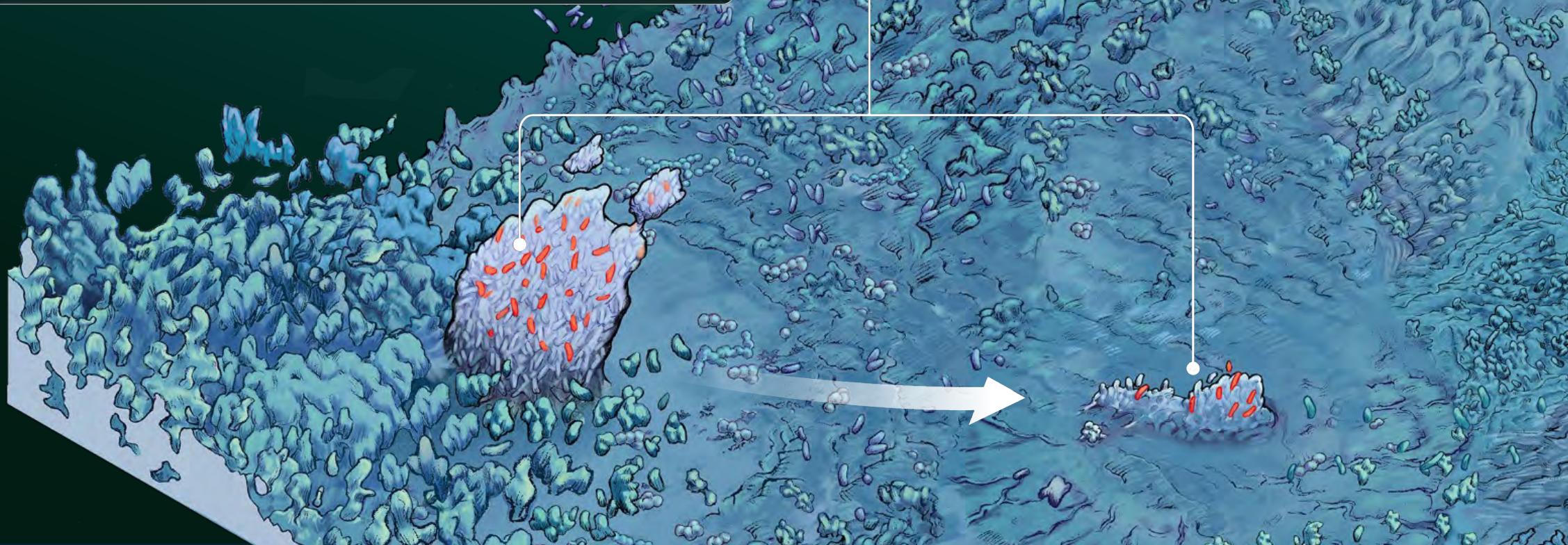
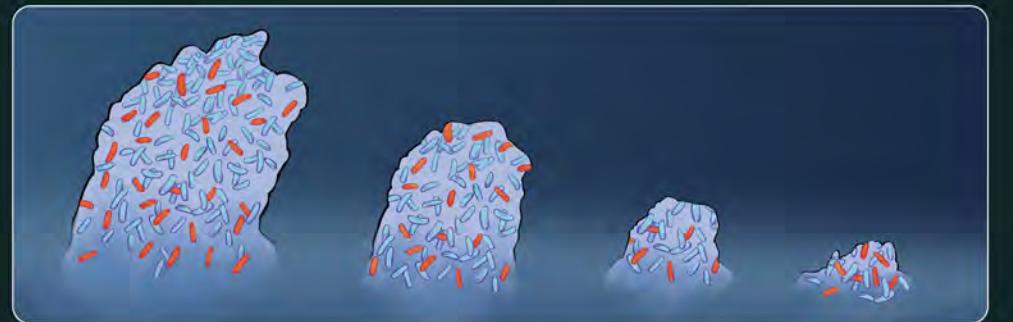


Pre-existing biofilm



Cleaning and control intervention

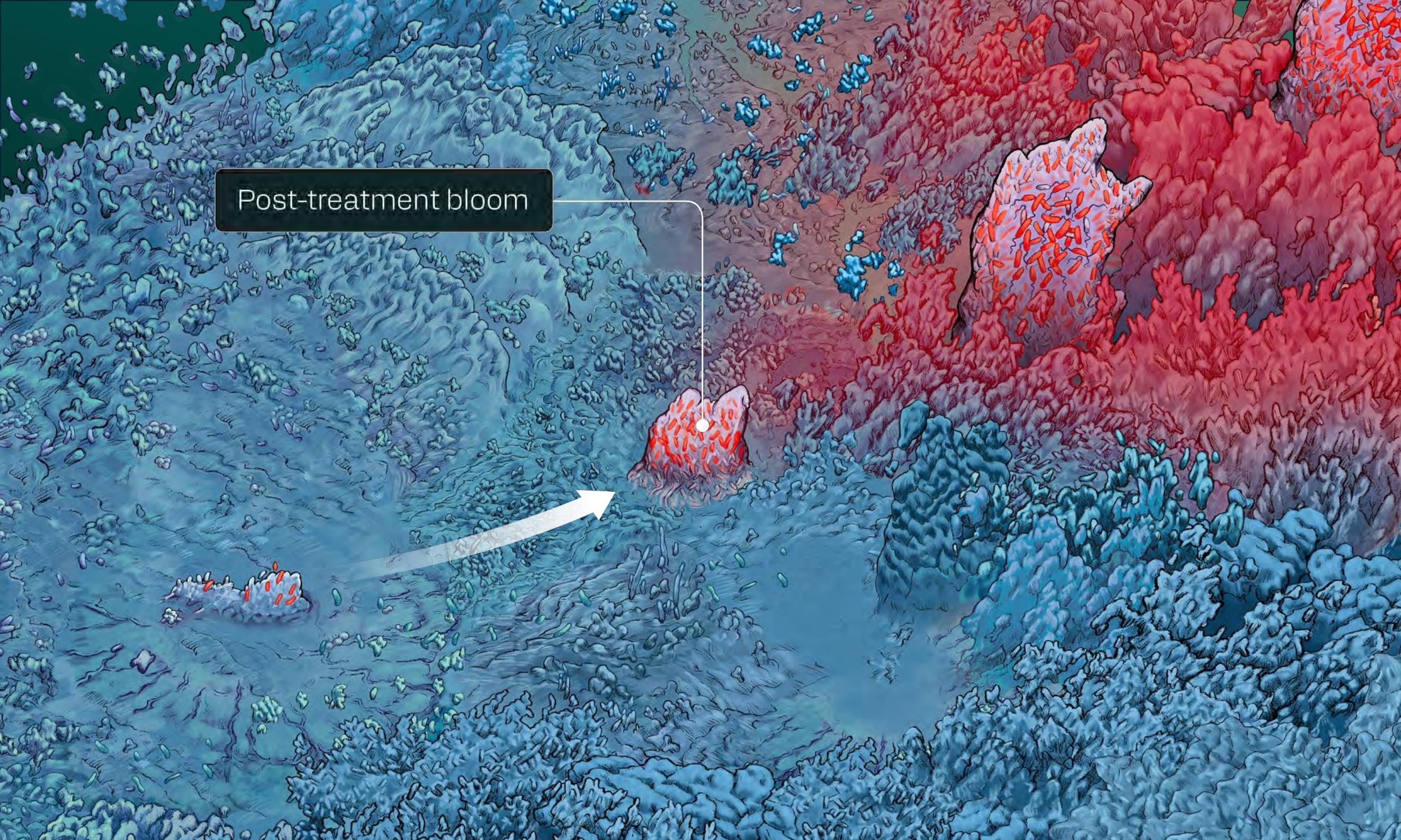
Schematic of a mature biofilm after surface treatment. The structure is only partially reduced due to biofilm resilience, driven by limited diffusion, metabolic specialization, and matrix-based protection.



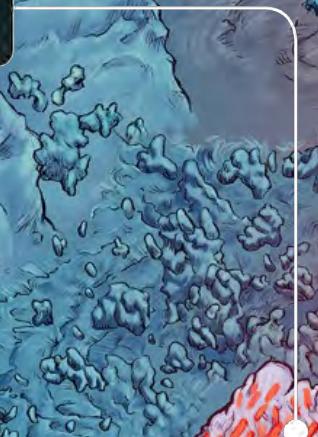


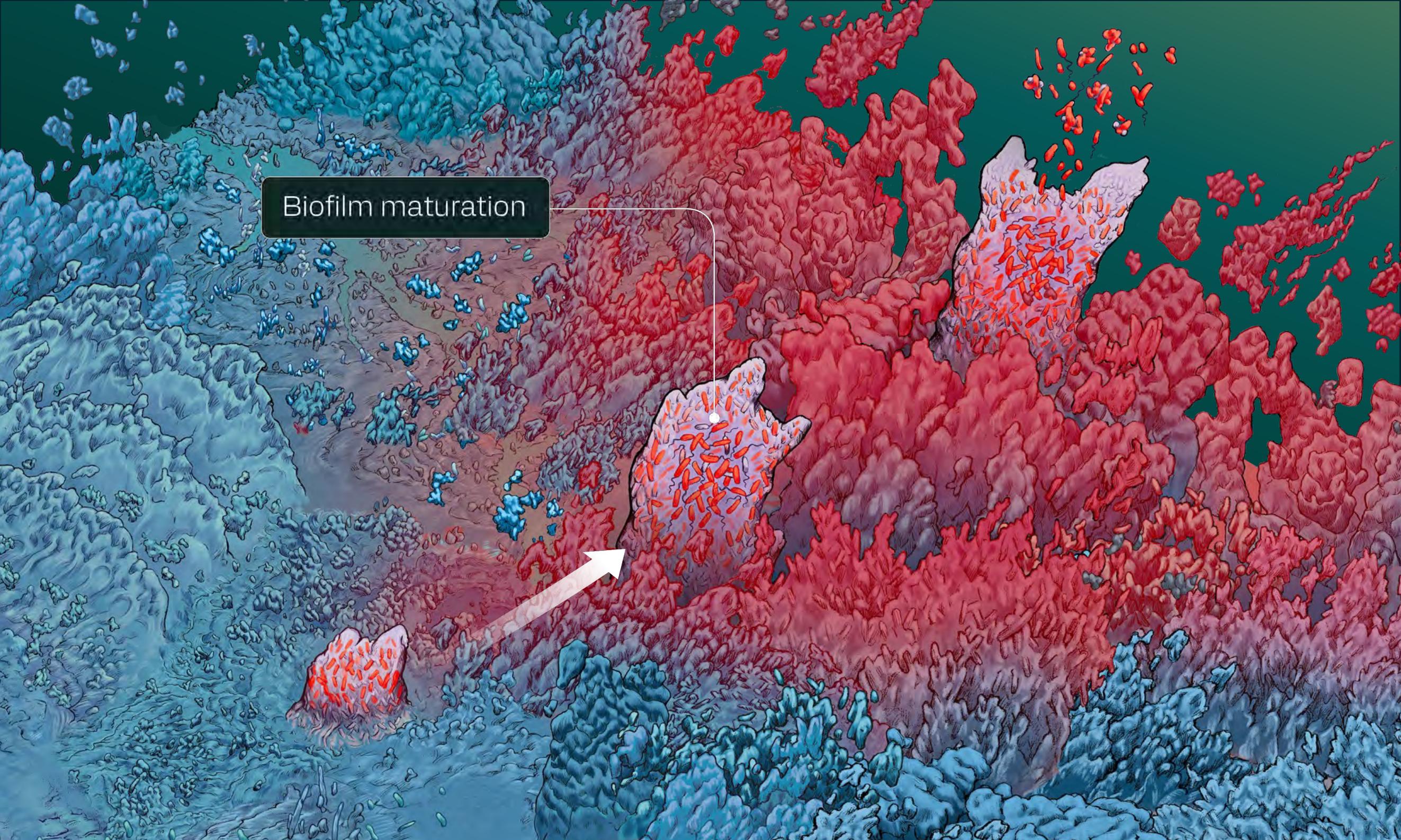
Residual biofilm survival

1



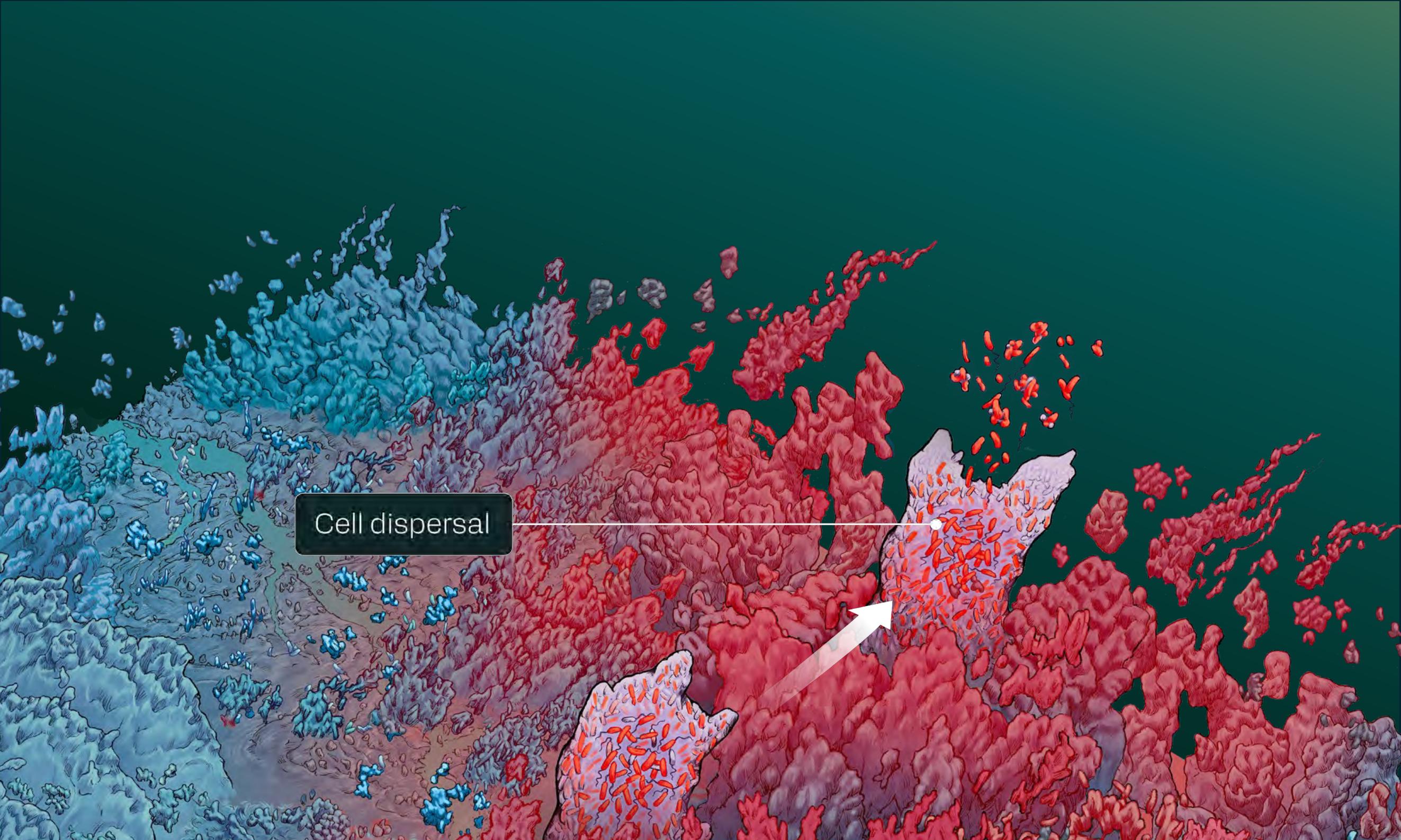
Post-treatment bloom



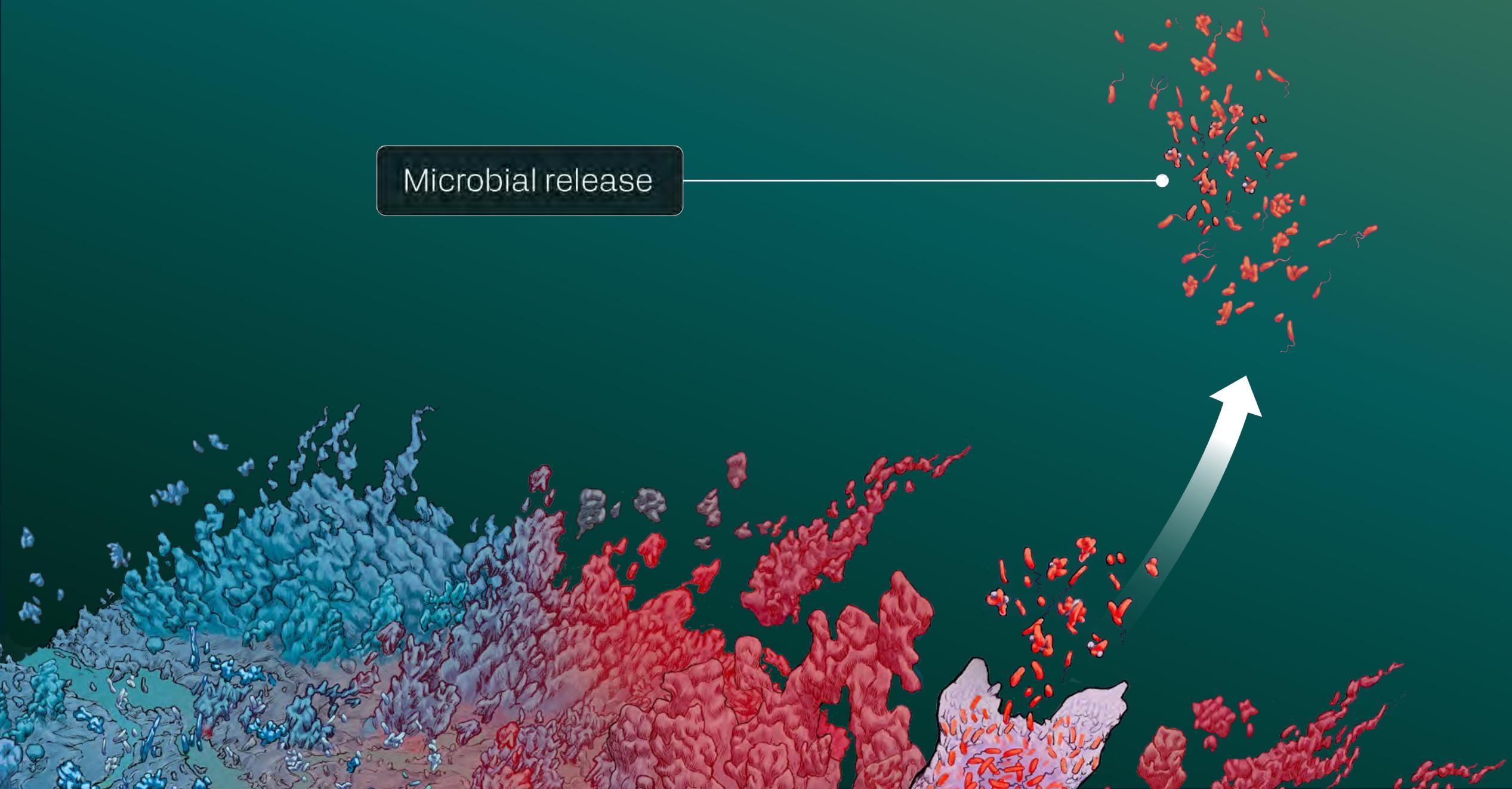


Biofilm maturation



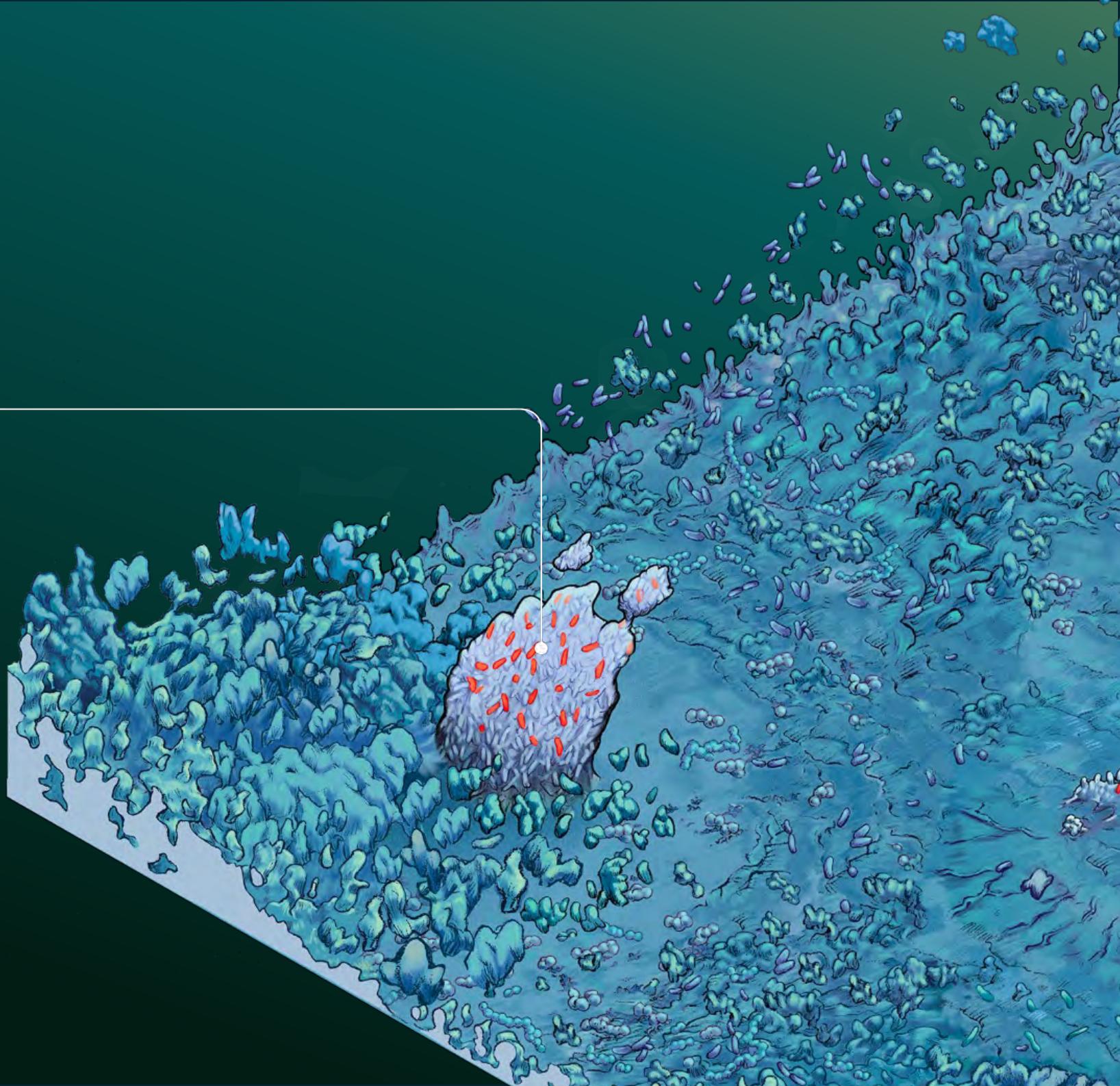


Cell dispersal

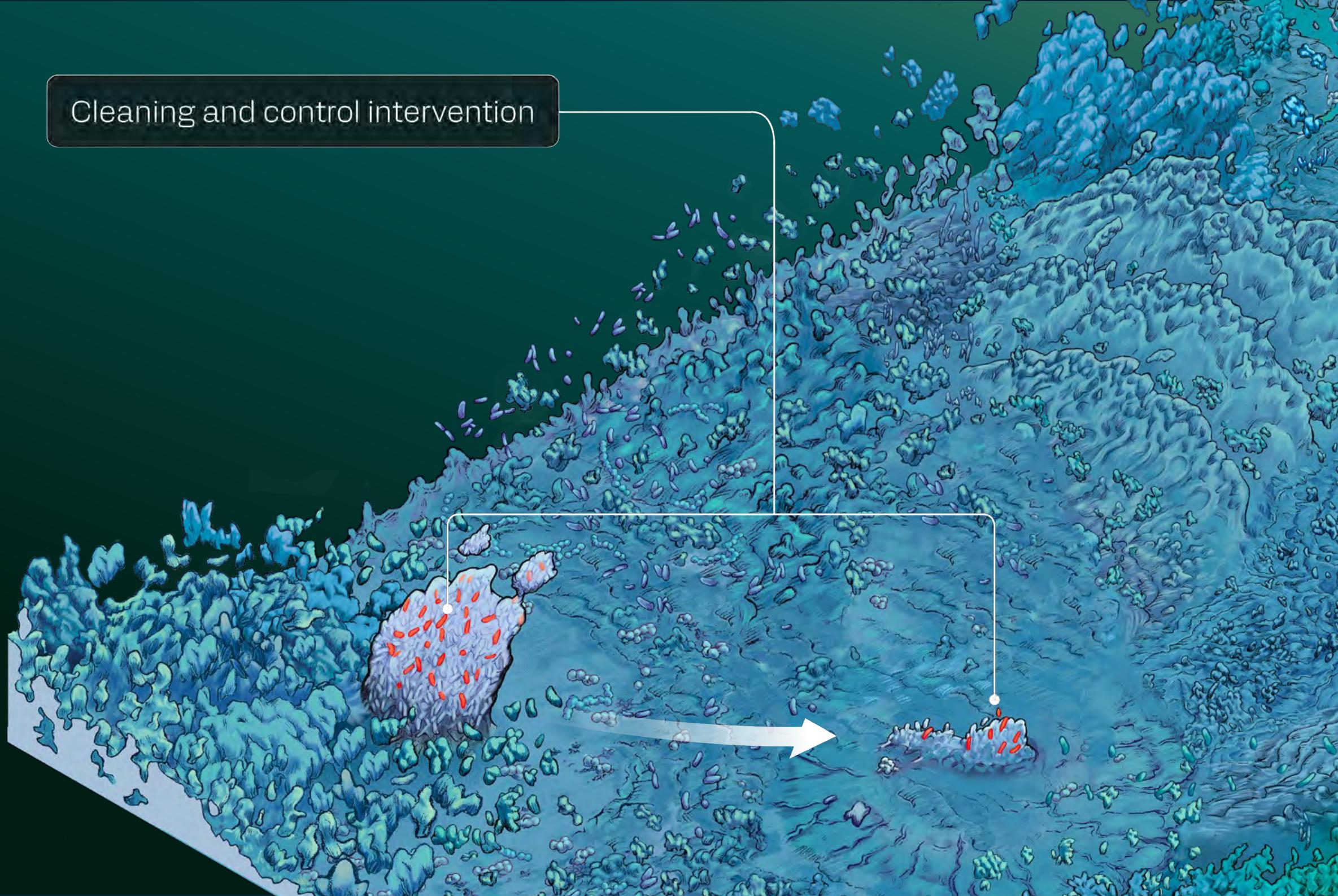


Microbial release

Pre-existing biofilm

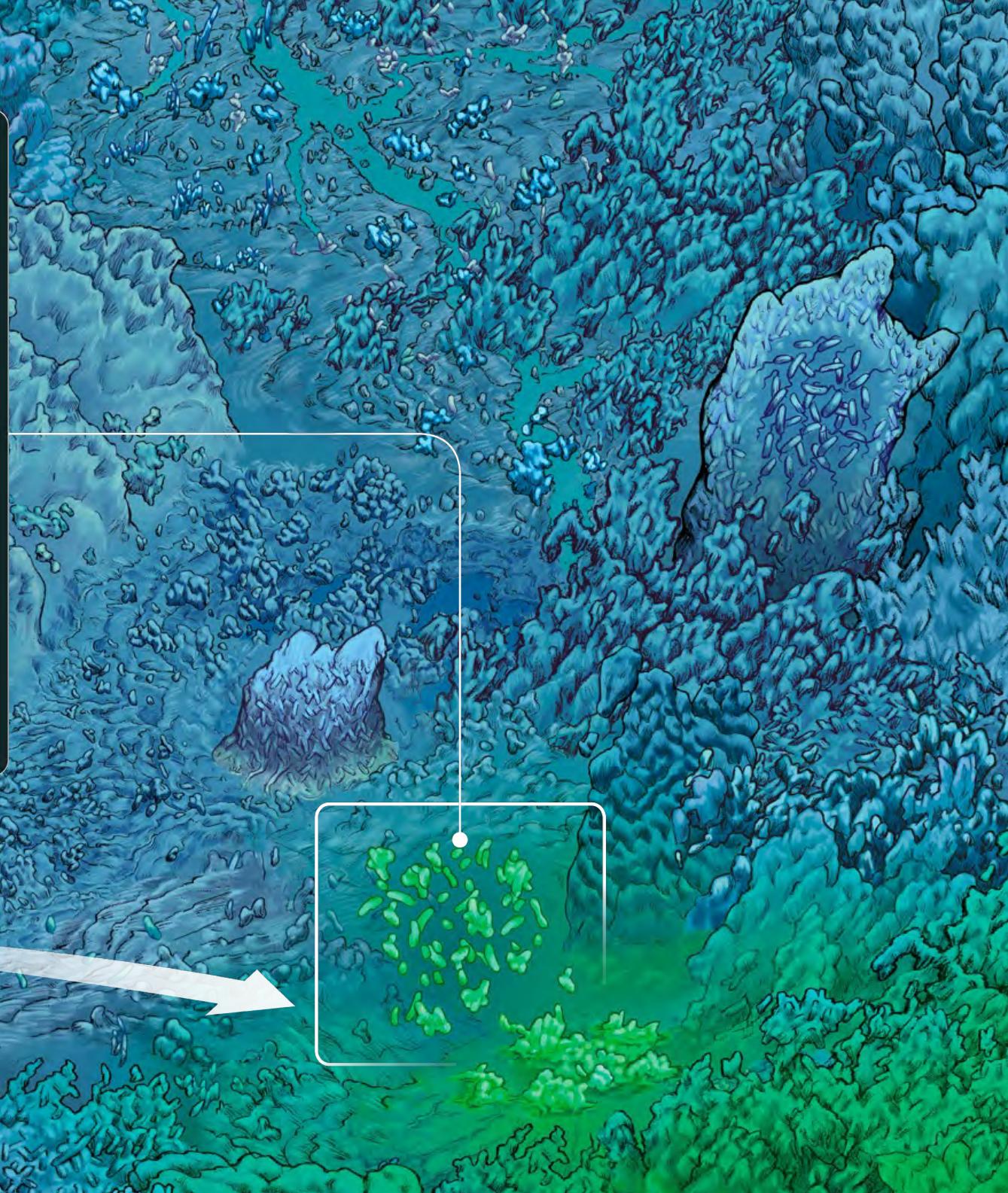


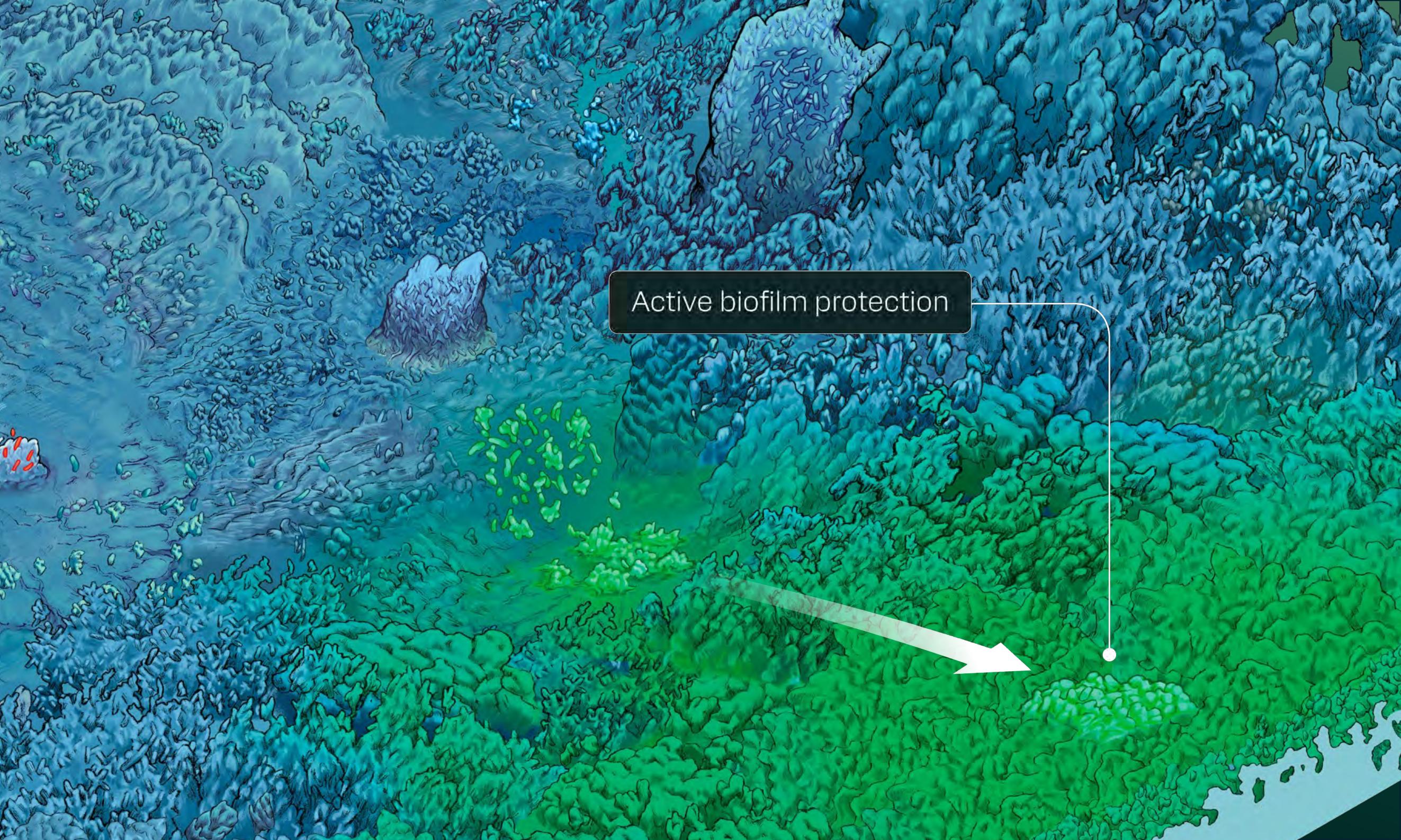
Cleaning and control intervention



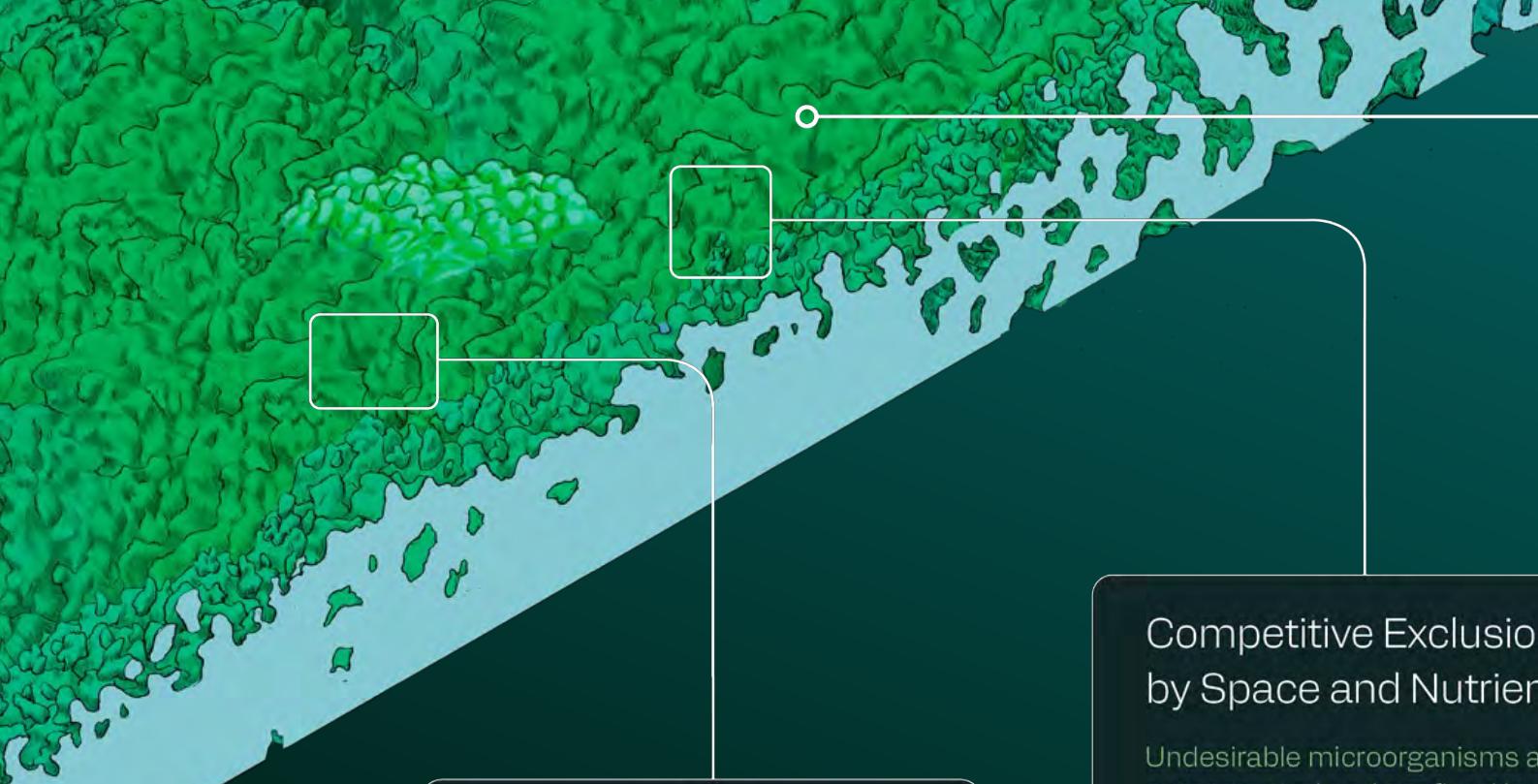
Establishment of a beneficial biofilm

Schematic showing the implantation of selected beneficial microbial consortia. Applied at high concentration to a sparsely colonized surface, they initiate a protective biofilm that prevents unwanted colonization.



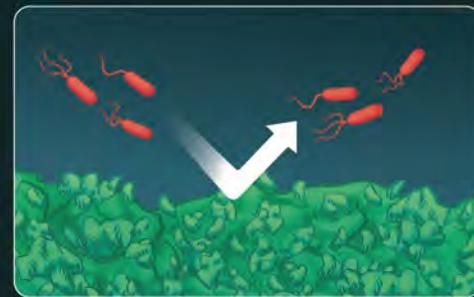


Active biofilm protection



Adhesion inhibition

Approaching pathogenic bacteria are repelled by adhesion-inhibitory molecules secreted by the protective biofilm, preventing surface attachment.



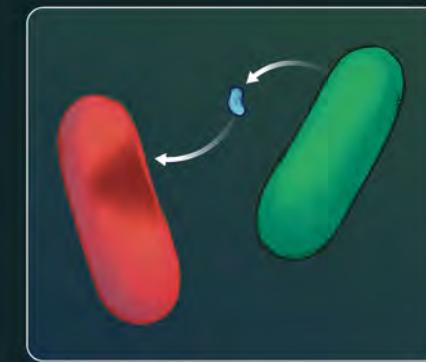
Competitive Exclusion by Space and Nutrients

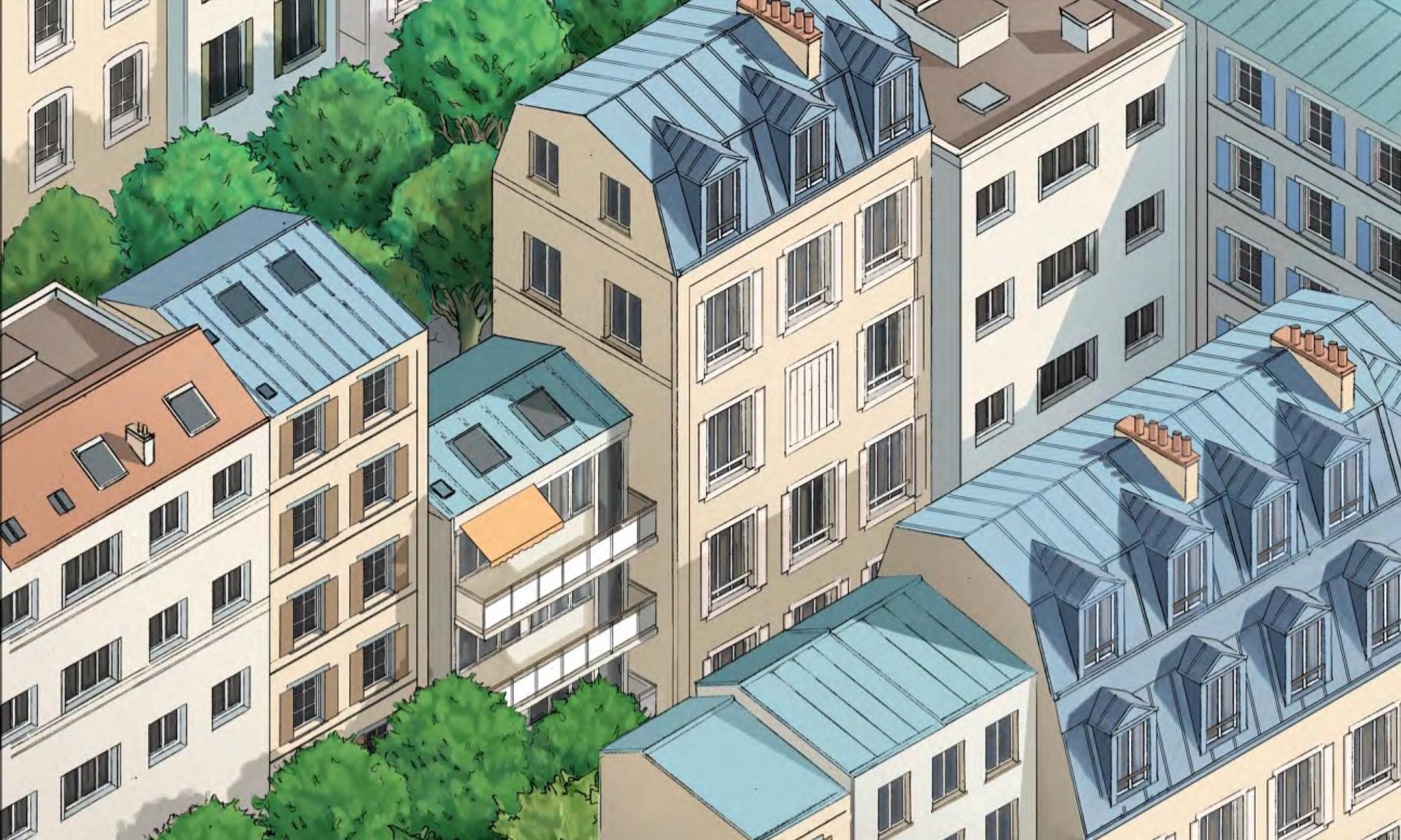
Undesirable microorganisms are outcompeted and confined by the dense beneficial population, which monopolizes both physical space and available nutrients within the biofilm.



Microbial interference

Beneficial bacteria interfere with unwanted populations by producing inhibitory metabolites that limit their growth or viability.





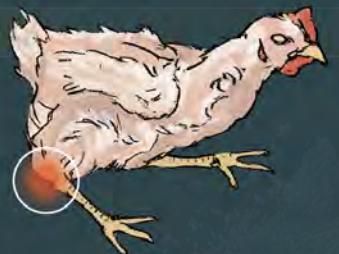


From contaminated surfaces to animal infections

Animals exposed to contaminated surfaces can develop localized infections, such as digestive tract colonization or joint infections, with potential propagation through cross-contamination between individuals.



Surfaces in livestock buildings are colonized by undesirable biofilms. These persistent structures on walls and equipment continuously release cells into the environment, increasing the risk of contamination and animal infection.

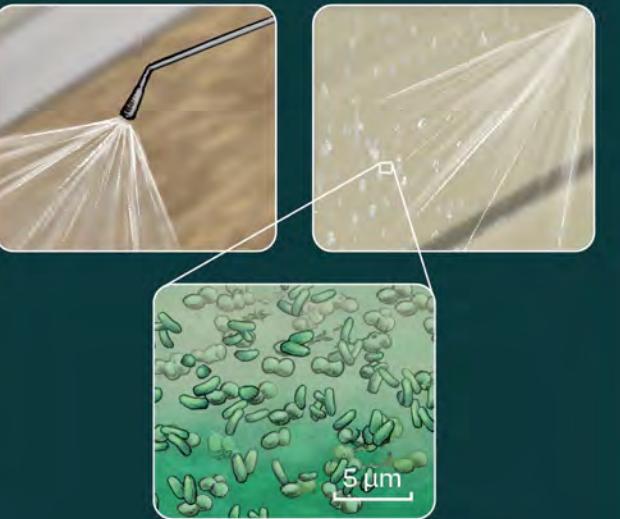


Salmonella enterica commonly colonizes the digestive tract in poultry, leading to enteric infections.

Enterococcus cecorum is frequently associated with joint infections in poultry, causing locomotor disorders.

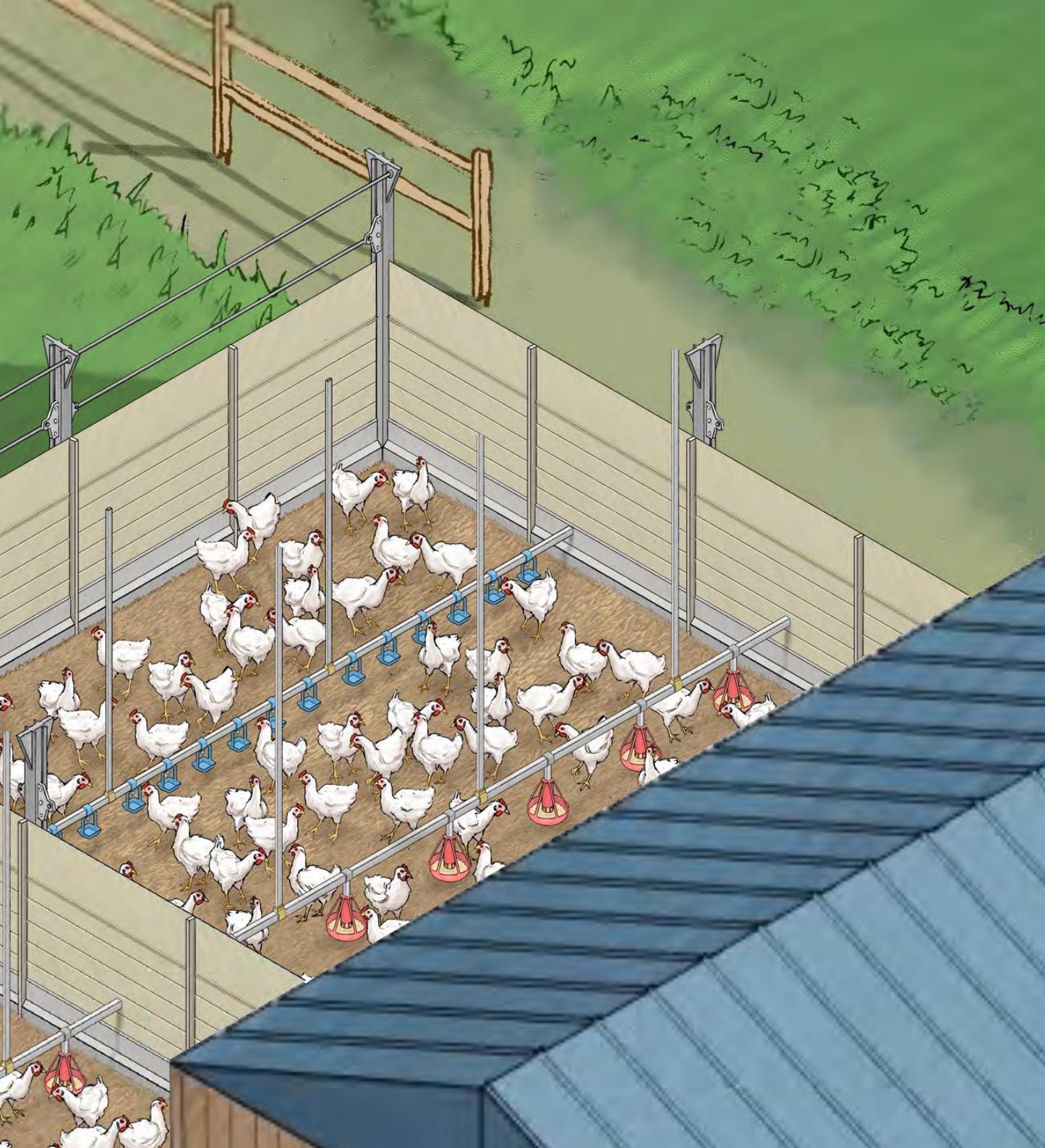


Establishment of a protective beneficial biofilm



A selected consortium of beneficial microorganisms is sprayed onto surfaces to establish a protective biofilm.

This active barrier competitively excludes undesirable microorganisms, preventing their attachment and proliferation.



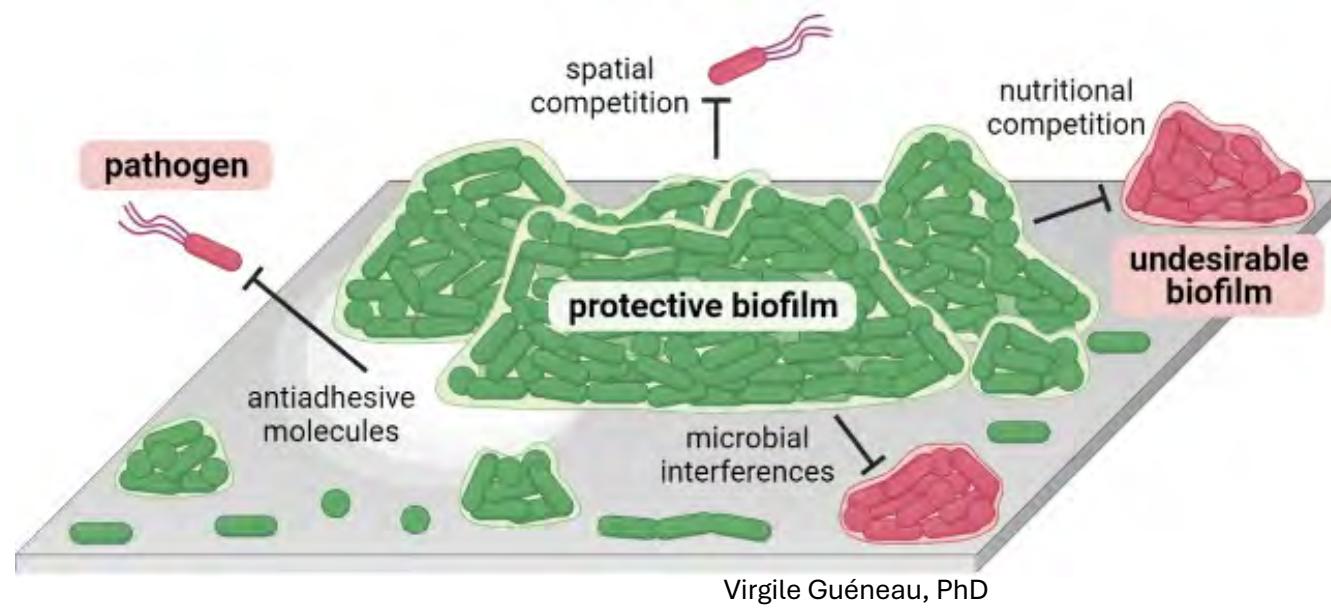




Graphic Design: ©Antoine Carlioz



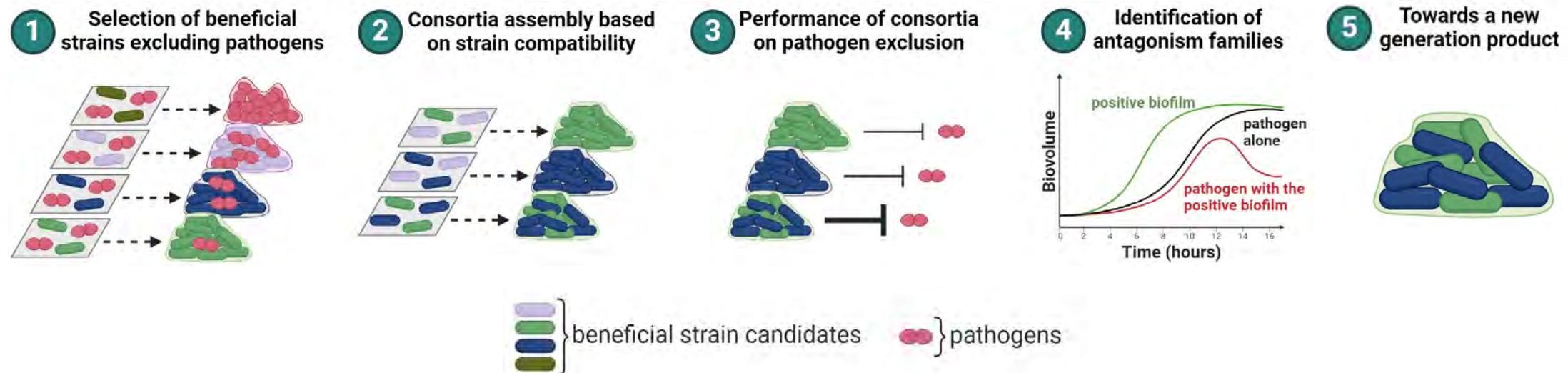
Exclusion of undesirable microorganisms by the settlement protective biofilms



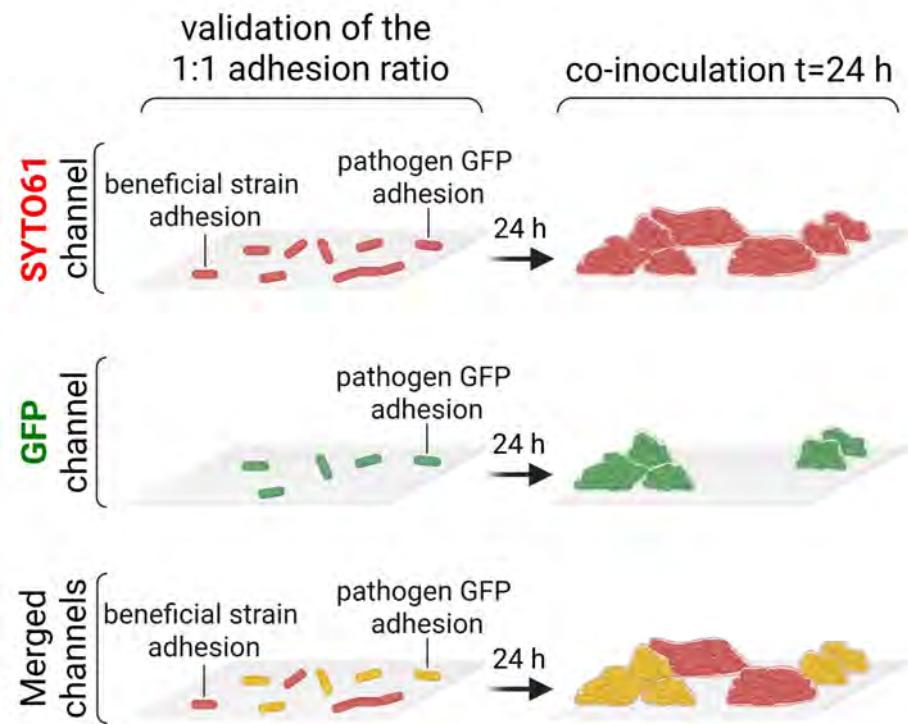


3D imaging-driven assembly of beneficial multi-species biofilms targeting undesirable bacteria

Virgile Guéneau, Laurent Guillier, Cécile Berdous, Marie-Françoise Noirot-Gros, Guillermo Jiménez, Julia Plateau-Gonthier, Pascale Serror, Mathieu Castex, Romain Briandet



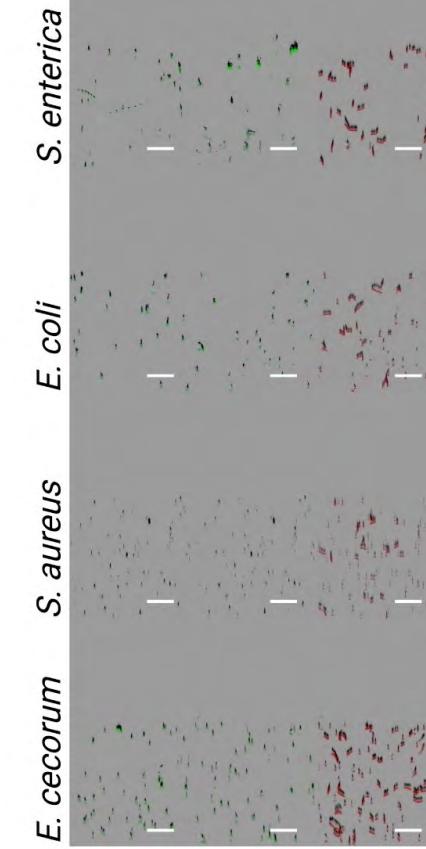
3D Imaging of co-culture models to investigate microbial interactions



validation of the 1:1 adhesion ratio

pahtogen *B. velezensis* ILPB8

GFP GFP SYTO61



E. coli

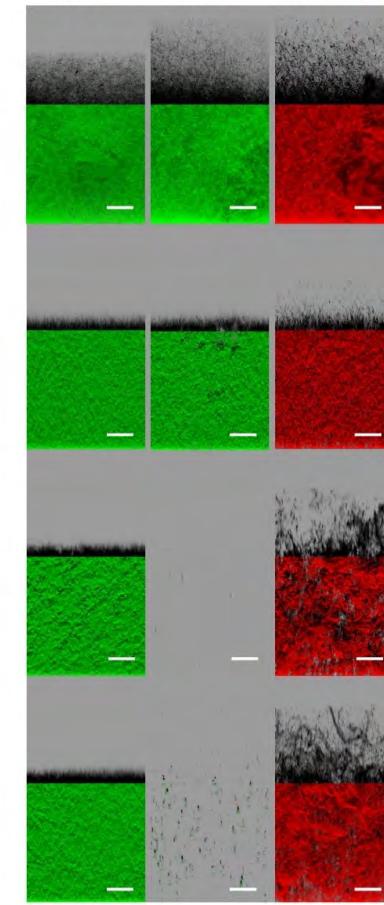
S. aureus

E. cecorum

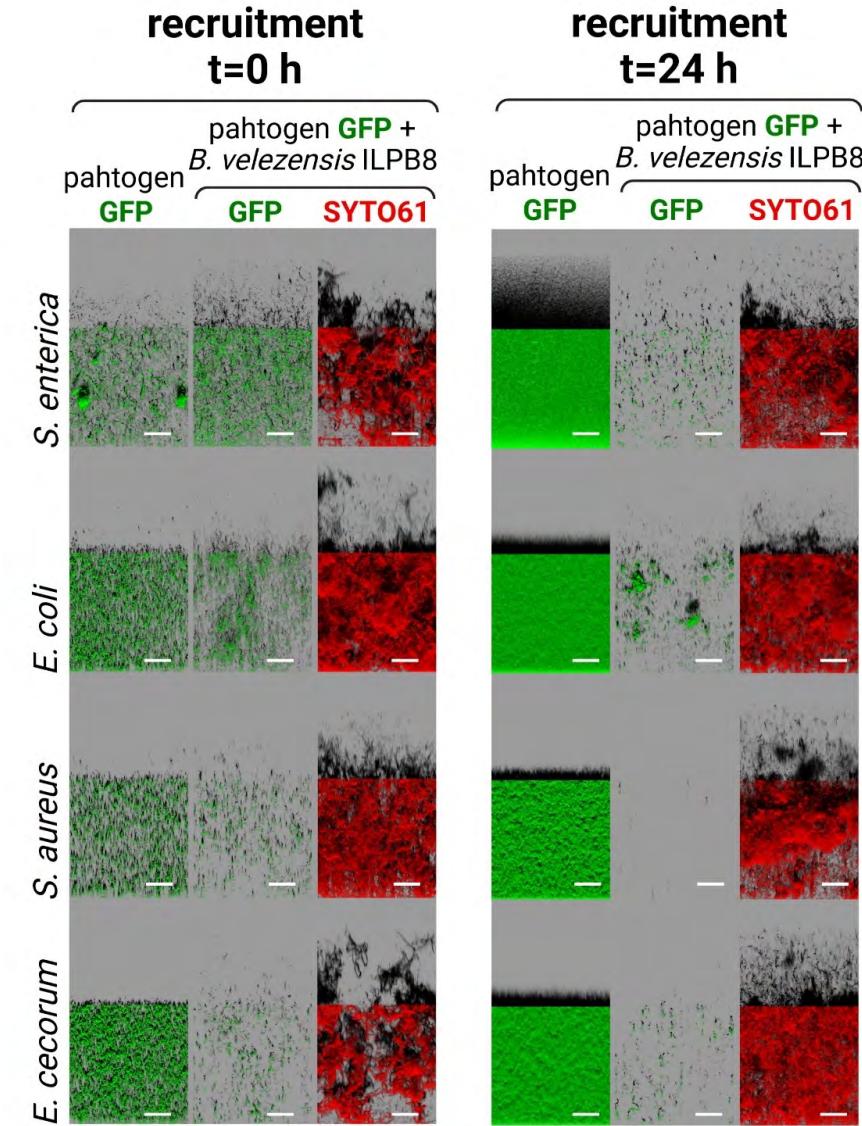
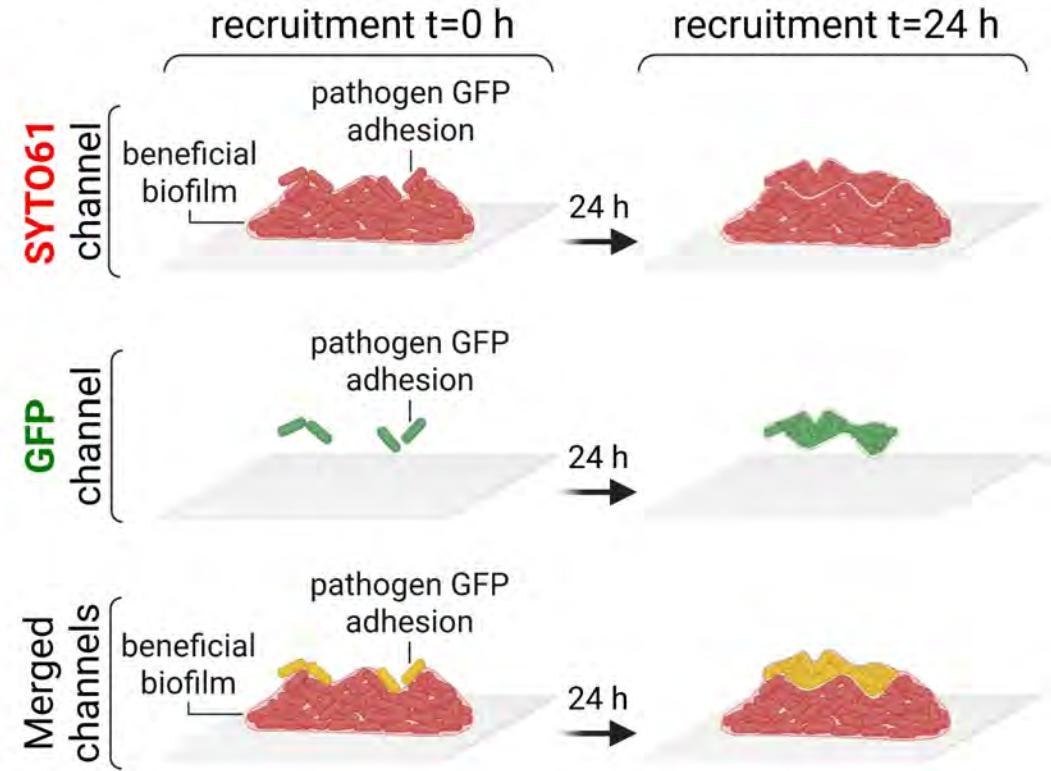
co-inoculation t=24 h

pahtogen *B. velezensis* ILPB8

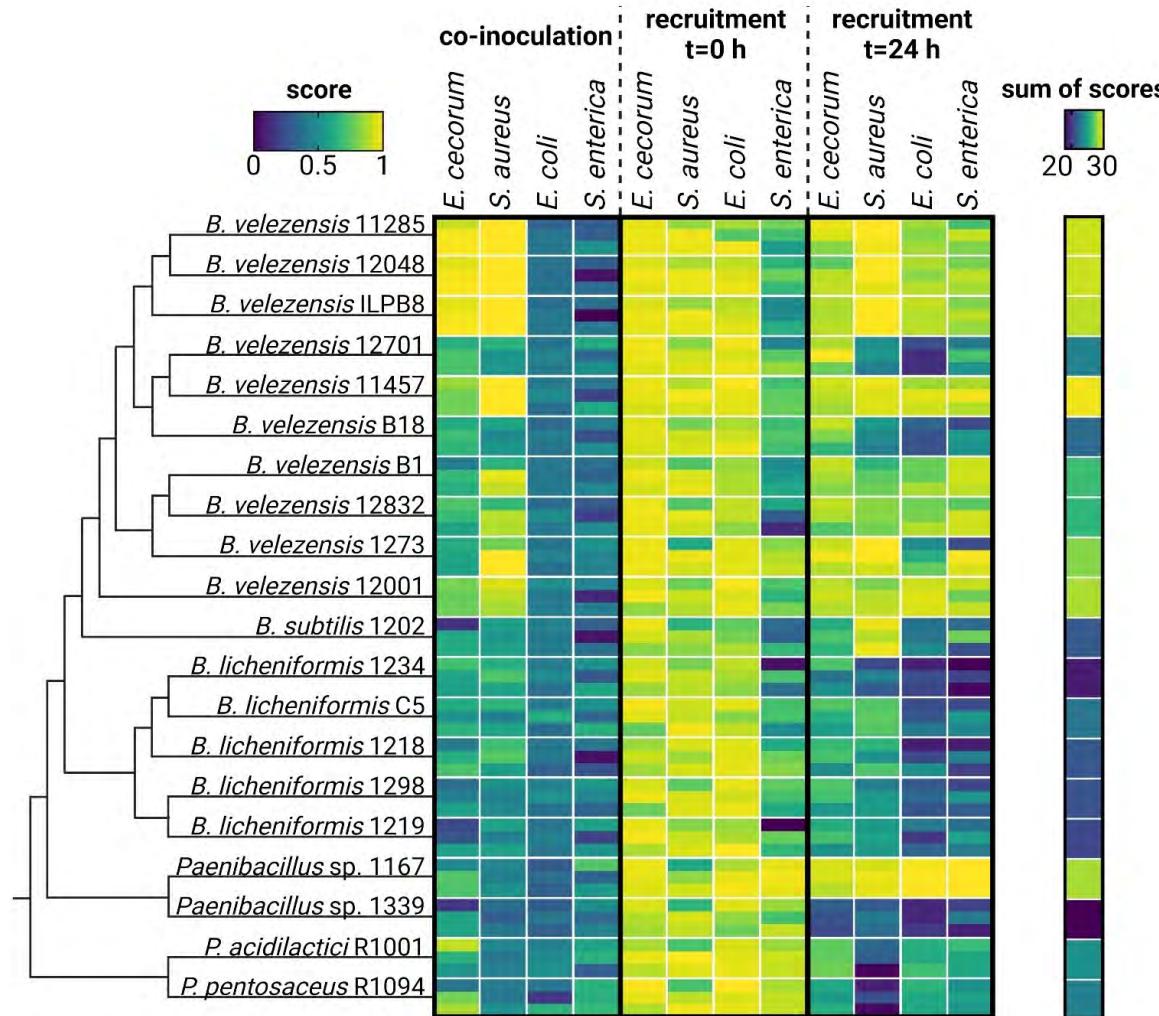
GFP GFP SYTO61



3D Imaging of co-culture models to investigate microbial interactions

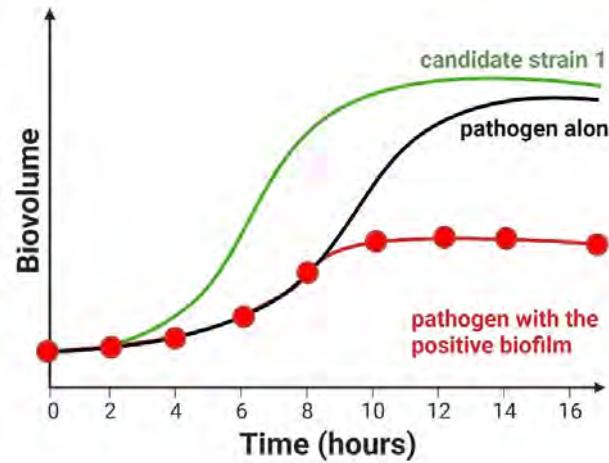


Assessing antagonistic activity against undesirable bacteria



- Diversity of responses
- Link between phylogenetic distance and exclusion
- Exclusion potential revealed when the biofilm is already formed
- Selection of best candidate strains

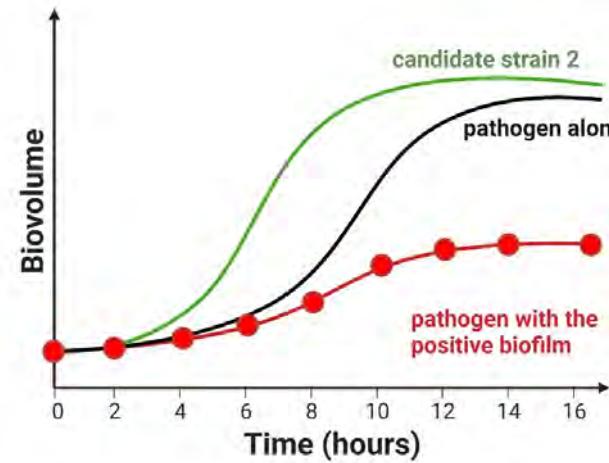
Time-resolved analysis of antagonistic mechanisms using 3D Imaging



Growth parameters:

- max biovolume: modified
- μ_{max} : not modified
- population decline: no

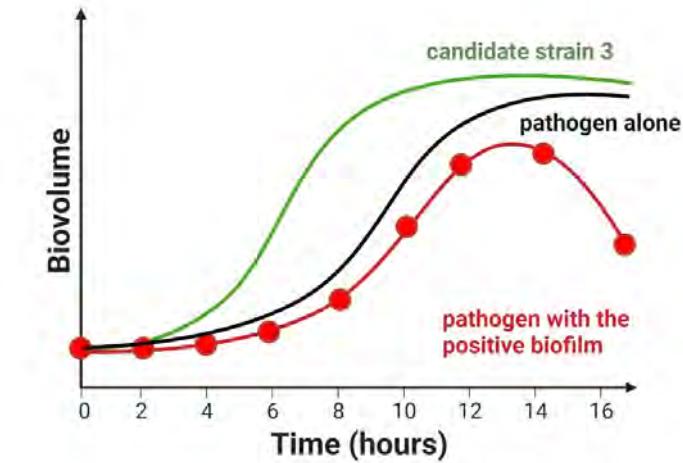
Nutritional and spatial competition described by the Jameson model



Growth parameters:

- max biovolume: modified
- μ_{max} : modified
- population decline: no

Nutritional and spatial competition described by the Jameson model + interference



Growth parameters:

- max biovolume: modified
- μ_{max} : modified or not
- population decline: yes

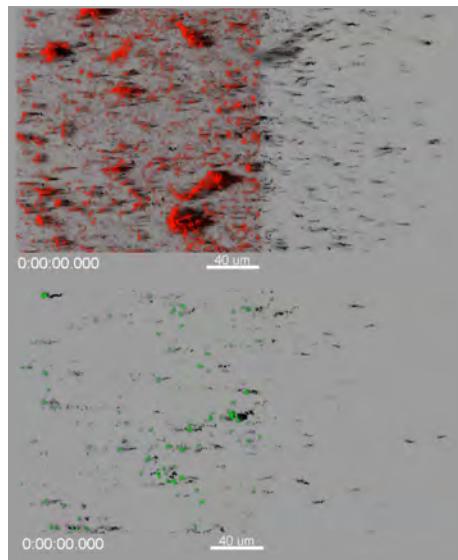
Prey-predator interaction described by the Lotka-Volterra model

Case studies of pathogen exclusion

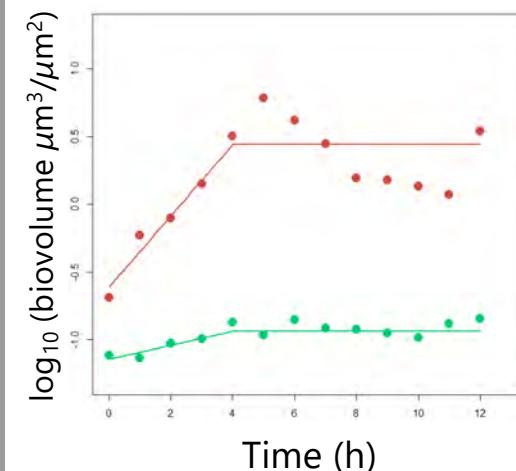
E. cecorum exclusion by *B. vellezensis*

Initial ratio in the co-inoculation and recruitment model:

E. cecorum < *B. vellezensis*

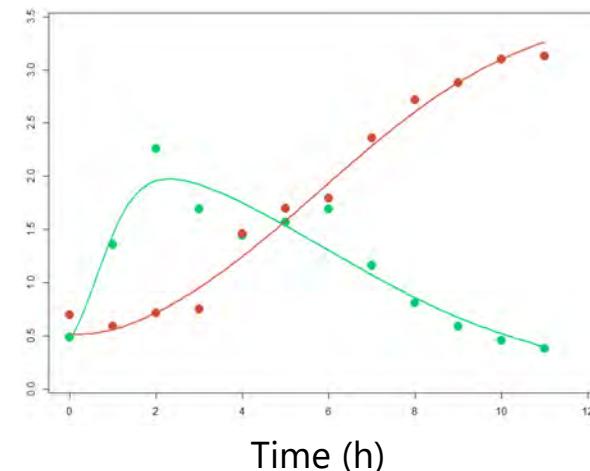
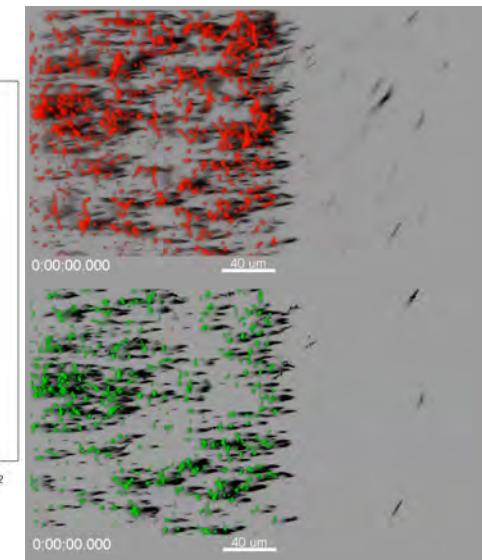


FM4-64 to label all the biofilm
GFP to label the pathogen



Initial ratio in the co-inoculation model:

E. cecorum > *B. vellezensis*



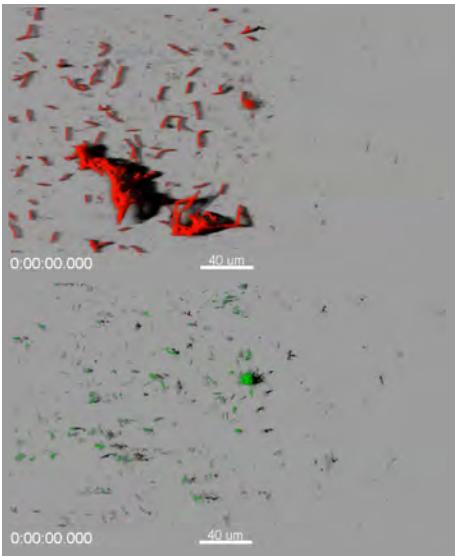
**Nutritional and spatial competition
described by the Jameson model
+ interference**

**Predation
described by the Lotka-Volterra model**

Case studies of pathogen exclusion

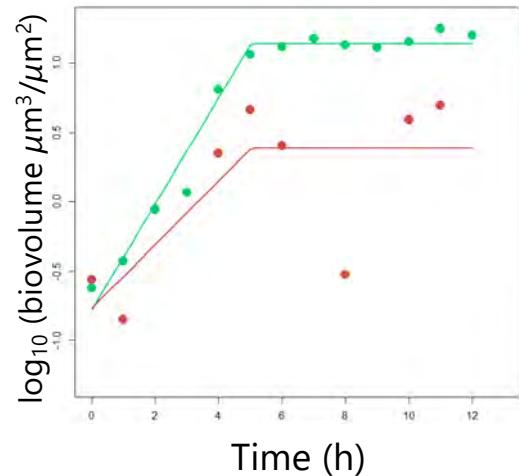
Pre-established *B. velezensis* biofilms inhibit *S. enterica* colonization

Co-inoculation model with all initial ratios:

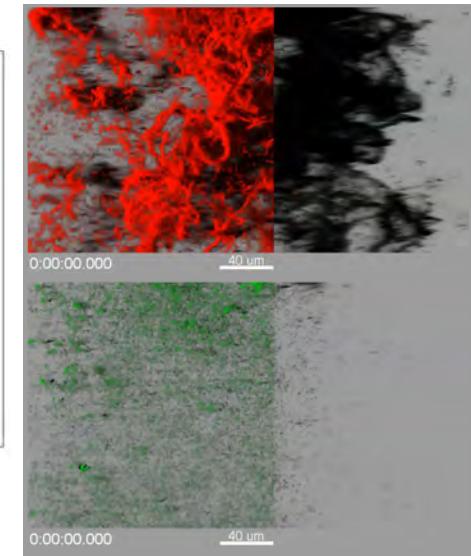
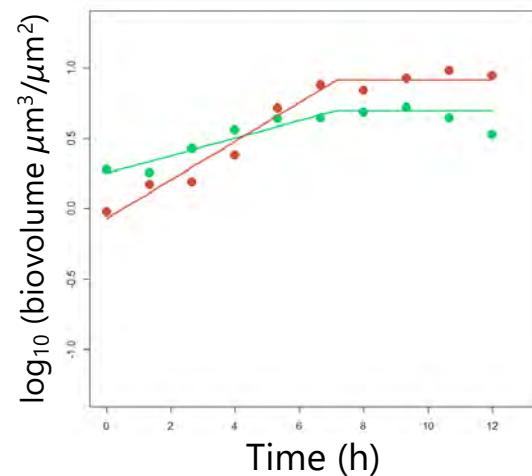


FM4-64 to label all the biofilm
GFP to label the pathogen

S. enterica growth is not affected



Recruitment model:

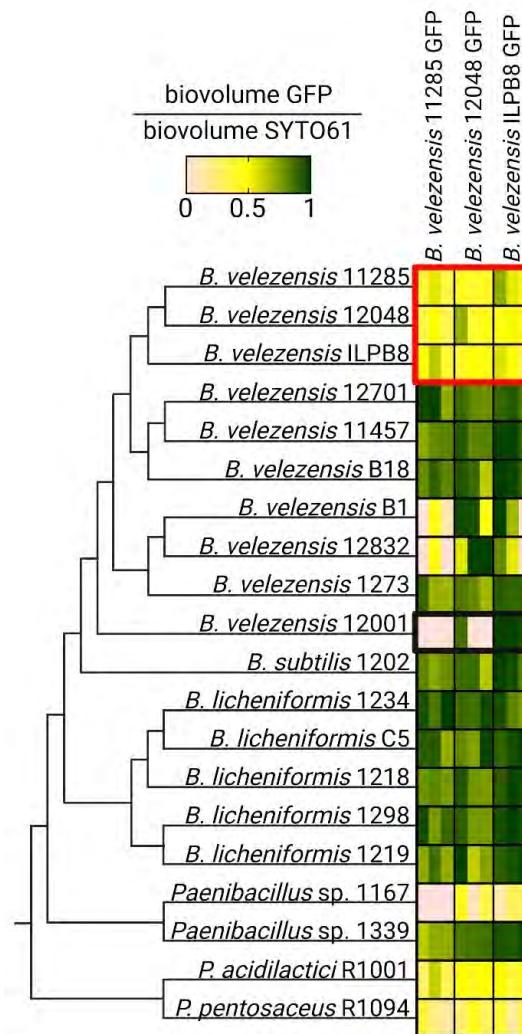


Nutritional and spatial competition
described by the Jameson model
+ interference

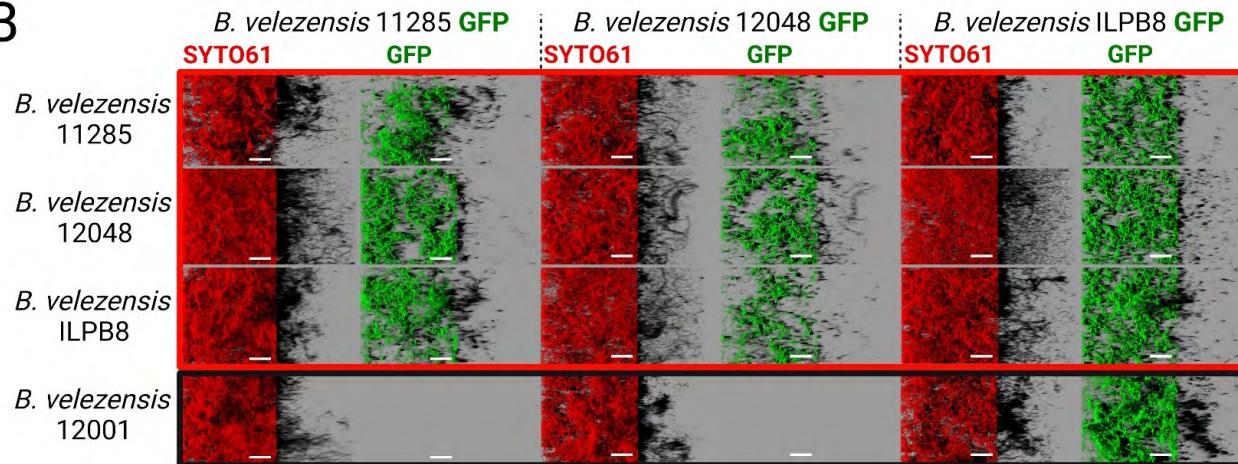
→ Generation time: 40 min to 5h

Design & compatibility of beneficial consortia

Evaluating compatibility in beneficial multispecies consortia



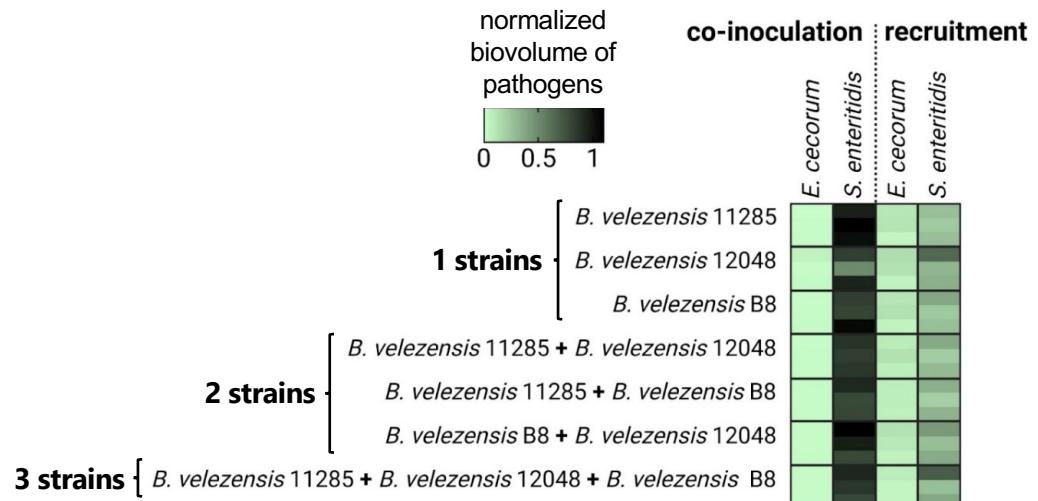
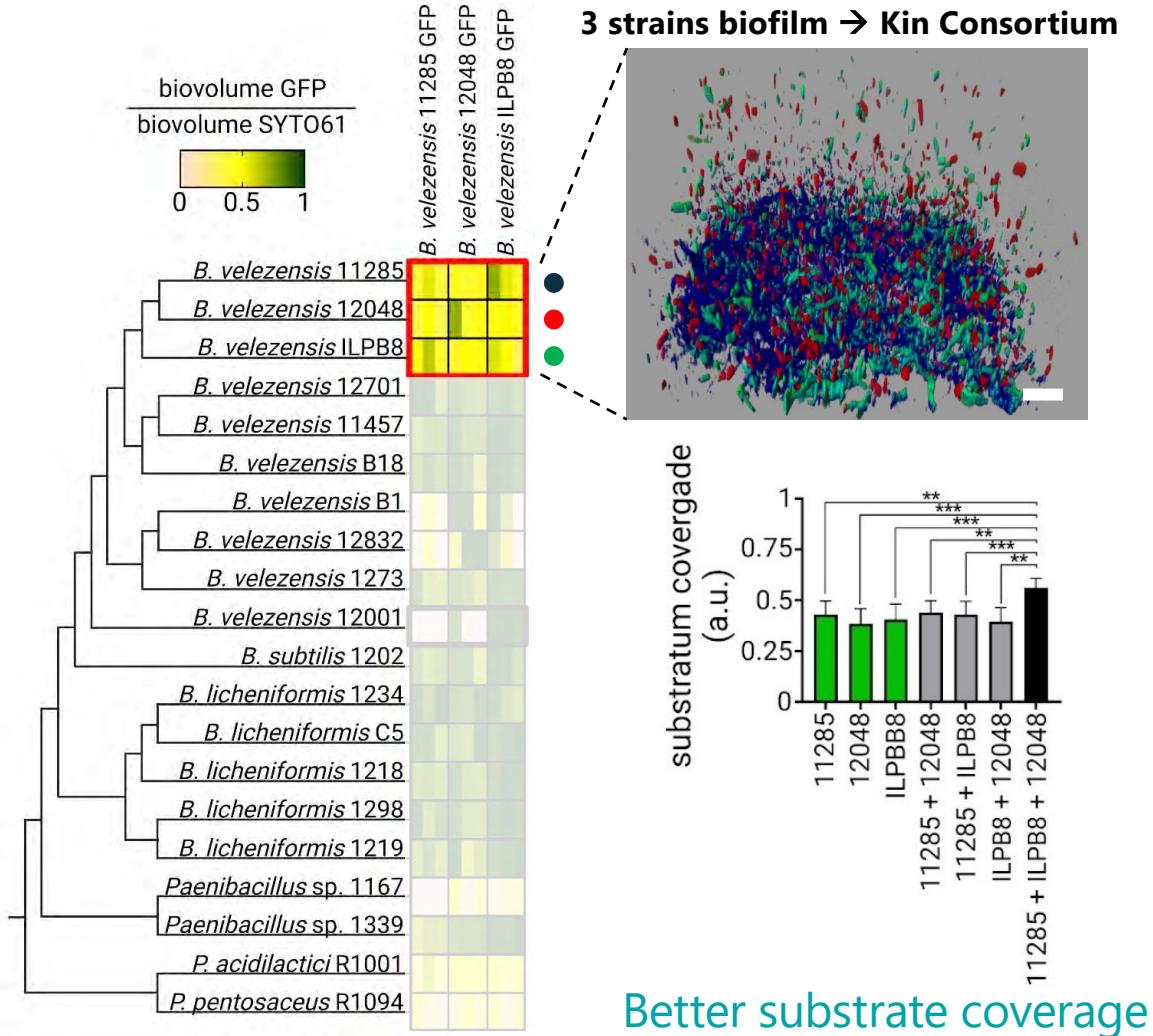
B



- **Very limited compatible consortia**
- Compatible strains are very close or very distant phylogenetically

Design & compatibility of beneficial consortia

Multi-strains *Bacillus velezensis* kin-consortium

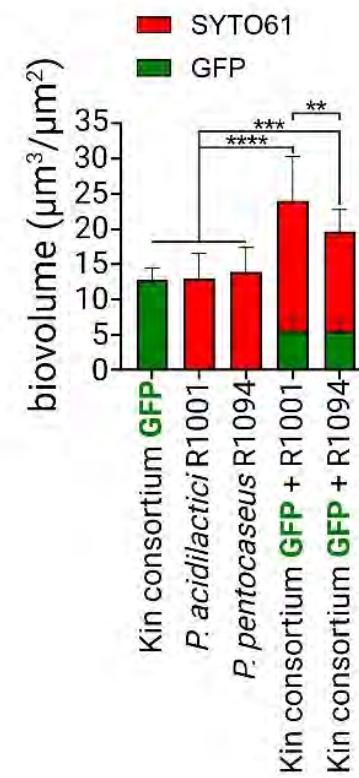
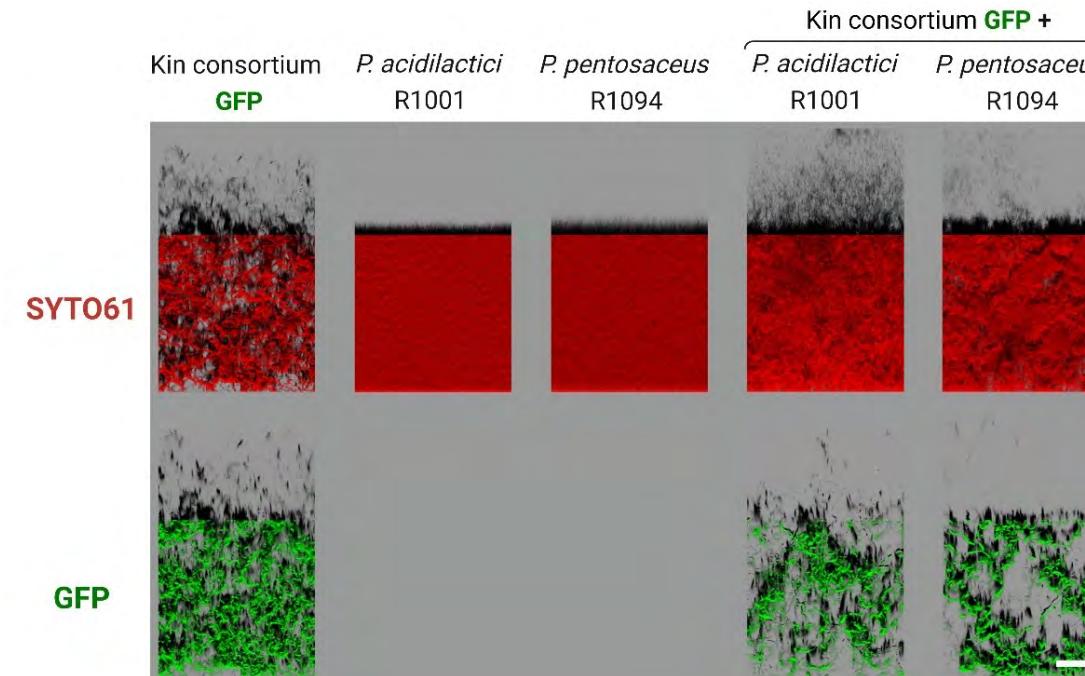
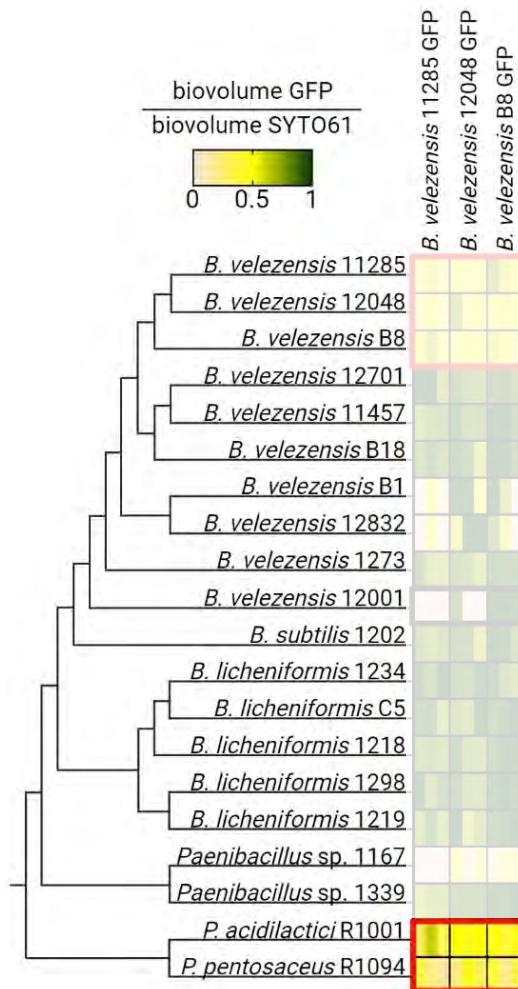


No additive effect of compatible SynComs on antagonism

→ Same antagonistic mechanisms?

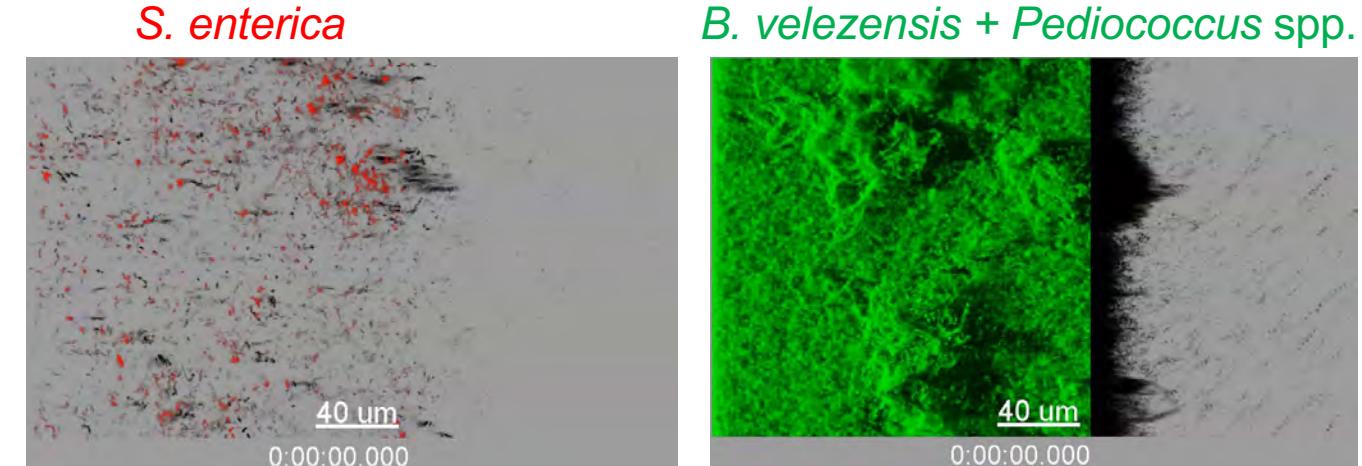
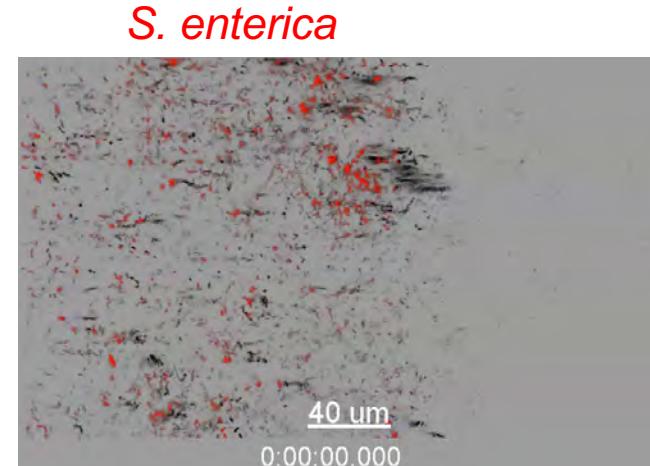
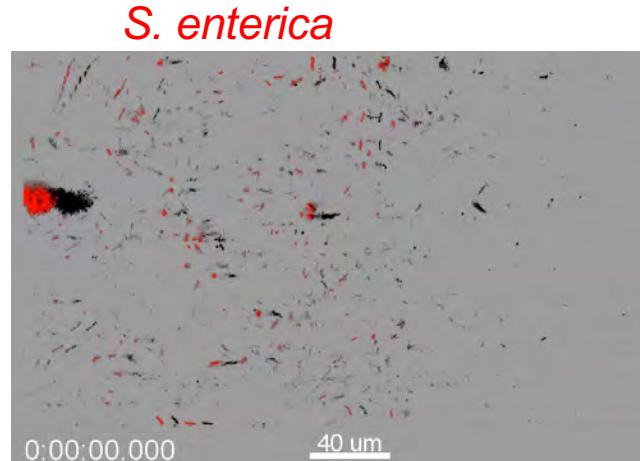
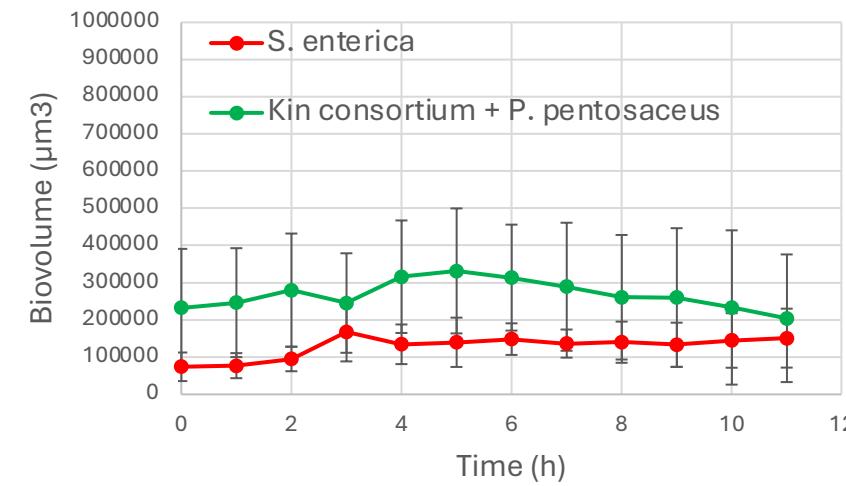
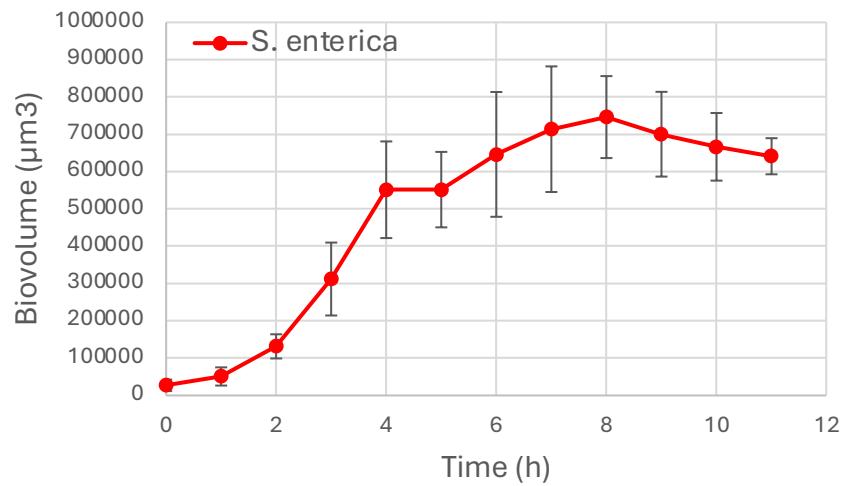
Design & compatibility of beneficial consortia

Multi-genus biofilms: enhancing protective effects through synergy?



Design & compatibility of beneficial consortia

Multi-Genus biofilms: enhancing protective effects through synergy?

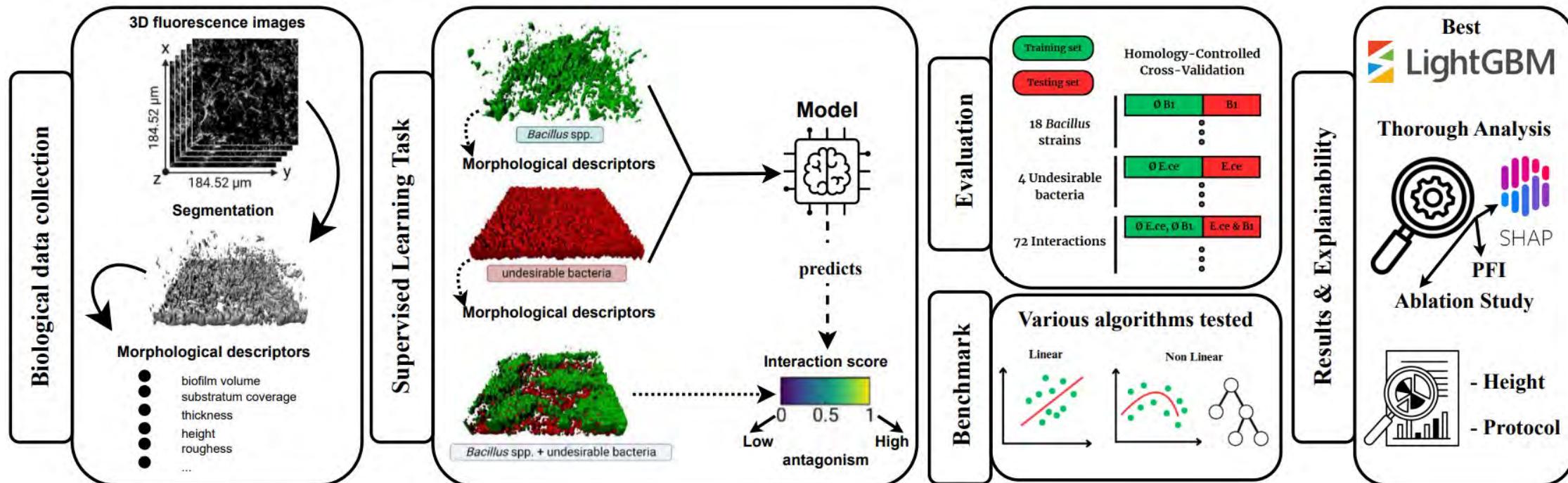


AI-guided strain selection

AI-assisted selection of beneficial strains for pathogen exclusion

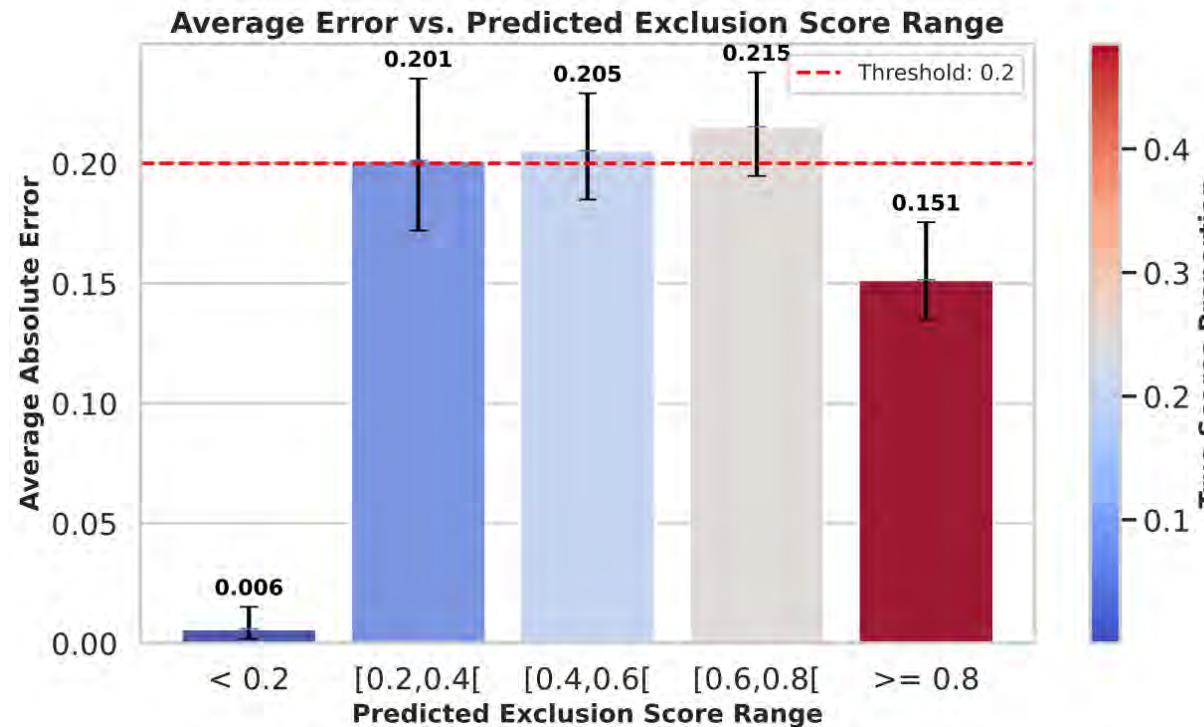
Strategy: use morphological descriptors of mono-strain biofilms to predict the interaction with pathogens

Objective: machine learning to reduce experiments for candidate strains selection



AI-guided strain selection

Accurate prediction of high exclusion potential using machine learning

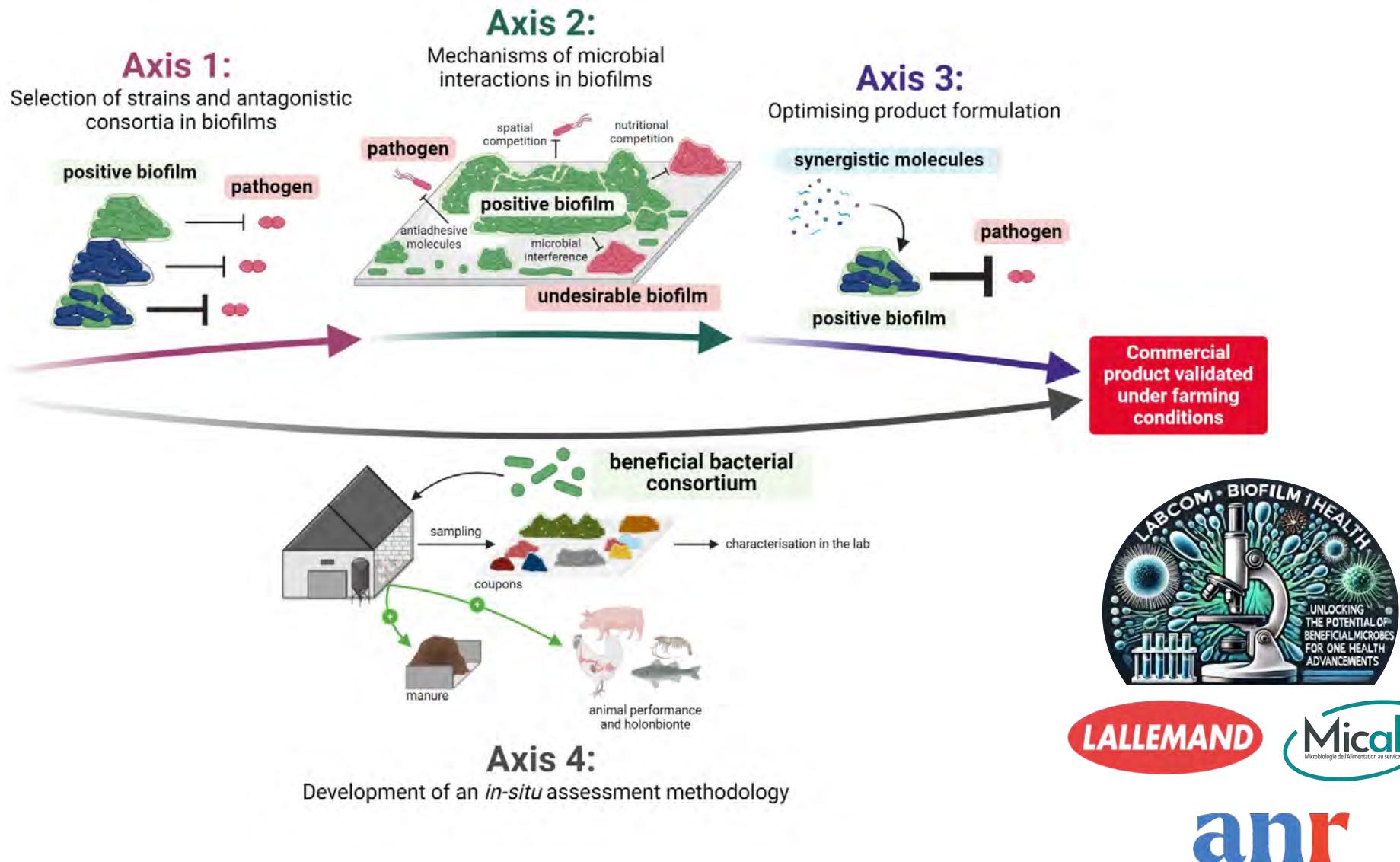


Many perspectives

- add genomes
- deep-learning on raw images
- other biofilm models
- add more data

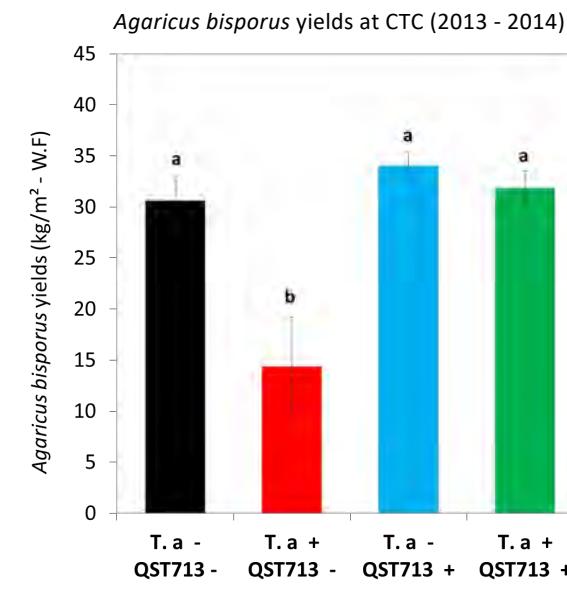
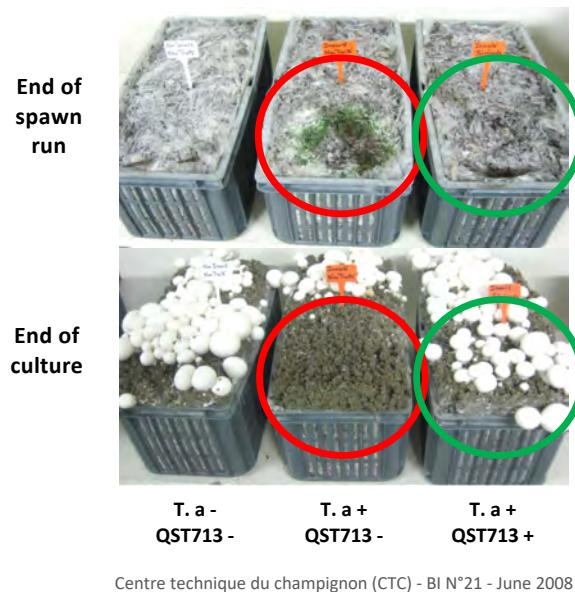
Project & Perspective: LabCom Biofilm1Health

Unlocking the potential of biofilm properties in beneficial microbes for One Health advancements



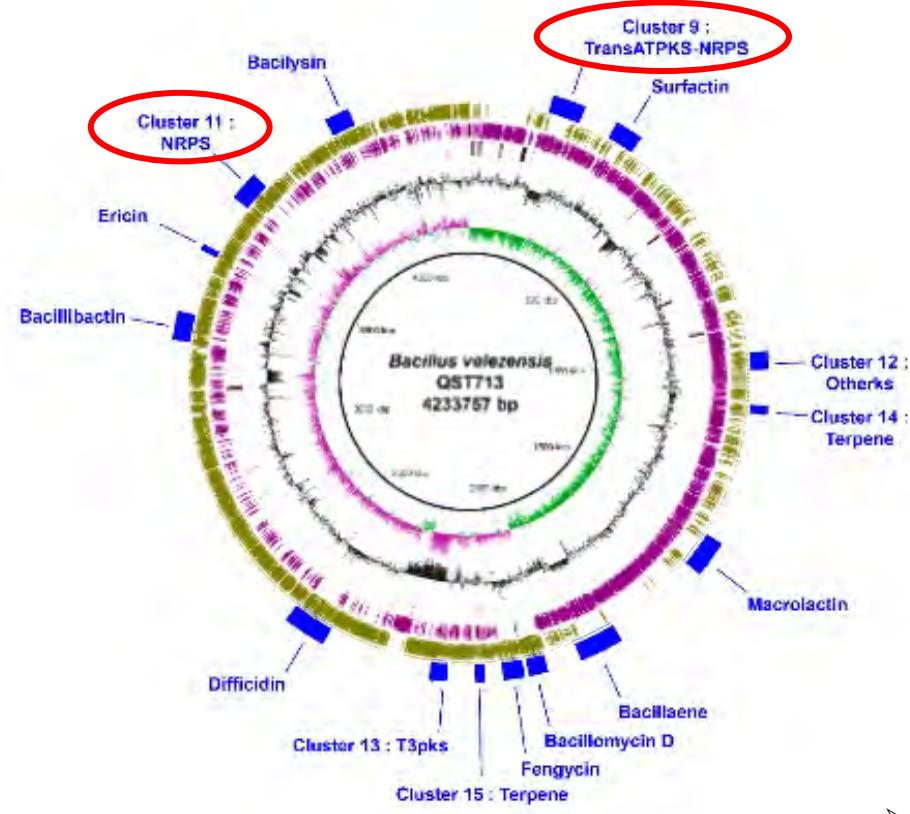
Protective biofilm in the French button mushroom sector

- > 25 % of annual losses of *Agaricus bisporus* → *Trichoderma aggressivum* (green mould)
- > Biocontrol used for 10 years in France → *Bacillus subtilis* vellezensis QST713



↗ 52 % Return to normal yield

Bacillus velezensis QST713



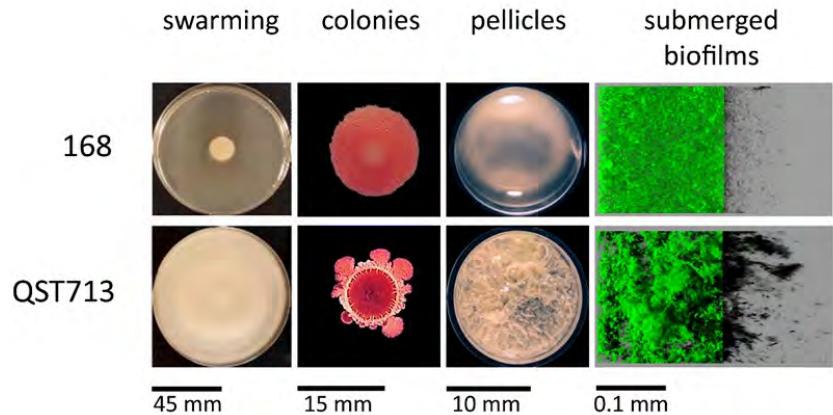
| <i>Bacillus velezensis</i> strain QST713 | |
|--|---------|
| Genome size (bp) | 4233757 |
| G + C content (%) | 45,9 |
| Total number of genes | 4263 |
| Coding sequences (CDSs) | 4056 |
| rRNAs genes | 25 |
| tRNAs genes | 79 |
| ncRNAs genes | 5 |
| Pseudo genes (total) | 98 |

Secondary metabolites: antiSMASH webserver

| Cluster number | Synthetase | Metabolites | bp | Bioactivity |
|----------------|------------------|------------------------|--------|----------------|
| 1 | NRPS | Surfactin | 65414 | Antimicrobial |
| 2 | TransATPKS | Macrolactin | 85893 | Antibacterial |
| 3 | TransATPKS-NRPS | Bacillaene | 102672 | Antimicrobial |
| 4 | NRPS | Bacillomycin D | 44669 | Antimicrobial |
| 5 | TransATPKS-NRPS | Fengycin | 49543 | Antimicrobial |
| 6 | TransATPKS | Difficidin | 100457 | Antibacterial |
| 7 | Bacteriocin-NRPS | Bacilibactin | 66790 | Antibacterial |
| 8 | NRPS | Bacilysin | 58252 | Antibacterial |
| 9 | TransATPKS-NRPS | - | 77736 | Not determined |
| 10 | Lantipeptide | Subtilin-like / Ericin | 15123 | Antibacterial |
| 11 | NRPS | - | 68430 | Not determined |
| 12 | OtherKS | - | 41245 | Not determined |
| 13 | T3PKS | - | 41101 | Not determined |
| 14 | Terpene | - | 20741 | Not determined |
| 15 | Terpene | - | 21884 | Not determined |

- *B. velezensis* QST713 = **12%** of the genome allocated to the biosynthesis, regulation and transport of antimicrobials
- *B. subtilis* = **4-5%** (Stein 2005; PMID:15853875)
- *B. velezensis* FZB42 = **8.5%** (Chen *et al.* 2009; PMID: 19041913)
- **2 potential new antimicrobials**

QST713 Abilities: biofilm formation, swarming motility & antibiosis



Phenotypic comparisons of *B. velezensis* QST713 with *Bacillus subtilis*

168

T. Aggressivum (T.

a)



168 vs T. a



QST713 vs T. a



Interaction tests on agar plates with *B. velezensis* QST713 or *B. subtilis* 168 with *T. aggressivum* at 25°C

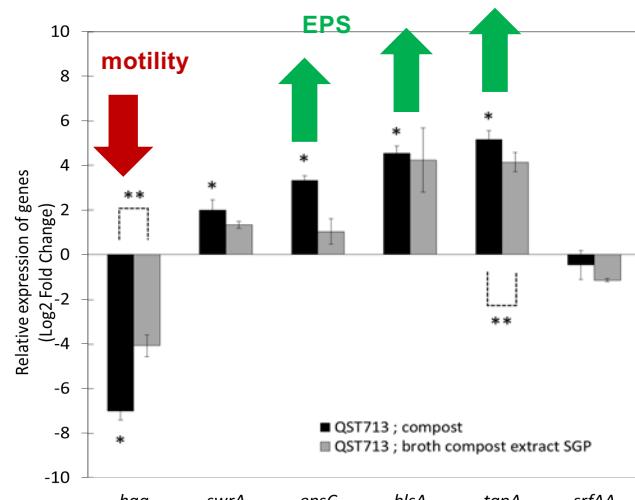
➤ ***B. velezensis* QST713 :**

- Swarming motility
- Complex macrocolony structure which is mucoid under a thick and dry surface layer
- Thick & solid pellicle
- Spatially organised submerged-biofilm

➤ **Pathogen inhibition test:**

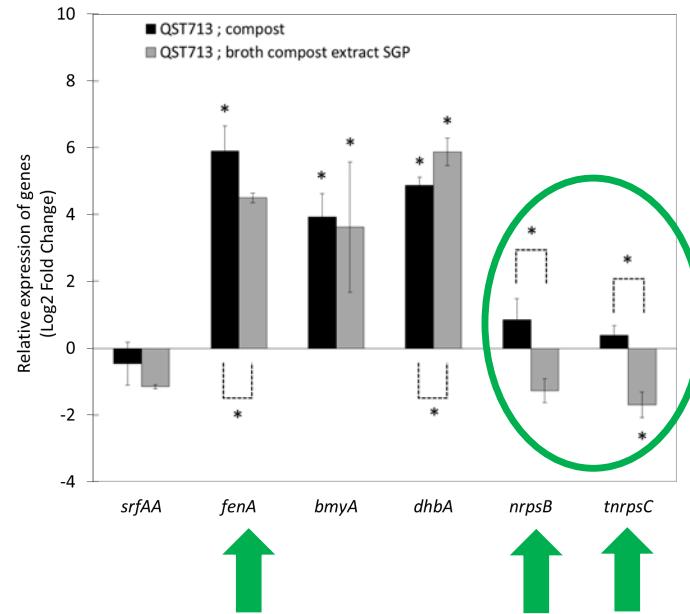
- No inhibition of *T. aggressivum* by *B. subtilis* 168
- Antibiosis of *T. aggressivum* by *B. velezensis* QST713 : growth inhibition and sporulation inhibition

In situ gene expression of the biocontrol agent



Control group: QST713 in broth compost extract (planktonic), exponential growth phase (EGP)

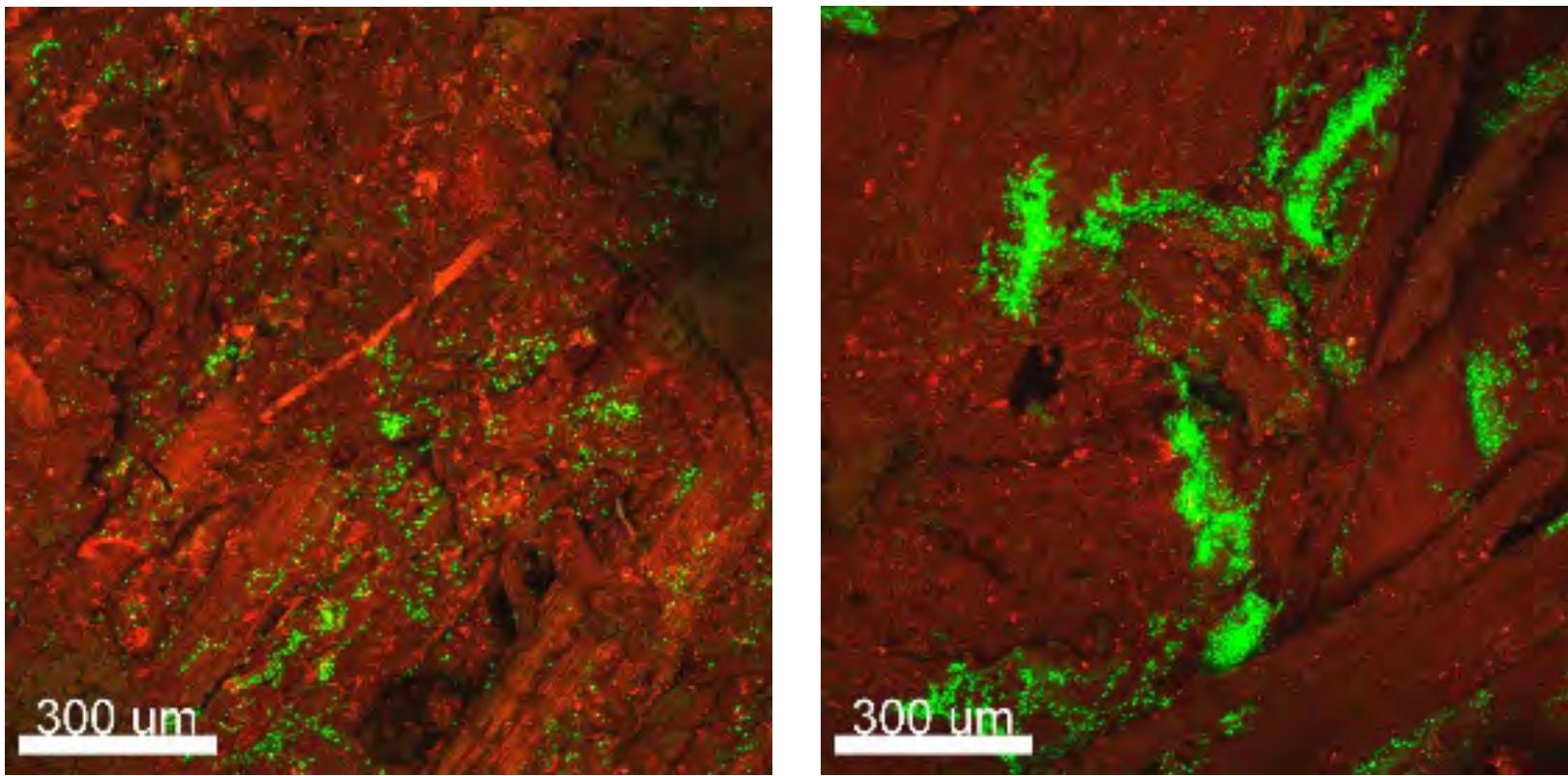
> biofilm is the preferential mode of life of the biocontrol agent in compost micromodels



> Several antimicrobials upregulated in the biofilm form

Direct visualisation of biofilm formation in the compost by fluorescent *B. velezensis* QST713 at 6 days of culture

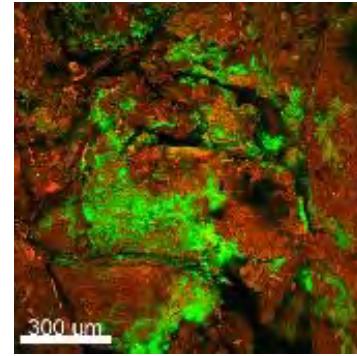
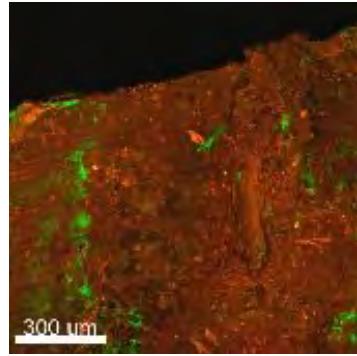
Confocal laser scanning microscopy visualisation



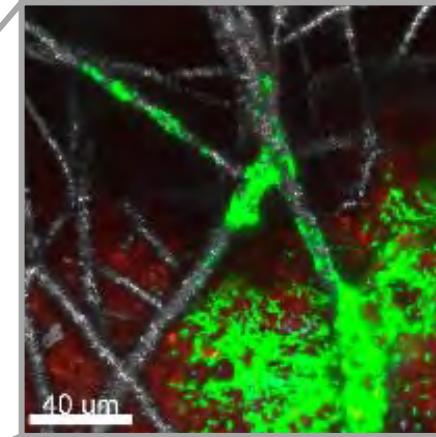
Biofilm formation by fluorescent *B. velezensis* in the compost

- Biocontrol strain *B. velezensis* FZB42 authorized in button mushroom field

FZB42



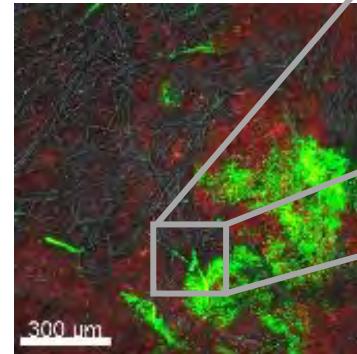
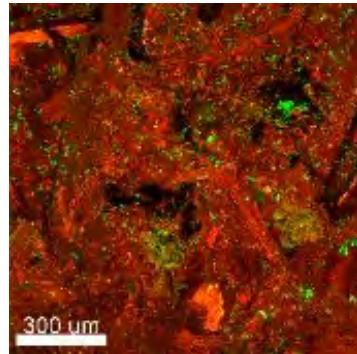
Proximity with *Agaricus* mycelium



Use of available biofilm and antimicrobial gene mutants of strain FZB42

- Other *B. velezensis* strain with biocontrol effect against fungi

SQR9



micro-aggregates were most frequently observed

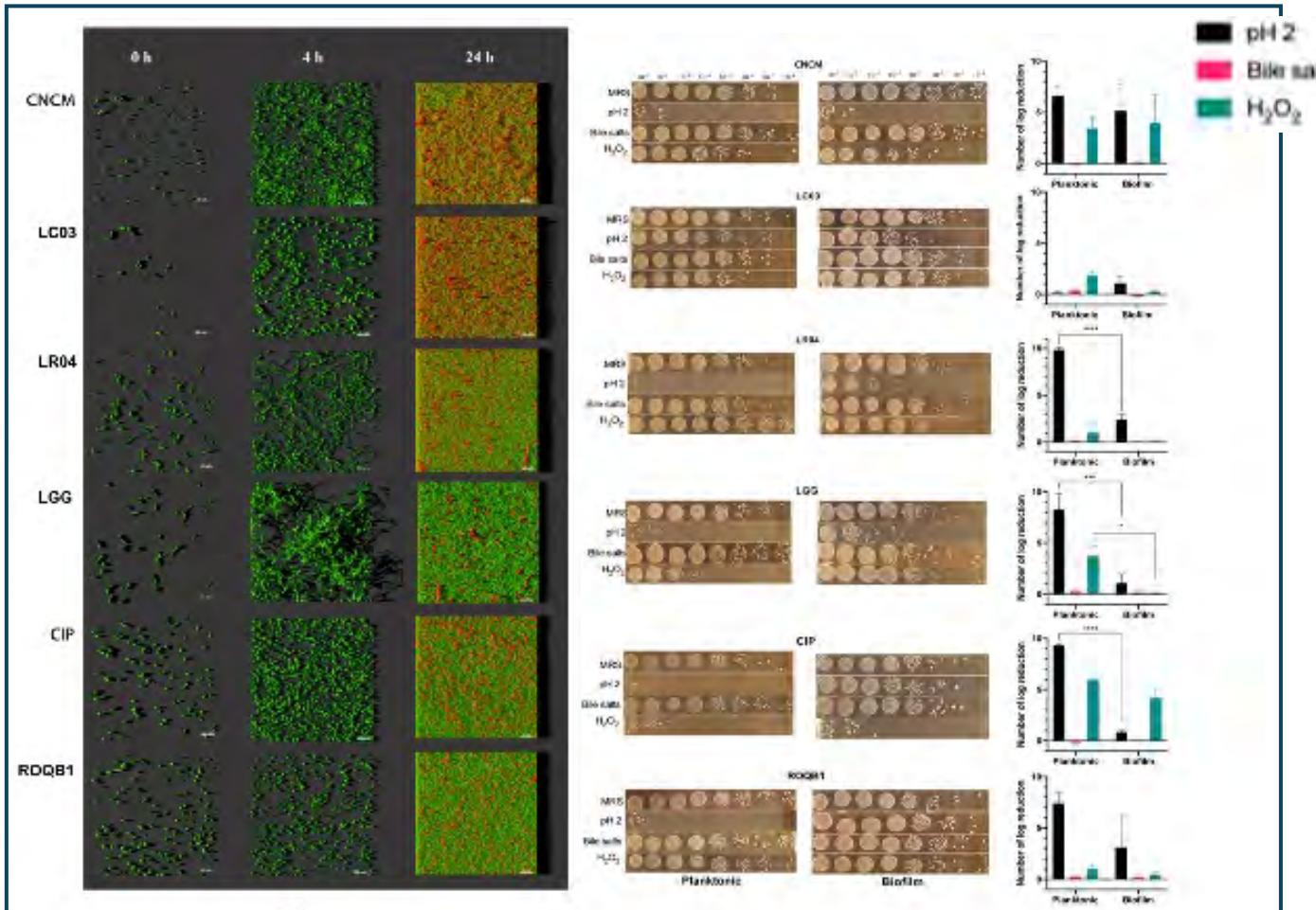
massive spatially organized biofilms were punctually observed

Bacteria use fungal « highways » to disperse?

Simon et al. 2015 (PMID:26432804)

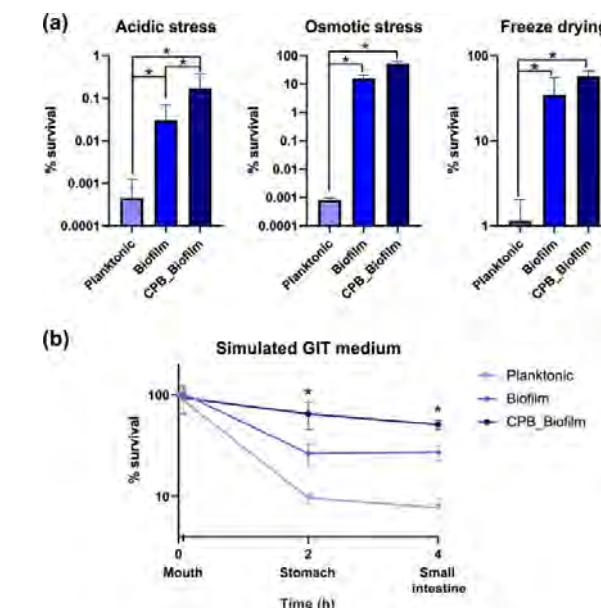
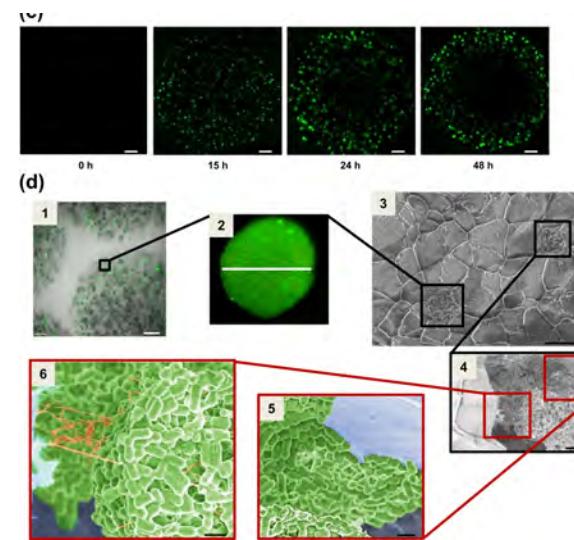
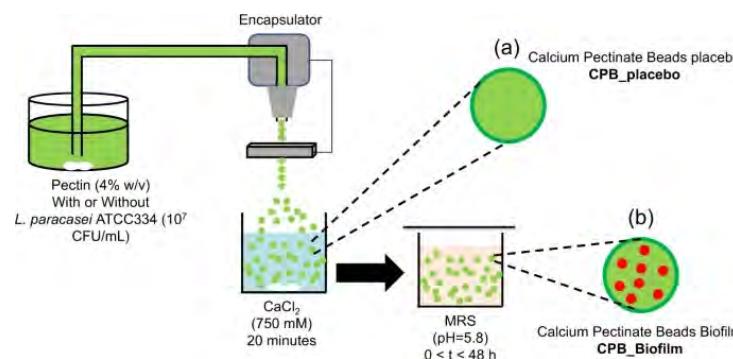
Confocal laser scanning microscopy visualisation

Potential probiotic strains tolerance to digestive stresses

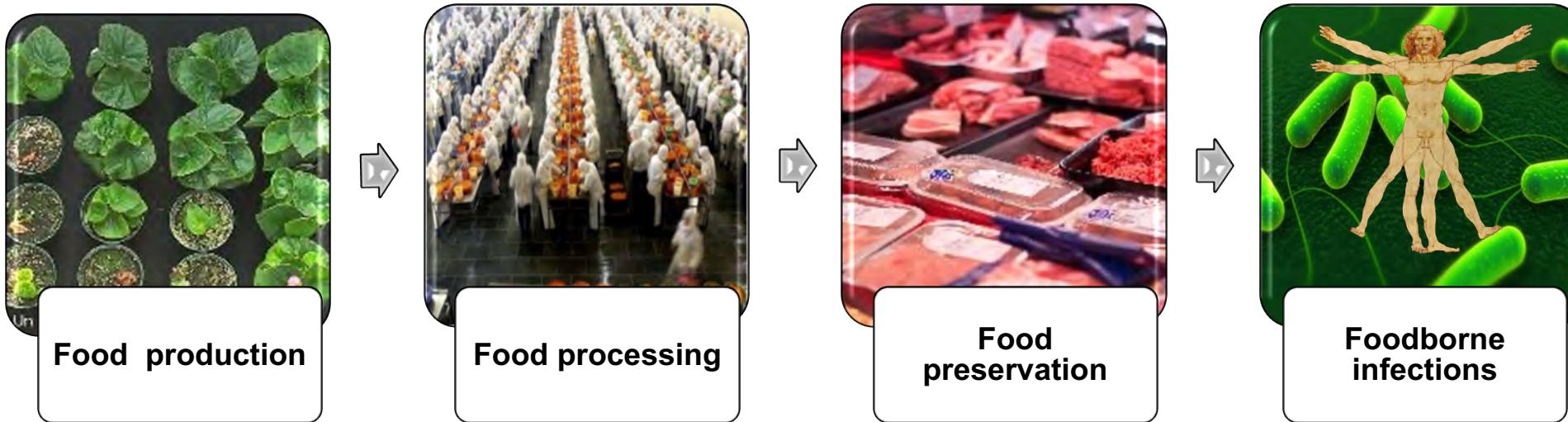


Chamignon et al. 2020. *Microorganisms*

Biofilm-like microcolonies encased in calcium-pectinate beads increases probiotic properties of *Lacticaseibacillus paracasei*



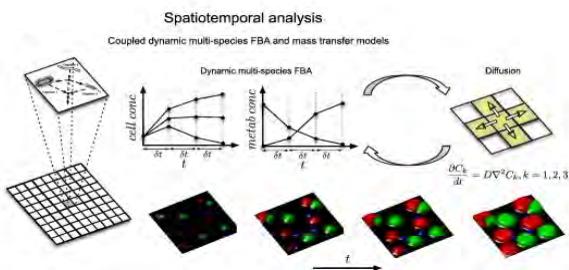
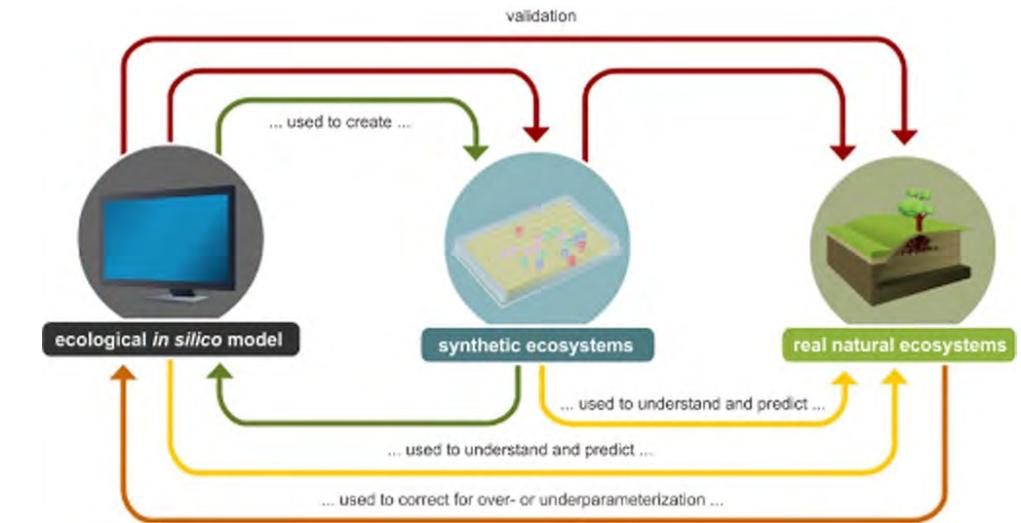
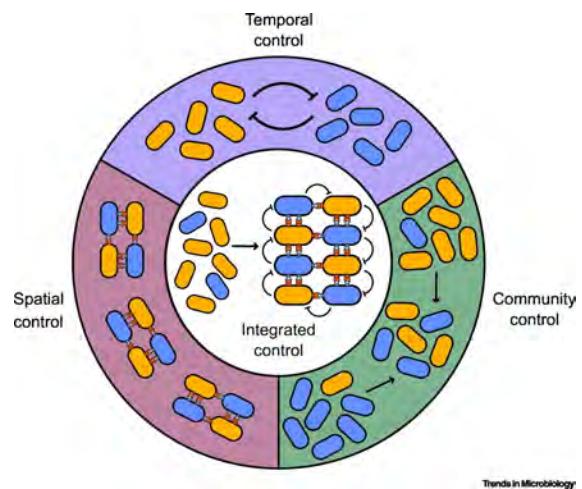
Toward microbial-based solution for biofilm control?



- > pesticides, disinfectants, preservatives, antibiotics
- > biocontrol, bioprotection, biopreservation, probiotics

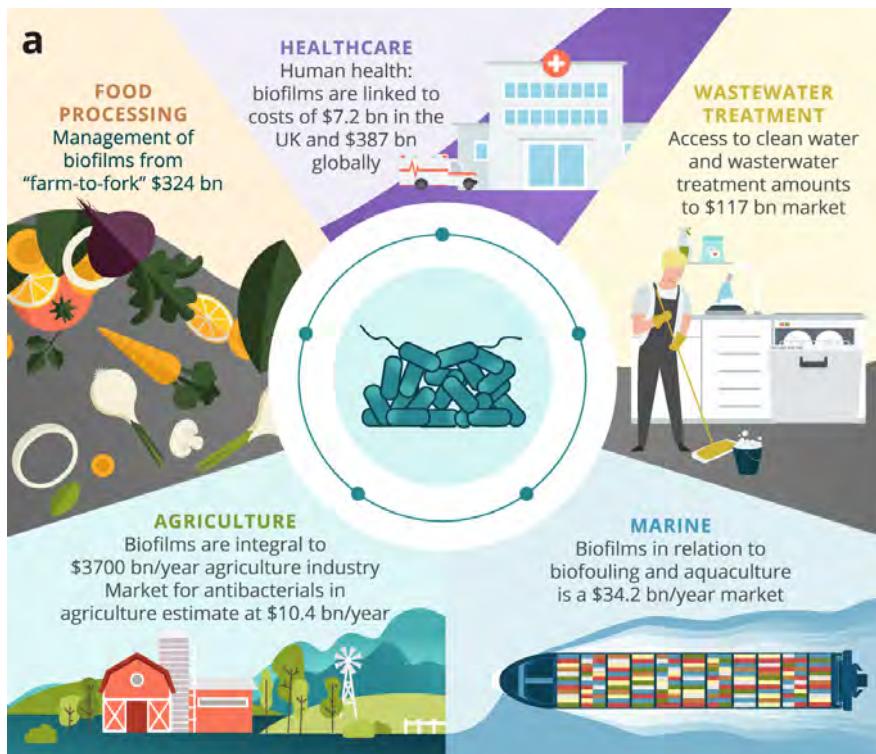
Future perspectives and challenges

> synthetic microbial ecology, artificial and programmable biofilms

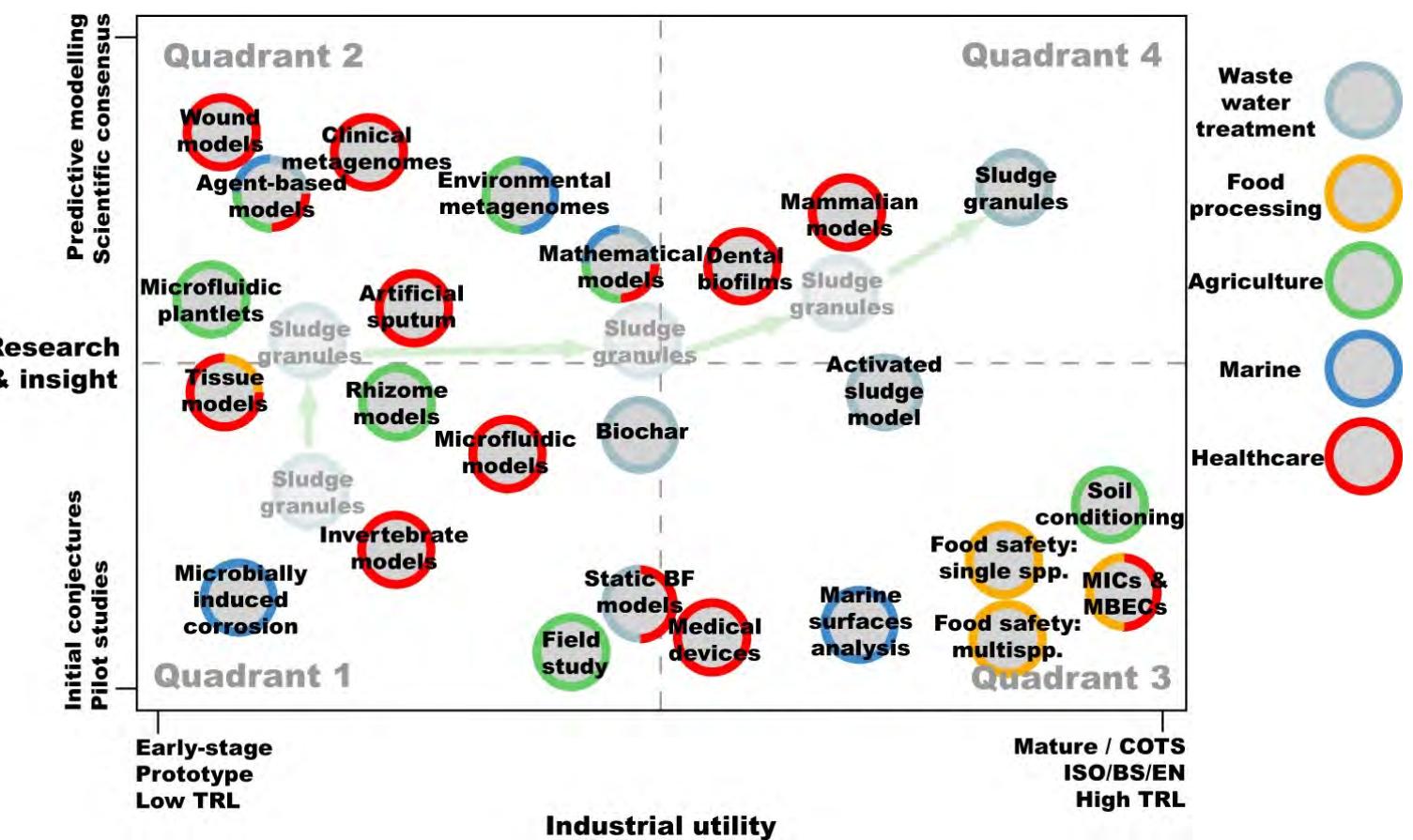


Future perspectives and challenges

>Translational challenges and opportunities in biofilm science



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Thank you !

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