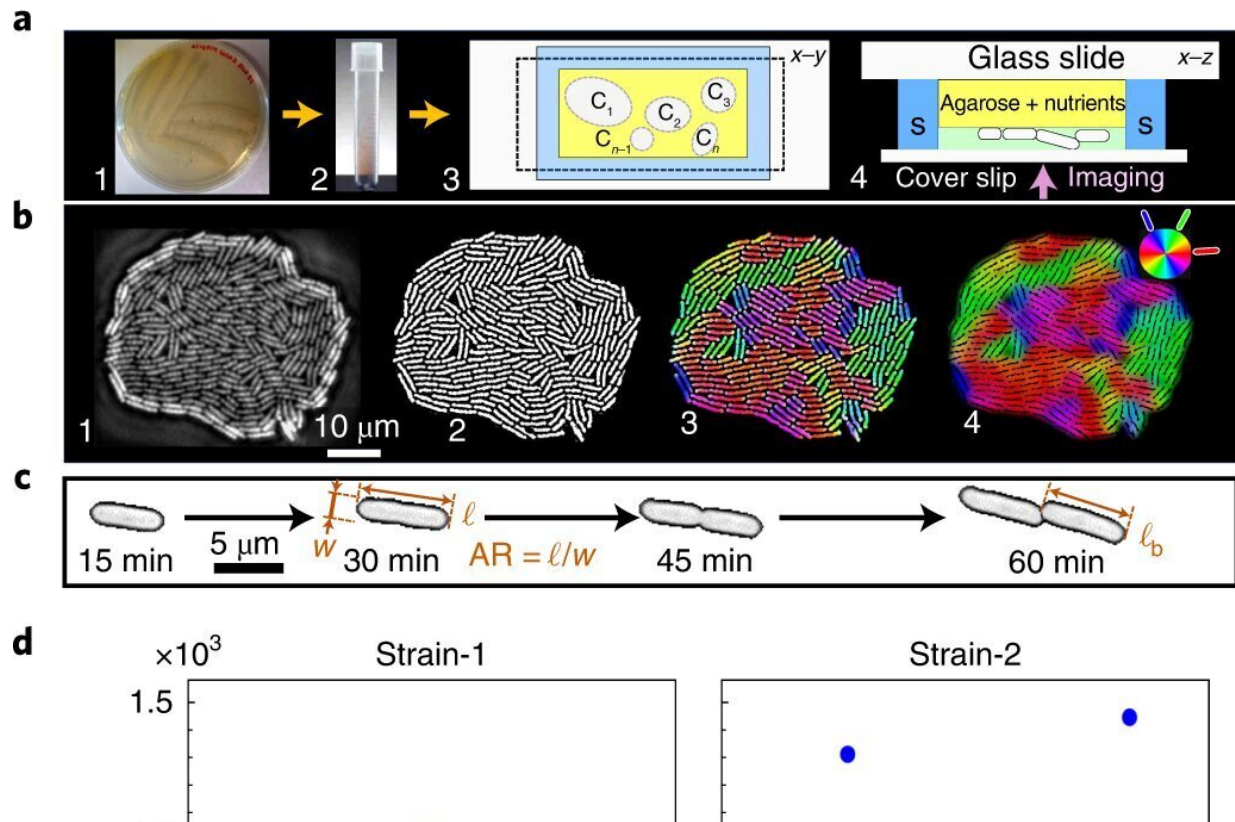


Physics meets biology: How bacteria synchronize to build complex structures

August 29 2022



From noisy phenotypes to a statistically precise mono-to-multilayer transition event. **a**, Cells streaked on nutrient-rich agar plate (1) are transferred in the liquid medium (2), then seeded within the microfluidic chamber, here C_i denotes individual colonies (3), for phase-contrast time-lapse imaging (4) in the swimming pool (S). **b**, Raw images of growing colonies (1) are binarized to extract phenotypic traits (2) and colour-coded to visualize local cell orientation (3,4). **c**, Single-cell geometric traits: AR and l_b . **d,e**, A_c is independent of growth temperature (**d**), as revealed both in our theory and experiments, whereas t_c is

temperature-dependent (e). Blue and red points indicate distinct biological replicates (including multiple technical replicates) for A_c and t_c experimental data. A_c has large variance across all T , whereas t_c has low variance, which minimizes further with the growth temperature. f, Trait-specific phenotypic noise, quantified as the normalized variance, $F = \text{var}(\dots) / \langle \dots \rangle^2$, across T : ℓ_b (blue squares), A_c (red triangles), cell length doubling time τ_{sc} (black diamonds), AR (magenta triangles) and t_c (green circles). Despite the high phenotypic noise at individual scales (ℓ_b , AR and τ_{sc}), t_c is statistically precise (corresponding noise is orders of magnitude lower across all T). The error bar denotes the standard deviation of F across colonies. Credit: *Nature Physics* (2022). DOI: 10.1038/s41567-022-01641-9

Bacteria collaborate and coordinate collectively as they form a shared structure called a biofilm, such as the dental plaque on our teeth or the microbiome associated with our gut. This self-organization in multiple complex layers—despite variations of cellular properties at individual level—requires that the living systems share common, yet precise time, which has now been uncovered by physicists from the University of Luxembourg.

Prof. Anupam Sengupta and his team explained this by zooming into the growth and development of nascent bacterial biofilms, wherein they discover emergent cross-talks between biophysical properties which enable precise timing of structural, topological and hydrodynamic transitions at the population scales. The groundbreaking findings appear in the recent edition of the journal *Nature Physics*.

Bacterial biofilms are found on nearly every natural and artificial surface we come across. They are crucial for human existence: from those in the gut regulating our physical well-being, to the ones regulating the progression of cancers and holding our ecosystem in balance. Thanks to the exquisite physico-chemical feedback mechanisms mediating their

prolific surface-associated lifestyles, biofilms are highly resilient and can colonize diverse systems.

Recent single-cell research by Prof. Sengupta and collaborators have revealed critical roles that single-cell geometry and growth dynamics play in shaping the dynamical properties of growing bacterial layers. Yet how variability in cell-level statistics—an entirely unaccounted player in existing living and active matter models—impact emergent collective phenomena in bacterial biofilms have remained a challenge, and thus largely unexplored.

Now, researchers from the Physics of Living Matter Group at the University of Luxembourg solve this long-standing conundrum: how biofilms—and living systems in general—regulate the timing of key developmental events emerging from individuals with highly variable and dynamic attributes. Taking a quantitative cross-disciplinary approach to spy within nascent bacterial biofilms, Prof. Sengupta's team provides the mechanistic underpinnings of how well-timed developmental steps emerge collectively due to the self-regulation of cell-level traits, which cross-talks to mitigate each other's noisy effects, ultimately enabling a [precise timing](#) of key developmental events.

The work dives deeper into rationalizing the results, providing a physiologically consistent picture based on the metabolic requirements during the early stages of [biofilm](#) development. The results demonstrate how self-regulation of phenotypic noise drive well-timed transitions in the structure, topology and active flows in bacterial colonies, highlighting the strategic role of physics in understanding [biological systems](#). The work fits broadly into the Physics Meets Biology initiative within the University of Luxembourg.

Noisy phenotypes drive well-timed developmental

events

Despite the variability in [phenotypic traits](#), such as cell geometry, growth rates, and surface association, bacterial biofilms undergo critical developmental steps in their lifecycle at precise time points, starting with the mono-to-multilayer transition (MTMT). Following this key structural transition, an emergent cascade is set off, synchronously timing the changes in topology and active flow fields within and in the vicinity of the nascent biofilms.

The catalog key phenotypic traits in *Escherichia coli* and *Serratia marcescens* growing under diverse conditions (nutrient levels and temperatures), allowing for careful control of the biological activity, and quantify how the cross-talks between phenotypic noises determine the timeliness of structural organization and emergence of active local transport. This work thus forges the first direct, mechanistic link between activity-dependent cell-level variability and population-scale emergent properties in living systems.

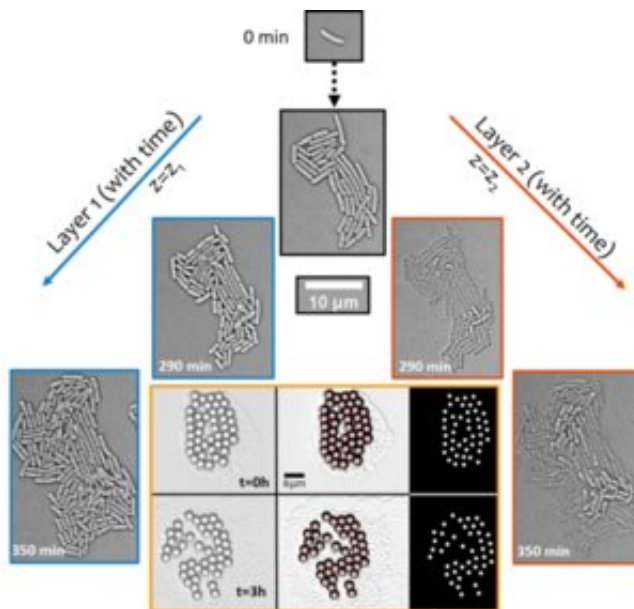


Figure 1. Structural transition of nascent biofilms from mono- to multi-layers drives emergent synchronous transport around the colony. The active transport phenomenon was uncovered by visualizing displacement of micron-sized particles over time (shown in the six-paneled image with orange border). Credit: University of Luxembourg

Leveraging a cross-scale and cross-disciplinary approach

Using a combination of single-cell time-lapse imaging, particle image velocimetry, numerical simulations and continuum modeling, the team quantifies cell-level variability in terms of phenotypic noise, and uncover trade-offs that ultimately suppress variability in emergent properties and render them synchronous. Specifically, a trade-off between the noise in the cell geometry and the growth rate self-regulate the timing of the mono-to-multilayer-transition (MTMT), a key step in biofilm initiation, setting off a synchronous active hydrodynamic cascade that ultimately drives enhanced local transport around the confluent bacterial colonies.

Remarkably, despite the species being non-motile in nature, the growth-dependent active hydrodynamics within confluent colonies trigger local flows in their surroundings which are strong enough to disrupt self-assembled particle clusters (used as tracers, see Figure 1) and transport them actively across the microbial environment. The enhancement in transport—in an otherwise diffusion-limited setting—by more than two orders of magnitude suggest biological functions of such active flows in the transport of molecular and micro-cargo during the early stages of biofilm development.

This work presents a breakthrough in the domain of physics of microbial active matter, and provides a new paradigm to understand how confluent bacterial populations may cope with environmental variations, including

those imposed by lifestyle and climate changes, by leveraging their individual-scale variability.

Perspectives on the synchronous active transport in nascent biofilms

The team demonstrates that sessile colonies can generate and harness active flows for shuttling sub-micron to micron sized biological cargo which are widely found associated with [bacterial colonies](#). Taken together the associated micro-cargo span dimensions ranging from tens of nanometers to a few microns. Typical micro-cargo include bacterial cells (different carrier and cargo species of micron size), fungal spores in micron scale; liposomes and extracellular vesicles delivering genetic or biochemical cargo (tens to hundreds of nanometers in size), hitchhiking bacteriophages (hundreds of nanometers), and synthetic beads and capsules relevant for drug delivery (sub-microns to tens of microns in size).

The biological significance of the time-synchronous nexus between structure-flow-transport is multi-pronged: Strict timing of the extrusion events presents MTMT as a biophysical proxy to quorum-like sensing between the colonies, with potential ramifications of and in synchronous selection for resistant cells (for instance, against antibiotics). Disrupting the well-defined MTMT timing (for instance, by appropriately tuning environmental factors) could offer future alternatives to inhibit quorum-sensing, and thereby regulate bacterial resistance to antibiotics.

Looking ahead, it will be critical to understand how sessile colonies harness the well-defined critical time to tune inter-colony communications under stressful environments (in the parameter space of intrinsic noise and structure-flow time-synchronicity). The noise-mediated spatio-temporal phenomena presented in this work provides

key missing insights into the developmental biophysics of morphogenesis in systems with higher complexity, including polymicrobial consortia observed in the human and plant microbiomes, and multi-cellular tissue systems relevant for embryonic and cancer progression.

This work opens new research paths in the fields of physics of microbial active matter, soft and biological physics, and cell biology, and—owing to the important role of bacterial biofilms in human microbiome and environmental ecology—also biomedical and biotechnological sciences. The structural and multifield topological outcomes presented here through Prof. Sengupta's vision will be consequential for diverse confluent systems spanning active tissue and cell systems, and a wide-range of organoid models.

Finally, this work provides a new experimental model system for active matter research where passive entities co-exist with active agents, and inspire a new class of theoretical models that incorporate cellular variability and trade-offs therein to understand the impact of noise, an intrinsic biophysical variable, on emergent properties in living systems.

More information: Jayabrata Dhar et al, Self-regulation of phenotypic noise synchronizes emergent organization and active transport in confluent microbial environments, *Nature Physics* (2022). [DOI: 10.1038/s41567-022-01641-9](https://doi.org/10.1038/s41567-022-01641-9)

Provided by University of Luxembourg

Citation: Physics meets biology: How bacteria synchronize to build complex structures (2022, August 29) retrieved 17 June 2024 from <https://phys.org/news/2022-08-physics-biology-bacteria-synchronize-complex.html>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.