



Microbiology

Célia Manaia is Full Professor at the Faculty of Biotechnology

The language of communities: microorganisms as an inspirational model

Communities can be conceptualized as concentric and overlapping rings that together form a complex, deeply interconnected assemblage.

In biology, a community is understood as a collection of diverse living organisms, each of which both gives to and receives from others. These interactions underpin the stability and functionality of the system as a whole. When considering the nature of microbial communities, it becomes almost inevitable to draw parallels with human societies and social communities. In both systems, collective stability and functionality emerge from shared organizing principles, including cooperation, communication, dynamic balance, the management of conflict, and the capacity for cohesion and resilience.

Studying microbial interactions can inspire approaches in human organizational design, from public health strategies to collaborative work environments.

It is through **cooperation** that microorganisms enhance their survival and functionality, establishing structured and persistent communities. Rather than acting as isolated individuals, microbes frequently engage in cooperative strategies such as metabolic cross-feeding, in which the by-products of one organism serve as resources for others. This division of labour allows communities to exploit complex substrates that no single species could process alone. Cooperation is also evident in the formation of biofilms, where cells produce shared extracellular matrices that provide protection against environmental stress, antimicrobial agents,

and host defences. While only some organisms invest in producing these structures, many benefit from them. Although cooperation entails costs at the individual level, it often yields collective benefits that increase community stability, efficiency, and resilience.

Just as microbial communities thrive through collective effort and division of labour, human societies depend on cooperation and shared responsibilities to achieve stability and progress.

Communication in microbial communities enables microorganisms to coordinate behaviour and respond collectively to external changes. This communication is primarily mediated through chemical signalling molecules that allow cells to sense the presence, density, and activity of neighbouring organisms. One well-studied example is quorum sensing, a mechanism through which microbes regulate gene expression in response to population size, synchronizing behaviours such as biofilm formation, virulence, and resource utilization. This is a very smart and efficient way of saving energy and resources. Communication also facilitates cooperation by aligning individual actions with community-level needs and by preventing premature or inefficient responses. Beyond intraspecies signalling, many microbial communities rely on interspecies communication, allowing diverse organisms to coexist and adapt within shared environments. Disruptions in communication can destabilize communities, leading to uncoordinated behaviour and reduced functionality.

The chemical signalling that coordinates microbial behaviour mirrors how effective communication underpins coordination and problem-solving in human communities.

Cooperation is essentially based on communication but strongly dependent on complementarity of needs and functions. This complementarity is essential to ensure an adequate **balance and equilibrium** of microbial communities. Diversity, dynamic regulation of activities, efficient resource use, and close interactions are key drivers that maintain a strong microbial community. Rather than remaining static, these communities maintain equilibrium through continuous feedback mechanisms that prevent any single species from dominating the system. Competition for nutrients, metabolic dependencies, and environmental constraints collectively shape population sizes and activity levels. Host-associated

microbial communities, such as the gut microbiota, further illustrate how balance is sustained through interactions with the host immune system and physiology. Disturbances in these communities, due to environmental stress, antibiotic exposure, or dietary changes, can disrupt equilibrium, leading to shifts in composition and function. Balanced microbial communities are often capable of reorganization, restoring key functions through adaptive interactions.

In both microbial ecosystems and human societies, maintaining balance through diversity, feedback, and mutual adaptation is essential for long-term stability.

No community is free of disruptive actors, and microbial communities are no exception. **Cheaters, exploiters, and pathogens** (microbes capable of causing disease) represent strategies that benefit individual organisms at the expense of collective function. The intention is not to cause harm to the community or the host; it is simply to prioritize one's own interests. Cheaters take advantage of shared resources, such as extracellular enzymes or protective matrices, without contributing to their production, reducing the overall efficiency of cooperation. Exploiters may manipulate communication systems or metabolic interactions to gain disproportionate access to resources. Pathogens, in turn, can disrupt community balance by prioritizing rapid growth or virulence over long-term stability. While these behaviours can undermine cooperation and equilibrium, they are also integral to community dynamics, stimulating regulatory mechanisms that mitigate potential adverse effects. As a reaction, microbial communities activate safeguards, such as spatial structuring, genetic memory, or host-mediated control, to limit exploitation. These evolutionary tensions refine microbial social systems and shape host–microbe interactions.

Just as cheaters, exploiters, and pathogens challenge microbial cohesion, human societies must manage individuals or groups whose self-interest can disrupt collective welfare.

The impairment of cheaters, exploiters, and pathogens, along with the successful functioning of cooperative members, is ultimately determined by the community's **cohesion and resilience**. These qualities emerge from the successful integration of cooperation, communication, balance, and controlled conflict. When these processes function effectively, individual

microorganisms are linked through networks of interaction that promote collective stability rather than fragmentation. Cohesive communities are better able to withstand environmental disturbances, such as nutrient fluctuations, physical stress, or antimicrobial challenges, because functional redundancy and coordinated responses allow the system to absorb shocks. Resilience is further reinforced by the capacity of the community to reorganize after disruption, restoring key functions even if species composition changes. Strength in microbial communities does not depend on uniformity, but on well-regulated diversity and interaction. When cooperative behaviours are maintained, communication remains effective, equilibrium is preserved, and exploitation is constrained, microbial communities emerge as robust, adaptable systems capable of long-term persistence.

The cohesion and resilience of microbial communities reflect how strong, well-connected human societies withstand crises and recover from disruption. Within these systems, diversity provides a buffer against environmental or social shocks, ensuring that both microbial and human communities can adapt and persist.

Both microbial communities and human societies develop forms of 'memory' that guide 'future' responses, whether through functional and genetic adaptation in microbes or institutional 'knowledge' in humans. This shared capacity for learning and adaptation underscores the importance of 'interconnectedness', 'cooperation', and 'resilience' across scales of life.
