



Phage transmission strategies: are phages farming their host?

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Extensive coevolution has led to utterly intricate interactions between phages and their bacterial hosts. While both the (short-term) intracellular molecular host-subversion mechanisms during a phage infection cycle and the (long-term) mutational arms race between phages and host cells have traditionally received a lot of attention, there has been an underestimating neglect of (mid-term) transmission strategies by which phages manage to cautiously spread throughout their host population. However, recent findings underscore that phages encode mechanisms to avoid host cell scarcity and promote coexistence with the host, giving the impression that some phages manage to ‘farm’ their host population to ensure access to host cells for lytic consumption. Given the tremendous impact of phages on bacterial ecology, charting and understanding the complexity of such transmission strategies is of key importance.

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Introduction

Bacterial viruses (also referred to as bacteriophages or phages) are ubiquitous in nature, and although they require a host cell to multiply, they often outnumber their hosts by an order of magnitude [1,2]. The life cycle and biology of phages has been extensively studied, and mainly became mapped along the lines of two distinct reproductive routes: lytic development (performed by

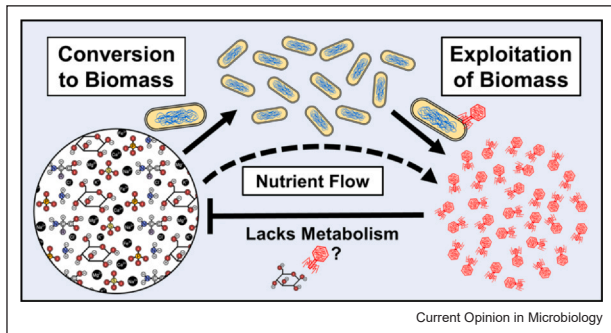
both lytic and temperate phages) and lysogenic development (only performed by temperate phages) [3]. In case of lytic development, the incoming phage chromosome enforces its replication and the production of new phage particles that are typically released by lysing the host and that enable further horizontal transmission throughout the host population. During this lytic development, phage proteins typically hijack host machinery and resources for massive replication of phage chromosomes and production of capsid proteins [4,5]. In case of temperate phages, the incoming phage chromosome can alternatively decide to lysogenically convert its host, thereby establishing itself as a seemingly more dormant prophage. The latter is typically integrated into the host chromosome, where it becomes stably replicated and segregated to ensure further vertical transmission [6]. However, when the host cell suffers stress such as DNA damage, the prophage can manage to exit this quiescent state and enter the lytic cycle [7,8].

Despite the above-perceived paradigms of phage transmission dynamics, the function and necessity of many phage-encoded proteins still remain obscure [9,10], suggesting higher-than-anticipated complexities in the way phages deal with their host population. In this context, we would like to focus on the possible ‘farming’ capacity of phages, being their presumed ability to intentionally cultivate and harvest host cells. Indeed, since phages can only indirectly access the nutritional resources from their environment via the metabolism and biomass of their host cells (Figure 1), it seems intuitive that phages have evolved mechanisms and strategies to avoid host cell scarcity and/or secure a steady supply of host cells for lytic consumption. In fact, recent literature is increasingly yielding examples that tend to incline toward this perspective and that showcase the resourcefulness of phages managing to stably coexist with their host.

Lytic phages manage to avoid host scarcity

The idea of strictly lytic phages inevitably consuming all of their host cells within the niche (or forcing them toward phage resistance) tends to conflict with actual observations in nature where lytic phages are found to stably coexist with their susceptible host cells [11,12]. Obviously, the heterogeneity of some natural environments or the

Figure 1



Phages need to exploit bacterial cells to gain access to the nutritional resources of their environment. While bacteria can directly access nutrients and convert them to biomass, phages need their host as an essential go-between to turn resources into phage particles. It therefore seems intuitive that phages have evolved strategies to avoid host cell scarcity and/or secure a steady supply of host cells for lytic consumption. These ("farming") strategies range from deferring from lytic development when host cells have poor growth prospects, up to supporting the emergence of transiently nonsusceptible subpopulations of host cells that can be lytically consumed ("harvested") afterward.

presence of biofilms can often provide spatially structured refuges that can serve as a continuous sink of fresh host cells [13,14]. However, also in homogeneous environments lacking such refuges, many lytic phages manage to coexist with their host without the need of being caught up in a mutational arms race, indicating lytic phages must have genetically wired strategies to avoid host depletion as well.

Indeed, some lytic phages (such as coliphage T4) seem to be able to defer their lytic development when infecting a starved host that likely signals a nutritionally poor and unproductive environment [15,16]. In this context, the chromosome of *Bacillus subtilis* phage $\phi 29$ even encodes binding sites for the host Spo0A protein (being the master regulator triggering the sporulation pathway) in order for its lytic cycle to become repressed and postponed when a hostile environment urges its host to commit to sporulation [17]. In the same vein, Φ Cbk-like phages of *Caulobacter* species seem to be able to monitor the cell-cycle state of its host, thereby distinguishing between its stalked versus swarmer phenotype. More specifically, swarmer cells of *Caulobacter* are characterized by low levels of activated CtrA developmental master regulator that coincide with active flagella and pili that are recognized by Φ Cbk-like phages to infect the cell. The phage chromosome, however, contains several CtrA-binding sites that are predicted to synchronize the cell-cycle state with lysis timing. High levels of active CtrA might trigger lysis and are only achieved when the swarmer cell switches to the stalker state in suitable environments. As such, the phage seems to defer lysis of the nondividing and more likely to be

a solitary swarmer cell up until this cell differentiates into a stalker cell under conditions that support a higher density of host cells [18].

Alternatively, some lytic phages seem to foster the emergence of a transiently resistant subpopulation that serves as a feedstock for lytic consumption later on. For example, *Bacteroides intestinalis* phage Φ crAss001 can only infect host cells with specific capsular polysaccharides (CPS). Since *B. intestinalis* displays stochastic phase variability in its CPS modifications, only the fraction of cells with susceptible CPS becomes lytically consumed, while cells with nonsusceptible CPS are left unharmed. These nonsusceptible cells can therefore grow out as a transiently resistant population that stochastically keeps spawning off susceptible cells that can be lytically consumed by Φ crAss001 [19]. Along similar lines, a more elaborate and circular use of phase variation was recently demonstrated with Fletchervirus F358, which uses its receptor-binding protein 1 (RBP1) to bind to specific CPS of its *Campylobacter jejuni* host. In this case, it was found that *C. jejuni* likewise displays stochastic phase variability in the modification of its CPS, leading to OFF/ON switching behavior of F358's receptor [20]. As such, infection by Fletchervirus F358 results in the lysis of receptor^{ON} cells (targeted by RBP1 of phage F358), which in turn allows receptor^{OFF} cells to take over the host population. However, phage F358 itself also displays phase variation in the expression of an alternative RBP2. If F358 has an active RBP2, it can circumvent the need for the CPS modification and infect and lyse receptor^{OFF} cells, leading receptor^{ON} revertants to take over the population [20]. It could perhaps be argued that many lytic phages have throughout evolution been selected to reach for phase-variable host receptors, as this would allow them to prevent host eradication and steer toward coexistence with their host.

Interestingly, the massive lysis of susceptible host cells on itself can also be a trigger toward securing a reserve population that can be lytically harvested later on. In an *Enterococcus faecalis* population, for example, massive lysis by phage Efs7 leads to the accumulation of phage-encoded endolysins in the environment, which in turn forces uninfected survivors into cell wall-deficient L-forms that lack phage receptors associated with the cell wall [21]. As such, these L-forms proliferate and remain phenotypically resistant to the phage until resynthesis of their cell wall can be reinitiated later on [21]. Similarly, upon phage SPP1 infection of its *Bacillus subtilis* host, a yet-to-be-determined signal is released and transmitted to neighboring noninfected cells that triggers expression of the host-encoded SigX sigma factor. SigX in turn activates the *dlt* operon, which encodes enzymes that modify the phage receptor (i.e. wall teichoic acids) and thereby raise a transiently resistant subpopulation [22]. As such, a reservoir of transiently uninfected cells is

formed that has the ability to revert back to an infectable state. Moreover, lysis by this same phage was also suggested to facilitate the spread of receptor-containing membrane vesicles that can fuse with nonsusceptible cells and hence make them susceptible for infection as well. As a result, massive lysis can even expand the host range toward previously unsusceptible cells [23].

Temperate phages manage to defy stable lysogeny

Contrary to lytic phages, temperate phages are assumed to have found the solution to host scarcity by integrating into the host chromosome as a prophage. But even though establishment of lysogeny is often seen as the path toward a stable vertical transmission and coexistence with the host, important phage (lytic) capacities escape selection and might become genetically eroded during lysogeny, as is evident from the countless examples of prophage remnants in bacterial genomes [24,25]. However, recent findings support the notion that lysogeny might be overrated as the (only) go-to strategy for temperate phages faced with host scarcity.

The classic coliphage Lambda model suggests that temperate phages tend to use a high multiplicity-of-infection (i.e. how many phage chromosomes simultaneously infect the same cell) as a proxy for imminent host depletion, and therefore as a cue to change the reproductive strategy from horizontal (lytic) toward vertical (lysogenic) transmission [26]. However, microscopic scrutiny of the dynamics of the temperate lambdoid P22 phage infecting its *Salmonella* Typhimurium host led to the observation that host scarcity can also lead to the establishment of a phage carrier state in which a polarly localized P22 episome is formed that segregates asymmetrically among daughter cells [27]. Although the cell inheriting the P22 episome eventually becomes lysogenized, its former P22-free daughter cells nevertheless cytoplasmically inherit superinfection exclusion factors (SEFs) produced by the P22 episome. This allows a transiently resistant P22-free subpopulation to emerge that — upon the gradual cytoplasmic dilution of these SEFs — again becomes susceptible to P22 infection. However, because of this gradual SEF dilution, the number of incoming P22 phages is heavily restricted, thereby relaying a high phage-to-host ratio (normally favoring establishment of lysogeny) into a low multiplicity-of-infection (favoring lytic consumption) [28]. Because of these carrier state and SEF dynamics, P22 seems to be able to partly defy lysogeny by continuously raising and lytically harvesting subpopulations of host cells.

In the same context, other temperate phages seem to have evolved intricate mechanisms to *wake up in time* and

reverse lysogeny when new susceptible host cells are detected and before potential crippling mutations accumulate. In this context, it was already previously discovered that SPbeta phages infecting *Bacillus subtilis* make use of a particular quorum sensing mechanism (rather than the multiplicity-of-infection) to switch from horizontal to vertical transmission. Indeed, during lytic consumption of a large part of the host population, a phage-encoded small peptide (AimP, also referred to as arbitrium peptide) progressively accumulates in the environment and eventually signals the remaining infections to incline toward establishment of lysogeny [29]. However, most recently, this system was also shown to affect the decision-making of the subsequently established SPbeta prophages. More specifically, at high lysogen concentrations, the accumulation of arbitrium peptide produced by the prophage will block the host DNA damage response and therefore restrict prophage induction. However, in environments where the lysogen again becomes surrounded by susceptible (nonlysogen) host cells, the correspondingly diluted arbitrium concentration will eventually enable the DNA damage response to activate prophages and release phage particles that can commit to lytic consumption of susceptible host cells [30–32]. As such, prophage dormancy can be alleviated as soon as new susceptible host cells arise that can again fuel horizontal transmission.

Alternatively, some prophages rather eavesdrop on host-derived quorum sensing signals for their wake up call [33,34]. As such, phage ARM81ld was found to sense the C4-HSL quorum signal of its *Aeromonas* host by using a phage-encoded LuxR homolog to decide upon prophage induction [33–35]. Furthermore, ARM81ld also seems to estimate the nutritional competition encountered by its host, since ARM81ld is also able to detect the C8-HSL quorum sensing signal of *Vibrio fischeri* cells (that it cannot infect). In fact, in cases where *Aeromonas* is being outcompeted by *V. fischeri*, ARM81ld will sense the high C8-HSL concentration and — given the poor prospects of its host — refrain from induction [35].

Finally, it is worth underscoring that filamentous phages have technically found a very fruitful merge between horizontal and vertical transmission. Indeed, their ability to continuously replicate within and egress from their host cell without actually lysing it by default circumvents host cell scarcity and secures continuous coexistence [36].

Conclusion

Many examples support the notion that phages are not only able to subvert individual host cells, but actually succeed in managing their host at the population level. Nevertheless, it still remains to be further established

whether all of the provided examples represent intentional strategies to secure coexistence. Indeed, while some mechanisms clearly appear to be supported by phage-borne functions, other dynamics might rather stem from intrinsic host cell behavior for which it is unclear whether it is truly exploited by the phage. However, with host cells as the essential intermediary and catalyst toward turning an environment's resources into phage particles, strategies to prevent host scarcity and raise or exploit transiently resistant subpopulations (i.e. 'farming' capacity) could well have been positively selected throughout evolution.

Typically, the mechanisms underlying these possible farming strategies depend on often subtle and transient interactions that need to be interpreted over different scales. In fact, biochemical interactions need to be put in a cellular context and variations (both cell-to-cell and over time) in individual cellular behavior need to be integrated at the level of populations in order to understand the relevant emergent properties and how they impact infection dynamics at large. For this, the live visualization, monitoring, and comparison of different phage–host interactions inside and between large numbers of cells within an infected population will become increasingly important, and will require further advancements in fluorescent reporter engineering (in both phages and bacteria) and fluorescence microscopy approaches. Finally, these efforts should further scale up to also take into account the full complexity imposed by ecologically relevant settings, such as differently structured microenvironments and the presence of various phages competing for the same host.

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Data Availability

No data were used for the research described in the article.

Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

References and recommended reading

Papers of particular interest, published within the period of review, have been highlighted as:

- of special interest
- 1. Comeau AM, Hatfull GF, Krisch HM, Lindell D, Mann NH, Prangishvili D: **Exploring the prokaryotic virosphere**. *Res Microbiol* 2008, **159**:306-313.
- 2. Stern A, Sorek R: **The phage-host arms race: shaping the evolution of microbes**. *BioEssays* 2011, **33**:43-51.
- 3. Hobbs Z, Abedon ST: **Diversity of phage infection types and associated terminology: the problem with 'Lytic or lysogenic'**. *FEMS Microbiol Lett* 2016, **363**:fnw047.
- 4. Dove WF: **Action of the lambda chromosome: I. Control of functions late in bacteriophage development**. *J Mol Biol* 1966, **19**:187-201.
- 5. Hood IV, Berger JM: **Viral hijacking of a replicative helicase loader and its implications for helicase loading control and phage replication**. *Elife* 2016, **5**:213.
- 6. Howard-Varona C, Hargreaves KR, Abedon ST, Sullivan MB: **Lysogeny in nature: mechanisms, impact and ecology of temperate phages**. *ISME J* 2017, **11**:1511-1520.
- 7. Silpe JE, Duddy OP, Bassler BL: **Induction mechanisms and strategies underlying interphage competition during polylysogeny**. *PLoS Pathog* 2023, **19**:e1011363.
- 8. Raya RR, H'bert EM: **Isolation of phage via induction of lysogens**. *Methods Mol Biol* 2009, **501**:23-32.
- 9. Yin Y, Fischer D: **Identification and investigation of ORFans in the viral world**. *BMC Genom* 2008, **9**:24.
- 10. Fremin BJ, Bhatt AS, Kyrpidis NC, Sengupta A, Sczyrba A, Maria da Silva A, Buchan A, Gaudin A, Brune A, Hirsch AM, et al.: **Thousands of small, novel genes predicted in global phage genomes**. *Cell Rep* 2022, **39**:110984.
- 11. Lourenço M, Chaffringeon L, Lamy-Besnier Q, Titécot M, Pédrón T, Sismeiro O, Legendre R, Varet H, Coppée JY, Bérard M, et al.: **The gut environment regulates bacterial gene expression which modulates susceptibility to bacteriophage infection**. *Cell Host Microbe* 2022, **30**:556-569.e5.
- 12. Aylward FO, Boeuf D, Mende DR, Wood-Charlson EM, Vislova A, Eppley JM, Romano AE, DeLong EF: **Diel cycling and long-term persistence of viruses in the ocean's euphotic zone**. *Proc Natl Acad Sci USA* 2017, **114**:11446-11451.
- 13. Winans JB, Wucher BR, Nadell CD: **Multispecies biofilm architecture determines bacterial exposure to phages**. *PLoS Biol* 2022, **20**:e3001913.
- 14. Attrill EL, Claydon R, Łapińska U, Recker M, Meaden S, Brown AT, Westra ER, Harding SV, Pagliara S: **Individual bacteria in structured environments rely on phenotypic resistance to phage**. *PLoS Biol* 2021, **19**:e3001406.
- This research combines advanced microfluidics with time-lapse microscopy to scrutinize phage-host infection dynamics within simulated structured environments. As such, the behaviour of a high number of individual cells can be continuously monitored over time.
- 15. Bryan D, El-Shibiny A, Hobbs Z, Porter J, Kutter EM: **Bacteriophage T4 infection of stationary phase E. coli: life after log from a phage perspective**. *Front Microbiol* 2016, **7**:1391.
- 16. Attrill EL, Łapińska U, Westra ER, Harding SV, Pagliara S: **Slow growing bacteria survive bacteriophage in isolation**. *ISME Commun* 2023, **3**:95.
- 17. Meijer WJJ, Castilla-Llorente V, Villar L, Murray H, Errington J, Salas M: **Molecular basis for the exploitation of spore formation as survival mechanism by virulent phage ϕ 29**. *EMBO J* 2005, **24**:3647-3657.
- 18. Mascolo E, Adhikari S, Caruso SM, deCarvalho T, Folch Salvador A, Serra-Sagrà J, Young R, Erill I, Curtis PD: **The transcriptional regulator CtrA controls gene expression in**

Alphaproteobacteria phages: evidence for a lytic deferral pathway. *Front Microbiol* 2022, **13**:918015.

19. Shkoporov AN, Khokhlova EV, Stephens N, Hueston C, Seymour S, Hryckowian AJ, Scholz D, Ross RP, Hill C: **Long-term persistence of crAss-like phage crAss001 is associated with phase variation in Bacteroides intestinalis.** *BMC Biol* 2021, **19**:163.

This research shows how the phase variability of a phage receptor can support long-term coexistence of a lytic phage with its host. Phenotypically resistant subpopulations emerge that continuously spawn off susceptible cells that fuel the infection.

20. Sørensen MCH, Vitt A, Neve H, Soverini M, Ahern SJ, Klumpp J, Brøndsted L: **Campylobacter phages use hypermutable polyG tracts to create phenotypic diversity and evade bacterial resistance.** *Cell Rep* 2021, **35**:109214.

This research shows how also phages can depend on phase variability of their RBPs to target different subpopulations of host cells.

21. Wohlfarth JC, Feldmüller M, Schneller A, Kilcher S, Burkolter M, Meile S, Pilhofer M, Schuppler M, Loessner MJ: **L-form conversion in Gram-positive bacteria enables escape from phage infection.** *Nat Microbiol* 2023, **8**:387-399.

This research shows how massive host cell lysis by a lytic phage can drive the remaining surviving host cells towards transient phenotypic resistance. In this case, the massive spread of endolysins converts the survivors into transiently unsusceptible L-forms.

22. Tzipilevich E, Pollak-Fiyaksel O, Shraiteh B, Ben-Yehuda S: **Bacteria elicit a phage tolerance response subsequent to infection of their neighbors.** *EMBO J* 2022, **41**:e109247.

This research shows how massive host cell lysis by a lytic phage can drive the remaining surviving host cells towards transient phenotypic resistance. In this case, the spread of a yet unknown signal triggers a genetic response making the survivors transiently unsusceptible for phage infection.

23. Tzipilevich E, Habusha M, Ben-Yehuda S: **Acquisition of phage sensitivity by bacteria through exchange of phage receptors.** *Cell* 2017, **168**:186-199.e12.

24. Fortier LC, Sekulovic O: **Importance of prophages to evolution and virulence of bacterial pathogens.** *Virulence* 2013, **4**:354-365.

25. Bobay LM, Touchon M, Rocha EPC: **Pervasive domestication of defective prophages by bacteria.** *Proc Natl Acad Sci USA* 2014, **111**:12127-12132.

26. Oppenheim AB, Kobiler O, Stavans J, Court DL, Adhya S: **Switches in bacteriophage lambda development.** *Annu Rev Genet* 2005, **39**:409-429.

27. Cenens W, Makumi A, Govers SK, Lavigne R, Aertsen A: **Viral transmission dynamics at single-cell resolution reveal transiently immune subpopulations caused by a carrier state association.** *PLoS Genet* 2015, **11**:e1005770.

28. Staes I, Bäcker LE, Simoens K, De Winter K, Marolt G, Cenens W, Wolput S, Vazquez AR, Goos P, Lavigne R, et al.: **Superinfection exclusion factors drive a history-dependent switch from vertical to horizontal phage transmission.** *Cell Rep* 2022, **39**:110804.

This research shows how phage-borne carrier state dynamics and cytoplasmic inheritance of SEFs allow temperate *Salmonella enterica* phage P22 to relay a high phage-to-host ratio towards a low multiplicity-of-infection, thereby deferring establishment of lysogeny and enabling the lytic consumption of subpopulations of host cells.

29. Erez Z, Steinberger-Levy I, Shamir M, Doron S, Stokar-Avihail A, Peleg Y, Melamed S, Leavitt A, Savidor A, Albeck S, et al.: **Communication between viruses guides lysis-lysogeny decisions.** *Nature* 2017, **541**:488-493.

30. Aframian N, Omer Bendori S, Kabel S, Guler P, Stokar-Avihail A, Manor E, Msaeed K, Lipsman V, Grinberg I, Mahagna A, et al.: **Dormant phages communicate via arbitrium to control exit from lysogeny.** *Nat Microbiol* 2022, **7**:145-153.

This research shows how a prophage can use its own quorum sensing signal to time activation of its lytic cycle when susceptible host cells become available. When mainly surrounded by similarly lysogenized host cells, the prophage refrains from activating its lytic cycle.

31. Brady A, Quiles-Puchalt N, Gallego del Sol F, Zamora-Caballero S, Felipe-Ruiz A, Val-Calvo J, Meijer WJJ, Marina A, Penadés JR: **The arbitrium system controls prophage induction.** *Curr Biol* 2021, **31**:5037-5045.e3.

32. Bruce JB, Lion S, Buckling A, Westra ER, Gandon S: **Regulation of prophage induction and lysogenization by phage communication systems.** *Curr Biol* 2021, **31**:5046-5051.e7.

33. Silpe JE, Duddy OP, Johnson GE, Beggs GA, Hussain FA, Forsberg KJ, Bassler BL: **Small protein modules dictate prophage fates during polylysogeny.** *Nature* 2023, **620**:625-633.

34. Silpe JE, Bassler BL: **Phage-encoded LuxR-type receptors responsive to host-produced bacterial quorum-sensing autoinducers.** *mBio* 2019, **10**:e00638-19.

35. Silpe JE, Duddy OP, Bassler BL: **Natural and synthetic inhibitors of a phage-encoded quorum-sensing receptor affect phage-host dynamics in mixed bacterial communities.** *Proc Natl Acad Sci USA* 2022, **119**:e2217813119.

This research shows how a prophage can read out the quorum sensing signal of non-host species that likely compete with its host. When the concentration of competing species is too high, the prophage refrains from activating its lytic cycle.

36. Hay ID, Lithgow T: **Filamentous phages: masters of a microbial sharing economy.** *EMBO Rep* 2019, **20**:e47427.