1	Prophage-dependent neighbor predation fosters horizontal gene transfer by natural
2	transformation
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13	Running Head: Prophage predation promotes lateral gene transfer
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## 22 ABSTRACT

23	Natural transformation is a broadly conserved mechanism of horizontal gene transfer
24	(HGT) in bacteria (1) that can shape their evolution through the acquisition of genes that
25	promote virulence, antibiotic resistance, and other traits (2). Recent work has established
26	that neighbor predation via Type VI secretion systems (3), bacteriocins (4) and virulent
27	phages (5), play an important role in promoting HGT. Here, we demonstrate that in chitin
28	estuary microcosms, Vibrio cholerae K139 lysogens exhibit prophage-dependent
29	neighbor predation of non-lysogens to enhance HGT. Through predation of non-lysogens,
30	K139 lysogens also have a fitness advantage in these microcosm conditions. The
31	ecological strategy revealed by our work provides a better understanding of the
32	evolutionary mechanisms used by bacteria to adapt in their natural setting and contributes
33	to our understanding of the selective pressures that may drive prophage maintenance in
34	bacterial genomes.

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#### 36 IMPORTANCE

37 Prophages are nearly ubiquitous in bacterial species. These integrated phage elements 38 have previously been implicated in horizontal gene transfer (HGT) largely through their 39 ability to carry out transduction (generalized or specialized). Here, we show that 40 prophage-encoded viral particles promote neighbor predation leading to enhanced HGT 41 by natural transformation in the water-borne pathogen Vibrio cholerae. Our findings 42 contribute to a comprehensive understanding of the dynamic forces involved in prophage 43 maintenance which ultimately drive the evolution of naturally competent bacteria in their 44 natural environment.

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### 46 TEXT

47	Several bacterial species have evolved to capture DNA (natural competence) as a
48	source of nutrients (6) or to incorporate into their genome to speed their evolution via a
49	process termed natural transformation (NT). V. cholerae is a genetically tractable and
50	well-established model organism to study NT. This human pathogen is usually found in
51	association with the chitinous carapaces of zooplankton in estuary and ocean waters (7).
52	V. cholerae can utilize chitin as a major carbon and nitrogen source and additionally, this
53	polymer is a required signal for induction of NT (8). In estuarine chitin microcosms, it
54	has been shown that this pathogen can take up multiple large DNA fragments when the
55	exogenous DNA concentration is high (3, 9). Much evidence points to NT and HGT
56	having contributed to the evolution of V. cholerae (10).
57	About 50% of <i>V. cholerae</i> clinical isolates carry the temperate kappa phage K139
58	(11). However, the role of K139 in the ecology of <i>V. cholerae</i> has been ill-defined. Here,
59	we explore whether the lytic replication of K139 affects the physiology of V. cholerae in
60	chitin microcosms, which mimics the aquatic reservoir for this facultative pathogen.
61	Using the K139 lysogen E7946, an O1 El Tor V. cholerae strain, we first
62	evaluated bacterial replication and production of viral particles over time in chitin
63	microcosms. V. cholerae growth was slow and bacterial numbers increased 100-fold by
64	24h and 1000-fold by 48h of incubation, while K139 plaque forming units (PFUs)
65	increased 3-logs by 6 hours, which is when bacteria were entering exponential growth
66	(Figure 1A). Interestingly, we found that insoluble chitin specifically increased K139
67	PFUs when compared to other carbon sources (Fig. S1). Based on these results, we

68	hypothesized that K139 might have a role in the ecology of V. cholerae in its
69	environmental reservoir by increasing the competitive fitness of lysogenic strains. To
70	evaluate if K139 is able to kill non-lysogenic strains in co-cultures, E7946 (lysogen) and
71	an isogenic E7946 $\Delta$ K139 mutant (non-lysogen) were competed in a chitin estuary
72	microcosm. After 24 h of incubation, the E7946 $\Delta$ K139 non-lysogen was outcompeted 8-
73	fold by the E7946 lysogenic strain (Figure 1B). To further test if K139 could help
74	lysogens outcompete non-lysogens in this environment, we competed E7946 with a panel
75	of diverse clinical isolates that naturally lack K139. E7946 was able to outcompete all of
76	these clinical isolates (Figure 1B). This effect was dependent on K139 viral production,
77	because a E7946 $\Delta$ K139 strain was unable to outcompete non-lysogens ( <b>Figure 1B</b> ).
78	These results strongly suggest that K139 plays an ecological role by providing a
79	competitive advantage for lysogenic strains in mixed populations containing non-
80	lysogens. By excising in a small fraction of lysogenic cells, and efficiently killing the
81	neighboring non-lysogenic cells via lytic replication, K139 may allow the remaining
82	lysogenic population to successfully compete for resources in a nutrient-limited estuary
83	environment.
84	It has previously been shown that some phages can lyse bacterial cells and release
85	intact DNA (5), while other phages degrade host DNA following lysis (12). Release of

86 intact DNA could aid in promoting horizontal gene transfer by NT (5, 13). Here, we have

87 found that K139 promotes neighbor predation in the same chitin microcosm conditions

that are required to induce NT. Therefore, we next wanted to test if K139-dependent

89 neighbor predation could promote HGT in a chitin microcosm. To that end, we co-

90 cultured a mixture of strains where each carried an unlinked selection marker at a neutral

site in the genome (9). The generation of strains with both selection markers indicated HGT. After 48 h of growth in chitin estuary microcosms, the co-culture containing E7946 (lysogen, wt) and E7946  $\Delta$ K139 (non-lysogen), where neighbor predation is expected to occur, showed the highest number of transformants, which was ~100-fold higher than a co-culture of two K139 lysogens or two non-lysogens where neighbor predation is not expected to occur (**Figure 1C**). These results are consistent with neighbor predation promoting HGT.

98 Prophages have mainly been linked to HGT by transduction (13). However, in 99 chitin microcosms, V. cholerae can also undergo HGT via NT. Thus, next we designed a 100 strategy to determine if the HGT observed is attributed to phage transduction or NT. To 101 distinguish between these, we inactivated dprA in our V. cholerae strains, a gene that is 102 essential for NT but is not required for transduction (14). After 48 h of growth in chitin 103 estuary microcosms, the co-culture containing a NT+ lysogen (E7946) and a NT-104 nonlysogen (E7946  $\Delta$ K139  $\Delta$ *dprA*) showed elevated rates of HGT similar to the co-105 culture containing both NT+ strains (Figure 1C). In contrast, a mixture containing an 106 NT- lysogen and an NT+ nonlysogen showed the basal levels of HGT seen in co-cultures 107 where no phage predation occurs (Figure 1C). This suggests that the HGT observed is 108 due to the transfer of DNA from the non-lysogen to the lysogen. This is the opposite of 109 what would be expected for phage transduction where a prophage excised from a 110 lysogenic strain would transduce DNA to the non-lysogen. Together, our results point to 111 an adaptative strategy used by lysogenic strains to induce prophage-dependent neighbor 112 predation in order to thrive and also to capture released DNA for HGT in its aquatic 113 environment.

- 114 To the best of our knowledge, our work is the first to establish an ecological role
- 115 for K139 in enhancing *V. cholerae* fitness. We show that K139 can enhance survival of
- 116 *V. cholerae* lysogens in chitin estuarine environments by neighbor predation of non-
- 117 lysogens and by driving evolution via HGT. More broadly, our results suggest a novel
- 118 mechanism by which prophages benefit their lysogenized hosts, which may contribute to
- 119 the maintenance of these genetic elements in bacterial genomes.

### 121 METHODS

### 122 Bacterial strains and growth conditions

123 See Table S1 for a list of all *V. cholerae* strains used in this study. All strains were

- 124 routinely grown in Luria-Bertani Miller (LB) Broth and agar at 30°C. Where necessary,
- media was supplemented with erythromycin ( $10 \mu g/mL$ ) or trimethoprim ( $10 \mu g/mL$ ).

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- 127 For chitin utilization experiments, cells were grown overnight in LB at 30°C shaking.
- 128 The next morning, cultures were washed and diluted in 0.7% Instant Ocean (Aquarium
- 129 Systems) or M9 minimal media. 10<sup>5</sup> CFU were inoculated in a 1% shrimp shell chitin
- 130 (Sigma) suspension in 0.7% Instant Ocean (chitin microcosm) and were incubated up to
- 131 48h at 30°C statically in 14 ml glass test tubes (Fisher Scientific). Lactate or glucose
- 132 were added to a final concentration of 0.2% and 0.5% when required. GlcNAc sugars
- 133 were added at 2mM for pentasaccharides, 3.33 mM for trisaccharides, 5 mM for
- 134 disaccharides, and 10 mM for monosaccharides. CFU counts were evaluated by serially
- 135 diluting and plating on LB agar plates.

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### 137 Bacteriophage assays

- 138 To assay phage titers, supernatants from bacterial cultures were filtered using 0.2 μm
- 139 filters (Costar). Filtered supernatants were serially diluted and tittered using E7946
- 140  $\Delta$ K139 (nonlysogenic, susceptible strain) as described in (15). Plates were incubated
- 141 overnight at 37°C and turbid plaques were counted the next morning.

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143 Competition and HGT assays

144	Competition assays were conducted in chitin microcosms for 24 h at 30°C statically.
145	Strains were distinguished by lacZ phenotype (using $lacZ$ + and $\Delta lacZ$ strain pairs) as
146	previously described (16). Cultures were mixed in a 1:1 ratio and plated for quantitative
147	culture on LB+Xgal. Competitive indices were calculated as previously described (16).
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149	HGT assays between strain pairs were conducted with differentially marked strains where
150	each contained an Ab <sup>R</sup> marker at a distinct neutral locus ( $\Delta VC1807$ ::Erm <sup>R</sup> or
151	$\Delta$ VCA0692::Tm <sup>R</sup> ). As indicated, strains were mixed in a 1:1 ratio in chitin microcosms
152	for 48 h at 30°C statically (9). After 48 h, a portion of each co-culture was diluted in LB
153	broth and outgrown for 2 hr prior to plating. Reactions were plated for quantitative
154	culture on Tm+Erm plates to quantify transformants, as well as on Erm and Tm alone to
155	quantify the abundance of each strain within the coculture. Transformation frequency is
156	expressed as CFU of transformants ( $\text{Erm}^{\text{R}} + \text{Tm}^{\text{R}}$ double resistant) / (CFU of $\text{Erm}^{\text{R}} +$
157	CFU of Tm <sup>R</sup> ).

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- 166 AUTHOR CONTRIBUTIONS
- 167 CAS-V, TND and RCM-Q performed experiments. CAS-V, RCM-Q, ABD and A.C.
- 168 designed experiments and provided materials and strains. CAS-V, ABD and RCM-Q
- 169 wrote the manuscript. All authors discussed the results and commented on the
- 170 manuscript.
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# 229 FIGURE LEGENDS

### 230 Figure 1. K139 promotes neighbor predation and enhances HGT by NT in chitin

231 microcosms. A, V. cholerae E7946 growth (CFU, full circles) and K139 phage titer in

culture supernatants (PFU, empty circles) were measured in chitin microcosms for 48 hrs.

- 233 B, Competition between V. cholerae strains mixed 1:1 was assessed following 24 hrs of
- growth in chitin microcosms. The strain indicated at the top of each graph was competed
- with each strain indicated on the X-axis. The competitive index is reported as the ratio of
- 236 (top strain) / (X-axis strain) in the output divided by the same ratio in the input. Data are
- from 4 independent experiments and the line within samples denotes the median. The
- dotted line indicates a CI of 1, which is the value expected if strains compete equally.
- 239 Statistical comparisons were made by Mann-Whitney test (\*p<0.05, \*\*\*\*p<0.0001). C,
- 240 The indicated variants of *V. cholerae* E7946 were co-cultured in chitin microcosms at 30°
- 241 C for 48 h to assess HGT, which is reported as the transformation frequency (see
- 242 Methods for details). Data represents three independent biological replicates shown as the
- 243 mean  $\pm$  SD. Statistical comparisons were made by one-way ANOVA with Tukey's post-

test on the log-transformed data (\*\*p<0.01, \*\*\*p<0.001).

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#### Figure S1. Growth on insoluble chitin specifically induces K139 lytic replication in

247 an estuarine environment. (A-C) *V. cholerae* E7946 was grown in chitin microcosms

with glucose (0.2%) and lactate (0.5%) added to growth reactions as indicated. Addition

- of these readily available carbon sources should diminish the reliance on chitin as a
- 250 carbon source in these assays. After 24 h of static growth A, K139 PFU in culture
- supernatant and **B**, *V. cholerae* CFU were determined. **C**, K139 PFU generated per viable

cell plotted as PFU/CFU using the data shown in A and B. (D-F) V. cholerae E7946

- grown in M9 minimal medium with the sole carbon sources indicated: chitin (insoluble),
- 254 glucose, lactate, the chitin monosaccharide N-acetyl glucosamine (GlcNAc), or soluble β
- 255 1,4-linked GlcNAc oligosaccharides. After 24 h of static growth **D**, K139 PFU in culture
- supernatant and E, V. cholerae CFU were determined. F, K139 PFU generated per viable
- 257 cell plotted as PFU/CFU using the data shown in **D** and **E**. Through these experiments,
- we found that the ratio of K139 viral particles per cell was highest during growth on
- 259 insoluble chitin, thus, *V. cholerae* may specifically induce K139 production while
- 260 utilizing insoluble chitin in microcosms. All data are from three independent experiments
- and plotted as the mean  $\pm$  SD. Statistical comparisons were made by Student's two-tailed
- t-test by comparing each condition to the control on chitin (p<0.05, p<0.01,

263 \*\*\*p<0.001, \*\*\*\*p<0.0001).

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