

Inhibition of phagocytic killing of Escherichia coli in Drosophila Hemocytes by RNA Chaperone Hfq

メタデータ	言語: eng 出版者: 公開日: 2017-10-04 キーワード (Ja): キーワード (En): 作成者: メールアドレス: 所属:
URL	http://hdl.handle.net/2297/46175

1 **Inhibition of Phagocytic Killing of *Escherichia coli* in *Drosophila* Hemocytes by**
2 **RNA Chaperone Hfq¹**

3
4 Running title: ROLE OF RNA CHAPERONE IN BACTERIAL EVASION OF HOST
5 IMMUNITY

6
7 Akiko Shiratsuchi,^{*,†,2} Mao Nitta,[†] Ayumi Kuroda,[†] Chiharu Komiyama,^{*} Mitsuko Gawasawa,[†]
8 Naoto Shimamoto,^{*} Tran Quoc Tuan,^{*} Teppei Morita,[‡] Hiroji Aiba,[‡] and Yoshinobu Nakanishi^{*,†,2}

9
10 ^{*}Graduate School of Medical Sciences and [†]School of Pharmacy, Kanazawa University, Kanazawa,
11 Ishikawa 920-1192, Japan; [‡]Faculty of Pharmaceutical Sciences, Suzuka University of Medical
12 Sciences, Suzuka, Mie 513-0816, Japan

13
14 Keywords: Other Animals; Bacterial Infection; Transcription Factors; Phagocytosis

15
16 Corresponding authors:

17 Dr. Akiko Shiratsuchi, Graduate School of Medical Sciences, Kanazawa University, Shizenken,
18 Kakuma-machi, Kanazawa, Ishikawa 920-1192, Japan

19 Phone number: +81 76 234 4424; Fax number: +81 76 264 6223; E-mail address:
20 ashira@staff.kanazawa-u.ac.jp; ORCID: 0000-0002-7811-4409

21 Dr. Yoshinobu Nakanishi, Graduate School of Medical Sciences, Kanazawa University, Shizenken,
22 Kakuma-machi, Kanazawa, Ishikawa 920-1192, Japan

23 Phone number: +81 76 234 4481; Fax number: +81 76 234 4480; E-mail address:
24 nakanaka@p.kanazawa-u.ac.jp; ORCID: 0000-0002-8767-3587

25

1 **Abstract**

2

3 An RNA chaperone of *Escherichia coli*, called host factor required for phage Q β RNA replication
4 (Hfq), forms a complex with small non-coding RNAs to facilitate their binding to target mRNA for
5 the alteration of translation efficiency and stability. Although the role of Hfq in the virulence and
6 drug resistance of bacteria has been suggested, how this RNA chaperone controls the infectious
7 state remains unknown. In the present study, we addressed this issue using *Drosophila*
8 *melanogaster* as a host for bacterial infection. When adult flies abdominally infected with *E. coli*
9 were analyzed, an *E. coli* strain with mutation in *hfq* was eliminated earlier, while flies survived
10 longer, compared to infection with a parental strain. The same was true with flies deficient in
11 humoral responses, but the mutant phenotypes were not observed when a fly line with impaired
12 hemocyte phagocytosis was infected. The results from an assay for phagocytosis *in vitro* revealed
13 that Hfq inhibits the killing of *E. coli* by *Drosophila* phagocytes after engulfment. Furthermore,
14 Hfq seemed to exert this action partly through enhancing the expression of σ^{38} , a stress-responsive
15 sigma factor that was previously shown to be involved in the inhibition of phagocytic killing of *E.*
16 *coli*, by a post-transcriptional mechanism. Our study indicates that the RNA chaperone Hfq
17 contributes to the persistent infection of *E. coli* by maintaining the expression of bacterial genes,
18 including one coding for σ^{38} , that help bacteria evade host immunity.

19

1 **Introduction**

2

3 Under hostile environmental conditions, bacteria alter their gene expression pattern for adaptation
4 and survival. One such mechanism involves the action of small non-coding RNAs that bind
5 mRNAs and change their translational efficacy and stability (1–6). A protein called ‘host factor
6 required for phage Q β RNA replication’ (Hfq) forms a complex with a variety of small non-coding
7 RNAs and facilitates their binding to target mRNAs (7–13). Hfq of *Escherichia coli* is a protein
8 of 102 amino-acid residues, and its orthologues exist in a number of Gram-negative and -positive
9 bacteria constituting the Hfq family of proteins (14–17). Hfq has been shown to play a role in the
10 virulence of uropathogenic *E. coli* (18) and other bacterial species (19–35). However, the
11 mechanism involved in Hfq-mediated changes in the virulence of bacteria remains unclear.

12 We approached this issue using *Drosophila melanogaster* as a host for bacterial infection.
13 This insect has been used as an animal model to clarify the mechanisms of a wide variety of
14 biological events occurring in higher animal species, including humans (36, 37). As for its
15 immune system, *Drosophila* has only innate immunity, which closely resembles that of humans
16 (38–40). Bacteria that have invaded the hemolymph of *Drosophila* are recognized by innate
17 immune receptors, which exist in a soluble form or at the surface of immune cells called hemocytes.
18 These receptors bind substances that are secreted from bacteria or present as components of the cell
19 wall and cell membrane of bacteria (38–40). Once bound by bacterial substances, the receptors
20 activate signaling pathways in immune cells to induce antibacterial reactions, including the
21 production of antimicrobial peptides and the phagocytosis of bacteria (38–40). *E. coli* is generally
22 considered to be non-pathogenic in *Drosophila*, but our previous study showed that a high load of *E.*
23 *coli* may kill flies (41). Under such conditions, the Imd pathway, which is responsible for the
24 production of antimicrobial peptides, and phagocytosis by hemocytes are required for the long
25 survival of adult flies after infection with *E. coli* (41), suggesting the involvement of both humoral

1 and cellular immune responses in the protection of *Drosophila* from pathogenic infection with *E.*
2 *coli*.

3 Phagocytosis is a major cellular response among the self-defense strategies of *Drosophila*,
4 which are employed to fight against infection with pathogenic bacteria, and its mechanism has been
5 investigated in adult flies abdominally injected with bacteria, a surrogate procedure for septic
6 infection (41–47). We previously reported that σ^{38} , a stress-responsive sigma factor of *E. coli*
7 (48–52), is involved in the bacterial evasion of phagocytic killing in *Drosophila* (41). RNA
8 polymerase of *E. coli* is a multi-subunit enzyme consisting of a complex of four ‘core’ subunits,
9 $\alpha_2\beta\beta'\omega$, and a promoter-recognizing subunit named σ (53–55). There are seven sigma factors in
10 *E. coli*, and σ^{38} forms a complex with the core subunits of RNA polymerase (54, 56–58) when *E.*
11 *coli* encounters hostile conditions (50–54). σ^{38} -encoding *rpoS* is included in the genes, the
12 expression of which is controlled by Hfq (13, 14, 51, 59, 60). The expression of *rpoS* is
13 transiently augmented when *E. coli* abdominally infects adult *Drosophila* (41). Anticipating the
14 involvement of Hfq in σ^{38} actions to modulate the susceptibility of *E. coli* to phagocytosis, we
15 conducted a series of biochemical and genetic experiments. The outcome suggested a link
16 between Hfq and σ^{38} in controlling the phagocytic killing of *E. coli* in *Drosophila*.

17

1 **Materials and Methods**

2

3 *Fly stocks, bacterial strains, and cell cultures*

4 The following fly lines were used in this study: *Oregon R* (Kyorin-Fly, Kyorin University, Tokyo,
5 Japan) as a wild-type line, w^{1118} , $y^1 w^{1118}$, *da-GAL4*, *pxn-GAL4 8.1* (a gift from Michael J. Galiko),
6 *UAS-rpr* (Bloomington Drosophila Stock Center, Indiana University, Bloomington, IN; stock
7 number 5824), *imd^l* (61) (a gift from Bruno Lemaitre), and *PGRP-LE¹¹²;PGRP-LC⁷⁴⁵⁴* (62) (a gift
8 from Shoichiro Kurata). Some of the fly lines were used after changing balancers. The *E. coli*
9 strains IT1568 (parental) and TM589 (deficient in *hfq* expression) (Δhfq) (63) were used throughout
10 the study. Δhfq was transfected with the plasmid pTWV-Hfq-HIS₆ for the expression of
11 His-tagged Hfq. To construct pTWV-Hfq-HIS₆, the DNA fragment coding for the entire Hfq,
12 fused to the His tag at the C-terminus, was prepared by PCR using the chromosomal DNA of
13 W3110*mlc* (IT1568) as a template and the DNA oligomers 5'–
14 CCCAAGCTTTAAACTTTAACGGAAGTGA–3' and 5'–
15 CGCGAATTCTTAGTGATGGTGATGGTGATGCGATCCTTCGGTTTCTTCGCTG–3' as
16 primers, and inserted into the plasmid pTWV228 (Takara Bio Inc., Ohtsu, Japan) after digestion
17 with *Hind*III and *Eco*RI. We have confirmed the functional complementation of the defect
18 observed with Δhfq by the expression of His-tagged Hfq. The *E. coli* gene *katG* was forcedly
19 expressed in Δhfq using the plasmid pCA24N and T5-*lac* promoter, which was obtained from
20 ASKA clone (National BioResource Project: National Institute of Genetics, Mishima, Japan).
21 Bacteria were cultured with Luria-Bertani liquid medium, supplemented with antibiotics if
22 necessary, at 37°C until the cultures reached the stationary phase of cell growth, after which they
23 were harvested by centrifugation, washed with PBS, and used in the experiments. Hemocytes
24 were isolated from third-instar larvae according to a standard procedure (64) with slight
25 modifications (45). l(2)mbn, a larval hemocyte-derived cell line, was maintained with Schneider's
26 *Drosophila* medium (Life Technologies Japan, Tokyo, Japan) containing 10% (v/v) heat-inactivated

1 FBS at 25°C, as described previously (45).

2

3 *Bacterial infection and assays for colony-forming bacteria and phagocytosis*

4 All procedures were performed as described previously (41, 45). In brief, male adult flies, 3~7
5 days after eclosion (15~20 flies per vial, 1~3 vials in each experiment), were anesthetized with CO₂
6 and abdominally injected with 50 nl of PBS containing a given number of bacteria (stated in each
7 figure) with the aid of a nitrogen gas-operated microinjector (Narishige, Tokyo, Japan). We varied
8 the number of bacteria when infecting flies so that about 80% of the flies that succumbed most
9 severely to infection died in 3-7 days. Flies that had received the injection were maintained at
10 29°C until analysis. To determine the level of persistent infection in flies with bacteria, five
11 randomly chosen viable flies were homogenized, inoculated onto agar-solidified Luria-Bertani
12 medium, and incubated at 37°C, followed by the determination of the number of colonies. Under
13 these conditions, lysates prepared from uninfected flies did not give rise to colonies (data not
14 shown), indicating that endogenous bacteria, commensal or environmental, do not influence this
15 assay. To determine the level of phagocytosis, larval hemocytes were incubated with
16 FITC-labeled bacteria (phagocytes:bacteria = 1:500) at 25°C for 10 min, supplemented with trypan
17 blue to extinguish the fluorescence from unengulfed bacteria, and microscopically examined for
18 hemocytes containing engulfed bacteria (41). For the determination of bacterial deaths in
19 phagocytes, 1(2)mbn cells were incubated with unlabeled bacteria as described above, washed with
20 serum-free medium to remove unengulfed bacteria, maintained in culture up to 1.5 h, osmotically
21 lysed with water, and subjected to an assay for colony-forming bacteria. We were unable to use
22 larval hemocytes for this experiment because they died when cultured longer than 30 min (41).
23 For the inhibition of hemocyte phagocytosis of bacteria, flies were induced to undergo hemocyte
24 apoptosis by the expression of the pro-apoptotic gene *reaper* using the GAL4-UAS system with the
25 hemocyte-specific GAL4 driver *pxn-GAL4 8.1*, the activity of which was confirmed in the
26 hemocytes of adult flies using the fly line *pxn-GAL4 UAS-GFP* (see Fig. 4A), or the administration

1 of latex beads (2 μ m in diameter; Life Technologies Japan) in the abdomen before the injection of
2 bacteria (41).

3

4 *Western blotting and RT-PCR*

5 All procedures were performed as described previously (41). Briefly, flies, l(2)mbn cells, and
6 bacteria were lysed in a buffer containing detergents, separated by SDS-PAGE (5 μ g of proteins
7 were analyzed unless otherwise stated in the corresponding figure legends), followed by Western
8 blotting analysis with anti-Hfq rabbit antiserum (63), anti- σ^{38} rabbit antiserum (a gift from Akira
9 Ishihama), or anti-GFP mouse monoclonal IgG (mFX75: Wako Pure Chemical Industries, Osaka,
10 Japan), in which signals were visualized by a chemiluminescence reaction. For RT-PCR and
11 RT-qPCR analyses of flies, cultured cells, and *E. coli*, total RNA was prepared using the
12 acid-phenol method with TRIzol (Life Technologies Japan), and used as a template for RT with
13 random primers, and the resulting cDNA was subjected to conventional semi-quantitative PCR or
14 qPCR using THUNDERBIRD SYBR qPCR Mix (Toyobo, Osaka, Japan) and Mx5005p (Agilent,
15 CA). The DNA oligomers used as the primers in PCR were: 5'–
16 ATGCAGGGTTCTGTGACAGA–3' (forward) and 5'–AGAATACGGCGCAGTGCGTT–3'
17 (reverse) for the mRNA of RNA polymerase α subunit (RpoA);
18 5'-GATGAGAACGGAGTTGAGGT-3' (forward) and 5'-ACGCAAGTTACTCTCGATCAT-3'
19 (reverse) for σ^{38} mRNA; 5'–GTTAACCACCGCTAATTGAT–3' (forward) and 5'–
20 GACTACACACAGCAATTCGT–3' (reverse) for RpoS regulator RNA A (RprA); 5'–
21 GCTGCGCAATCGCTTCTACT–3' (forward) and 5'–TGGTGGAGTGGGCTTCATG–3'
22 (reverse) for dipteracin-A mRNA; 5'–CCCGGAGTGAAGGATG–3' (forward) and 5'–
23 GTTGCTGTGCGTCAAG–3' (reverse) for the mRNAs of attacin-A and attacin-B; and 5'–
24 GACGCTTCAAGGGACAGTATCTG–3' (forward) and 5'–AAACGCGGTTCTGCATGAG–3'
25 (reverse) for the mRNA of ribosomal protein 49 (Rp49). The levels of bacterial and fly mRNA
26 are shown relative to those of RpoA and Rp49, respectively, which were analyzed as internal

1 controls.

2

3 *Data processing and statistical analysis*

4 The results from quantitative analyses are expressed as the mean \pm SD of the data from at least
5 three independent experiments, unless otherwise stated in the corresponding figure legends.

6 Statistical analyses were performed using the log-rank test (Kaplan-Meier method) or two-tailed
7 Student's *t* test. A *p* value < 0.05 was considered significant.

8

1 **Results**

2

3 *Hfq-mediated persistent infection of E. coli in Drosophila*

4 As the first step toward clarifying how Hfq participates in controlling the persistent infection and
5 pathogenicity of *E. coli* in *Drosophila*, we examined how an *E. coli* strain deficient in the
6 expression of *hfq* (Δhfq) behaves, in comparison with its parental strain, when infecting adult flies.
7 When these bacteria were maintained in Luria-Bertani liquid medium, Δhfq showed a lower growth
8 rate with almost the same level of culture density at the stationary phase compared with the parental
9 strain (Fig. 1A). We then abdominally infected adult flies with these bacterial strains at two
10 different doses and examined the rate of fly death and the bacterial load. We first analyzed the
11 pathogenic effect of bacteria. Under conditions where the parental strain killed nearly 70% of flies
12 in 2 days, about 80% of flies were still alive even 4 days after the injection with Δhfq (Fig. 1B).
13 For the amount of bacteria present in flies, we homogenized infected flies and inoculated the
14 resulting lysates onto agar-solidified culture medium to determine the number of colony-formable
15 bacteria. The data showed that Δhfq was eliminated more rapidly from flies than the parental
16 strain (Fig. 1C). These results suggest the involvement of Hfq in the persistent infection of *E. coli*
17 in flies. To confirm that the lowered persistence observed in the mutant strain was solely due to
18 the loss of Hfq, we expressed *hfq* in Δhfq as a protein fused to 'His tag' (Fig. 2A) and examined its
19 effect on the two phenotypes. The growth rate of Δhfq in a liquid medium became almost the
20 same as that of the parental strain after expression of His-tagged Hfq (Fig. 2B), suggesting that the
21 tagged Hfq was functional in *E. coli*. We found that the expression of His-tagged Hfq increased
22 the rate of fly deaths after infection with Δhfq (Fig. 2C) and augmented the level of persistent
23 infection of Δhfq in adult flies (Fig. 2D), to the levels comparable with the data obtained using the
24 parental *E. coli* strain. From these findings, we concluded that Hfq is required for *E. coli* to
25 achieve a high level of persistent infection in *Drosophila*.

1

2 *Observation of hfq phenotype in Drosophila deficient in humoral immune response*

3 We then investigated whether the function of Hfq, in terms of the control of bacterial persistence, is
4 related to the humoral immunity of the host. Humoral immune responses in *Drosophila* against
5 Gram-negative bacteria are induced through the recognition of invaders by peptidoglycan
6 recognition proteins (PGRPs) and the subsequent activation of the Imd pathway leading to the
7 production of antimicrobial peptides such as diptericin and attacin (61, 62, 65–68). First, we
8 determined the level of mRNA of these peptides in flies infected with Δhfq and the parental strain,
9 but found no significant difference between flies infected with the two *E. coli* strains (Fig. 3A).
10 Next, we used mutant *Drosophila* lines having a defect in the humoral response as a host for
11 infection with *E. coli*. The fly line *imd*¹ that lacks Imd, a central component in the Imd pathway,
12 succumbed to *E. coli* infection more severely than the control fly line (left panel in Fig. 3B),
13 indicating a reduced level of immunity in those flies. We found that Δhfq showed decreased levels
14 of killing effect and persistent infection in this mutant fly line (middle and right panels in Fig. 3B,
15 respectively), as observed in the experiment using wild-type flies (see Figs. 1B and 1C). To
16 confirm this further, we used another mutant fly line deficient in the expression of both PGRP-LE
17 and PGRP-LC, peptidoglycan-recognizing receptors responsible for the detection of *E. coli* and the
18 subsequent induction of the Imd pathway (left panel in Fig. 3C), and again found that Δhfq killed
19 flies at a lower rate (middle panel in Fig. 3C) and was less persistent (right panel in Fig. 3C) than
20 the parent strain. Collectively, these results indicate that Hfq is required for the host-killing effect
21 and persistent infection of *E. coli* in flies deficient in the Imd pathway as observed using wild-type
22 flies as a host. It is most probable that Hfq-mediated persistence of *E. coli* in adult flies is
23 achieved independent of humoral immune responses.

24

25 *Involvement of cellular immunity in the control of bacterial persistence in Drosophila by Hfq*

26 Next, we asked if phagocytosis, a main cellular immune response, is related to the Hfq-mediated

1 control of the persistence of *E. coli*. To generate flies defective in phagocytosis, we expressed the
2 pro-apoptotic gene *reaper* (69) specifically in hemocytes, the major phagocytes in *Drosophila*,
3 using the GAL4-UAS system. Flies that possessed both *UAS-rpr* and hemocyte-specific *pxn-Gal4*
4 contained a smaller number of hemocytes compared to control flies having only *pxn-Gal4* (left
5 panel in Fig. 4A). Those flies were more vulnerable to infection with the parental *E. coli* strain
6 than control flies (middle panel in Fig. 4A), suggesting impaired phagocytosis of bacteria. We
7 found that these flies died at almost the same rate after infection with the parental and Δhfq strains
8 (right panel in Fig. 4A). Similar results were obtained with wild-type flies that had been
9 pre-injected with latex beads (Fig. 4B), another method of inhibiting hemocyte phagocytosis (41,
10 70). Furthermore, Δhfq no longer showed a lowered persistence phenotype when it infected the
11 bead-injected flies (Fig. 4C). To further confirm the above findings, we used flies deficient in
12 both the Imd pathway and hemocyte phagocytosis as a host for bacterial infection. *imd¹* flies were
13 injected with latex beads, abdominally infected with *E. coli*, and subjected to assays for fly survival
14 and bacterial persistence. We found no difference in either assay between the parental and Δhfq *E.*
15 *coli* strains (Fig. 4D), as observed using flies with impaired phagocytosis. The data obtained so far
16 indicate that Hfq is no longer required for the host-killing effect and persistent infection of *E. coli* in
17 flies deficient in hemocyte phagocytosis. It is thus likely that Hfq-mediated persistence of *E. coli*
18 in adult flies is related to phagocytosis.

19 A simple explanation for the above observations is that Hfq-lacking *E. coli* undergoes
20 phagocytic killing more efficiently than the parental strain. Next, we tested this hypothesis by
21 conducting an assay for the phagocytosis of *E. coli in vitro* using hemocytes isolated from
22 *Drosophila* larvae. The extent of phagocytosis was assessed by determining the ratio of
23 hemocytes that had accomplished phagocytosis as well as the number of *E. coli* engulfed by a given
24 number of hemocytes, but we observed no significant difference between Δhfq and the parental
25 strain in either index (Fig. 5A). Then, we examined a possible difference in the rate of killing of *E.*
26 *coli* in hemocytes after engulfment. For this examination, l(2)mbn cells, a cell line derived from

1 larval hemocytes, were used as phagocytes because larval hemocytes spontaneously lysed and could
2 not be maintained in culture for a period long enough to analyze the killing of engulfed bacteria.
3 l(2)mbn cells and *E. coli* were co-incubated, unengulfed *E. coli* were washed out, and phagocytes,
4 after engulfment, were maintained in culture further for 1.5 h. Whole-cell lysates of l(2)mbn cells
5 at 0 and 1.5 h of incubation were inoculated onto agar-solidified medium to determine the number
6 of bacteria capable of forming colonies that existed in the phagocytes. We found that Δhfq
7 decreased more rapidly than its parental strain (Fig. 5B), showing that the Hfq-lacking strain is
8 more effectively killed in phagocytes after engulfment. These results collectively indicated that
9 Hfq helps *E. coli* evade killing by *Drosophila* phagocytes after engulfment.

10 The observation that flies deficient in the Imd pathway (Figs. 3B and 3C) or hemocyte
11 phagocytosis (Figs. 4A and 4B) succumbed more severely to infection with *E. coli* compared to
12 control flies indicates the roles of both humoral and cellular immunity in the protection of
13 *Drosophila* from bacterial infection. Thus, we examined whether these two immune responses are
14 related. To do so, flies were abdominally injected with latex beads or left untreated, infected with
15 the parental *E. coli* strain, and analyzed for the level of antimicrobial peptide mRNA. Also,
16 hemocytes isolated from larvae of Imd-lacking and control flies were tested for the phagocytosis of
17 the parental *E. coli* strain *in vitro*. We found that impaired phagocytosis did not influence the
18 production of antimicrobial peptide mRNA in flies after infection, and the compromised humoral
19 response did not alter the hemocyte phagocytosis of *E. coli* (Supplemental Fig. 1). These results
20 suggest that humoral and cellular responses are independently involved in the antibacterial
21 immunity of *Drosophila*. A possible explanation for the observation that the inhibition of either
22 immune response causes earlier fly deaths after bacterial infection could be that two responses are
23 induced at different times: phagocytosis is an immediate reaction for the direct elimination of
24 invading pathogens followed by the production of antimicrobial peptides to kill the remaining
25 pathogens.

26

1 σ^{38} -mediated Hfq function in controlling *E. coli* persistence

2 *E. coli* genes whose expression is controlled by Hfq include *rpoS*, which codes for the
3 stress-responsive sigma factor σ^{38} (13, 49, 50), and the phenotype of Δhfq described above
4 resembles what is seen in the *rpoS* mutant, as we previously reported (41). To test the possible
5 involvement of σ^{38} in the Hfq-mediated control of the persistence and killing effect of *E. coli* in
6 *Drosophila*, we examined whether the forced expression of *rpoS* restores the phenotypes of *E. coli*
7 observed after a loss of Hfq. When the lysates of *E. coli* were analyzed by Western blotting using
8 Ab specific to σ^{38} , the signal was less intense in Δhfq than in the parental strain (left panel in Fig.
9 6A), confirming the Hfq control of *rpoS* expression. The forced expression of *rpoS* in Δhfq , which
10 was confirmed by Western blotting (right panel in Fig. 6A), partly recovered a host-killing effect
11 (Fig. 6B). Also, *rpoS*-expressing Δhfq a higher level of persistent infection in adult flies compared
12 to Δhfq that harbored vector alone (Fig. 6C). Furthermore, Δhfq became more resistant to killing
13 in l(2)mbn cells, after engulfment, by the expression of *rpoS* (Fig. 6D). These results indicated
14 that *rpoS* partly accounts for the Hfq-dependent persistent infection of *E. coli* in *Drosophila*, and
15 suggested the presence of other Hfq-controlled genes besides *rpoS* that explain the entire actions of
16 Hfq. Target genes of σ^{38} include *katE* and *katG* that code for an enzyme detoxifying reactive
17 oxygen species. In our previous study (41), forced expression of *katE* or *katG* recovered lowered
18 persistence and host-killing effect seen in σ^{38} -lacking *E. coli*. We thus examined if the defective
19 phenotypes of Hfq-lacking *E. coli* were rescued by the expression of *katG* (Fig. 6E). The data
20 indicated that overexpression of *katG* (left panel) restored, though partly, host-killing effect (middle
21 panel) and persistence of Δhfq (right panel). It is thus likely that the Hfq-mediated inhibition of
22 phagocytic killing is achieved through augmented expression of *katG*.

23

24 Post-transcriptional increase of σ^{38} mRNA by Hfq during infection

25 Finally, we examined the mechanism of the control of *rpoS* expression by Hfq. We first

1 determined the level of σ^{38} in the whole-cell lysates of Δhfq by Western blotting and found that σ^{38}
2 increased after the expression of His-tagged Hfq (Fig. 7A), confirming a positive control of *rpoS*
3 expression by Hfq. Hfq as an RNA chaperone is thought to influence the level of mRNA
4 independent of transcription. The strength of the *rpoS* promoter was then analyzed using GFP as a
5 reporter, and the level of GFP determined in Western blotting was almost the same between the
6 parental and Hfq-lacking *E. coli* strains (Fig. 7B), suggesting no influence of Hfq on the promoter
7 activity of *rpoS*. We found that σ^{38} mRNA increased in adult flies infected with the parental but
8 not Hfq-lacking *E. coli* strains (Fig. 7C), indicating the Hfq-mediated expression of *rpoS* after
9 infection. The expression of *rpoS* is post-transcriptionally enhanced by RprA (71–73), a small
10 non-coding RNA controlled by Hfq, and Hfq remains abundant in *E. coli* regardless of the phase of
11 its growth (11). We thus determined a possible change in the level of this RNA by RT-qPCR after
12 infection. The level of RprA was higher in the parental, but not RpoS-lacking, strains in adult flies
13 (Fig. 7D), suggesting the presence of a positive feedback loop in the expression of *rpoS* and
14 RprA-encoding genes. Collectively, an elevated level of RprA in the parental *E. coli* most likely
15 accounts for the Hfq-dependent increase of σ^{38} after infection.

16

1 **Discussion**

2

3 There are a number of reports describing the role of Hfq in the virulence and drug resistance of
4 bacteria (14–17, 74, 75). However, the mechanism by which Hfq is involved in the control of
5 bacterial behavior under infectious conditions has yet to be clarified. In the present study, we
6 showed that Hfq is required for the maintenance of the persistent infection of *E. coli* in *Drosophila*.
7 Hfq seemed to inhibit the killing of bacteria in hemocytes, a major phagocytic cell type in
8 *Drosophila* (76, 77), after engulfment, which should be a mechanistic explanation for the
9 Hfq-mediated persistent infection of *E. coli*. There was no difference in the efficacy of
10 phagocytosis of *E. coli* between the parental and Hfq-lacking strains, while the Hfq-lacking strain
11 was more rapidly killed in phagocytes after engulfment than the parent strain. Therefore, it is
12 possible that the Hfq-lacking strain is more efficiently engulfed by *Drosophila* phagocytes than the
13 parental strain.

14 Hfq is known as an RNA chaperone that binds small non-coding RNAs to facilitate their
15 actions in post-transcriptionally controlling the stability and translational efficacy of mRNAs (3, 12,
16 13). One of the target RNAs of Hfq is RprA, which positively controls the expression of *rpoS*
17 coding for the stress-inducing sigma factor σ^{38} (71–73). The levels of RprA and σ^{38} mRNA rose
18 in *E. coli* after infection of adult flies. Furthermore, we showed that the forced expression of *rpoS*
19 or *katG* restored, though partly, the *E. coli* phenotype of lowered persistence caused by the lack of
20 Hfq. These findings allowed us to propose the following mechanism for the Hfq- and
21 σ^{38} -mediated persistent infection of *E. coli* in *Drosophila*: the expression of RprA is augmented
22 after infection; Hfq forms a complex with RprA; Hfq-bound RprA stabilizes σ^{38} mRNA; the
23 production of σ^{38} is enhanced; σ^{38} induces the transcription of genes including *katG*; killing of *E.*
24 *coli* engulfed by hemocytes is inhibited; and the persistent infection of *E. coli* is achieved. As a
25 result, an increase in the abundance of bacteria in *Drosophila* brings about early death of the host
26 organism. Rescue of the defective phenotype in Hfq-lacking *E. coli* by the forced expression of

1 *rpoS* and *katG* was partial. This suggests that other genes besides *rpoS* are induced by Hfq and
2 involved in the control of *E. coli* persistence in *Drosophila*, and that other genes in addition to *katG*
3 are induced by σ^{38} and responsible for the inhibition of the phagocytic killing of *E. coli* in
4 *Drosophila* hemocytes. Further study is required for identifying and characterizing such genes.

5 The expression of *hfq* seems to remain unchanged regardless of the phase of *E. coli* cell
6 growth (11). We showed an increase of RprA, a small non-coding RNA that stabilizes σ^{38} mRNA,
7 in adult flies. It is therefore likely that the post-transcriptional control of *rpoS* expression depends
8 on increased RprA levels after infection. Thus, another important issue to be resolved is the
9 mechanism behind the augmented expression of the RprA-encoding gene in *Drosophila*. The
10 two-component signal transduction system (TCS) senses environmental changes and causes the
11 alteration of gene expression so that bacteria adapt to the new environment and survive (78–80).
12 The TCS consists of two distinct proteins, a membrane protein called the sensor kinase that serves
13 as a receptor, and a DNA-binding transcription factor called the response regulator that is
14 phosphorylated by the sensor kinase for activation (79). It is probable that the TCS control of
15 gene expression is related to the levels of persistence and pathogenicity of bacteria in host
16 organisms. Indeed, we previously found that EnvZ-OmpR, a TCS of *E. coli*, is required for
17 bacteria to maintain low pathogenicity in *Drosophila* (81). More specifically, positive control of
18 the expression of RprA-encoding gene by RcsC/RcsD-RcsB and CpxA-CpxR was reported
19 previously (82). To examine the possible involvement of such TCSs in the control of *rprA*
20 expression under infectious conditions, further investigation is required.

21

22 **Acknowledgements**

23 We are grateful to Dr. Akira Ishihama of Hosei University for antibody. Dr. Shoichiro Kurata of
24 Tohoku University, Dr. Bruno Lemaitre of the École Polytechnique Fédérale de Lausanne, Dr.
25 Michael J. Galko of the University of Texas, Bloomington *Drosophila* Stock Center, and
26 Kyorin-Fly are thanked for fly lines. We also thank the National BioResource Project for plasmid.

1

2 **Disclosures**

3 The authors have no financial conflict of interest.

4

1 **References**

2

- 3 1. Gottesman, S. 2004. The small RNA regulators of *Escherichia coli*: roles and mechanisms. *Annu.*
4 *Rev. Microbiol.* 58: 303–328.
- 5 2. Waters, L. S., and G. Storz. 2009. Regulatory RNAs in bacteria. *Cell* 136: 615–628.
- 6 3. Gottesman, S., and G. Storz. 2011. Bacterial small RNA regulators: versatile roles and rapidly
7 evolving variations. *Cold Spring Harb. Perspect. Biol.* 3: a003798.
- 8 4. Moon, K., and S. Gottesman. 2011. Competition among Hfq-binding small RNAs in *Escherichia*
9 *coli*. *Mol. Microbiol.* 82:1545–1562.
- 10 5. Papenfort, K., and J. Vogel. 2009. Multiple target regulation by small noncoding RNAs rewires
11 gene expression at the post-transcriptional level. *Res. Microbiol.* 160: 278–287.
- 12 6. Fröhlich, K. S., and J. Vogel. 2009. Activation of gene expression by small RNA. *Curr. Opin.*
13 *Microbiol.* 12: 674–682.
- 14 7. Kajitani, M., and A. Ishihama. 1991. Identification and sequence determination of the host factor
15 gene for bacteriophage Q β . *Nucleic Acids Res.* 19: 1063–1066.
- 16 8. Muffler, A., D. Fischer, and R. Hengge-Aronis. 1996. The RNA-binding protein HF-I, known as
17 a host factor for phage Qbeta RNA replication, is essential for *rpoS* translation in *Escherichia*
18 *coli*. *Genes Dev.* 10: 1143–1151.
- 19 9. Møller, T., T. Franch, P. Højrup, D. R. Keene, H. P. Bächinger, R. G. Brennan, and P.
20 Valentin-Hansen. 2002. Hfq: a bacterial Sm-like protein that mediates RNA-RNA interaction.
21 *Mol. Cell* 9: 23–30.
- 22 10. Aiba, H. 2007. Mechanism of RNA silencing by Hfq-binding small RNA. *Curr. Opin.*
23 *Microbiol.* 10: 134–139.
- 24 11. Brennan, R. G., and T. M. Link. 2007. Hfq structure, function and ligand binding. *Curr. Opin.*
25 *Microbiol.* 10: 125–133.
- 26 12. Vogel, J., and B. F. Luisi. 2011. Hfq and its constellation of RNA. *Nat. Rev. Microbiol.* 9: 578–

- 1 589.
- 2 13. De Lay, N., D. J. Schu, and Gottesman, S. 2013. Bacterial small RNA-based negative
3 regulation: Hfq and its accomplices. *J. Biol. Chem.* 288: 7996–8003.
- 4 14. Chao, Y., and J. Vogel. 2010. The role of Hfq in bacterial pathogens. *Curr. Opin. Microbiol.* 13:
5 24–33.
- 6 15. Liu, Y., N. Wu, J. Dong, Y. Gao, X. Zhang, C. Mu, N. Shao, and G. Yang. 2010. Hfq is a global
7 regulator that controls the pathogenicity of *Staphylococcus aureus*. *PLoS ONE* 5: e13069.
- 8 16. Sauer, E. 2013. Structure and RNA-binding properties of the bacterial LSm protein Hfq. *RNA*
9 *Biol.* 10: 610–618.
- 10 17. Van Assche, E., S. Van Puyvelde, J. Vanderleyden, and H. P. Steenackers. 2015. RNA-binding
11 proteins involved in post-transcriptional regulation in bacteria. *Front. Microbiol.* 6:141.
- 12 18. Kulesus, R. R., K. Diaz-Perez, E. S. Slechta, D. S. Eto, and M. A. Mulvey. 2008. Impact of the
13 RNA chaperone Hfq on the fitness and virulence potential of uropathogenic *Escherichia coli*.
14 *Infect. Immun.* 76: 3019–3026.
- 15 19. Boudry, P., C. Gracia, M. Monot, J. Caillet, L. Saujet, E. Hajnsdorf, B. Dupuy, I.
16 Martin-Verstraete, and O. Soutourina. 2014. Pleiotropic role of the RNA chaperone protein Hfq
17 in the human pathogen *Clostridium difficile*. *J. Bacteriol.* 196: 3234–3248.
- 18 20. Sittka, A., V. Pfeiffer, K. Tedin, and J. Vogel. 2007. The RNA chaperone Hfq is essential for
19 the virulence of *Salmonella typhimurium*. *Mol. Microbiol.* 63: 193–217.
- 20 21. Meng, X., X. Meng, C. Zhu, H. Wang, J. Wang, J. Nie, P. R. Hardwidge, and G. Zhu. 2013. The
21 RNA chaperone Hfq regulates expression of fimbrial-related genes and virulence of *Salmonella*
22 *enterica* serovar *Enteritidis*. *FEMS Microbiol. Lett.* 346: 90–96.
- 23 22. Meibom, K. L., A. L. Forslund, K. Kuoppa, K. Alkhuder, I. Dubail, M. Dupuis, A. Forsberg,
24 and A. Charbit. 2009. Hfq, a novel pleiotropic regulator of virulence-associated genes in
25 *Francisella tularensis*. *Infect. Immun.* 77: 1866–1880.
- 26 23. Dietrich, M., R. Munke, M. Gottschald, E. Ziska, J. P. Boettcher, H. Mollenkopf, and A.

- 1 Friedrich. 2009. The effect of hfq on global gene expression and virulence in *Neisseria*
2 *gonorrhoeae*. *FEBS J.* 276: 5507–5520.
- 3 24. Fantappiè, L., M. M. Metruccio, K. L. Seib, F. Oriente, E. Cartocci, F. Ferlicca, M. M. Giuliani,
4 V. Scarlato, and I. Delany. 2009. The RNA chaperone Hfq is involved in stress response and
5 virulence in *Neisseria meningitidis* and is a pleiotropic regulator of protein expression. *Infect.*
6 *Immun.* 77: 1842–1853.
- 7 25. Schiano, C. A., and W. W. Lathem. 2012. Post-transcriptional regulation of gene expression in
8 *Yersinia* species. *Front. Cell. Infect. Microbiol.* 2: 129.
- 9 26. Subashchandrabose, S., R. M. Leveque, R. N. Kirkwood, M. Kiupel, and M. H. Mulks. 2013.
10 The RNA chaperone Hfq promotes fitness of *Actinobacillus pleuropneumoniae* during porcine
11 pleuropneumonia. *Infect. Immun.* 81: 2952–2961.
- 12 27. Hempel, R. J., D. J. Morton, T. W. Seale, P. W. Whitby, and T. L. Stull. 2013. The role of the
13 RNA chaperone Hfq in *Haemophilus influenzae* pathogenesis. *BMC Microbiol.* 13: 134.
- 14 28. Gangaiah, D., M. Labandeira-Ray, X. Zhang, K. R. Fortney, S. Ellinger, B. Zwickl, B. Baker, Y.
15 Liu, D. M. Janowicz, B. P. Katz, C. A. Brautigam, R. S. Munson Jr., E. J. Hansen, and S. M.
16 Spinola. 2014. *Haemophilus ducreyi* Hfq contributes to virulence gene regulation as cells enter
17 stationary phase. *mBio* 5: e01081–13.
- 18 29. Lenz, D. H., K. C. Mock, B. N. Lilley, R. V. Kulkarni, N. S. Wingreen, and B. L. Bassler. 2004.
19 The small RNA chaperone Hfq and multiple small RNAs control quorum sensing in *Vibrio*
20 *harveyi* and *Vibrio cholerae*. *Cell* 118: 69–82.
- 21 30. Christiansen, J. K., J. S. Nielsen, T. Ebersbach, P. Valentin-Hansen, L. Sjøgaard-Andersen, B. H.
22 Kallipolitis. 2006. Identification of small Hfq-binding RNAs in *Listeria monocytogenes*. *RNA*
23 12: 1383–1396.
- 24 31. McNealy, T. L., V. Forsbach-Birk, C. Shi, and R. Marre. 2005. The Hfq homolog in *Legionella*
25 *pneumophila* demonstrates regulation by LetA and RpoS and interacts with the global regulator
26 CsrA. *J. Bacteriol.* 187: 1527–1532.

- 1 32. Bibova, I., K. Skopova, J. Masin, O. Cerny, D. Hot, P. Sebo, and B. Vecerek. 2013. The RNA
2 chaperone Hfq is required for virulence of *Bordetella pertussis*. *Infect. Immun.* 81: 4081–4090.
- 3 33. Robertson, G. T., and R. M. Roop Jr. 1999. The *Brucella abortus* host factor I (HF-I) protein
4 contributes to stress resistance during stationary phase and is a major determinant of virulence in
5 mice. *Mol. Microbiol.* 34: 690–700.
- 6 34. Mitobe, J., T. Morita-Ishihara, A. Ishihama, and H. Watanabe. 2008. Involvement of
7 RNA-binding protein Hfq in the post-transcriptional regulation of *invE* gene expression in
8 *Shigella sonnei*. *J. Biol. Chem.* 283: 5738–5747.
- 9 35. Zeng, Q., R. R. MaNally, and G. W. Sundin. 2013. Global small RNA chaperone Hfq and
10 regulatory small RNAs are important virulence regulators in *Erwinia amylovora*. *J. Bacteriol.*
11 195: 1706–1717.
- 12 36. Wang, L., I. Kounatidis, and P. Ligoxygakis. 2013. *Drosophila* as a model to study the role of
13 blood cells in inflammation, innate immunity and cancer. *Front. Cell. Infect. Microbiol.* 3: 113.
- 14 37. Fuchs, Y., and Steller, H. 2011. Programmed cell death in animal development and disease. *Cell*
15 147: 742–758.
- 16 38. Elrod-Erickson, M., S. Mishra, and D. Schneider. 2000. Interactions between the cellular and
17 humoral immune responses in *Drosophila*. *Curr. Biol.* 10: 781–784.
- 18 39. Brennan, C. A., and K. V. Anderson. 2004. *Drosophila*: the genetics of innate immune
19 recognition and response. *Annu. Rev. Immunol.* 22: 457–483.
- 20 40. Lemaitre, B., and J. Hoffmann. 2007. The host defense of *Drosophila melanogaster*. *Annu. Rev.*
21 *Immunol.* 25: 697–743.
- 22 41. Shiratsuchi, A., N. Shimamoto, M. Nitta, T. Q. Tuan, A. Firdausi, M. Gawasawa, K. Yamamoto,
23 A. Ishihama, and Y. Nakanishi. 2014. Role for σ^{38} in prolonged survival of *Escherichia coli* in
24 *Drosophila melanogaster*. *J. Immunol.* 192: 666–675.
- 25 42. Elrod-Erickson, M., S. Mishra, and D. Schneider. 2000. Interactions between the cellular and
26 humoral immune responses in *Drosophila*. *Curr. Biol.* 10: 781–784.

- 1 43. Ulvila, J., L. M. Vanha-aho, A. Kleino, M. Vähä-Mäkilä, M. Vuoksio, S. Eskelinen, D.
2 Hultmark, C. Kocks, M. Hallman, M. Parikka, and M. Rämet. 2011. Cofilin regulator 14-3-3 ζ is
3 an evolutionarily conserved protein required for phagocytosis and microbial resistance. *J.*
4 *Leukoc. Biol.* 89: 649–659.
- 5 44. Kocks, C., J. H. Cho, N. Nehme, J. Ulvila, A. M. Pearson, M. Meister, C. Storm, S. L. Conto, C.
6 Hetru, L. M. Stuart, T. Stehle, J. A. Hoffmann, J. M. Reichhart, D. Ferradon, M. Rämet, R. A.
7 Ezekowitz. 2005. Eater, a transmembrane protein mediating phagocytosis of bacterial pathogens
8 in *Drosophila*. *Cell* 123: 335–346.
- 9 45. Hashimoto, Y., Y. Tabuchi, K. Sakurai, M. Kutsuna, K. Kurokawa, T. Awasaki, K. Sekimizu, Y.
10 Nakanishi, and A. Shiratsuchi. 2009. Identification of lipoteichoic acid as a ligand for Draper in
11 the phagocytosis of *Staphylococcus aureus* by *Drosophila* hemocytes. *J. Immunol.* 183: 7451–
12 7460.
- 13 46. Shiratsuchi, A., T. Mori, K. Sakurai, K. Nagaosa, K. Sekimizu, B. L. Lee, and Y. Nakanishi.
14 2012. Independent recognition of *Staphylococcus aureus* by two receptors for phagocytosis in
15 *Drosophila*. *J. Biol. Chem.* 287: 21663–21672.
- 16 47. Nonaka, S., K. Nagaosa, T. Mori, A. Shiratsuchi, and Y. Nakanishi. 2013. Integrin
17 α PS3/ β v-mediated phagocytosis of apoptotic cells and bacteria in *Drosophila*. *J. Biol. Chem.*
18 288: 10374–10380.
- 19 48. Kazmierczak, M. J., M. Wiedmann, and K. J. Boor. 2005. Alternative sigma factors and their
20 roles in bacterial virulence. *Microbiol. Mol. Biol. Rev.* 69: 527–543.
- 21 49. Hengge, R. 2009. Proteolysis of σ^S (RpoS) and the general stress response in *Escherichia coli*.
22 *Res. Microbiol.* 160: 667–676.
- 23 50. Dong, T., and H. E. Schellhorn. 2010. Role of RpoS in virulence of pathogens. *Infect. Immun.*
24 78: 887–897.
- 25 51. Battesti, A., N. Majadani, and S. Gottesman. 2011. The RpoS-mediated general stress
26 response in *Escherichia coli*. *Annu. Rev. Microbiol.* 65: 189–213.

- 1 52. Bak, G., K. Han, D. Kim, and Y. Lee. 2013. Roles of *rpoS*-activating small RNAs in pathways
2 leading to acid resistance of *Escherichia coli*. *Microbiol. Open* 3: 15–28.
- 3 53. Ishihama, A. 2010. Prokaryotic genome regulation: multifactor promoters, multitarget
4 regulators and hierarchic networks. *FEMS Microbiol. Rev.* 34: 628–645.
- 5 54. Ishihama, A. 2012. Prokaryotic genome regulation: A revolutionary paradigm. *Proc. Jpn. Acad.,
6 Ser. B* 88: 485–508.
- 7 55. Ishihama, A. 2000. Functional modulation of *Escherichia coli* RNA polymerase. *Annu. Rev.
8 Microbiol.* 54: 499–518.
- 9 56. Jishage, M., and A. Ishihama. 1995. Regulation of RNA polymerase sigma subunit synthesis in
10 *Escherichia coli*: intracellular levels of σ^{70} and σ^{38} . *J. Bacteriol.* 177: 6832–6835.
- 11 57. Hengge-Aronis, R. 2002. Signal transduction and regulatory mechanisms involved in control of
12 the σ^S (RpoS) subunit of RNA polymerase. *Microbiol. Mol. Biol. Rev.* 66: 373–395.
- 13 58. Hengge, R. 2008. The two-component network and the general stress sigma factor RpoS (σ^S) in
14 *Escherichia coli*. *Adv. Exp. Med. Biol.* 631: 40–53.
- 15 59. Hengge, R. 2009. Proteolysis of σ^S (RpoS) and the general stress response in *Escherichia coli*.
16 *Res. Microbiol.* 160: 667–676.
- 17 60. Soper, T., P. Mandin, N. Majalani, S. Gottesman, and S. A. Woodson. 2010. Positive regulation
18 by small RNAs and the role of Hfq. *Proc. Natl. Acad. Sci. USA* 107: 9602–9607.
- 19 61. Lemaitre, B., E. Kromer-Metzger, L. Michaut, E. Nicolas, M. Meister, P. Georgel, J.-M.
20 Reichhart, and J. A. Hoffmann. 1995. A recessive mutation, immune deficiency (*imd*), defines
21 two distinct control pathways in the *Drosophila* host defense. *Proc. Natl. Acad. Sci. USA* 92:
22 9465–9469.
- 23 62. Takehana, A., T. Yano, S. Mita, A. Kotani, Y. Oshima, and S. Kurata. 2004. Peptidoglycan
24 recognition protein (PGRP)-LE and PGRP-LC act synergistically in *Drosophila* immunity.
25 *EMBO J.* 23: 4690–4700.
- 26 63. Morita, T., K. Maki, and H. Aiba. 2005. RNase E-based ribonucleoprotein complexes:

- 1 mechanical basis of mRNA destabilization mediated by bacterial noncoding RNAs. *Genes Dev.*
2 19: 2176–2186.
- 3 64. Avet-Rochex, A., E. Bergeret, I. Attree, M. Meister, and M.-O. Fauvarque. 2005. Suppression
4 of *Drosophila* cellular immunity by directed expression of the ExoS toxin GAP domain of
5 *Pseudomonas aeruginosa*. *Cell. Microbiol.* 7: 799–810.
- 6 65. Choe, K. M., T. Wemer, S. Stöven, D. Hultmark, and K. V. Anderson. 2002. Requirement for a
7 peptidoglycan recognition protein (PGRP) in Relish activation and antibacterial immune
8 response in *Drosophila*. *Science* 296: 359-562.
- 9 66. Gottar, M., V. Gobert, T. Michel, M. Belvin, G. Duyk, J. A. Hoffmann, D. Ferrandon, and J.
10 Royet. 2002. The *Drosophila* immune response against Gram-negative bacteria is mediated by a
11 peptidoglycan recognition protein. *Nature* 416: 640-644.
- 12 67. Rämet, M., P. Manfruelli, A. Pearson, B. Mathey-Prevo, and R. A. Ezekowitz. 2002.
13 Functional genomic analysis of phagocytosis and identification of a *Drosophila* receptor for *E.*
14 *coli*. *Nature* 416: 644-648.
- 15 68. Myllymäki, H., S. Valanne, and M. Rämet. 2014. The *Drosophila* Imd signaling pathway. *J.*
16 *Immunol.* 192: 3455–3462.
- 17 69. White, K., E. Tahaoglu, and H. Steller. 1996. Cell killing by the *Drosophila* gene reaper.
18 *Science* 271: 805–807.
- 19 70. Ganesan, S., I. Aggarwal, N. Paquette, and N. Silverman. 2011. NF-κB/Rel proteins and the
20 humoral immune responses of *Drosophila melanogaster*. *Curr. Top. Microbiol. Immunol.* 349:
21 25–60.
- 22 71. Argaman, L., R. Hershberg, J. Vogel, G. Bejerano, E. G. Wanger, H. Margalit, and S. Altuvia.
23 2001. Novel small RNA-encoding genes in the intergenic regions of *Escherichia coli*. *Curr. Biol.*
24 11: 941–950.
- 25 72. Majdalani, N., S. Chen, J. Murrow, K. St John, and S. Gottesman. 2001. Regulation of RpoS by
26 a novel small RNA: the characterization of RprA. *Mol. Microbiol.* 39: 1382–1394.

- 1 73. Majdalani, N., D. Hernandez, and S. Gottesman. 2002. Regulation and mode of action of the
2 second small RNA activator of RpoS translation, RprA. *Mol. Microbiol.* 46: 813–826.
- 3 74. Lalaouna, D., A. Eyraud, S. Chabelskaya, B. Felden, and E. Massé. 2014. Regulatory RNAs
4 involved in bacterial antibiotic resistance. *PLoS Pathog.* 10:e1004299.
- 5 75. Nishino, K., S. Yamasaki, M. Hayashi-Nishino, and A. Yamaguchi. 2011. Effect of
6 overexpression of small non-coding DsrA RNA on multidrug efflux in *Escherichia coli*. *J.*
7 *Antimicrob. Chemother.* 66: 291-296.
- 8 76. Underhill, D. M., and A. Ozinsky. 2002. Phagocytosis of microbes: complexity in action. *Annu.*
9 *Rev. Immunol.* 20: 825–852.
- 10 77. Vlisidou, I., and W. Wood. 2015. *Drosophila* blood cells and their role in immune responses.
11 *FEBS J.* 282: 1368–1382.
- 12 78. Stock, A. M., V. L. Robinson, and P. N. Goudreau. 2000. Two-component signal transduction.
13 *Annu. Rev. Biochem.* 69: 183–215.
- 14 79. Mascher, M., J. D. Helmann, and G. Uden. 2006. Stimulus perception in bacterial
15 signal-transducing histidine kinases. *Microbiol. Mol. Biol. Rev.* 70: 941–938.
- 16 80. Gao, R., T. R. Mack, and A. M. Stock. 2007. Bacterial response regulators: versatile regulatory
17 strategies from common domains. *Trends Biochem. Sci.* 32: 225–234.
- 18 81. Pukklay, P., Y. Nakanishi, M. Nitta, K. Yamamoto, A. Ishihama, and A. Shiratsuchi. 2013.
19 Involvement of EnvZ-OmpR two-component system in virulence control of *Escherichia coli* in
20 *Drosophila melanogaster*. *Biochem. Biophys. Res. Commun.* 438: 306–311.
- 21 82. Vogt, S. L., A. D. Evans, R. L. Guest, and T. L. Raivio. 2014. The Cpx envelope stress response
22 regulates and is regulated by small noncoding RNAs. *J. Bacteriol.* 196: 4229–4238.

23

1 **Footnotes**

2

3 ¹This work was supported by Grants-in-Aid for Scientific Research from the Japan Society for the
4 Promotion of Science (grant numbers 23570160 and 26440049 to A.S., and grant number 25291021
5 to Y.N.), a grant from The Institute for Fermentation (to A.S.), and an institutional research grant
6 from Kanazawa University (to A.S.).

7

8 ²Address correspondence and reprint requests to Dr. Akiko Shiratsuchi or Dr. Yoshinobu Nakanishi,
9 Graduate School of Medical Sciences, Kanazawa University, Shizenken, Kakuma-machi,
10 Kanazawa, Ishikawa 920-1192, Japan. E-mail addresses: ashira@staff.kanazawa-u.ac.jp (A.S.)
11 and nakanaka@p.kanazawa-u.ac.jp (Y.N.)

12

13 ³Abbreviations used in this paper: Hfq, host factor required for phage Q β RNA replication; PGRP,
14 peptidoglycan recognition protein(s); Rp49, ribosomal protein 49; RpoA, α subunit of *E. coli* RNA
15 polymerase; RpoS, σ^{38} subunit of *E. coli* RNA polymerase; RprA, RpoS regulator RNA A; TCS,
16 two-component regulatory system.

17

1 **Figure Legends**

2

3 **FIGURE 1.** Decreases in the persistent infection and host killing effect of *E. coli* with the loss of
4 Hfq. (A) Growth rates of Hfq-lacking (Δhfq) and parental strains of *E. coli* were determined in a
5 Luria-Bertani liquid medium. Representative results from three independent experiments with
6 similar results are shown. (B) The ratio of live *Oregon R* flies was determined at the indicated
7 time points after abdominal infection with Δhfq and parental strains. Data were analyzed using the
8 log-rank test. Results from one of three independent experiments that gave similar results are
9 shown. (C) The level of colony-forming bacteria was determined in lysates of *Oregon R* flies at
10 the indicated time points after abdominal infection with Δhfq and parental strains. Data from five
11 (parent) and three (Δhfq) independent experiments are shown after analysis by Student's *t* test.

12

13 **FIGURE 2.** Recovery of the persistence and host killing effect of Δhfq by the forced expression
14 of His-tagged Hfq. The mutant *E. coli* strain Δhfq , which had been transfected with a plasmid for
15 *hfq* expression (*hfq*) or a vector alone (vector), was analyzed. (A) The levels of Hfq, endogenous
16 Hfq and ectopically expressed His-tagged Hfq, in lysates of the indicated bacteria were determined
17 by Western blotting with anti-Hfq Ab. Representative data from one of two independent
18 experiments with similar results are shown. (B) The growth rate of the indicated bacteria was
19 determined in a Luria-Bertani liquid medium. Data from one of three independent experiments
20 with similar results are shown. (C) The survival of *Oregon R* flies was analyzed after abdominal
21 infection with bacteria. Data from one of three independent experiments are shown after analysis
22 by the log-rank test. (D) The level of colony-forming bacteria was determined in lysates of
23 *Oregon R* flies at the indicated time points after abdominal infection with bacteria. Results from
24 three (parent) and four (Δhfq) independent experiments are shown after analysis by Student's *t* test.

25

26 **FIGURE 3.** No relationship of the control of *E. coli* persistence by Hfq with host humoral

1 immunity. (A) mRNA levels of dipteracin and attacin were determined by RT-qPCR with RNA
2 prepared from *Oregon R* flies infected with the indicated *E. coli* strains for 3 h. Values for the
3 mRNAs of attacin and dipteracin are shown after normalization with those of Rp49 mRNA.
4 Results from 13 (attacin) and 11 (dipteracin) independent experiments are shown after analysis by
5 Student's *t* test. NS, difference not significant. (B) Levels of the persistence and killing effect of
6 *E. coli* were determined using the fly line *imd^l* defective in the Imd pathway as the host. (left) The
7 fly lines *imd^l* and *w¹¹¹⁸*, as a control with the same genetic background, were abdominally infected
8 with the parental *E. coli* strain, and the proportion of live flies was determined at the indicated time
9 points. Data were analyzed by the log-rank test. Representative results from one of three
10 independent experiments that gave similar results are shown. (middle) The ratio of viable *imd^l*
11 flies was determined at the indicated time points after infection with Δhfq or the parental strain.
12 Data were analyzed by the log-rank test. Representative results from one of three independent
13 experiments with similar results are shown. (right) Level of colony-forming activity in lysates of
14 *imd^l* flies, which had been infected with Δhfq or the parental strain, was determined at the indicated
15 time points after infection. *Oregon R* flies infected with the parental strain were similarly
16 analyzed as a control. Data from three (*Oregon R*) and four (*imd^l*) independent experiments are
17 shown after analysis by Student's *t* test. (C) The killing effect of *E. coli* was determined using the
18 fly line *PGRP-LE¹¹²;PGRP-LC⁷⁴⁵⁴* lacking both PGRP-LC and PGRP-LE as the host. (left) The
19 fly lines *PGRP-LE¹¹²;PGRP-LC⁷⁴⁵⁴* and *y¹ w¹¹¹⁸*, as a control with the same genetic background,
20 were abdominally infected with the parental *E. coli* strain, and the proportion of live flies was
21 determined at the indicated time points. Data were analyzed by the log-rank test. Representative
22 results from one of three independent experiments that gave similar results are shown. (middle)
23 The proportion of live *PGRP-LE¹¹²;PGRP-LC⁷⁴⁵⁴* flies was determined at the indicated time points
24 after infection with Δhfq or the parental strain. Data were analyzed by the log-rank test.
25 Representative results from one of three independent experiments that gave similar results are
26 shown. (right) The level of colony-forming activity in lysates of *PGRP-LE¹¹²;PGRP-LC⁷⁴⁵⁴* flies,

1 which had been infected with Δhfq or the parental strain, was determined at the indicated time
2 points after infection. *Oregon R* flies infected with the parental strain were similarly analyzed as a
3 control. Data from three (*Oregon R*) and four (*PGRP-LE¹¹²;PGRP-LC⁷⁴⁵⁴*) independent
4 experiments are shown after analysis by Student's *t* test.

5
6 **FIGURE 4.** Involvement of phagocytosis in Hfq-mediated persistence of *E. coli*. (A) (left)
7 Cells contained in the hemolymph of adult flies that possessed the trans-gene *pxn-GAL4 UAS-GFP*
8 or *pxn-GAL4 UAS-rpr UAS-GFP* were examined by fluorescence microscopy. Phase contrast and
9 fluorescence views of the same microscopic fields are shown. Scale bar, 20 μm . (middle) Flies
10 possessing the trans-gene *pxn-GAL4 UAS-rpr*, *pxn-GAL4* or *UAS-rpr* were abdominally infected
11 with the parental *E. coli* strain, and the proportion of live flies was determined at the indicated time
12 points. Data were analyzed by the log-rank test. Representative results from one of three
13 independent experiments that gave similar results are shown. (right) The proportion of live flies
14 possessing *pxn-GAL4 UAS-rpr* was determined after infection with the indicated *E. coli* strains.
15 Data were analyzed using the log-rank test. NS, difference not significant. Representative
16 results from one of three independent experiments that gave similar results are shown. (B) (left)
17 *Oregon R* flies, which had been pre-injected with latex beads or PBS alone followed by the
18 infection of the parental *E. coli* strain, were subjected to a survival assay. Data were analyzed by
19 the log-rank test. Representative results from one of three independent experiments that gave
20 similar results are shown. (right) The proportion of live *Oregon R* flies, which had been
21 pre-injected with latex beads followed by the infection with the indicated *E. coli* strains, was
22 determined at the indicated time points. Data were analyzed by the log-rank test. NS, difference
23 not significant. Representative results from one of three independent experiments that gave
24 similar results are shown. (C) The level of colony-formable bacteria was determined in the
25 whole-fly lysates of *Oregon R*, which had been pre-injected with latex beads, at the indicated time
26 points after abdominal infection with the parental or Δhfq *E. coli* strain. Flies not treated with

1 latex beads and infected with the parental bacteria were similarly analyzed as a control. Results
2 from three (no bead injection) and five (the others) independent experiments are shown after
3 analysis by Student's *t* test. NS, difference not significant. **(D)** Levels of the killing effect (left)
4 and persistence (right) of *E. coli* were determined using *imd^l* flies that had been pre-injected with
5 latex beads as a host. Data were analyzed by the log-rank test (left, representative data from one
6 of three independent experiments shown) or Student's *t* test (right, results from five independent
7 experiments analyzed). NS, difference not significant.

8

9 **FIGURE 5.** Hfq inhibition of *E. coli* phagocytosis by *Drosophila* phagocytes. **(A)** Hemocytes
10 obtained from third-instar larvae of *Oregon R* flies were incubated with the indicated *E. coli* strains,
11 and the level of phagocytosis was determined using two indices. Data from four independent
12 experiments are shown after analysis by Student's *t* test. NS, difference not significant. **(B)** The
13 hemocyte cell line l(2)mbn that had been incubated with the indicated *E. coli* strains was
14 maintained in culture further for 1.5 h, and their lysates were subjected to an assay of
15 colony-forming bacteria. Data from three independent experiments are shown after analysis by
16 Student's *t* test.

17

18 **FIGURE 6.** σ^{38} -mediated function of Hfq in maintaining the persistent infection of *E. coli*. **(A)**
19 Lysates of parental *E. coli* and Δhfq , which had been transfected with *rpoS*-expressing plasmid
20 (*rpoS*) or an empty vector, were subjected to Western blotting with anti- σ^{38} Ab. Data from one of
21 two independent experiments that gave similar results are shown. **(B)** *Oregon R* flies were
22 infected with Δhfq possessing *rpoS*-expressing plasmid (*rpoS*) or an empty vector, and the
23 proportion of live flies was determined at the indicated time points. Flies infected with the
24 parental bacteria were similarly analyzed as a control. The data from one of four independent
25 experiments that yielded similar results are shown after analysis by the log-rank test. **(C)** Lysates
26 of *Oregon R* flies that had been infected with the indicated bacterial strains were analyzed for the

1 level of colony-forming bacteria at the indicated time points after infection. Data from three
2 independent experiments are shown after analysis by Student's *t* test. (D) l(2)mbn cells that had
3 been incubated with the indicated *E. coli* strains were maintained in culture further for 1.5 h, and
4 their lysates were subjected to an assay of colony-formable bacteria. Data from three (parent) and
5 four (Δhfq) independent experiments are shown after analysis by Student's *t* test. (E) The killing
6 effect and persistence of Δhfq in adult flies were determined after forced expression of *katG*. (left)
7 Proteins in whole-cell lysates of the indicated bacteria were separated by SDS-PAGE followed by
8 staining with Coomassie brilliant blue. The arrowhead points to the position of KatG.
9 Representative data from one of three independent experiments with similar results are shown.
10 (middle) Killing effect of the indicated bacteria was determined using *Oregon R* flies as a host.
11 Data were analyzed by the log-rank test. Representative results from one of three independent
12 experiments that gave similar results are shown. (right) Persistence of the indicated bacteria in
13 *Oregon R* flies was determined. Data from three independent experiments are shown after
14 analysis by Student's *t* test.

15
16 **FIGURE 7.** Increases of σ^{38} mRNA and RprA in *E. coli* after infection. (A) Lysates prepared
17 from parental *E. coli* and Δhfq , which had been transfected with a plasmid for the expression of
18 His-tagged Hfq (*hfq*) or a vector alone, were analyzed for the level of σ^{38} by Western blotting with
19 anti- σ^{38} Ab. Data from one of three independent experiments with similar results are shown. (B)
20 Lysates from parental *E. coli* and Δhfq harboring plasmid for GFP expression driven by *rpoS*
21 promoter were analyzed by Western blotting with anti-GFP Ab. Data from one of three
22 independent experiments with similar results are shown. (C) RNA prepared from *Oregon R* flies
23 that had been infected with the parental or Δhfq *E. coli* strain for 5 min together with the RNA of
24 the same *E. coli* strains was subjected to RT-qPCR analysis for the level of the mRNA of σ^{38} and
25 RpoA. Values for σ^{38} mRNA are shown after normalization with those of RpoA mRNA.

1 Results from 15 (parent), eight (Δhfq , no infection), and seven (Δhfq , infection) independent
2 experiments are shown after analysis by Student's t test. NS, difference not significant. **(D)**
3 RNA prepared from *Oregon R* flies, after the infection with parental or σ^{38} -lacking ($\Delta rpoS$) *E. coli*
4 strain for 5 min together with RNA of the same *E. coli* strains, was subjected to RT-qPCR for the
5 level of RprA and RpoA mRNA. Values of RprA were normalized with those of RpoA mRNA,
6 and results from nine (left panel: parent, no infection), six (left panel: parent, infection), three (right
7 panel: parent, no infection), eight ($\Delta rpoS$, no infection), and seven ($\Delta rpoS$, infection) independent
8 experiments analyzed by Student's t test are shown.

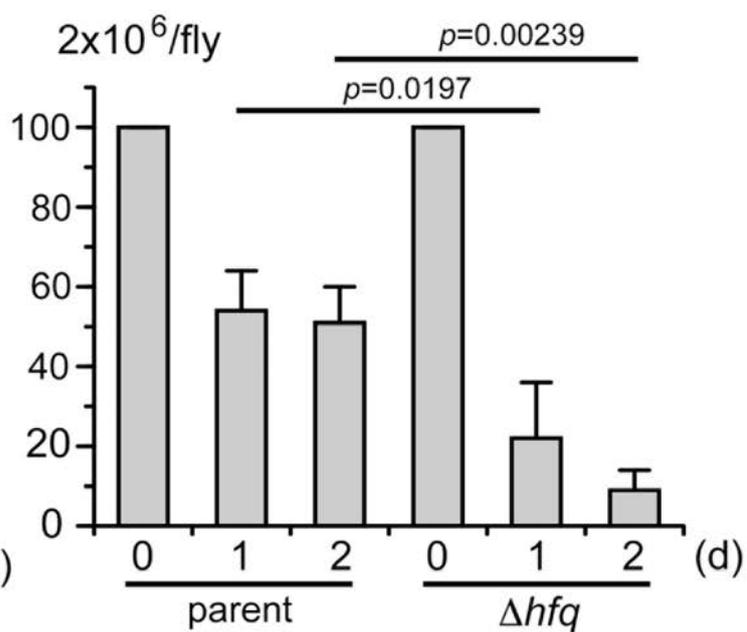
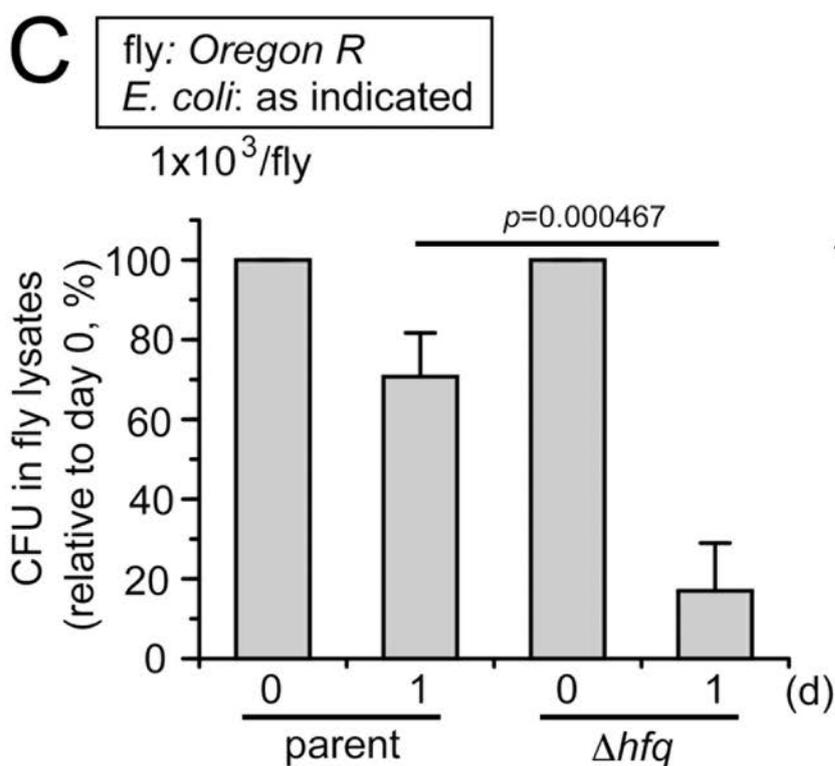
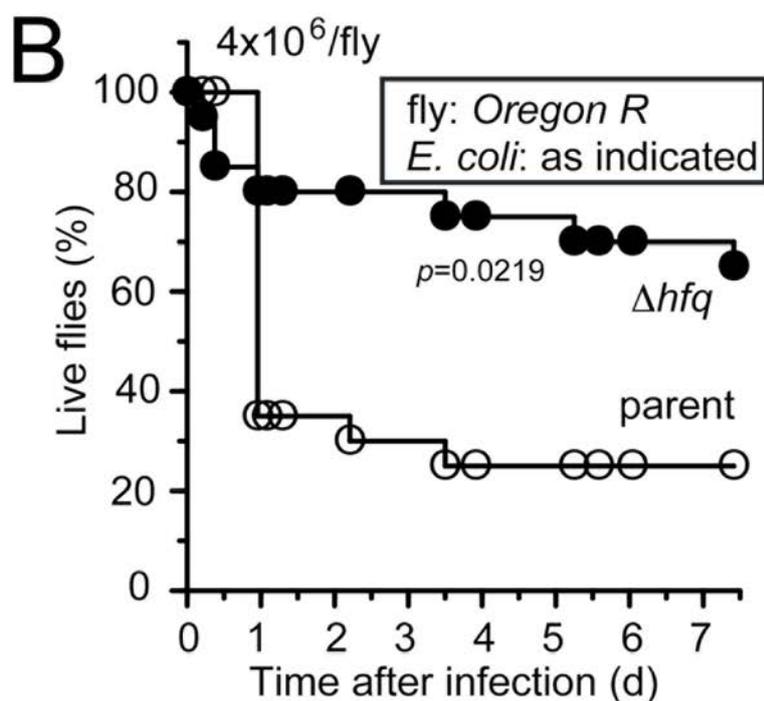
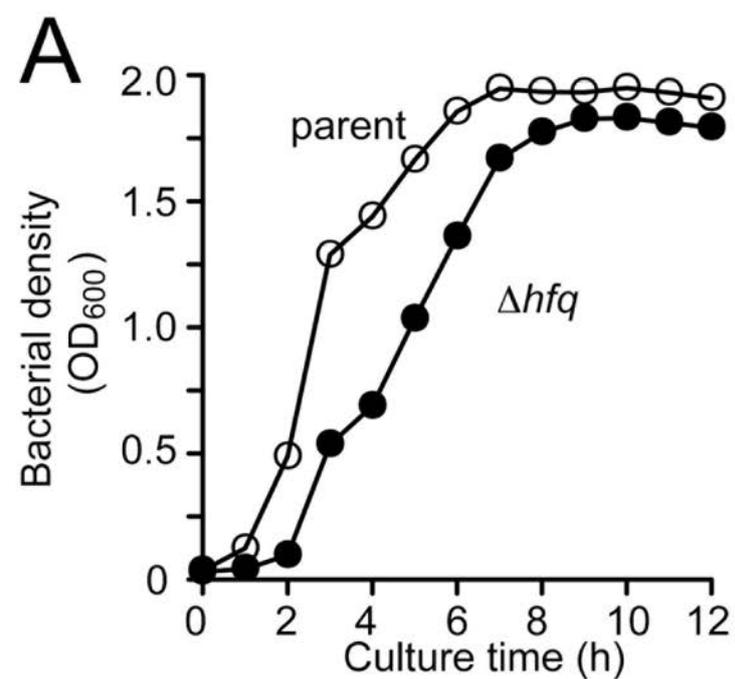


Fig. 1

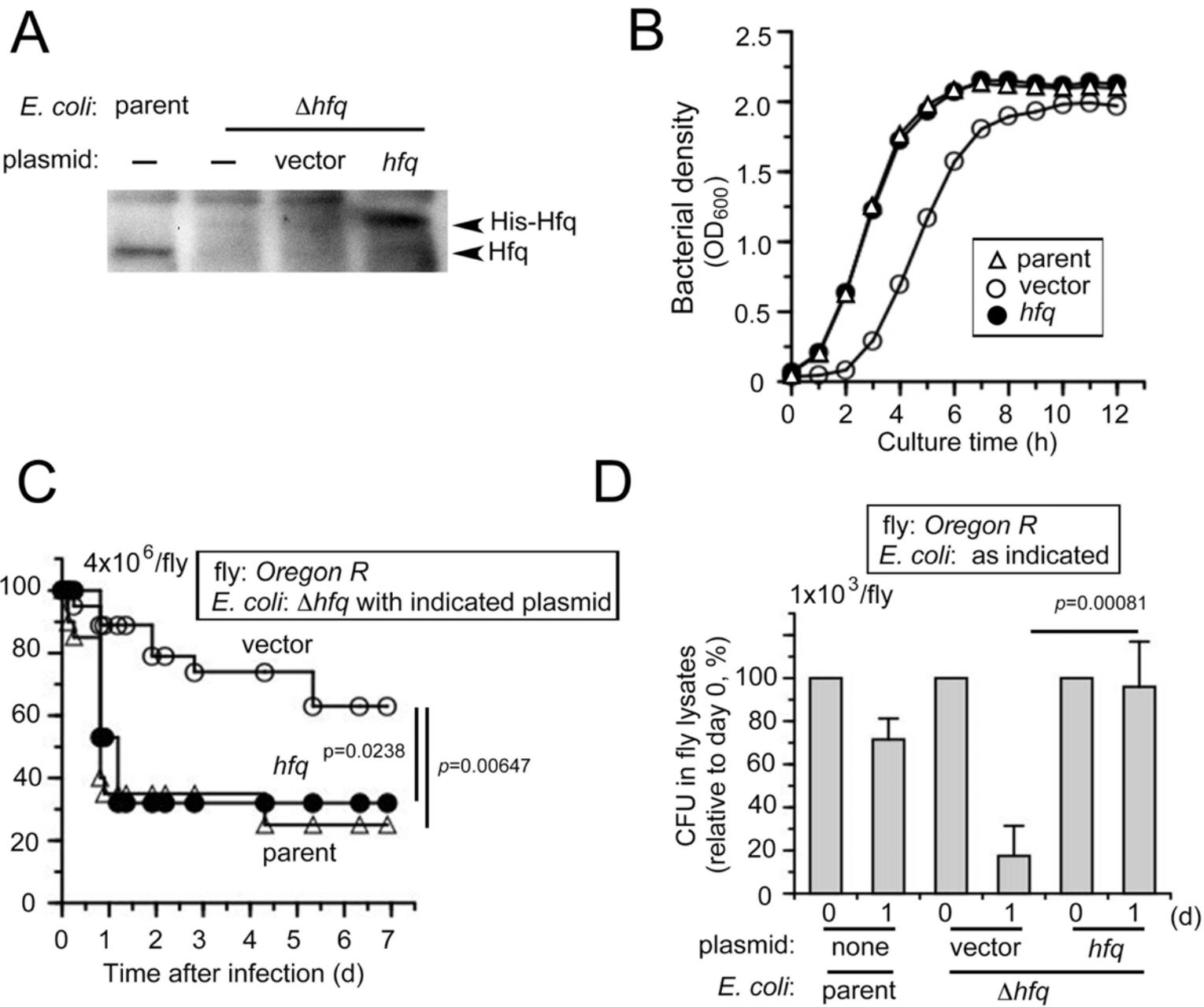


Fig. 2

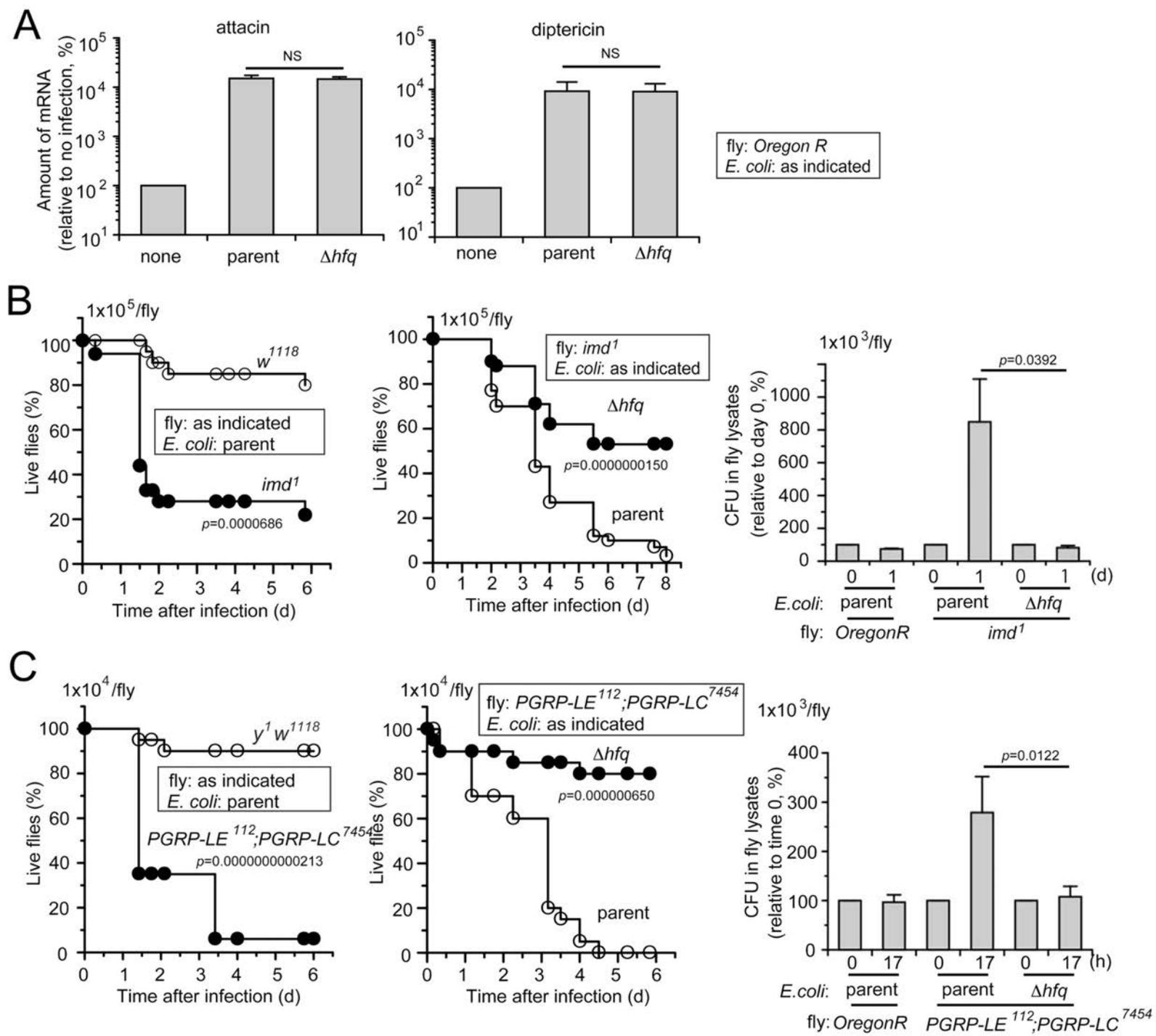
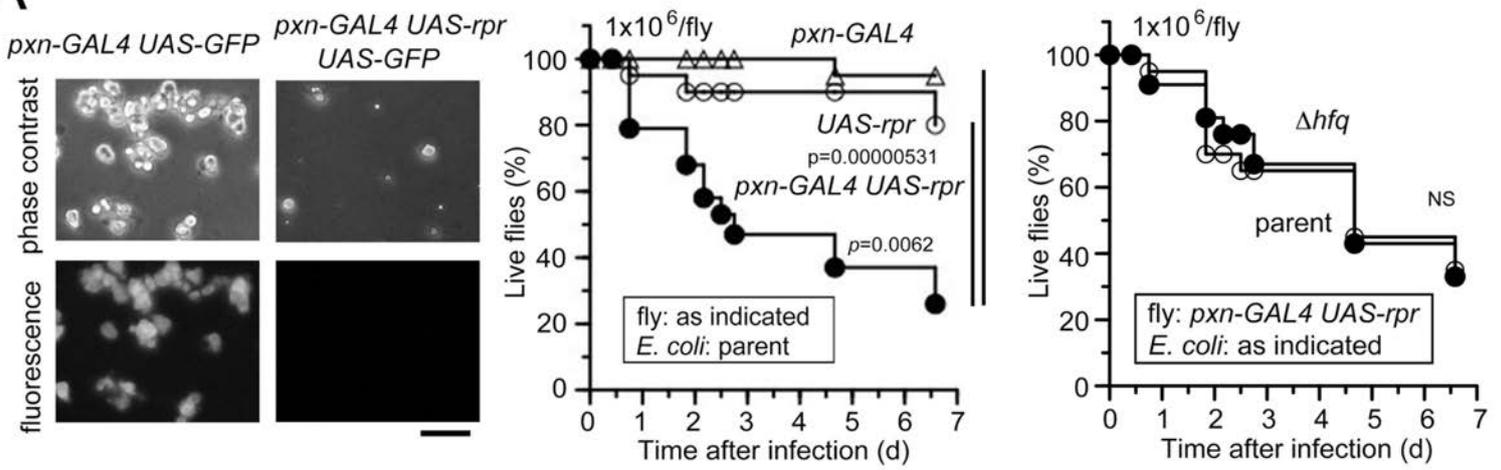
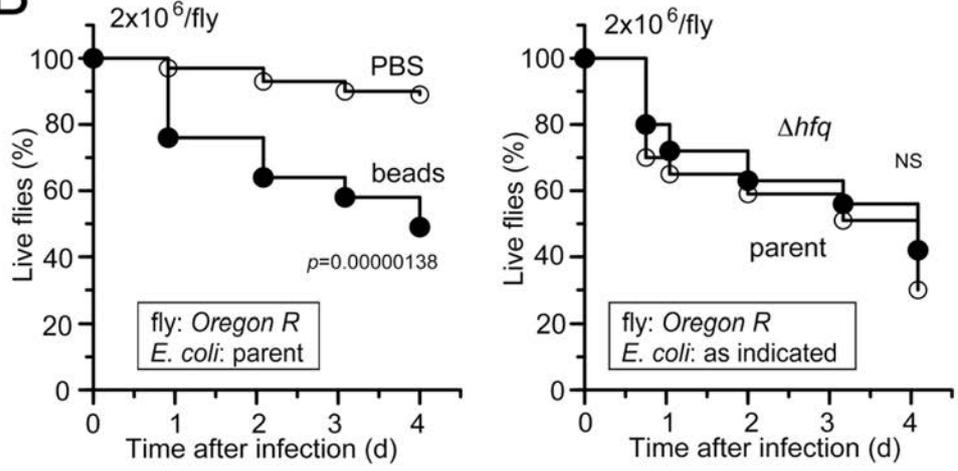
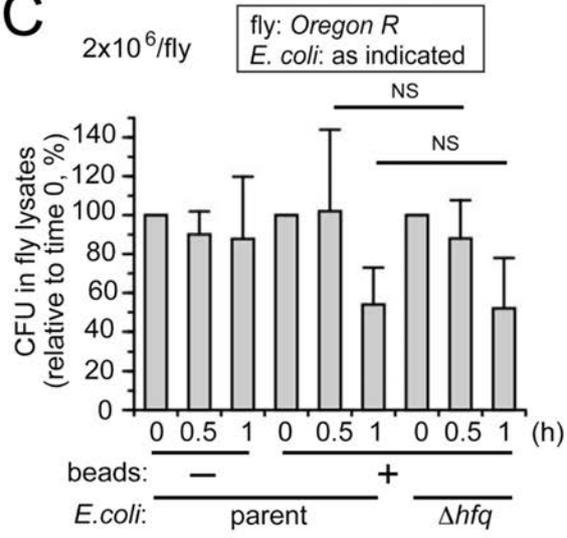
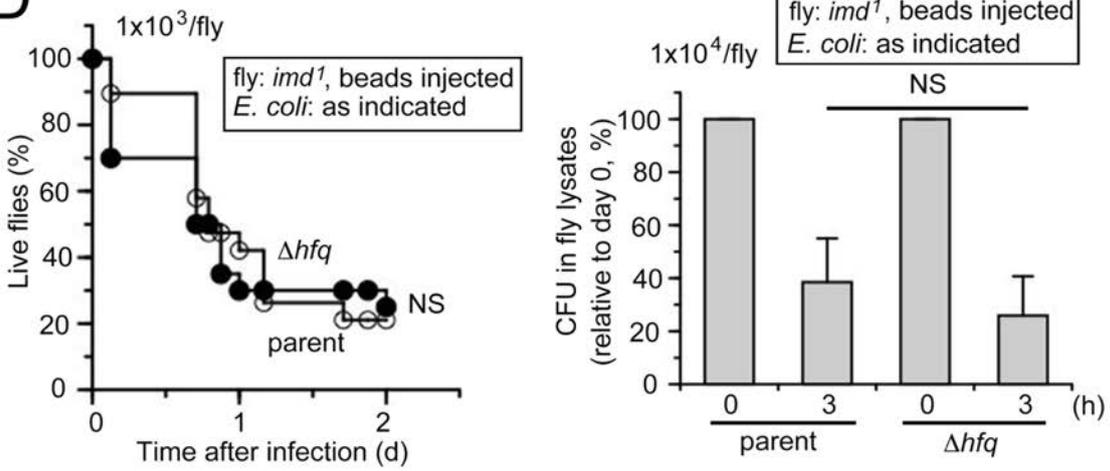


Fig. 3

A**B****C****D****Fig. 4**

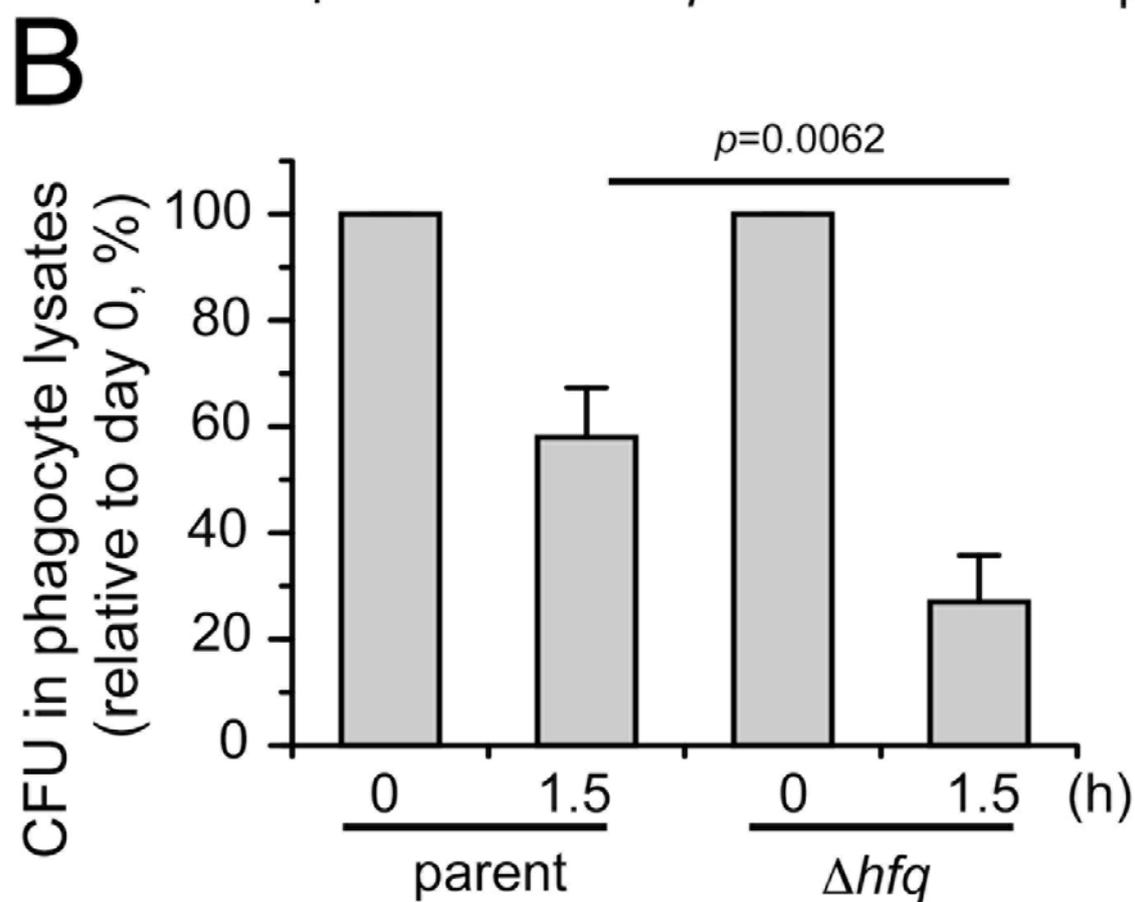
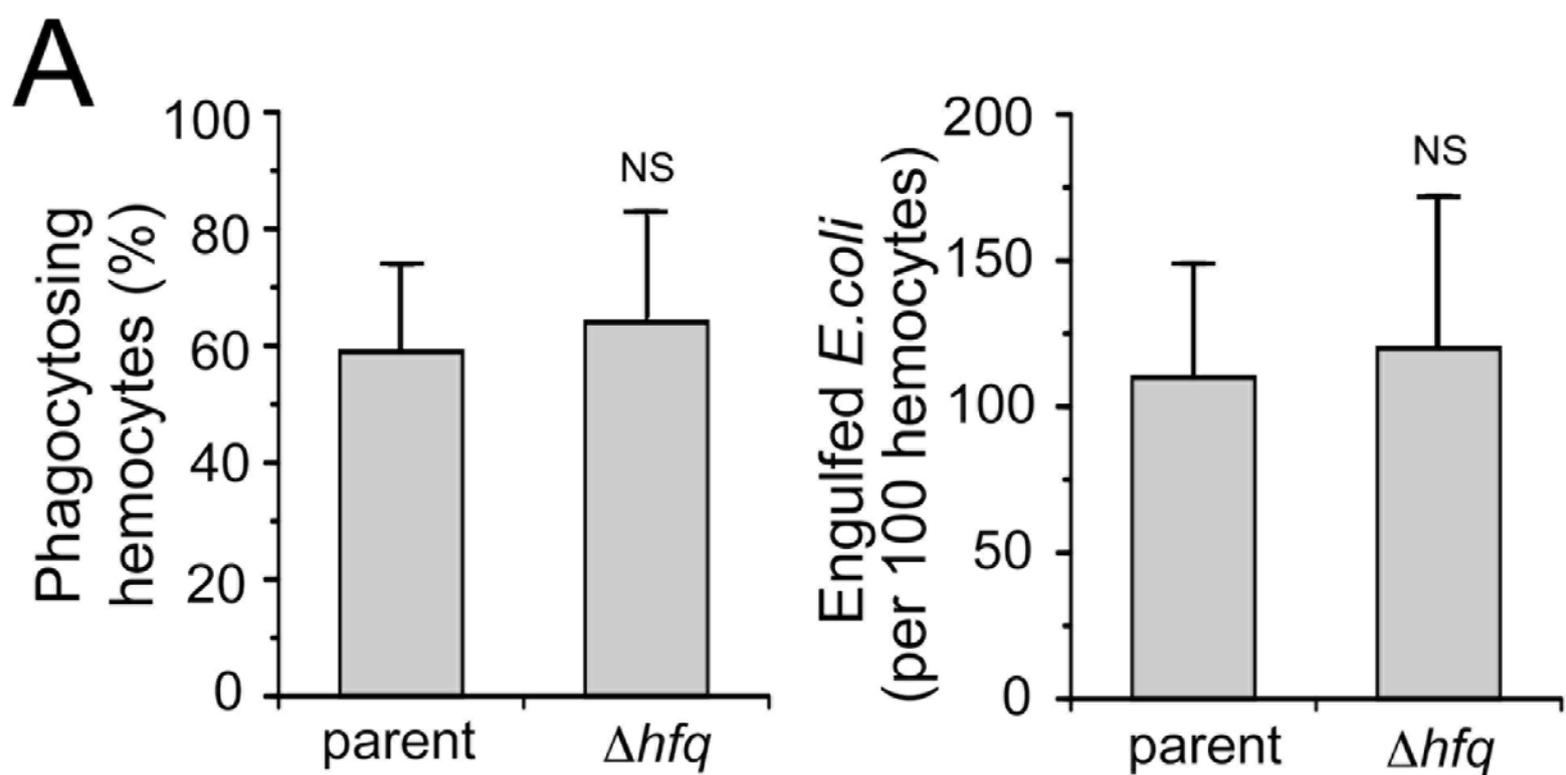


Fig. 5

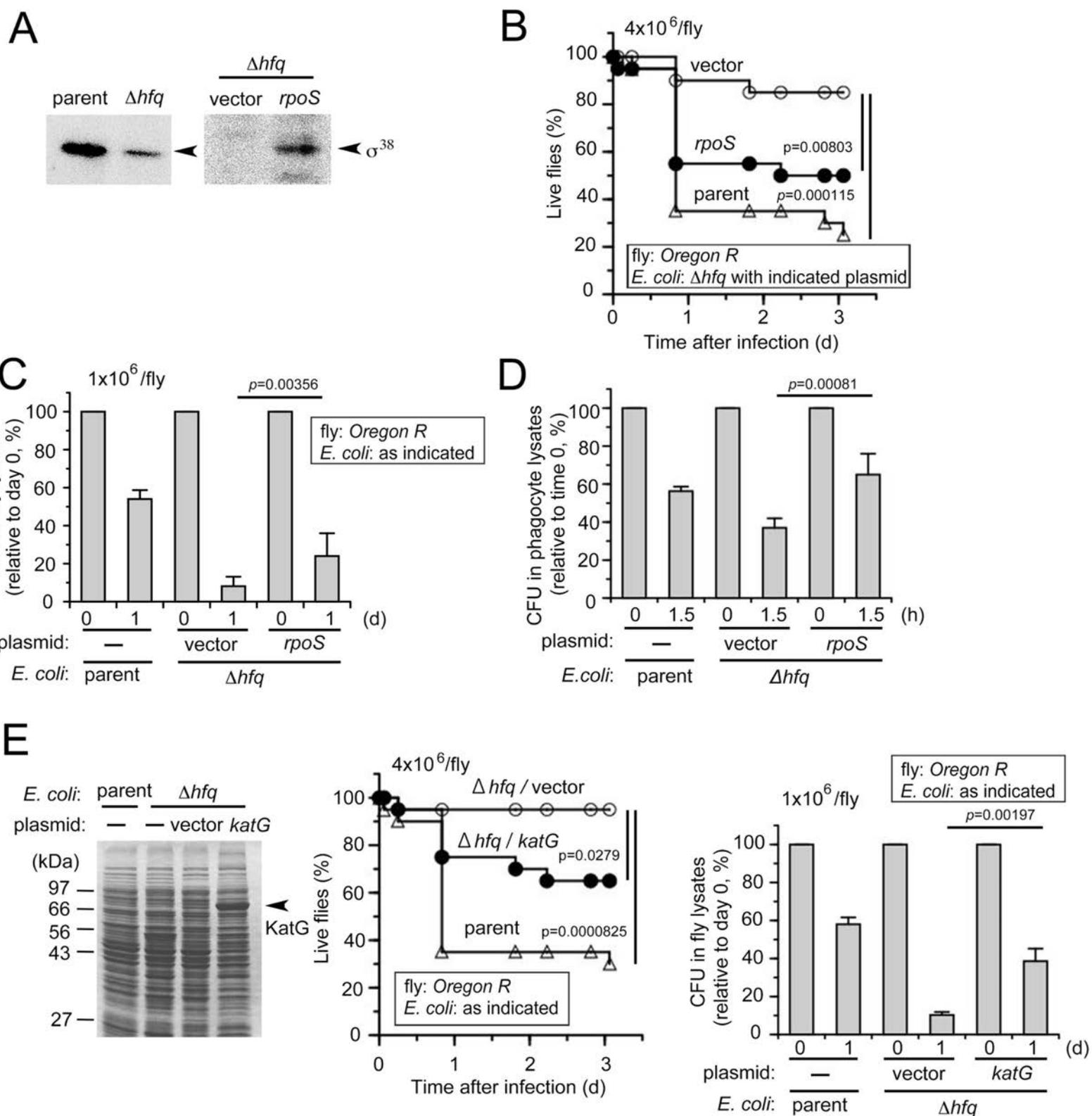
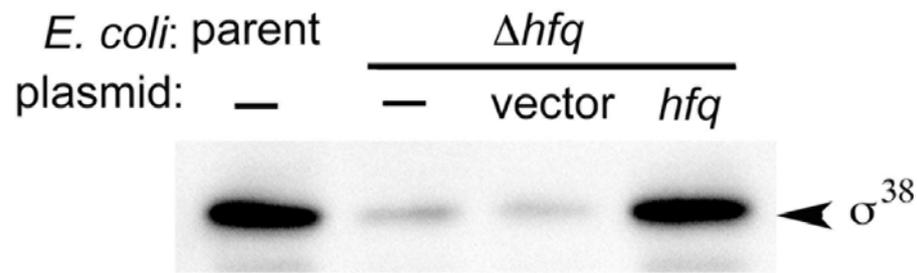
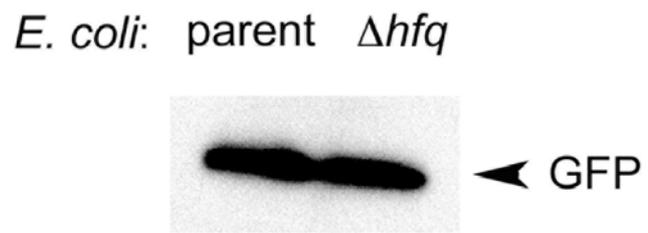
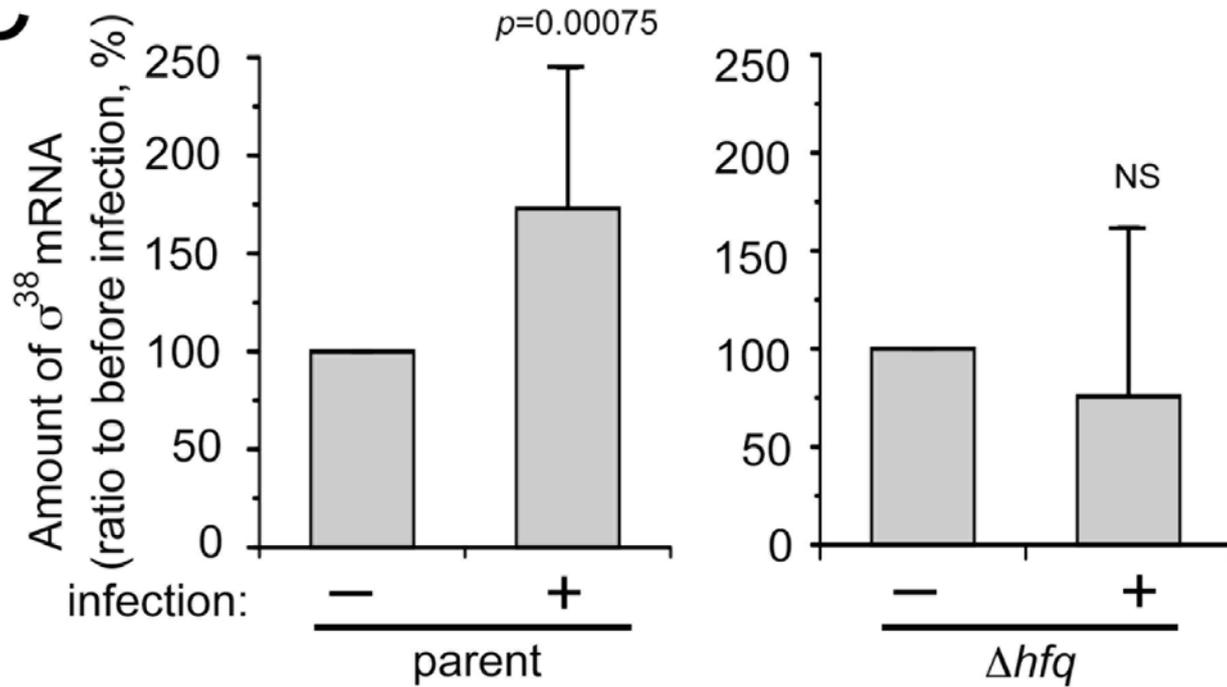
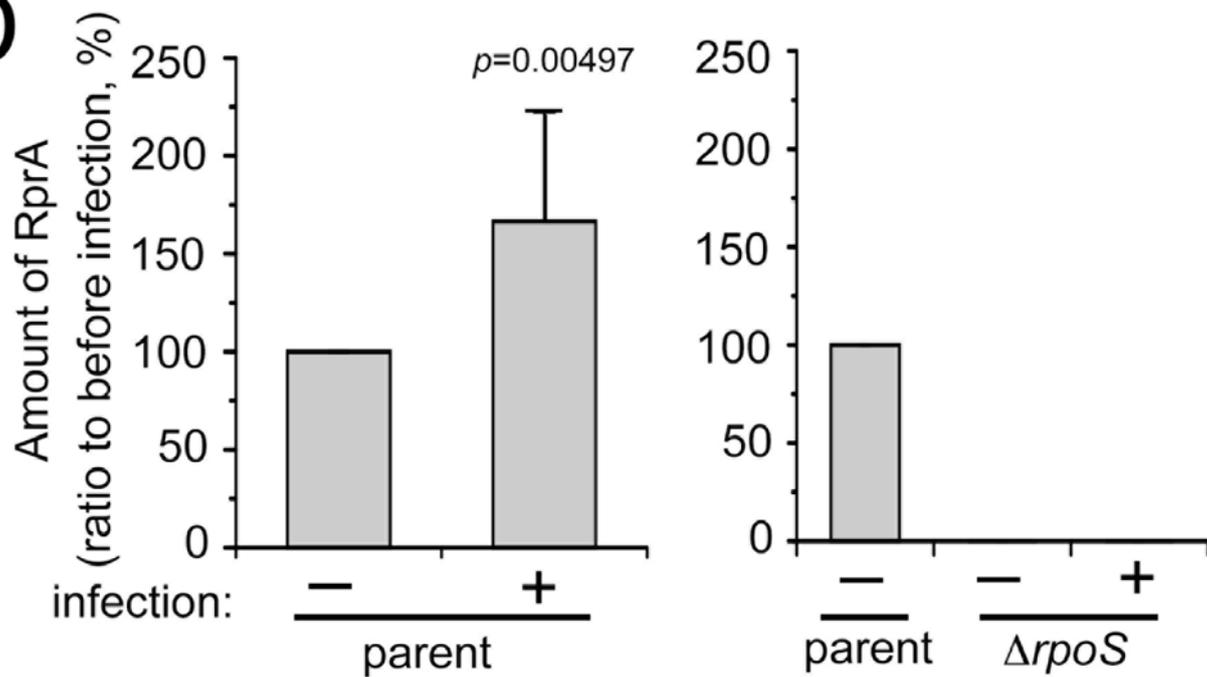
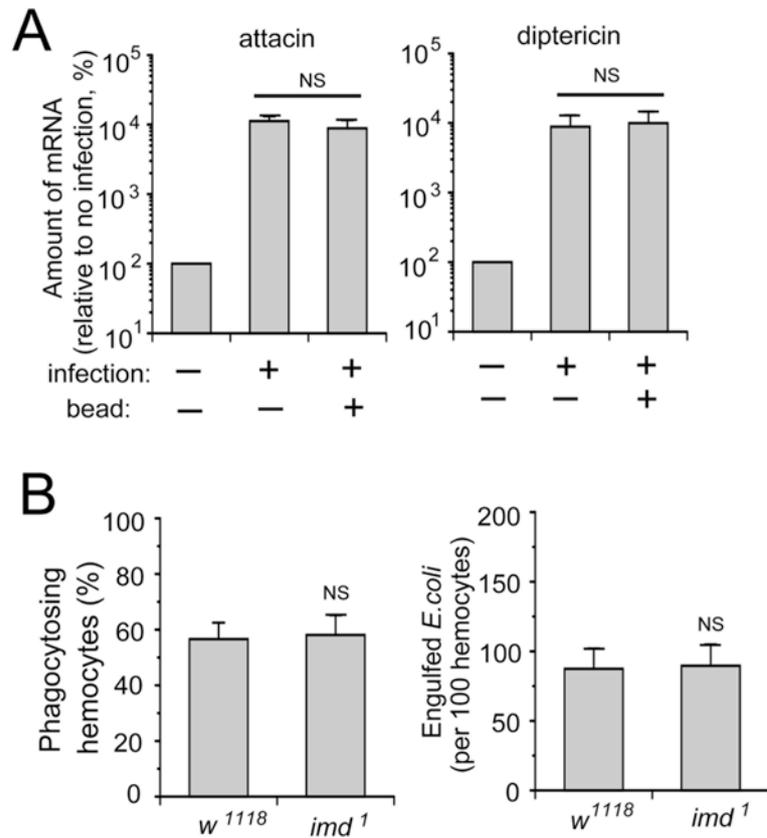


Fig. 6

A**B****C****D****Fig. 7**



Supplemental Figure 1

No relation between the production of antimicrobial peptide mRNA and the phagocytosis of *E. coli* by hemocytes. **(A)** Adult flies of *Oregon R* were injected with latex beads or PBS prior to the abdominal infection with the parental *E. coli* strain. After 3 h, RNA was prepared from those flies as well as the *E. coli* strain before infection, and subjected to RT-qPCR for the mRNAs of attacin, dipteracin, and Rp49. The values for the mRNAs of attacin and dipteracin were normalized with those of Rp49 mRNA, and the data from independent experiments are shown as the mean \pm SD after analysis by Student's *t* test. NS, difference not significant. The numbers of replicated experiments are: (in the analysis of attacin mRNA, from left to right) four, seven, and eight; (in the analysis of dipteracin mRNA, from left to right) five, three, and eight. **(B)** Hemocytes were isolated from third-instar larvae of the indicated fly lines and used as phagocytes in the phagocytosis reaction *in vitro* with the parental *E. coli* strain as the target. Data from three independent experiments are shown as the mean \pm SD after analysis by Student's *t* test. NS, difference not significant.