Bacteria Endosymbiont, *Wolbachia*, Promotes Parasitism of Parasitoid Wasp *Asobara japonica*

Shunsuke Furihata*, Makiko Hirata*, Hitoshi Matsumoto, Yoichi Hayakawa*

Department of Applied Biological Sciences, Saga University, Saga, Japan

*These authors contributed equally to this work.

**hayakayo@cc.saga-u.ac.jp**

**Abstract**

*Wolbachia* is the most widespread endosymbiotic bacterium that manipulates reproduction of its arthropod hosts to enhance its own spread throughout host populations. Infection with *Wolbachia* causes complete parthenogenetic reproduction in many Hymenoptera, producing only female offspring. The mechanism of such reproductive manipulation by *Wolbachia* has been extensively studied. However, the effects of *Wolbachia* symbiosis on behavioral traits of the hosts are scarcely investigated. The parasitoid wasp *Asobara japonica* is an ideal insect to investigate this because symbiotic and aposymbiotic strains are available: *Wolbachia*-infected Tokyo (TK) and noninfected Iriomote (IR) strains originally collected on the main island and southwest islands of Japan, respectively. We compared the oviposition behaviors of the two strains and found that TK strain females parasitized *Drosophila melanogaster* larvae more actively than the IR strain, especially during the first two days after eclosion. Removing *Wolbachia* from the TK strain wasps by treatment with tetracycline or rifampicin decreased their parasitism activity to the level of the IR strain. Morphological and behavioral analyses of both strain wasps showed that *Wolbachia* endosymbionts do not affect development of the host female reproductive tract and eggs, but do enhance host-searching ability of female wasps. These results suggest the possibility that *Wolbachia* endosymbionts may promote their diffusion and persistence in the host *A. japonica* population not only at least partly by parthenogenesis but also by enhancement of oviposition frequency of the host females.

**Introduction**

Many parasitoid wasps harbor endosymbiotic bacteria *Wolbachia*, a single bacterial lineage in the alpha-group of the Proteobacteria. *Wolbachia* is generally a facultative reproductive parasite in arthropods, and invades the host population by reproductive manipulations such as cytoplasmic incompatibility, male killing, feminization, or thelytokous parthenogenesis [1–5]. *Wolbachia*-induced parthenogenesis is most commonly found in haplodiploid organisms, such
as Hymenoptera [6–8]. In uninfected haplodiploid organisms, fertilized eggs develop into diploid daughters and unfertilized eggs develop into haploid sons, while infection with Wolbachia causes diploidization of the haploid eggs by alteration of meiotic and/or mitotic processes, resulting in the production of daughters from unfertilized eggs [9–11]. Since an asexual female produces only daughters, all of her offspring will contribute to the next generation. Therefore, an asexual population can grow faster than a sexual population and when they are in competition, the asexual population should outcompete the sexual one [12].

Aside from such reproductive effects of Wolbachia infection, recent findings in Diptera indicate that Wolbachia may also modify the host’s physiologies such as metabolism and immunity. In D. melanogaster, Wolbachia may play a role as a nutritional mutualist by affecting iron utilization by the hosts [13,14]. Wolbachia infection has also been shown to generate oxidative stress in one Aedes aegypti cell line, which reacts by the overexpression of host antioxidant genes [15]. This finding must be closely related with the fact that reactive oxygen species (ROS) are known to play a major role in immune response to pathogens [16,17]. Furthermore, Wolbachia is known to induce resistance against RNA viral infection in D. melanogaster and D. simulans [18,19], and against several arboviruses including dengue, yellow fever, and Chikungunya viruses in the mosquito A. aegypti, by priming the innate immune system [20–22].

Although increasing evidence is emerging on the phenotypic effects of Wolbachia infection on host reproduction and physiologies in non-Hymenopteran species, it is unknown whether it exerts similar effects on the physiologies of parasitoid wasps. Furthermore, it remains ambiguous whether Wolbachia causes other phenotypic effects such as behaviors of host insects, parasitoid wasps, in a particular symbiotic association in which Wolbachia affects parasitism processes through its effect on wasp oviposition. The objective of this study was to clarify the effect of Wolbachia on behavioral traits of the host female wasps. Here, we compared symbiotic and aposymbiotic parasitoid wasps in terms of parasitism activities of females using Asobara japonica because Wolbachia-infected Tokyo (TK) and noninfected Iriomote (IR) strains, which are originally collected on the main and southwest islands of Japan, respectively, are available [23–26]. Behavioral analyses demonstrated that symbiotic TK strain females parasitize D. melanogaster larvae significantly more actively than the aposymbiotic IR strain especially during an early adult stage. A series of analysis revealed that Wolbachia-induced parasitism activity of female wasps is primarily due to the enhancement of their host-searching ability.

Materials and Methods

Animals

Wolbachia-infected (Tokyo, TK) and uninfected (Iriomote, IR) strains of Asobara japonica were originally collected from Tokyo and Iriomote-jima island, respectively [27]. These strains were kindly provided by M.T. Kimura (Hokkaido University) and subsequently maintained under laboratory conditions as follows. Each generation, 5–10 female wasps were allowed to parasitize about 200–300 larvae of D. similans in glass vials (35 x 200 mm) at 23°C under 16L:8D regime. For experiments, yellow-white (yw) strain of D. melanogaster was used as a host [23,24].

Antibiotic treatment

Removal of Wolbachia was performed basically according to the procedure of Dedeine et al. [28] as follows. Parasitoid eggs and larvae were exposed to tetracycline or rifampicin through hemolymph of the hosts which had been fed Drosophila medium containing 2 mg/ml tetracycline or 1 mg/ml rifampicin for 24 h. After pupariation occurred, host pupae were taken into a
different vial and wasps emerged from hosts were collected. Removal of Wolbachia was confirmed by PCR analysis as follows.

**PCR, RT-PCR, and quantitative real-time PCR**

PCR analysis was conducted using genome DNA samples prepared from whole bodies of A. japonica wasps according to the previously described procedure [29]. RT-PCR was conducted essentially according to the previously described procedure [30] as follows. Total RNAs were prepared from test tissues of parasitoid wasps using TriPure Isolation reagent (Roche Applied Science, USA). First-strand cDNA was synthesized with oligo(dT)\textsubscript{12-18} primer using ReverTra Ace RT-PCR kit (Toyobo, Japan) according to the manufacturer’s protocol. The cDNAs for target genes were amplified with a specific primer pair indicated in Table 1. PCR was conducted under the following conditions: 25–35 cycles at 95°C for 30 s, 52–55°C for 30 s, and 72°C for 45 s.

Quantitative real-time PCR was carried out with the cDNAs in a 20 ml reaction volume of LightCycler Fast DNA Master SYBR Green I (Roche Applied Science, USA) using the LightCycler 1.2 instrument and software (Roche Applied Science) [31]. The PCR cycling conditions were denaturation at 95°C for 10 min, followed by 45 cycles of heating at 95°C for 10 s, annealing at 55°C for 5 s, and extension at 72°C for 15 s. Using the second derivative maximum method provided in the LightCycler software (version 3.5), a standard curve was generated by plotting the external standard concentration against threshold cycle. The software automatically calculated PCR product concentration for each tissue sample. All samples were analyzed in duplicate, and assay variation was typically within 10%. Data were normalized according to the expression level of rp49 determined in duplicate by reference to a serial dilution calibration curve.

**Table 1. List of PCR primers used in this study.**

| Primers | Forward | Reverse |
|---------|---------|---------|
| **rp49** | FW | AAAGGTATYGAYAACAGAGT |
| | RV | TATTCSTTCYCCYCARATCG |
| **Wsp** | FW | TGGTCCAATAAGTGATGAGAAAAC |
| | RV | AAAATTAAACGCTACTCCA |
| **WP1** | FW | TTGTAAGCCTGTATGTTAACT |
| | RV | GAATAGGTATGATTTTCATGT |
| **Orco** | FW | AAYATCKCACAAGCCATCC |
| | RV | CTGGTGGMACKACGGTGC |
| **Common RACE Primers** | Qt | CCAGTGACGAGCAGTAGGAGGACTCGAGCTCAAGGCTTTTTTTTTTTTTTTTTTTT |
| | Qo | CCAGTGACGAGCAGTGACG |
| | Qi | GAGGACTCGAGCTCAAGC |
| **Primers for 5’-RACE** | Aj-Orco-R1 | AAGAGCGGCAGCCGAGCAGAGAGAAGC |
| | Aj-Orco-R2 | TGATTCCGGTCCAGCAGAAGT |
| **Primers for 3’-RACE** | Aj-Orco-F1 | CCATCAATAAGAAGCTC |
| | Aj-Orco-F2 | GGTTTCCAAATCTCTGGC |
Activity of parasitism

To measure parasitism activity of test wasp females, one female of TK (or antibiotic-treated TK strains) or one pair of IR strain was put in a 30 ml glass vial with Drosophila medium containing host D. melanogaster larvae. IR strain females can mate within a few hours after eclosion. Therefore, we used only mated IR females (even day 0) in all experiments to eliminate a possible effect of mating on parasitism behavior. Furthermore, we have also demonstrated that the presence of males did not cause any change of oviposition rates of IR strain wasps. Detailed conditions were indicated in each experiment.

Y-tube olfactory assay

The olfactory responses of A. japonica female wasps to odors were tested in a Y-tube olfactometer. The olfactometer consisted of a glass Y-tube (base 4.5 cm long; Y-arms each 4.5 cm long; 10 mm inner diameter). The Y-tube apparatus was modified after the design of Carroll et al. [32]. The base of the Y-tube was connected to a Teflon tube of similar size that was attached directly to the vacuum source. Odor sources, 50 μg of Drosophila medium with or without D. melanogaster larvae, were placed at either end of the Y-arms and the odors were extracted through the base arm at a flow rate of 2L/min by a vacuum pump to ensure a steady flow. Ice-anesthetized test wasps were introduced individually by disconnecting the Y-tube at its base and allowing the wasp to move into the olfactometer. After wasp recovered from anesthesia, the tube was reconnected to reestablish the airflow from the odor sources through the arms and out at the base towards the vacuum pump. A choice and consumed time were recorded when a test wasp reached either end of Y-arms. Test wasps were recorded as ‘not moved’ when they remained in the base arm. Each test was terminated after 30 min from the initiation of airflow. To avoid positional bias, odor chambers were switched after every replicate.

Morphological analysis of lateral oviduct

Paired lateral oviducts were dissected from A. japonica female wasps and the magnification images were taken with a microscope with camera attachment. Sizes (length and width) of the tissues were measured using Image J National Institute of Health, USA. The number of eggs harbored in the oviducts was also counted under a microscope.

Sequencing of A. japonica Orco gene

RT-PCR was carried out using cDNA prepared from antennae of day 0 females and degenerate primers designed from conserved sequences of Apis mellifera odorant receptor 2 (Or2), transcript variant X1, mRNA (gi 571501583) and Microplitis demolitor odorant receptor coreceptor (LOC103571567), mRNA (gi 665785588). Amplified fragments of A. japonica Orco gene were sequenced by using BigDye Terminator v3.1 (Applied Biosystems, USA) with 3130 Genetic Analyzer (Applied Biosystems, USA) [33]. Alignment of sequences was carried out on BioEdit Sequence Alignment Editor 7.0.9.0. (Ibis Biosciences, USA).

To obtain full-length cDNA, 5’ rapid amplification of cDNA ends (5’RACE) was performed using 5’ RACE kit (Invitrogen, Carlsbad, CA) as described previously [30]. All identified cDNA sequences were subjected to computer-assisted sequence analysis using GENETYX-MAC Ver. 13.1.7 (Software Development Co., Tokyo, Japan).

Statistical analyses

For comparison of parasitism activity, development of ovary and preference for host odor, Tukey’s HSD tests were carried out. As a result of normality test, Shapiro-Wilk test, it was
found that each data set does not deviate from the normality. All statistical analyses were performed using JMP 9.0.2 (SAS Institute).

Results
Frequencies of parasitism by TK and IR strains
We first confirmed that the TK strain flies harbored *Wolbachia pipientis* but the IR strain did not by PCR with two primer sets specific to *W. pipientis* genes (Fig 1A). To roughly compare parasitism activity by both strains, one young female of each strain was put into a 30 ml glass tube containing a *D. melanogaster* colony with *Drosophila* medium. All wasps were removed 15 h later, and *D. melanogaster* colonies were maintained until all wasp offspring emerged from the hosts. The number of the TK strain offspring was higher than those of the IR strain (Fig 1B). Although we did not count the number of eggs oviposited by both strain females, these results were interpreted to imply a possibility that TK strain female wasps with *Wolbachia* possess higher parasitic activity than IR strain animals.

Effect of *Wolbachia* on oviposition rates of host wasps
To seek detailed differences in parasitism behavior between TK and IR strains, we counted the number of eggs in *D. melanogaster* larvae that had been oviposited by both strain females during first three days after eclosion (Fig 2A). On day 0, the number of eggs oviposited by TK strain females was 2~3 times higher than that of IR strains. The difference in oviposition rates between both strains decreased on day 1 and disappeared on day 2 (Fig 2A). These data suggested the possibility that *Wolbachia* enhances parasitism behavior especially during early adult stages of the wasp. To examine this, we eliminated *Wolbachia* by using *D. melanogaster* larvae fed an antibiotic (tetracycline (tet) or rifampicin (rif)) as the wasp’s hosts. This treatment succeeded in the elimination of *Wolbachia* from TK strain wasps (Fig 2B). TK strain female wasps without *Wolbachia* showed significantly lower oviposition rates on day 0 compared to the normal TK strain (Fig 2C). Furthermore, tet-treated IR strain wasps did not show any significant change in their oviposition rates, indicating that tet-treatment itself did not affect on the parasitism behavior of wasps (Fig 2C). Based on these results, it is reasonable to assume that the active parasitism in young adults of the TK strain wasps is due to the presence of symbiont *Wolbachia*.

Effect of *Wolbachia* on development of host wasp reproductive tissues
To examine whether *Wolbachia* affects development of the reproductive tracts and eggs in female hosts, we dissected the lateral oviducts and measured their sizes. Neither the widths nor the lengths of TK, IR, and tet-treated TK (tet-TK) strains were different even on day 0 and day 2 of adult stages (Fig 3A and 3B). Furthermore, the number of eggs in the oviducts was not different among the three strains. Therefore, *Wolbachia* symbiosis did not affect reproductive maturation rates of the parasitoid wasps.

Effect of *Wolbachia* on parasitism behavior of host wasps
To find the main cause of high oviposition rates of young TK strain female wasps, we compared behavioral traits of both strains of female wasps. First, we measured the frequency of ovipositor insertion into an artificial diet containing host larvae by female wasps of both strains. TK strain females inserted their ovipositors much more frequently than IR strain and tet-TK strain females did on day 0, but afterwards, the frequencies of ovipositor insertion of IR and tet-TK strains increased to a level almost equivalent to that of the TK strain on day 2 (Fig 4A). The number of eggs oviposited by the three strains of females also showed the same tendencies...
Fig 1. (A) RT-PCR of *Wolbachia* specific genes, *Wolbachia* surface protein gene (*wsp*) and *Wolbachia* protein 1 gene (*WP1*), and *A. japonica* ribosomal protein 49 gene (*rp49*) in whole bodies of *A. japonica* wasps. (B) The number of offspring wasps emerged from parasitized female. One female wasp randomly selected from day 0 to day 2 adults of TK or IR strain colony was put in a glass vials (35 x 120 mm) containing 200–300 *D. melanogaster* larvae and allowed to freely parasitize for 15 h. In the case of IR strain, mated females were used. Each value represents the mean ± S.D. for 16 independent determinations. Ten wasps of each strain were used for each determination. P = 0.056 (Tukey’s HSD).

doi:10.1371/journal.pone.0140914.g001
as the ovipositor insertion frequencies: the TK strain laid eggs much more actively than IR and tet-TK strains did on day 0, but the difference in oviposition activities between TK and other two strains was decreased on day 2 (Fig 4B).

Recognition of host *Drosophila* larvae by three strains of female wasps was determined in a Y-tube olfactometer. TK strain females were attracted to the smell of the host larvae much more strongly than IR and tet-TK strains were on day 0, but the behavioral differences between
TK and other strains were reduced on day 2 (Fig 5A). Furthermore, TK strain females reached an artificial diet containing the host larvae much more quickly than IR and tet-TK strains did on day 0 (Fig 5B). The difference in host searching behavior between TK and IR strains was preliminarily demonstrated not to be changed when we had switched the host Drosophila species from D. similans to D. melanogaster for maintaining wasp colonies.

**Effect of Wolbachia on olfactory receptor gene expression in host wasps**

Finally, we tested whether the behavioral differences between TK and IR (or tet-TK) strains were due to a difference in olfactory abilities in these strains. Since the antennae have been reported to be the sensorial organs used by female parasitoid wasps to locate their hosts, we cut off the antennae of both strains and observed oviposition behaviors of female wasps. This treatment strongly destroyed their host searching ability and removed the difference in TK and IR strain females, (Fig 6A).

In Drosophila, odorant receptor co-receptor (Orco, Or83b) is expressed in almost all olfactory receptor neurons and found to form heteromers with other odorant receptor proteins [34], indicating that Orco is essential for Drosophila to sense all odorants. Since it also has been demonstrated that other insect species, such as silkworm and honeybee, possess an Orco family protein [35], we made a degenerate Orco primer set in the conserved region of reported genes and sequenced by RT-PCR (S1 Fig). The full length of Orco genes was amplified by repeated 5′- and 3′-RACE reactions using cDNAs prepared from both strain wasp antennae. The homologies of the nucleotide sequences and the deduced amino acid sequences of both strain Orco genes were found to be 99.5% and 99.8%, respectively, indicating that the structures of Orco genes in both strains are almost identical (S2 and S3 Figs). We then performed quantitative real-time PCR by using cDNA expressed in the antennae. Expression levels of the A. japonica Orco orthologous gene were unexpectedly higher in IR than those in TK strains both on day 0 and day 2. Furthermore, the low gene expression in TK strain was not elevated by tet-treatment, indicating that the better host searching ability of TK strain cannot be simply explained by expression levels of Orco (Fig 6B).

**Discussion**

All populations of A. japonica collected on the main islands of Japan have been reported to harbor Wolbachia. Although uninfected populations were exceptionally found on the small southern islands, the total infection of the main island populations suggests that Wolbachia infection has an ecological advantage for this wasp species to extend its habitat. Wolbachia-induced parthenogenesis is most generally thought to be beneficial for host insects to increase their population; in the case of haplodiploid organisms such as wasps, diploidization of the haploid eggs caused by Wolbachia produces daughters from unfertilized eggs [36]. Such a reproductive manipulation allows Wolbachia to increase infected wasp populations, which increases their transmission [37].

In this study, we sought additional positive effects of Wolbachia infection for A. japonica wasps other than the reproductive manipulation, and found that the oviposition rates of Wolbachia-infected (TK strain) females were clearly higher than those of uninfected (IR strain) females on day 0 of the adult stage. Therefore, Wolbachia infection enhances the oviposition behavior of A. japonica wasps during an early adult stage. This is certainly beneficial for the
Wolbachia-infected stain because life spans of both strains are not significantly different in glass culture tubes containing host larvae and their artificial diet: life spans of TK and IR strains.

Fig 4. Frequencies of ovipositor insertion behavior (A) and oviposition by wasp (B) of A. japonica TK strain, IR strain, or tet-TK strain. In the case of IR strain, mated females were used. Each value represents the mean ± S.D. for 6–7 independent determinations. Significant difference indicated by Tukey’s HSD (*P<0.05, **P<0.01, ***P<0.001). Two female wasps of each strain were put in an assay plate containing twenty D. melanogaster larvae with their medium, and each test wasp was allowed to parasitize for 1 h during each determination.

doi:10.1371/journal.pone.0140914.g004

Wolbachia-infected stain because life spans of both strains are not significantly different in glass culture tubes containing host larvae and their artificial diet: life spans of TK and IR strains.
Fig 5. (A) Rate of *A. japonica* TK, IR, and tet-TK strain wasps attracted to *Drosophila* medium with *Drosophila* larvae. Each value represents the mean of tested wasps; 11 (day 0) and 21 (day 2) TK, 7 (day 0) and 9 (day 2) IR, and 7 (day 0) and 8 (day 2) tet-TK strains were used for the determination. In the case of IR strain, mated females were used. Although TK and IR strain wasp colonies were
were calculated to be 4.33 ± 1.22 days (n = 9) and 4.60 ± 2.22 (n = 10), respectively, by our preliminary experiments. Furthermore, we observed that A. japonica females of TK strain could parasitize soon after the emergence from the host in the field as well as in the laboratory. Given that their average life span must be shorter under natural conditions than in the culture tube due to the presence of predators, infection, and various physical accidents, it is reasonable to think that this behavioral trait must become more beneficial for wasps under natural conditions. To reveal the mechanism underlying the elevation of the oviposition rates of young Wolbachia-infected females, we compared the sizes of the reproductive tracts and number of eggs in the oviducts of TK strain and IR strain young wasps. Neither the size of the lateral oviduct nor the number of eggs was different between both strains of wasps. However, we found that the oviposition activity of the TK strain was significantly higher than that of the IR strain during the young adult stage: the numbers of oviposited eggs in host larvae, as well as frequency of ovipositor insertion to the artificial diet containing host larvae, were significantly higher in the TK strain compared to the IR strain on day 0. This difference was demonstrated to be primarily due to Wolbachia-mediated enhancement of host-searching ability of the TK strain. The better host searching ability of TK strain was verified irrespective of host Drosophila species: switching the host from D. melanogaster to D. simulans did not change the behavioral difference between TK and IR strain wasps (data not shown).

It has been reported that the sensorial organs used by female parasitoid wasps to locate and evaluate their hosts are mostly present on their antennae where olfactory chemosensory sensilla have been described. Many olfactory receptors are present in the sensilla of antennae. Although they are highly diverse in insect species, the Orco family is exceptional: members share a highly conserved gene sequence among different species and play an important role in regulating insect behavior [38]. Therefore, we examined whether Wolbachia infection affects expression of A. japonica Orco. The expression levels of the ortholog of the Drosophila Orco gene were unexpectedly higher in IR than in TK strain wasps, suggesting at least that the high host-searching ability of TK strain is not due to a high expression level of Orco under the effect of Wolbachia infection. Although it is known that Orco plays an essential general roles in olfaction of Drosophila [39], there has not been reported the precise relationship between Orco expression levels and olfactory sensitivities. The present observation that tet-treatment of TK strain did not elevate the Orco expression level negated one possibility that extra-expression of Orco negatively affects olfaction or expression level of Orco does not affect any substantial olfactory ability. However, this result is not surprising because it is reasonable to consider that host-searching as well as oviposition behaviors cannot be controlled in a simple way by a sole gene. The fact that cutting off the antennae of both strain females drastically decreased their host searching abilities (Fig 6A) could be explained by postulating that both strain females cannot normally move or act without olfactory stimuli through the sensorial organ. Further investigation is required to clarify the relationship between expression levels of olfactory receptor genes such as Orco in the antenna and the olfactory sensitivities.

Pannebakker et al. [40] recently compared transcriptomes of the resting and ovipositing female parasitoid Nasonia vitripennis using a “DeepSAGE” gene expression approach. They identified 332 tags that were significantly differentially expressed between the two treatments, and found that nine of the most abundant differentially expressed tags showed greater expression in
Wolbachia Promotes Parasitism of Its Host Parasitoid Wasp

A

Rate of wasps attracted to host larvae

|          | Intact | Removed | Intact | Removed |
|----------|--------|---------|--------|---------|
| TK       | 100    | 0       | 0      | 0       |
| IR       | 75     | 0       | 75     | 0       |

B

Relative expression level (Drc/osp/p4/9)

|        | Day 0 | Day 2 |
|--------|-------|-------|
| TK     | 0.01  | 0.02  |
| tet-TK | 0.03  | 0.02  |
| IR     | 0.02  | 0.02  |
ovipositing females, including the genes purity-of essence (poe) (associated with behavioral phenotypes in Drosophila melanogaster) and glucose dehydrogenase. The poe protein is an evolutionarily conserved, large membrane protein containing a motif that affects behavior and synaptic transmission of D. melanogaster. Since it has been reported that two mutants of poe in Drosophila cause both increased nervous excitability and reduced motor function [41] and mutants also affect peripheral nerve morphology [42], it is reasonable to assume that Wolbachia infection might cause enhancement of such transcription factors in the A. japonica female wasp, resulting in the change of their oviposition behavior. Furthermore, approximately three-quarters of the changes involve greater expression in resting females, and enrichment analysis suggests that these down-regulated genes are more likely to be involved in various metabolic processes than expected by chance, suggesting that as female wasps move from resting to ovipositing, aspects of their metabolism are down-regulated, focusing gene expression on other processes [40]. These results can be interpreted to indicate that oviposition behavior is associated with changes in a variety of gene expressions in female whole bodies. Another recent comparative analysis of gene expression in ovaries between Wolbachia-infected and the uninfected parasitoid wasp Asobara tabida, a close species to A. japonica, showed that Wolbachia might interfere with numerous biological processes such as oogenesis, programmed cell death, and immunity [43]. These reports together with our unexpected result on A. japonica Orco expression imply that Wolbachia-induced enhancement of host-searching ability in the TK strain is not driven by a change of a single olfactory receptor gene that directly controls sensing of the host smell.

Supporting Information

S1 Fig. Comparison of nucleotide sequences of 5 insect species odorant receptor coreceptor genes (Orco). The fragment of Orco cDNA was prepared from antennae of A. japonica female wasps. The base sequence was analyzed as described in Materials and Methods. (PDF)

S2 Fig. Comparison of nucleotide sequences of TK and IR strain odorant receptor coreceptor genes (Orco). Whole Orco cDNAs were prepared from antennae of both strain female wasps by RT-PCR. Start and stop codons were red-colored. (PDF)

S3 Fig. Comparison of deduced amino acid sequences of TK and IR strain odorant receptor coreceptor genes (Orco). Whole Orco cDNAs were prepared from antennae of both strain female wasps by RT-PCR. Only one amino acid, Lue402, is replaced with Phe. (PDF)

Author Contributions

Conceived and designed the experiments: YH SF. Performed the experiments: SF MH HM YH. Analyzed the data: SF MH HM YH. Contributed reagents/materials/analysis tools: SF MH HM. Wrote the paper: YH.
References

1. Bouchon D, Rigaud T, Juchault P. Evidence for widespread Wolbachia infection in isopod crustaceans: molecular identification and host feminization. Proc Biol Sci. 1998; 265: 1081–1090. PMID: 9864374

2. Huigens ME, Luck RF, Klaassen RH, Maas MF, Timmermans MJ, Stouthamer R. Infectious parthenogenesis. Nature. 2000; 405: 178–179. PMID: 10821272

3. Jiggins FM, Hurst GD. The butterfly Danaus chrysippus is infected by a male-killing Spiroplasma bacterium. Parasitology. 2000; 120: 439–446. PMID: 10840973

4. Kageyama D, Nishimura G, Hoshizaki S, Ishikawa Y. Feminizing Wolbachia in an insect, Ostrinia furnacalis (Lepidoptera: Crambidae). Hereditas (Edinb). 2002; 88: 444–449.

5. Negri I, Pellecchia M, Mazzoglio PJ, Patetta A, Alma A. Feminizing Wolbachia in Zyginaida pullula (Insecta, Hemiptera), a leafhopper with an XX/X0 sex-determination system. Proc Biol Sci. 2006; 273: 2409–2416. PMID: 16928646

6. Stouthamer R, Breeuwer JA, Hurst GD. Wolbachia pipientis: microbial manipulator of arthropod reproduction. Annu Rev Microbiol. 1999; 53: 71–102. PMID: 10547866

7. Werren JH. Biology of Wolbachia. Annu Rev Entomol. 1997; 42: 587–609. PMID: 15012323

8. Werren JH, Baldo L, Clark ME. Wolbachia: master manipulators of invertebrate biology. Nat Rev Microbiol. 2006; 6: 741–751. doi: 10.1038/nrmicro1969 PMID: 18794912

9. Stouthamer R, Kazmer DJ. Cytogenetics of microbe-associated parthenogenesis and its consequences for gene flow in Trichogramma wasps. Heredity. 1994; 73: 317–327.

10. Stouthamer R, Luck RF, Hamilton WD. Antibiotics cause parthenogenetic Trichogramma (Hymenoptera/Trichogrammatidae) to revert to sex. Proc Natl Acad Sci U S A. 1990; 87: 2424–2427. PMID: 11607070

11. Weeks AR, Breeuwer JA. Wolbachia-induced parthenogenesis in a genus of phytophagous mites. Proc Biol Sci. 2001; 268: 2245–2251. PMID: 11674872

12. Reumer BM, van Alphen JJM, Kraaijeveld K. Ecology, Wolbachia infection frequency and mode of reproduction in the parasitoid wasp Tetrastichus coerules (Hymenoptera: Eulophidae). Mol Ecol. 2010; 19: 1733–1744. doi: 10.1111/j.1365-294X.2010.04599.x PMID: 20345674

13. Brownlie JC, Cass BN, Riegler M, Witsenburg JJ, Iturbe-Ormaetxe I, McGraw EA. Evidence for metabolic provisioning by a common invertebrate endosymbiont, Wolbachia pipientis, during periods of nutritional stress. PLoS Pathog. 2009; 5: e1000368. doi: 10.1371/journal.ppat.1000368 PMID: 19343208

14. Kremer N, Voronin D, Charif D, Mavingui P, Mollereau B, Vavre F. Wolbachia interferes with ferritin expression and iron metabolism in insects. PLoS Pathog. 2009; 5: e1000630. doi: 10.1371/journal.ppat.1000630 PMID: 19851452

15. Brennan LJ, Keddie BA, Braig HR, Harris HL. The endosymbiont Wolbachia pipientis induces the expression of host antioxidant proteins in an Aedes albopictus cell line. PLoS One. 2008; 3: e2083. doi: 10.1371/journal.pone.0002083 PMID: 18461124

16. Beutler B. Innate immunity: an overview. Mol Immunol. 2004; 40: 845–859. PMID: 14698223

17. Molina-Cruz A, DeJong RJ, Charles B, Gupta L, Kumar S, Jaramillo-Gutierrez G, et al. Reactive oxygen species modulate Anopheles gambiae immunity against bacteria and Plasmodium. J Biol Chem. 2008; 283: 3217–3223. PMID: 18065421

18. Osborne SE, Leong YS, O'Neill SL, Johnson KN. Variation in antiviral protection mediated by different Wolbachia strains in Drosophila simulans. PLoS Pathog. 2009; 5: e1000656. doi: 10.1371/journal.ppat.1000656 PMID: 19911047

19. Teixeira L, Ferreira A, Ashburner M. The bacterial symbiont Wolbachia induces resistance to RNA viral infections in Drosophila melanogaster. PLoS Biol. 2008; 6: e2. doi: 10.1371/journal.pbio.1000002 PMID: 19223034

20. Biao G, Xu Y, Yu P, Xie Y, Xi Z. The endosymbiotic bacterium Wolbachia induces resistance to dengue virus in Aedes aegypti. PLoS Pathog. 2010; 6: e1000833. doi: 10.1371/journal.ppat.1000833 PMID: 20368968

21. Moreira LA, Iturbe-Ormaetxe I, Jeffery JA, Li Q, Pyke AT, Hedges LM, et al. A Wolbachia symbiont in Aedes aegypti limits infection with dengue, Chikungunya, and Plasmodium. Cell. 2009; 139: 1268–1278. doi: 10.1016/j.cell.2009.11.042 PMID: 20064373

22. van den Hurk AF, Hall-Mendelin S, Pyke AT, Frentiu FD, McElroy K, Day A, et al. Impact of Wolbachia on infection with chikungunya and yellow fever viruses in the mosquito vector Aedes aegypti. PLoS Negl Trop. 2012; 6: e1892.
23. Furihata SX, Kimura MT. Effects of Asobara japonica venom on larval survival of host and nonhost Drosophila species. Physiol Entomol. 2009; 34: 292–295.

24. Furihata SX, Matsumoto H, Kimura MT, Hayakawa Y. Venom components of Asobara japonica impair cellular immune responses of host drosophila melanogaster. Arch Insect Biochem Physiol. 2013; 83: 86–100. doi: 10.1002/arch.21093 PMID: 23606512

25. Kremer N, Charif D, Henri H, Bataille M, Prevost G, Kraaijeveld K, et al. A new case of Wolbachia dependence in the genus Asobara: evidence for parthenogenesis induction in Asobara japonica. Heredity (Edinb). 2009; 103: 248–256.

26. Reumer BM, van Alphen JJ, Kraaijeveld K. Occasional males in parthenogenetic populations of Asobara japonica (Hymenoptera: Braconidae): low Wolbachia titer or incomplete coadaptation? Heredity (Edinb). 2012; 108: 341–346.

27. Mitsu H, Van Achterberg K, Nordlander G, Kimura MT. Geographical distributions and host associations of larval parasitoids of frugivorous Drosophilidae in Japan. J Nat Hist. 2007; 41: 1731–1738.

28. Dedeine F, Vavre F, Fleury F, Loppin B, Hochberg ME, Bouletreau M. Removing symbiotic Wolbachia bacteria specifically inhibits oogenesis in a parasitic wasp. Proc Natl Acad Sci U S A. 2001; 98: 6247–6252. PMID: 11353833

29. Hayakawa Y, Noguchi H. Growth-blocking peptide expressed in the insect nervous system. Cloning and functional characterization. Eur J Biochem. 1998; 253: 810–816. PMID: 9654083

30. Tsuzuki S, Ochiai M, Matsumoto H, Kurata S, Ohsishi A, Hayakawa Y. Drosophila growth-blocking peptide-like factor mediates acute immune reactions during infectious and non-infectious stress. Sci Rep. 2012; 2: 210. doi: 10.1038/srep00210 PMID: 22355724

31. Tsuzuki S, Matsumoto H, Furihata S, Ryuda M, Tanaka H, Sung EJ, et al. Switching between humoral and cellular immune responses in Drosophila is guided by the cytokine GBP. Nat Commun. 2014; 5: 4628. doi: 10.1038/ncomms5628 PMID: 25130174

32. Carroll MJ, Schmelz EA, Meagher RL, Teal PE. Attraction of Spodoptera frugiperda larvae to volatiles from herbivore-damaged maize seedlings. J Chem Ecol. 2006; 32: 1911–1924. PMID: 16902828

33. Oda Y, Matsumoto H, Kurakake M, Ochiai M, Ohnishi A, Hayakawa Y. Adaptor protein is essential for insect cytokine signaling in hemocytes. Proc Natl Acad Sci U S A. 2010; 107: 15862–15867. doi: 10.1073/pnas.1003785107 PMID: 20798052

34. Vosshall LB, Amrein H, Morozov PS, Rzhetsky A, Axel R. A spatial map of olfactory receptor expression in the Drosophila antenna. Cell. 1999; 96: 725–736. PMID: 10089887

35. Nakagawa T, Sakurai T, Nishioka T, Touhara K. Insect sex-pheromone signals mediated by specific combinations of olfactory receptors. Science. 2005; 307: 1638–1642. doi: 10.1126/science.1107361 PMID: 1592016

36. Pannebakker BA, Pijnacker LP, Zwaan BJ, Beukeboom LW. Cytology of Wolbachia-induced parthenogenesis in Leptopilina clavipes (Hymenoptera: Figitidae). Genome. 2004; 47: 299–303. PMID: 15060582

37. Serbus LR, Casper-Lindley C, Landmann F, Sullivan W. The genetics and cell biology of Wolbachia-host interactions. Annu Rev Genet. 2008; 42: 683–707. doi: 10.1146/annurev.genet.41.110306.130354 PMID: 18713031

38. Pitts RJ, Fox AN, Zwiebel LJ. A highly conserved candidate chemoreceptor expressed in both olfactory and gustatory tissues in the malaria vector Anopheles gambiae. Proc Natl Acad Sci U S A. 2004; 101: 5058–5063. PMID: 15037749

39. Larsson MC, Domingos AI, Jones WD, Chiappe ME, Amrein H, Vosshall LB. Or83b encodes a broadly expressed odorant receptor essential for Drosophila olfaction. Neuron. 2004; 47: 299–303. PMID: 15339651

40. Pannebakker BA, Trivedi U, Blaxter MA, Watt R, Shuker DM. The transcriptomic basis of oviposition behaviour in the parasitoid wasp Nasonia vitripennis. PLoS One. 2013; 8: e68608. doi: 10.1371/journal.pone.0068608 PMID: 23894324

41. Richards S, Hillman T, Stern M. Mutations in the Drosophila pushover gene confer increased neuronal excitability and spontaneous synaptic vesicle fusion. Genetics. 1996; 142: 1215–1223. PMID: 8446899

42. Yager J, Richards S, Hekmat-Scafe DS, Hurd DD, Sundaresan V, Capretta DR, et al. Control of Drosophila perineural glial growth by interacting neurotransmitter-mediated signaling pathways. Proc Natl Acad Sci U S A. 2001; 98: 10445–10450. PMID: 11517334

43. Kremer N, Delphine C, Helene H, Gavory F, Wincker P, Patrick M, et al. Influence of Wolbachia on host gene expression in an obligatory symbiosis. BMC Microbiology. 2012; 12: S7. doi: 10.1186/1471-2180-12-S1-S7 PMID: 22376153