HOST MICROBE INTERACTIONS

Symbiotic Fungal Flora in Leaf Galls Induced by *Illiciomyia yukawai* (Diptera: Cecidomyiidae) and in Its Mycangia

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Abstract We investigated the association between a gall midge, Illiciomyia yukawai, and its symbiotic fungi on Japanese star anise, Illicium anisatum. The number of fungal species isolated from the galls increased with development of the galls, whereas those from the leaves showed a different trend. Botryosphaeria dothidea was dominant in the galls from June to October, and after that Phomopsis sp. 1, Collectrichum sp., and Pestalotiopsis sp. became dominant. Although B. dothidea was not isolated from the leaves, it was detected from mycangia (abdominal sternite VII) of egg-laying adults at a high isolation frequency (>90%). However, B. dothidea was not isolated from mycangia of adults emerging from galls that were enclosed by plastic bags. This indicates that I. yukawai is closely associated with B. dothidea and that its newly emerged adults do not take the fungus into mycangia

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T. Kubono e-mail: kubono@ffpri.affrc.go.jp directly from the galls where they had developed. Also, the fungus from the fungal layers of ambrosia galls has less ability to propagate on artificial media despite the presence of its mycelial mass in mature galls.

Introduction

Insect-fungal symbioses exist widely in nature, and their mechanisms and interactions have been of considerable interest in ecology. For example, ambrosia beetles have specific fungi in their mycangia, which are structures specialized to carry fungal spores, and their larvae mainly feed on the fungi which are introduced into wood from the mycangia [3, 18]. Wood wasps inoculate *Amylostereum* fungi into wood with their eggs. Their larvae can make use of cellulose decomposed by digestive enzymes which are derived from their fungal symbionts [20]. Fungus-growing ants cultivate basidiomycetous fungi as a main food source in their nest (fungus garden) into which they carry various substrates [8, 21, 24].

"Ambrosia galls" are induced by gall midges (especially species in the tribes Asphondyliini, Lasiopterini, and Alycaulini, [28]) and are characterized by endomycetophagous larval feeding [4, 22]. Ambrosia galls have a fungal layer instead of nutrient plant tissue which is usually present in galls [40]. The gall midge larvae are thought to be dependent on the fungal mycelium growing on the inner surface of the galls as a food source [4, 16]. Female adults have mycangia, as in ambrosia beetles and wood wasps, which are located in the abdominal sternite VII in Asphondyliini and the abdominal segment VIII or IX in Lasiopterini and Alycaulini [5, 26, 27]. In addition, mycangial conidia are found with eggs in or on host plants [4, 26]. Therefore, relationships between gall midges inducing ambrosia galls and associated fungi are considered as mutualisms [1, 4, 13, 25].

Previously, fungal symbionts have been determined only by the isolation of fungi from galls [2, 14] or by morphological identification based on mycangial conidia [5]. Adair et al. [1] and Janson et al. [17] investigated fungal symbionts by molecular basis and identified fungi associated with distinct tribe gall midges as a single species Botryosphaeria dothidea, but the primary fungal symbionts are not yet understood in many gall midge species responsible for ambrosia galls. Therefore, we focused on the gall midge Illiciomvia vukawai, which is not yet examined to isolate symbiotic fungi, although fungal layers can be obviously seen inside its mature galls. Because the genus Illiciomyia is situated at one of the most basal groups of the subtribe Asphondyliina [35, 36], this species is critically important in understanding the evolutionary process of gall midges in relation to acquisition of their associated fungi.

I. yukawai induces hemiglobular and single-chambered galls on leaves of *Illicium anisatum* (Illiciaceae), a broad-leaved evergreen tree, and the galls remain on the tree throughout the year [34]. In order to understand the nature of gall midge–fungus relationships, fungal floras in galls and mycangia as well as their seasonal changes should be compared intensively.

In this study, we investigated associations between the gall midge *I. yukawai* and its fungal symbionts from both spatial and temporal points of view. We isolated fungi from the galls and mycangia and revealed successions of fungal floras in them. We also focused on micro-space within the gall by separating it into inner (fungal) and outer (plant tissue) layers and then compared fungal floras between them to reveal interaction between gall midge larvae and fungal symbionts within the galls.

Materials and Methods

Study Insect

I. yukawai is a monotypic species of the tribe Asphondyliini [34]. In May, adults emerge from galls on the tree in the morning and females lay eggs into new leaves. Larvae induce galls, overwinter at the third (final) instar, and pupate in the following spring [34, 39]. Although *I. yukawai* is fundamentally univoltine, some individuals exhibit prolonged diapause at the first instar stadium and require 2 years to complete a generation [34, 39].

The study was conducted in an I. anisatum plantation at

Kameyama City (34°50' N, 136°18' E, 200 m asl), Mie

Study Sites

Prefecture, central Japan, and in secondary forests consisting of *I. anisatum*, *Cryptomeria japonica*, and *Pieris japonica*, *Eurya japonica* in Taiki Town (34°18′ N, 136°18′ E, 800 m asl) and Komono Town (35°01′ N, 136°26′ E, 300 m asl), Mie Prefecture, central Japan, and Shimanto Town (33°12′ N, 133°02′ E, 450 m asl), Kochi Prefecture, western Japan. Five galled trees (approximately 3 m in height) of *I. anisatum* were randomly selected as sampling trees at each site.

Fungal Isolation from Galls and Leaves

From May 2006 to April 2007, five galled shoots (including about 40 galls) were randomly collected at 2– 4-week intervals in Kameyama, Taiki, and Shimanto. Collected galls were surface-sterilized in 70% ethanol for 1 min and then dipped for 1 min in sterile distilled water. Afterwards, the galls were aseptically dried on sterile filter paper, and they were then dissected and placed on four kinds of agar media (ten plates each of MA, YM, MPY, MM; "Electronic supplementary material"). The leaves, from which the galls were removed, were also used as the control for fungal isolation. After surface sterilization, leaf discs (approximately 2×2 mm) were aseptically cut and placed on the agar plates.

Fungal Isolation from Inner and Outer Layers of Galls

In November 2007, when the larvae were fully grown, the galls were collected from Kameyama (n=36), Komono (n=23), and Shimanto (n=26). They were surfacesterilized in 70% ethanol for 1 min and 1% NaClO for 2 mins and then dipped for 1 min in sterile distilled water. After the same process of drying, the galls were separated into inner layer (fungal tissue) and outer layer (lignified plant tissue) (Fig. 1a). Each layer was placed on MPY and MM agar media (each about half).

Fungal Isolation from Female Adults

In Kameyama and Komono, some galled shoots were enclosed in plastic bags $(340 \times 240 \text{ mm})$ before the emergence of *I. yukawai*. In May 2007, the shoots were observed almost daily in order to collect female adults that emerged within plastic bags. In total, 16 female adults (enclosed adults) have emerged from the enclosed galls. At the same time, 15 female adults that were laying eggs into new host leaves (egg-laying adults) were collected from the census sites.

In the laboratory, abdomens of the collected adults were aseptically pressed on agar plates to isolate fungi adherent to the body surface. Then, without surface sterilization, conidia in the mycangia on the abdominal

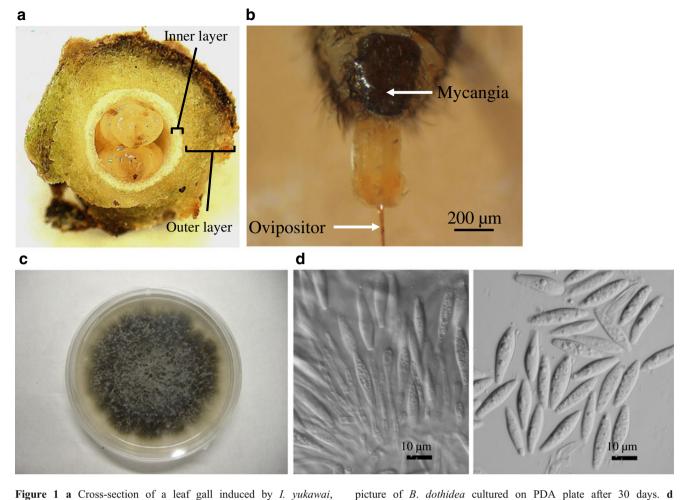


Figure 1 a Cross-section of a leaf gall induced by *I. yukawai*, showing inner (fungal) and outer (plant tissue) layers. b Mycangia on the abdominal sternite VII and retracted ovipositor of *I. yukawai*. c A

Conidiophores (*left*) and conidia (*right*) of *B. dothidea* induced on WA with pine needles

sternite VII (Fig. 1b) were inoculated into agar plates with a sterilized insect pin. MPY was used for both surface and mycangial fungal isolations of enclosed and egg-laying adults.

Identification

All cultures were incubated at 15°C. After fungal colonies were purified, isolated fungi were classified based on morphology and rDNA informations (LSU D1/D2 and/or ITS sequence data). Cultural morphologies and fungal structures on the media were observed under an optical microscope. Spore production on the representative isolates of each fungus was induced on water agar with pine needles under BLB light. If fungal structures were found on the media, they were picked up and mounted on glass slides and observed with a stereomicroscope.

The DNA of each isolate was extracted by the highthroughput method of Kikuchi et al. [19]. It was amplified using polymerase chain reaction (PCR) on a GeneAmp 9700 thermal cycler (Perkin-Elmer Applied Biosystems, Foster City, CA, USA). The oligonucleotide primer pairs, NL1 and NL4, ITS5 and ITS4 [38], were used for PCR. Then, 50 µl of reaction mixture containing 25 µl Qiagen GoTaq premix and 10 pmol each of primer and distilled water were added to the template. The reactions were initiated with 3 min of denaturation at 95°C, followed by 40 cycles of two-step PCR, consisting of 20 s at 95°C and 60 s at either 56°C for the primer pairs NL1 and NL4 or 54°C for ITS5 and ITS4, with a final extension for 10 min at 72°C. Amplification products were purified by QIAquick PCR Purification Kit (QIAGEN) and used for sequencing with a Big Dye Terminator Cycle Sequencing FS Ready Reaction kit ver. 3.1 and ABI PRISM 3100 genetic analyzer (Perkin-Elmer Applied Biosystems). Both strands of a fragment were sequenced. Blast homology search against the sequence data obtained was conducted (http://blast.ddbj.nig.ac.jp/top-j. html). We identified using the results of homology search together with morphological observations. The representative isolates (FFPRI411066-FFPRI411073) were deposited on

FFPRI culture collection (Forestry & Forest Products Research Institute, Japan). Selected sequence data (accession no. AB645744-AB645759) obtained in this study were deposited on DDBJ (DNA Data Bank of Japan).

Data Analysis

For the number of species and isolation frequency, data were pooled in each experiment because there were only small differences among the media kinds. Isolation frequencies of mycangial fungi were also combined irrespective of study sites.

The isolation frequency of each fungal species was calculated according to the following formula:

Isolation frequency(%) = $(N_i/N_t) \times 100$

where N_i and N_t are the numbers of samples from which the fungus was isolated and the total number of samples examined, respectively.

For the number of fungal species isolated from the inner and outer layers of galls, statistical significance was examined using Mann–Whitney *U*-test. Fisher's exact test was used to compare the isolation frequency of each fungus between inner and outer layers. All data were analyzed by SPSS ver. 11.5.1 j.

Results

Over 30 species of fungi were isolated in this study. They included plant-parasitic fungi such as B. dothidea, Colletotrichum sp., Phomopsis spp., and Pestalotiopsis sp. and saprophytic fungi such as *Cladosporium* spp. and Penicillium spp. Among the obtained fungal species, B. dothidea (Fig. 1c) was most frequently isolated from mycangia and ambrosia galls and was identified based on sequence data and its anamorphic characteristics. Its conidiomata were often produced on the medium, showing multilocular and eustromatic opening through several non-papillate pores. Conidiogenous cells were initially holoblastic, becoming enteroblastic, integrated, hyaline, smooth, $8-18 \times 2-5$ µm, cylindrical producing one or more conidia apically, often proliferating percurrently to produce conidia at successively higher levels on annellate conidiogenous cells or proliferating at the same level, resulting in periclinal thickenings. Conidia were fusiform to elliptical with a subobtuse apex and a truncate base bearing a minute basal frill, 20.4-25.6×4.9-6.5 µm in size (average of 84 conidia, $23.0 \times 5.7 \mu m$, L/W ratio=4.0), hyaline, aseptate, thin-walled, rarely becoming darker, 1septate, and well fitted with the description of its anamorph by Slippers et al. [30] (Fig. 1d).

Seasonal changes in the number of fungal species isolated from the galls and leaves are shown in Figs. 2 and 3, respectively. From June to October, several species of fungi were detected from the galls in all census sites (Fig. 2). The numbers tended to increase gradually from June to December and sharply during winter. The maximum numbers of fungal species detected were 14, 13, and 15 in Kamevama, Taiki, and Shimanto, respectively. In contrast, the fungi isolated from the leaves fluctuated from three to eight species after November, although the numbers reached peaks from September to October, in Taiki and Shimanto. In Kameyama, the number of fungal species fluctuated from five to ten through the study period (Fig. 3). Thus, there were clear differences in the isolation trend between the galls and the leaves rather than between the three study sites.

B. dothidea predominated among all of the isolates from May to September when the galls were still immature, and the isolation frequency was 10-62.5%. However, the frequency decreased to 0-35% following the development of galls (Fig. 4). Other common fungi at the three sites were Phomopsis sp. 1, Colletotrichum sp., and Pestalotiopsis sp. In contrast to *B. dothidea*, the isolation frequencies of these fungi were low from May to September, but they increased gradually during winter. Thus, they were not isolated frequently from the immature galls and then tended to increase gradually. The frequency of Phomopsis sp. 1 eventually reached about 90% and 40% in Kameyama and Shimanto, respectively, and that of Colletotrichum sp. about 60% in Taiki. Pestalotiopsis sp. was less dominant and its isolation frequency did not exceed 15% in all study sites throughout the survey.

In the leaves, *B. dothidea*, which is the dominant fungus in the galls, was seldom detected (< 10%) in all study sites throughout the survey. Although *Phomopsis* sp. 1, *Colletotrichum* sp., and *Pestalotiopsis* sp. were also

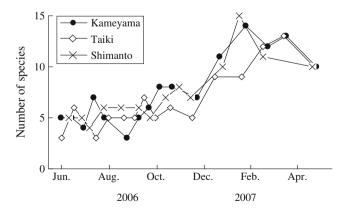


Figure 2 Seasonal changes in the number of fungal species isolated from galls induced by *I. yukawai*

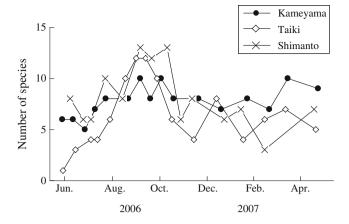


Figure 3 Seasonal changes in the number of fungal species isolated from *I. anisatum* leaves

isolated from the leaves, their frequencies were less than 50% (Fig. 5).

Table 1 summarizes the number of fungal species isolated from the inner and outer layers of the galls at each site (mean \pm SE, per gall). In all study sites, the number of species isolated was significantly larger in the outer layer than in the inner layer (Mann–Whitney *U*-test, p < 0.01).

Figure 6 illustrates the isolation frequency of three fungal species isolated from the inner and outer layers of the galls dominant at each site, including *B. dothidea*. Though the isolation frequency of *B. dothidea* showed no significant differences between the layers at all study sites (Fisher's exact test, p > 0.05), that of *Phomopsis* sp. 1 was significantly more frequent in the outer layer than in the inner layer at all sites. Similar results and trends were obtained for *Colletotrichum* sp. in Komono (Fisher's exact test, p < 0.05), respectively. An unidentified fungal species (unidentified sp. 1) was isolated in Kameyama, but the frequency was very rare and no significant differences were detected between the layers (Fisher's exact test, p > 0.05).

From mycangia of egg-laying adults, *B. dothidea* was the dominant species in immature galls with a frequency of 93.3%, but other fungi were seldom detected (Table 2). However, on the surface of the abdomen, the isolation frequency of *B. dothidea* was much lower (6.7%) than in mycangia, and that of the other fungi was higher (33.3%). In contrast to egg-laying adults, *B. dothidea* was not isolated at all from both parts of enclosed adults (both 0%), although other fungi were isolated in some cases (12.5% in mycangia, 75.0% on body surface).

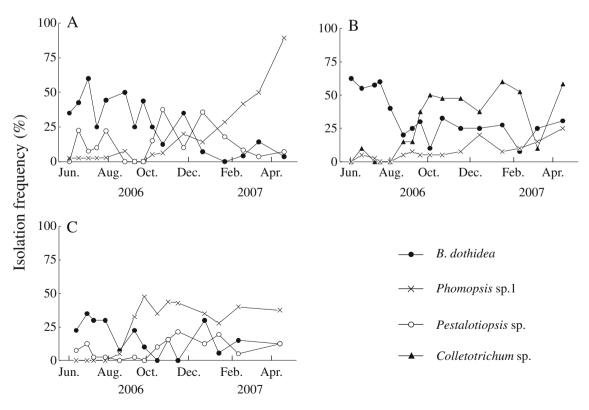


Figure 4 Seasonal changes in the isolation frequency of three dominant fungal species from galls induced by *I. yukawai*. A Kameyama, **b** Taiki, **c** Shimanto

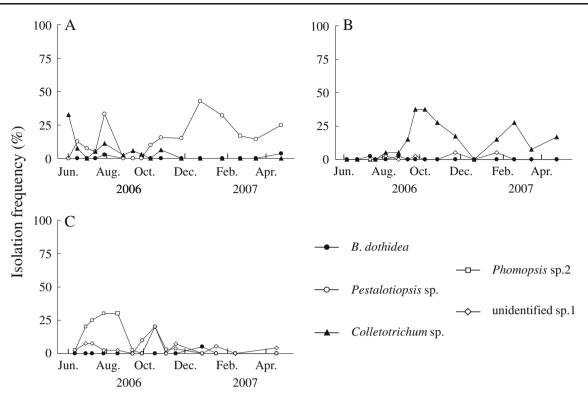


Figure 5 Seasonal changes in the isolation frequency of three dominant fungal species including *B. dothidea* from *I. anisatum* leaves. **a** Kameyama, **b** Taiki, **c** Shimanto

Discussion

Fungal flora in the galls and the leaves changed spatiotemporally. The number of fungal species isolated from the leaves increased from May to September or October and then decreased or fluctuated (Fig. 3). Generally, it is well known that fungal flora in leaves fluctuate with time [7, 12]. On the contrary, the number of fungal species isolated from the galls increased consistently with the development of galls (Fig. 2). Because the isolation frequency of each fungus was also different between leaves and galls (Figs. 4 and 5), galls are considered to have unique chemical and physical properties as fungal habitats distinct from the leaves, even though the gall tissue is a part of the leaf.

B. dothidea, *Phomopsis* sp. 1, *Colletotrichum* sp., and *Pestalotiopsis* sp. were dominant fungi commonly isolated from the galls at all study sites (Fig. 4). These fungi are known as endophytes of various plant species [6, 11, 23, 29, 31]. Among them, *B. dothidea* and *Phomopsis* sp. 1 showed different trends in temporary occurrence. *B.*

Table 1	The number	of fungal s	pecies ((mean±SE,	per gall)	isolated	from	inner	and	outer	layer	of gall	s
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		Study sites			
Layer	Kameyama	Komono	Shimanto		
	(<i>n</i> = 36)	(<i>n</i> = 23)	(<i>n</i> = 26)		
Outer	1.56 ± 0.09	1.78 ± 0.19	1.50 ± 0.10		
	**	**	**		
Inner	0.25 ± 0.07	0.09 ± 0.06	0.19 ± 0.08		

**P<0.01 (Mann–Whitney U-test)

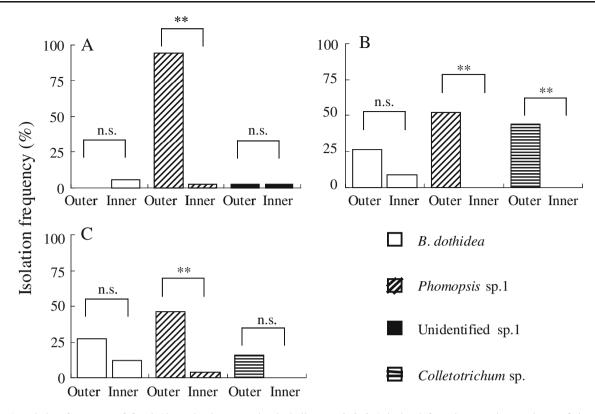


Figure 6 Isolation frequency of fungi (three dominant species including *B. dothidea*) isolated from inner and outer layer of the galls. **a** Kameyama (n=36), **b** Komono (n=23), **c** Shimanto (n=26). Fisher's exact test, **P < 0.01

dothidea was the most abundant during the immature-gall period but the isolation frequency became low after the galls were fully developed. In contrast, the isolation frequency of *Phomopsis* sp. 1 increased steadily with gall development and was often higher than that of *B. dothidea*. Although *Phomopsis* sp. 1 was also isolated from the leaves in some degree, *B. dothidea* was rarely detected from them throughout the year. These results suggest that *B. dothidea* is a symbiotic fungus particularly associated with the gall.

 Table 2
 List of fungi isolated from mycangia and body surface of enclosed and egg-laying female adults in Kameyama and Komono (data are combined)

	Number of cases, <i>n</i>	Mycangia	Surface
Enclosed adults ^a	16	Cladosporium sp. (1) Penicillium sp. (1)	Cladosporium spp. (5) Penicillium spp. (3) Phomopsis sp. 1 (2) Unidentified sp. 3 (1) Unidentified sp. 4 (1)
Egg-laying adults	15	B. dothidea (14) Colletotrichum sp. (1)	Cladosporium sp. (2) Penicillium sp. (2) B. dothidea (1)

^a Adults emerged from galls that were enclosed with plastic bag

We can propose a hypothesis why *B. dothidea* had low isolation frequency at mature-gall period (Fig. 4). Although the fungal inner layer existed inside the gall used for fungal isolation, the number of species and the isolation frequency of the isolated fungi including *B. dothidea* were extremely low in the inner layer (Table 1; Fig. 6). Generally, fungi should be isolated at high frequency from the mature gall with huge mycelia, but not in this study. Therefore, *B. dothidea* may have less ability to propagate on artificial media, showing physiological (trophic mode) change. This change is possibly caused by biological effects such as larval secretions of the gall midge. Manipulation of fungal symbionts by gall midges was also suggested in the case of *Asteromyia carbonifera*, goldenrod-galling midge, which belongs to the tribe Alycaulini [13, 17].

The dominant fungus in the galls, *B. dothidea*, was found frequently in mycangia of egg-laying adults, but not on their body (Table 2). Thus, we would strongly conclude that *B. dothidea* is the primary symbiotic fungus for *I. yukawai*. The enclosed adults, just emerging from the galls, had no *B. dothidea* in their mycangia or body surface (Table 2) as in previous reports [1, 4, 5, 13, 14]. This fact demonstrates that gall midges cannot take the symbiotic fungus into mycangia from the galls where they had developed. In other words, *B. dothidea* is not inherited via vertical (mother-to-infant) transmission as indicated by

Janson et al. [17]. Females of gall midge could collect conidia from the environment such as old non-natal galls in the leaf litter [4], incomplete galls where midge or parasitoids have died [1], and other parts of host plant [26]. Future studies are needed to clarify how and when the females search for conidia of symbiotic fungus and where they collect them. These are important study subjects to understand biological interactions between the gall midge and fungus.

The genus *Botryosphaeria* is a common endophyte in many plants [31, 32] and is also recognized as an economically important pathogen affecting various crops [15, 33]. Some species, including *B. dothidea*, are known to have a wide host range [31]. In this study, *B. dothidea* showed low isolation frequency from *I. anisatum* leaves (Fig. 5); however, it may possibly attack other parts, such as branches and stems of *I. anisatum* trees and/or other plants.

Bissett and Borkent [4] have predicted that fungal symbionts of gall midge species belonging to the tribe Asphondyliini and the supertribe Lasiopteridi may be Botryosphaeria and its related anamorphs (Fusicoccum, Diplodia, Dothiorella, and Macrophoma). For example, Adair et al. [1] suggested that fungal associates of some Asphondylia species distributed in Australia and South Africa are B. dothidea. Our study clarified that the symbiotic fungus associated with Illiciomyia was also Botryosphaeria. Asphondylia is the largest and radiating genus in tribe Asphondyliini [41] and contains nearly 300 described species [9, 10]. In contrast, Illiciomvia comprises only one species, as mentioned earlier, and is situated at one of the most basal clades of the subtribe Asphondyliina [35, 36]. To verify their prediction, we have to examine the symbionts of many other genera and species of gall midges further. However, the currently available evidence indicates that Botryosphaeria fungi appear to be common among Asphondyliini midges, all of which can provide a great opportunity for the fungi to disperse its conidia. Janson et al. [17] suggested that fungal symbionts of A. carbonifera are B. dothidea and identical to free-living populations with lack of genetical specialization. In our isolates of the symbionts of I. yukawai, further detailed genetic analysis is required to discuss this possibility.

We propose two hypotheses about the benefits of symbiotic fungus for *I. yukawai*. First, the fungus is directly utilized as a food source as predicted in ambrosia gall midges by Bissett and Borkent [4]. Furthermore, Heath and Stireman [13] showed that fungal proliferation and gall development are dependent on larvae of the gall midge *A. carbonifera*, indicating direct gall midge–fungal interaction. In the case of *I. yukawai*, full-grown larvae were obviously covered with fungal layers in galls. So, the larvae probably rely upon the fungus as the food resource. This observation

supports the first hypothesis, although we will need to detect fungal DNA from larval gut contents. Secondly, gall midge-associated fungus may function as a biological defense against antagonistic organisms [37], especially fungi. For example, mortality of the gall midge, *Contarinia* sp., which does not associate with symbiotic fungi, was higher on Douglas fir leaves infected by endophytes than on those uninfected [6]. Actually, many fungi other than *B. dothidea* existed in the leaves of *I. anisatum* and the outer layer of galls induced by *I. yukawai* (Figs. 5 and 6). Thus, *B. dothidea* might protect the larva from attack by antagonistic fungi.

Throughout the survey, we revealed the succession of fungal flora on galls, host leaves, and female adults of *I*. *yukawai* and identified the primary symbiotic fungus as *B*. *dothidea*. This is the first report to investigate the symbiosis between a gall midge and associated fungi from both spatial and temporal points of view.

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