
Diversity of fungi associated with the mountain pine beetle, *Dendroctonus ponderosae* and infested lodgepole pines in British Columbia

Sangwon Lee¹, Jae-Jin Kim² and Colette Breuil^{1*}

¹Department of Wood Science, University of British Columbia, 2424 Main Mall, Vancouver, B.C., V6T 1Z4, Canada

²Division of Environmental Science & Ecological Engineering, Korea University, Seoul 136-701, Korea

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The mountain pine beetle (*Dendroctonus ponderosae*), one of the most destructive bark beetles, has damaged large areas of lodgepole pine forests in British Columbia (Canada). It has been suggested that the beetle has a mutually beneficial relationship with the fungi that it carries. In this work, the fungal associates of the mountain pine beetle were extensively investigated. Fungi were isolated from the beetles, galleries and sapwood of infested lodgepole pines (*Pinus contorta* var. *latifolia*) at six epidemic sites in British Columbia. The isolated fungal species were more diverse than previously reported. We identified a total of 1042 isolates that belong to nine species. Among these, *Ophiostoma clavigerum*, an *Ophiostoma minutum*-like species, and *Ophiostoma montium* were frequently isolated. Unexpectedly, the *Ophiostoma minutum*-like species was found at high frequency on the mountain pine beetle. *Leptographium longiclavatum*, *Entomocorticium dendroctoni* and an unidentified species of *Entomocorticium* also appeared to be specifically associated with the mountain pine beetle.

Key words: *Dendroctonus ponderosae*, diversity, mountain pine beetle, ophiostomatoid, *Pinus contorta*

Introduction

Lodgepole pine (*Pinus contorta* var. *latifolia*) forests represent 25% of the total growing stock in British Columbia (COFI, 2005). However, large number of mature lodgepole pines has been killed by the mountain pine beetle, *Dendroctonus ponderosae*, and its fungal associates. Although outbreaks of the mountain pine beetle have occurred in British Columbia in the past, the present infestation is the most severe infestation ever recorded. The current mountain pine beetle epidemic started near Tweedsmuir Park and spread to northern

*Corresponding author: Colette Breuil; e-mail: colette.breuil@ubc.ca

British Columbia (Houston and Prince George) and the Alberta border. As of 2004, it has spread over 7 million ha, infesting 283 million m³ of lodgepole pine (British Columbia Ministry of Forests, 2005).

Usually, the mountain pine beetle attacks its host in July and August. At this time, trees are often stressed by water deficiency. The mountain pine beetle feeds on fungi and tree tissues (Harrington, 2005). New adults graze on fungi and acquire fungal spores in their mycangia and guts and on their exoskeletons before they emerge from the pupal chambers and attack new hosts. It appears that the mountain pine beetle and its fungal associates have a mutually beneficial relationship (Whitney, 1982; Paine *et al.*, 1997; Six, 2003a; Harrington, 2005).

The known fungal associates of the mountain pine beetle are ascomycetes in the genera *Ophiostoma* and *Leptographium*. They produce asexual and sexual spores in slimy masses that attach to insect bodies and are dispersed to new hosts that represent fresh nutrient sources (Harrington, 1993). Two fungal species, *Ophiostoma montium* and *O. clavigerum* have been consistently isolated from the mycangia and exoskeletons of the mountain pine beetle as well as from infested pines (Robinson, 1962; Whitney and Farris, 1970; Solheim, 1995; Six, 2003b; Kim *et al.*, 2005). Recently, *Leptographium longiclavatum* associated with the mountain pine beetle has also been reported (Lee *et al.*, 2005). In comparison, *Entomocorticium dendroctoni*, *Ophiostoma minutum* and *O. minus* have been occasionally found in the mountain pine beetle galleries and their association with the mountain pine beetle has been suggested (Robinson, 1962; Whitney *et al.*, 1987). However, most of the previous work was carried out with a limited number of isolation sites, and fungi were isolated only from the beetle or infested trees. Recent outbreaks have occurred over a wide range in British Columbia, which includes areas that have not been affected by mountain pine beetle epidemics in the past century and which have different climates from the ones previously studied. Therefore, the mycoflora in the current epidemic area in British Columbia might be different from what has been reported.

To obtain accurate information on the fungal diversity involved in the current mountain pine beetle outbreak, we investigated the fungi collected at six sites in British Columbia. We examined, first, which fungal species were consistently associated with the mountain pine beetle across the large epidemic area in British Columbia, and second, whether the fungal species and their frequencies as isolated from the beetles, beetle galleries and infested trees were different or not.

Materials and methods

Sampling strategies

Six epidemic sites (Fort St. James, Houston, Kamloops, Princeton, Tweedsmuir Park, and Williams Lake) were selected to cover the current outbreak regions in Canada (Fig. 1). Samplings were conducted before the emergence of teneral adult beetles from the hosts. A total of 23 trees (3-5 trees/site), which were attacked by the mountain pine beetle in the same or previous summer of the sampling year (early green and late green phase of trees, respectively), were harvested in 2001 and 2002 (Table 1). One bolt (50-80 cm in length) from each tree was cut at breast height and placed in a plastic bag in the field and then transported to the laboratory. The bolts were kept at 4°C for 1-3 days before fungal isolations were conducted.

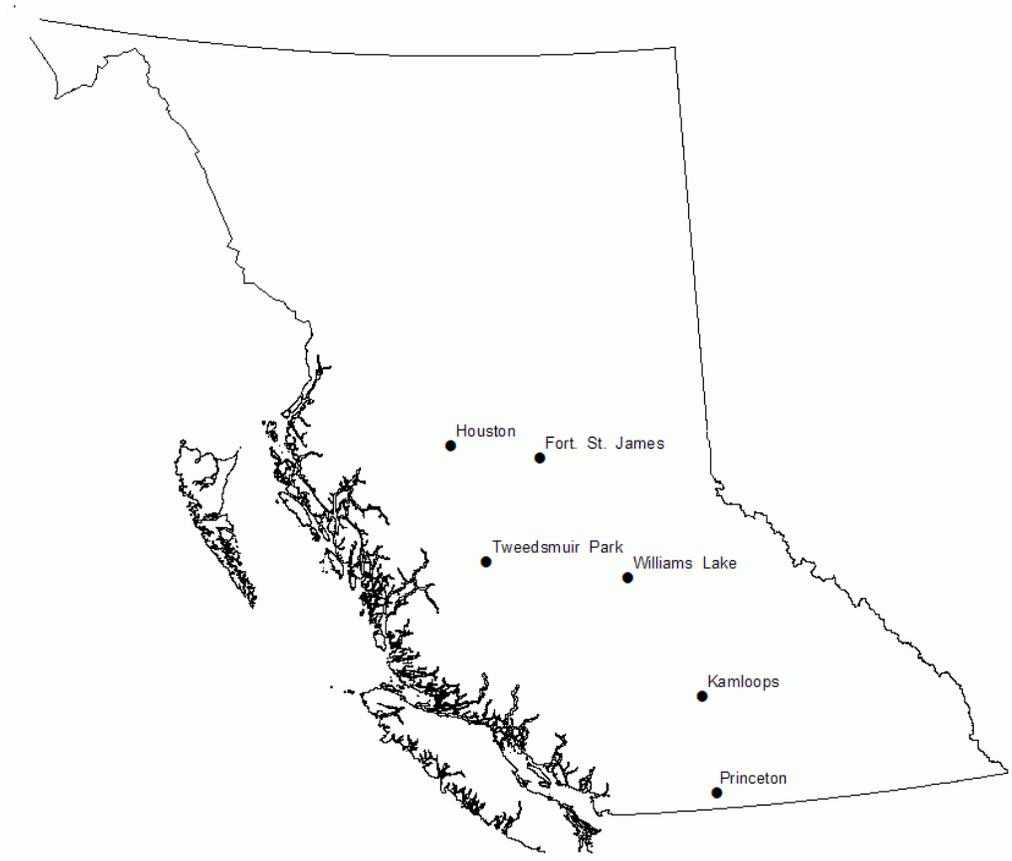


Fig. 1. Map of sampling sites in British Columbia, Canada.

Table 1. Characteristics of the mountain pine beetle-infested trees used for samplings.

Site	No. of trees	Location (latitude, longitude)	Date sampled	Tree phase*
Kamloops	2	N 50° 65', W 120° 36'	June 6, 2001	Late green
	1	N 50° 45', W 120° 31'	August 16, 2001	Early green
	1	N 50° 45', W 120° 31'	August 31, 2001	Early green
	1	N 50° 45', W 120° 31'	May 20, 2002	Late green
Williams Lake	3	N 52° 26' 21", W 122° 03' 15"	July 9, 2001	Late green
Princeton	3	N 49° 14' 58", W 120° 34' 43"	July 13, 2001	Late green
Tweedsmuir Park	4	N 52° 43' 41", W 125° 30' 23"	July 19, 2001	Late green
Houston	4	N 54° 08', W 126° 40'	July 17, 2002	Late green
Fort St. James	4	N 54° 38' 66", W 124° 25' 14"	August 8, 2002	Late green

*'Late green' indicates that trees were attacked in the previous summer and 'early green' means that the trees were attacked in the same summer of the sampling year (Kim *et al.*, 2005)

The trees harvested ranged in age from 75 to 140 years, with diameters varying from 20.5-35 cm. Most of the trees were heavily attacked with 70-160 pitch tubes/m², and we often observed that the mountain pine beetle larval galleries of adjacent broods overlapped at their ends. Generally, the entire sapwood was stained and secondary beetles such as *Ips* and ambrosia were commonly found in most of the trees.

Isolation of fungi from mountain pine beetle, galleries and sapwood

Bolts were debarked, and beetles were collected from galleries that were not intermingled with other galleries. Fungi on body surfaces of larvae, pupae, and adults were isolated using serial dilution with 0.01% (v/v) Tween-20, and were cultured on 2% malt extract agar (MEA) with ampicillin (50 µg/mL) as described by Lee *et al.* (2005). Fungi were sampled from the same galleries from which the beetles were collected, by aseptically transferring small amounts of frass onto 2% MEA with ampicillin. After 3-30 days of incubation at room temperature, either hyphal tips or the conidial mass on top of conidiophores were sub-cultured and grown on 2% MEA for identification. Fungi were also isolated from sapwood as described by Kim *et al.* (2005).

Fungal mycelia removed from the edge of a young colony were placed in micro-tubes (Sarstedt, Montreal, Québec) with either 1 mL of water for storage at 4°C, or 1 mL of 20% (v/v) glycerol (Sigma-Aldrich, Oakville, ON) solution for storage at -80°C.

Fungal identification

The filamentous fungal isolates were identified by the morphology of asexual structures (Rumbold, 1941; Batra, 1967; Robinson-Jeffrey and Davidson, 1968; Upadhyay, 1981; Jacobs and Wingfield, 2001). Identification was confirmed by sequencing the ribosomal DNA (rDNA) or β -tubulin gene of representative isolates (Table 2). Fungal DNA extraction and PCR were conducted following Lee *et al.* (2003), and sequencing was carried out as described by Lim *et al.* (2005). The identification of *O. clavigerum* was confirmed using an *O. clavigerum*-specific PCR-RFLP marker (Lee *et al.*, 2003). Fungal growth rates were measured at 23°C on 2% MEA as described by Lee *et al.* (2005).

Statistical analyses

As in our previous work (Kim *et al.*, 2005), the Simpson's diversity index (Simpson, 1949) was used to indicate fungal diversity because sample sizes in this study were relatively small (Mouillot and Leprêtre, 1999). The index is defined as

$$C = 1 - \sum_{i=1}^{i=S} p_i^2,$$

where P_i is the relative abundance of a species i , and S is the species richness, which is defined as the number of competing species present in the community. Fungal dominance was determined by Camargo's index ($1/S$) (Camargo, 1992), where S represents species richness. A species was defined as dominant if $P_i > 1/S$.

Results

A total of 1042 fungal isolates were obtained from mountain pine beetle adults, pupae, larvae, galleries, and sapwood collected from 23 lodgepole pines. Nine fungal species were isolated: *Ophiostoma montium*, *O. clavigerum*, an *O. minutum*-like species, an *O. nigrocarpum*-like species, *Leptographium longiclavatum*, *L. terebrantis*, *Entomocorticium dendroctoni*, and two unknown fungi, an *Entomocorticium* species and an *Ambrosiella* species.

Fungal diversity on the mountain pine beetle

From the exoskeletons of the mountain pine beetle adults we obtained 516 fungal isolates comprising eight species (Table 3). The most dominant species was *Ophiostoma montium*, whose average isolation frequency was

Table 2. GenBank accession numbers and growth rates of the fungi isolated from *D. ponderosae*.

Taxon	Strain	Site*	Primer used for amplification [†]	Accession no. [‡]	Closest match in BLAST	Accession of match	Identity [§] %	Growth rate mm/day
<i>Entomocorticium dendroctoni</i> Whitney	SL-A69	TP	ITS5/ITS4	DQ118419	<i>E. dendroctoni</i>	AF119506	99.6	0.7 ± 0.2
<i>Entomocorticium</i> sp.	SL-P44	P	ITS5/ITS4	DQ118418	<i>E. dendroctoni</i>	AF119506	99.6	0.5 ± 0.2
	SL-A3	TP	ITS5/ITS4	DQ118416	<i>Entomocorticium</i> sp. H	AF119512	99.1	3.6 ± 0.3
	SL-W002	WL	ITS5/ITS4	DQ118417	<i>Entomocorticium</i> sp. H	AF119512	99.1	3.5 ± 0.2
<i>Leptographium longiclavatum</i> Lee <i>et al.</i>	SL-K215	K	T10/BT12	AY288931	--	--	--	6.3 ± 0.9
	SL-W001	WL	T10/BT12	AY288936	--	--	--	7.1 ± 0.8
<i>Leptographium terebrantis</i> Barras & Perry	SL-A57	TP	T10/BT12	DQ118421	<i>L. terebrantis</i>	AY263192	100	12.8 ± 0.4
<i>Ophiostoma clavigerum</i> (Robins. Jeff. & Davids.) Harrington	SL-K1	K	T10/BT12	AY263210	<i>O. clavigerum</i>	AY263194	100	14.3 ± 0.7
<i>Ophiostoma minutum</i> -like sp.	SL-K70	K	ITS5/ITS4	DQ128175	<i>O. minutum</i>	DQ128173	93.1	1.5 ± 0.3
	SL-W15	WL	ITS5/ITS4	DQ128174	<i>O. minutum</i>	DQ128173	92.9	1.9 ± 0.3
<i>Ophiostoma montium</i> (Rumbold) von Arx	SL-K77	K	ITS1-F/ITS4	AY194942	<i>O. montium</i>	AY194941	100	6.7 ± 0.3
<i>Ophiostoma nigrocarpum</i> -like sp.	SL-A54	TP	ITS5/ITS4	DQ118420	<i>O. nigrocarpum</i> -like	AF484452	99.8	1.6 ± 0.2
<i>Pichia capsulata</i> (Wick.) Kurtzman	SL-WY2	WL	LR0R/LR3	DQ128167	<i>P. capsulata</i>	U70178	99.8	--
<i>Pichia holstii</i> (Wick.) Kurtzman	SL-W2Y4	WL	LR0R/LR3	DQ128171	<i>P. holstii</i>	U75722	99.2	--
<i>Pichia scolyti</i> (Phaff & Yoney.) Kreger	SL-W2Y3	WL	LR0R/LR3	DQ128172	<i>P. scolyti</i>	U45788	99.8	--
	SL-PY1	P	LR0R/LR3	DQ128170	<i>P. scolyti</i>	U45788	99.8	--
Unidentified yeast	SL-WY4	WL	LR0R/LR3	DQ128168	<i>P. ofunaensis</i>	U45829	98.3	--
	SL-W2Y1	WL	LR0R/LR3	DQ128169	<i>P. ofunaensis</i>	U45829	96.8	--

*K: Kamloops, P: Princeton, TP: Tweedsmuir Park, WL: Williams Lake

[†]ITS1-F (Gardes and Bruns, 1993), ITS4 and ITS5 (White *et al.*, 1990), LR0R and LR3 (Vilgalys and Hester, 1990), T10 and BT12 (Lee *et al.*, 2003).

[‡]Accession numbers in bold were sequenced during this work.

[§]Identity (%) was derived from the pairwise alignment of each isolate sequence with the closest BLAST match in GenBank or a reference strain (DQ128173, *O. minutum* CBS 145.59; AF484452, *O. nigrocarpum*-like C 314)

Table 3. Number of fungal isolates from *D. ponderosae* and number of beetles yielding each fungal species at six sites in British Columbia.

Taxon	Number of Isolates*						Total isolates
	Fort St. James	Houston	Kamloops	Princeton	Tweedsmuir Park	Williams Lake	
<i>Ambrosiella</i> sp.	--	3 (2)	--	--	--	--	3 (2)
<i>Entomocorticium dendroctoni</i> Whitney	--	--	--	17 (2)	3 (1)	--	20 (3)
<i>Entomocorticium</i> sp.	--	--	1 (1)	2 (2)	8 (2)	--	11 (5)
<i>Leptographium longiclavatum</i> Lee <i>et al.</i>	--	--	2 (1)	--	--	--	2 (1)
<i>Leptographium terebrantis</i> Barras & Perry	--	--	--	--	1 (1)	--	1 (1)
<i>Ophiostoma clavigerum</i> (Robins. Jeff. & Davids.) Harrington	11 (3)	20 (6)	9 (3)	--	2 (2)	2 (2)	44 (16)
<i>Ophiostoma minutum</i> -like sp.	23 ^a (6)	25 (5)	29 ^a (7)	--	2 (2)	5 (2)	84 ^a (22)
<i>Ophiostoma montium</i> (Rumbold) von Arx	27 ^a (5)	57 ^a (11)	91 ^a (16)	77 ^a (10)	24 ^a (4) [†]	75 ^a (12)	351 ^a (58)
No. of total fungal isolates	61	105	132	96	40	82	516
No. of total MPB	7	13	20	10	6	12	68
Species richness (<i>S</i>)	3	4	5	3	6	3	8
Camargo's index ($1/S$)	0.33	0.25	0.20	0.33	0.17	0.33	0.13
Simpson's index of diversity (<i>C</i>)	0.63	0.58	0.47	0.33	0.59	0.16	0.50

*The total number of fungi on the beetle = the number of fungal isolates written in the table $\times 10^4$.

[†]Values in parentheses are the number of MPB yielding each fungal species.

^aDominant species. Species was considered as dominant if $P_i > 1/S$, where P_i is the relative abundance of a species i and S is the species richness, which is the number of competing species present in the community (Camargo, 1992).

68% (from 44% at Fort St. James to 92% at Williams Lake) of the total number of fungal isolates. In total, 85% of the beetles (from 67% at Tweedsmuir Park to 100% at Princeton and Williams Lake) yielded *O. montium*. Unexpectedly, the second-most dominant isolate was the *O. minutum*-like species, which was isolated from 32% of the beetles with an average isolation frequency of 16% (from 0% at Princeton to 38% at Fort St. James). *Ophiostoma clavigerum* was also commonly isolated. However, it was isolated from fewer beetles (24%) than the *O. minutum*-like species, and its average frequency was lower (9%). In contrast to the *O. minutum*-like species, which was dominant at two sites, *O. clavigerum* was not dominant at any site. When the data were pooled, the dominant species on the mountain pine beetle exoskeletons were *O. montium* and the *O. minutum*-like species. *Entomocorticium* sp. and *E. dendroctoni* were often isolated, while *Leptographium longiclavatum*, *Ambrosiella* sp., and *Leptographium terebrantis* were found occasionally. At the sites where larvae, pupae and adults were found together (Kamloops, Tweedsmuir Park, and Williams Lake), the fungi obtained at each developmental stage were similar (Table 4).

Often, more than one filamentous fungal species was isolated from one beetle. Many beetles yielded two species (Tweedsmuir Park 83%, Houston 39%, Kamloops 29%, and Williams Lake 25% of beetles), or even three (Houston 15%, Princeton 10%, and Williams Lake 1% of beetles).

Yeasts were present in higher numbers than filamentous fungi. At Princeton, Williams Lake, and Kamloops, the average number of yeast colonies per adult beetle was approximately 3×10^5 , 5×10^5 , and 7×10^5 , respectively. Yeasts were obtained at all beetle developmental stages, but were more abundant on eggs (data not shown). Similarly to the filamentous fungi, more than one yeast species was isolated from most beetles. The 26S rDNA of the yeasts isolated in this study had high sequence identity (>99.2%) with those of *Pichia capsulata*, *P. holstii*, and *P. scolyti* (Table 2).

Fungal diversity in the beetle galleries and the stained sapwood

A total of 274 isolates were collected from galleries and sapwood (Table 5). *Ophiostoma montium* and *O. clavigerum* were frequently isolated from both galleries and sapwood. In contrast to the results from the beetle exoskeletons, the *O. minutum*-like species was isolated at low frequency, while *Leptographium longiclavatum* was often isolated. Other species, including *Entomocorticium* sp., *Ambrosiella* sp., and *Leptographium terebrantis*, were occasionally isolated in the sapwood. However, *Entomocorticium dendroctoni* was not found in sapwood. Yeasts were often found in galleries and occasionally in sapwood (data not shown).

Table 4. Number of fungal isolates from larvae, pupae and adults of *D. ponderosae**.

Taxon	Number of Isolates*											
	Kamloops			Tweedsmuir Park			Williams Lake			Total		
	Larvae	Pupae	Adults	Larvae	Pupae	Adults	Larvae	Pupae	Adults	Larvae	Pupae	Adults
<i>Entomocorticium dendroctoni</i> Whitney	2	--	--	--	--	3	--	--	--	2	--	3
<i>Entomocorticium</i> sp.	2	--	1	9	9	8	--	--	--	11	9	9
<i>Ophiostoma clavigerum</i> (Robins. Jeff. & Davids.) Harrington	10	--	9	--	4	2	--	--	2	10	4	13
<i>Ophiostoma minutum</i> -like sp.	8	4	29	3	18	2	--	3	5	11	25	36
<i>Ophiostoma montium</i> (Rumbold) von Arx	96	26	91	5	15	24	14	23	75	115	64	190
<i>Ophiostoma nigrocarpum</i> -like sp.	--	--	--	--	1	--	--	--	--	--	1	--
<i>Leptographium longiclavatum</i> Lee <i>et al.</i>	--	--	2	--	--	--	--	--	--	--	--	2
<i>Leptographium terebrantis</i> Barras & Perry	--	--	--	--	--	1	--	--	--	--	--	1
No. of <i>D. ponderosae</i>	14	4	20	2	5	6	2	3	12	18	12	38
No. of fungi	118	30	132	17	47	40	14	26	82	149	103	254

*The total number of fungi = the number of fungal isolates written in the table $\times 10^4$.

Table 5. Fungal isolates from gallery and wood at six sites in British Columbia.

Taxon	Number of Isolates													
	Fort St. James		Houston		Kamloops		Princeton		Tweedsmuir Park		Williams Lake		Total isolate	
	G*	W†	G	W	G	W	G	W	G	W	G	W	G	W
<i>Ambrosiella</i> sp.	--	--	--	--	--	--	--	2	--	--	--	--	--	2
<i>Entomocorticium</i> sp.	--	--	--	1	--	1	--	--	--	--	--	--	--	2
<i>Leptographium longiclavatum</i> Lee et al.	3	--	1	--	4	7	--	--	--	--	--	--	8	7
<i>Leptographium terebrantis</i> Barras & Perry	--	--	--	--	--	1	--	--	--	--	--	1	--	2
<i>Ophiostoma clavigerum</i> (Robins. Jeff. & Davids.) Harrington	8 ^a	13 ^a	5 ^a	2	27 ^a	54 ^a	2	6 ^a	7 ^a	3	4	8 ^a	53 ^a	86 ^a
<i>Ophiostoma minutum</i> -like sp.	2	--	1	--	--	1	--	--	--	--	--	1	3	2
<i>Ophiostoma montium</i> (Rumbold) von Arx	7 ^a	4	4 ^a	5 ^a	25 ^a	32 ^a	4 ^a	5 ^a	1	14 ^a	5 ^a	3	46 ^a	63 ^a
No. of total fungal isolates	20	17	11	8	56	96	6	13	8	17	9	13	110	164
Species richness (S)	4	2	4	3	3	6	2	3	2	2	2	4	4	7
Camargo's index (1/S)	0.25	0.50	0.25	0.33	0.33	0.17	0.50	0.33	0.50	0.50	0.50	0.25	0.25	0.14
Simpson's index of diversity (C)	0.69	0.36	0.65	0.53	0.56	0.57	0.44	0.49	0.22	0.29	0.49	0.56	0.59	0.56

*gallery, †wood

^aDominant species. Species was considered dominant if $P_i > 1/S$, where P_i is the relative abundance of a species i and S (Species richness) is the number of competing species present in the community (Camargo, 1992).

Discussion

Through an extensive survey, we isolated more diverse fungal associates of the mountain pine beetle than previously reported. While *Ophiostoma montium* and *O. clavigerum* were frequently isolated from the mountain pine beetle in accordance with previous studies (Robinson, 1962; Whitney and Farris, 1970; Six, 2003b), we also isolated an *O. minutum*-like species, *Leptographium longiclavatum*, *Entomocorticium* sp., and *E. dendroctoni*. The isolation methods in each study might have affected the species found. In this work, the beetles were washed and the diluted washes were plated onto media. In previous research the beetles were either streaked onto or allowed to walk on the surface of the media; with such methods, it is less likely that all spores in the cavities of the beetle exoskeletons would be removed. Since sampling fungi in mycangia without cross contamination from fungi present on the exoskeletons was difficult, we only isolated fungi from beetle body surfaces.

Unexpectedly, the frequency of the *Ophiostoma minutum*-like species on the mountain pine beetle exoskeletons was high; often this fungus was isolated more frequently than *O. clavigerum*. The *O. minutum*-like species appears to belong to a complex phylogenetic group that is often reported as *Ophiostoma minutum* Siem., or as *Ceratocystiopsis minuta* (Siem.) Upadhyay & Kendrick (Hausner *et al.*, 1993, 2003). To our knowledge, *Ophiostoma minutum* has been isolated occasionally from infested lodgepole pines (Robinson, 1962), but not from the mountain pine beetle exoskeletons. It has also been isolated from phoretic mites carried by other bark beetles (Moser and Macias-Samano, 2000). We often observed mites in the mountain pine beetle galleries, but further work needs to be done to determine whether the mountain pine beetle is associated with phoretic mites carrying the *O. minutum*-like species.

Consistent with Robinson's (1962) data, we isolated *O. montium* more frequently than *O. clavigerum* from the beetle exoskeletons. This was also the case when *O. clavigerum* was more prevalent than *O. montium* in the galleries, even though, for such cases, the beetles had more opportunity to contact *O. clavigerum* than *O. montium*. The lower frequency of *O. clavigerum* on beetles could be due to its large clavate spores, which might not be able to adhere to the beetles as stably as the small conidia of *O. montium*.

Leptographium longiclavatum was isolated from the mountain pine beetle exoskeletons and infested sapwood. In previous work it has been also found in the mountain pine beetle mycangia (Lee *et al.*, 2005). This species appeared to be affected by the moisture content of its environment, as it was isolated more often from early green phase trees than from the drier late green phase trees that had been infested in the previous summer (Kim *et al.*, 2005).

Like *Ophiostoma clavigerum*, *Leptographium longiclavatum* has long conidia, and may be more easily grazed by the beetles and carried preferentially in the mycangia rather than on the exoskeleton (Harrington and Zambino, 1990; Hsiau and Harrington, 1997; Six, 2003b).

In this work we isolated two basidiomycetes. *Entomocorticiium dendroctoni* was isolated from the mountain pine beetle exoskeletons, although it has been only reported in beetle galleries (Whitney *et al.*, 1987). The *Entomocorticiium* sp. had a faster growth rate than *E. dendroctoni* and its rDNA showed only 97.8-98.1% sequence identity to that of *E. dendroctoni*. *Entomocorticiium* species have been suggested to be good nutritional sources for the mountain pine beetle and other *Dendroctonus* species (Barras and Perry, 1972; Whitney and Cobb, 1972; Whitney *et al.*, 1987).

In contrast to all the above fungal species, which appear to be specifically associated with the mountain pine beetle, *Leptographium terebrantis*, *Ambrosiella* sp., and the *Ophiostoma nigrocarpum*-like species seem to be incidental associates. The presence of these fungi on the mountain pine beetle is likely due to cross-contamination with fungal associates of other cohabiting Scolytid beetles (e.g. *Ips* and ambrosia beetles), which were frequently observed in sampled trees. At this point, the information available on fungal associates of such other beetles is limited. *Ophiostoma minus* and *O. huntii*, which have been found in old mountain pine beetle galleries (Robinson, 1962; Robinson-Jeffrey and Grinchenko, 1964), were not isolated in this work.

The fungi that we isolated can be grouped into fast growing sapstaining fungi and slow growing non-staining fungi. It is likely that non-staining fungi, such as the *O. minutum*-like species, *Entomocorticiium* species and yeasts, found mainly on the beetle and rarely in sapwood, might be better nutritional sources for the mountain pine beetle than staining fungi, which contain melanin, a phenolic derivative. Whitney *et al.* (1987) has shown that the sapstaining fungi *Ophiostoma montium* and *O. clavigerum* were less efficient in supporting beetle reproduction than *E. dendroctoni*. Yeasts, commonly found on the mountain pine beetle in current and previous studies (Whitney, 1971; Lim *et al.*, 2005), may also be an important nutritional source. In addition, they may contribute to successful brood development by preventing excess numbers of beetles in individual trees, since some yeasts can convert the mountain pine beetle aggregation pheromone, *trans*-verbenol, into the anti-aggregation pheromone, verbenone (Hunt and Borden, 1990). Similarly, pathogenic species *Ophiostoma clavigerum*, *O. montium* and *Leptographium longiclavatum*, by invading the sapwood and interrupting water transportation, decrease the moisture content in the tree and may provide a better environment

for beetle brood development (author's unpublished data for *L. longiclavatum*; Webb and Franklin, 1978; Strobel and Sugawara, 1986; Yamaoka *et al.*, 1995).

In conclusion, in contrast to previous work, we showed that more diverse fungal species were associated with the mountain pine beetle. We also found that species dominant on the mountain pine beetle exoskeletons differed from those dominant in galleries and sapwood; with the difference mainly due to the *Ophiostoma minutum*-like species. Comparison of fungal frequencies from different parts of the mountain pine beetle, such as the exoskeleton, mycangia and gut, would suggest which fungi might be more preferentially consumed by the beetle. Further work will be needed to clarify interactions among fungi and the impact of each fungal species on the mountain pine beetle fitness and on host defence mechanisms.

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