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# Ecological aspects of myxomycetes associated with white and brown wood rot on coarse woody debris in subalpine coniferous forests in **Central Japan**

Kazunari Takahashi<sup>a\*</sup>, Yuichi Harakon<sup>b</sup>, Yu Fukasawa<sup>c</sup>

<sup>a</sup> Okayama Ryukoku High School, 874 Kasaoka, Kasaoka City, Okayama 714-0081, Japan

<sup>b</sup> Hiroshima Municipal Hiroshima Secondary School, 1-14-1 Miirihigasi, Asakitaku, Hiroshima City, Hiroshima 731-0212, Japan

<sup>c</sup> Laboratory of Forest Ecology, Graduate School of Agriculture Science, Tohoku University, Naruko, Osaki, Miyagi 898-6711, Japan

## ABSTRACT

We investigated factors affecting the community composition of lignicolous myxomycetes in dead wood with white and brown rot through summer and autumn surveys in a subalpine forest in Central Japan. In both seasons, wood had decayed to a softer state under brown rot than under white rot. The pH of wood with white rot was nearly neutral, while wood with brown rot was weakly acidic. Wood pH was lower in summer than in autumn. Forty-two myxomycetes taxa in 19 genera were identified in 302 fruiting-body colonies; white rot yielded 31 taxa and brown rot 24 taxa. Species diversity was higher on wood with white rot than on wood with brown rot. The effect of wood hardness on species composition depended on season. Several species exhibited a preference for one of the rot types. The substrate conditions associated with brown rot limit myxomycetes species diversity.

Keywords: brown rot, seasonality, white rot, wood pH, wood hardness

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In myxomycetes, a group of fungus-like protists, the reproductive stage comprises an amoeba and the developmental stage comprises spore-containing fruiting bodies (Novozhilov et al., 2017). Myxomycetes typically inhabit coarse woody debris and prey on decomposers such as bacteria and fungi (Clark & Haskins, 2015; Madelin, 1984), and their nutritional preferences may vary between species (Hoppe & Kutschera, 2015). Wood-inhabiting (lignicolous) myxomycetes are associated with the state of decay of dead wood (Novozhilov et al., 2017; Takahashi & Hada, 2009; Xavier de Lima & de Holanda Cavalcanti, 2015), but the factors influencing their association with white and brown rot remain to be clarified.

Fungi decompose the lignocellulose in wood, although the process differs depending on the type of wood rot (Fukasawa, 2013). This difference may impact lignicolous myxomycete community composition (Fukasawa et al., 2015). The relationships between the amount of white and brown rot present in the forest and myxomycete community dynamics remain incompletely understood. To address this gap in knowledge, we examined the responses of myxomycete communities to the state of wood decay and substrate pH associated with brown and white wood rot in a subalpine coniferous forest of Japan.

\* Corresponding author. Okayama Ryukoku High school, 874 Kasaoka, Kasaoka City, Okayama 714-0081, Japan.

E-mail address: kumakusu03@yahoo.co.jp (K. Takahashi)

The survey site was located in the subalpine forest of the Yatsugatake in Nagano Prefecture, Central Japan, beside Shirakoma-ike (2,132 m altitude, Sakuho-machi Minamisaku-gun, 36.04752121° N, 138. 36402719° E). The forest, comprising oldgrowth conifers, is dominated by Tsuga diversifolia Mast. and a mixture of Abies veitchii Lindley and Abies mariesii Mast. During our survey, large deadwood logs were abundant on the forest floor (Fig. 1A). The predicted mean annual temperature was 2.3 °C and mean annual precipitation >1,271.5 mm for 1999-2020, according to the neighboring Japanese Meteorological Agency station (Haramura; 35.5801200° N, 138.1301200° E, 1,017 m asl).

The surveys were performed over 3y during two field seasons (summer and autumn), on Jul 23, 2017; Jul 25, 2018; Oct 28, 2018; and Oct 28, 2019, at Shirakoma-ike. The process of data collection was as follows. First, we located portions of logs exhibiting myxomycete fruiting bodies (Fig. 1B). We identified the rot type in those portions based on macroscopic features. White rot is distinguished by the separation of the wood into fibers, whereas brown rot is distinguished by cracks along the length and breadth of the rotten wood (Araya, 1993). We then measured wood hardness (depth, in mm) using a soil hardness tester (No. 351; Fujiwara Scientific, Nishigahara Kita-ku, Tokyo, Japan), and pH using a hand-held pH meter (Shinwa code 72724, measurement range of 4.0-7.0), following Fukasawa et al. (2015). Surveys were conducted on ca. 200 fallen logs along a trekking path on the forest floor (ca.  $800 \text{ m} \times 30$ 



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**Fig. 1** – Survey site in a subalpine coniferous forest on Mt. Yatugatake, and myxomycete fruiting bodies. A: Fallen logs on the forest floor. B: Colony of myxomycete fruiting bodies on a portion of a log exhibiting white rot. C: Fruiting bodies of *Lamproderma columbinum*. D: Fruiting bodies of *Cribraria macrocarpa*.

m) using the naked eye and a magnifying glass. During each survey, searching continued until ca. 100 colonies of fruiting bodies were located and recorded. The average wood hardness and substrate pH were calculated for each species and for white rot and brown rot separately.

For each myxomycete species, the number of colonies was recorded for each season and wood rot type. Myxomycete fruiting bodies were partly sampled and glued to specimen boxes then identified to species level via microscopic observation in the laboratory. Myxomycete nomenclature was according to Yamamoto (1998), and the most recent published scientific names were used (Lado, 2005–2022). Both species and varieties were considered when assessing species richness. Voucher specimens were deposited in the Tottori Prefectural Museum herbarium (TRPM).

To describe the myxomycete assemblage, abundance (colony count) was compared by season and rot type. Survey accuracy was estimated using the index of exactitude, as [observed number of species  $(S_{obs})$  in each assemblage] / [estimated number of species  $(S_{oss})$ ] × 100;  $S_{est}$  was calculated using the Chao1 method (Chao, 1984). Myxomycete assemblage species diversity was calculated using the Shannon–Wiener index (H') (Shannon & Weaver, 1963) and the equitability (J') index (Pielou, 1996), as in previous studies (Stephenson, 1989). These indices were estimated using PAST (Hammer et al., 2001; https://past.en.lo4d.com/windows). Species composition was calculated using the relative abundance of each

species (as %), as [number of colonies for a given species] / [total colonies]  $\times$  100. Species preference for each rot type was determined using an independent sample *t*-test in ESUMI Excel Statistics 5.0 (ESUMI Co. Ltd., Tokyo, Japan).

In total, 302 myxomycete colonies were identified, 222 on white rot and 80 on brown rot. Considering wood with myxomycete colonies, in terms of wood hardness, the depth for white rot was  $21.0 \pm 7.0$  mm in summer and  $20.6 \pm 7.8$  mm depth in autumn; for brown rot, the depth was  $18.4 \pm 7.1$  mm in summer and  $16.0 \pm 8.0$  mm in autumn (Table 1). The wood was decayed to a significantly softer state under brown rot than under white rot in both seasons (p < 0.05). Overall, wood pH ranged from 4.4 to 7.0, and differed seasonally; mean pH was significantly lower in summer (pH = 5.9) than in autumn (pH = 6.7) (*t*-test, p < 0.01, Table 1). The rot types were significantly influenced by the pH: wood with brown rot was more acidic (pH 5.4  $\pm$  0.5 in summer, 6.4  $\pm$  0.3 in autumn) (p < 0.01).

In contrast, considering wood without myxomycete colonies in summer, wood with white rot exhibited hardness of  $14.9 \pm 6.5$  mm depth and pH of  $6.2 \pm 0.4$  (n = 66), while wood with brown rot exhibited hardness of  $14.7 \pm 6.9$  mm depth and pH of  $5.2 \pm 0.5$  (n = 71); the two rot types did not differ significantly in wood hardness, whereas brown rot exhibited significantly lower pH than white rot (*p* < 0.01). These findings indicate that rot type influenced wood

**Table 1.** Number of colonies, wood hardness, and pH at which myxomycete fruiting colony occurred on a portion of log according to the season and decay type. Hardness and pH are indicated by the mean value with its standard deviation (\*p < 0.05, \*\*p < 0.01).

		White rot	Brown rot	Total
Summer				
	Colonies	135	55	190
	Hardness	$21.0 \pm 7.0$	$18.4^{*} \pm 7.1$	$20.2 \pm 7.1$
	pH	$6.2 \pm 0.4$	$5.4^{**} \pm 0.5$	$5.9 \pm 0.6$ **
Autumn				
	Colonies	87	25	112
	Hardness	$20.6 \pm 7.8$	$16.0$ $^{*} \pm 8.0$	$19.6 \pm 8.0$
	pН	$6.8 \pm 0.2$	$6.4^{**} \pm 0.3$	$6.7 \pm 0.3$

pH, which was nearly neutral for wood with white rot, whereas wood with brown rot exhibited weak acidity.

Accuracy for the entire survey was 85.7%, based on the Chao1-estimated number of species (Table 2). In total, 42 myxomycete taxa belonging to 19 genera were identified. In summer, 27 taxa in 12 genera were identified in 190 colonies of fruiting bodies; in autumn, 16 taxa in 10 genera were identified in 112 colonies (Table 3). The species fruited seasonally, with some fruiting in summer and some in autumn, except for one taxon, Lycogala epidendrum (L.) Fr. The dominant species in summer was Ceratiomyxa fruticulosa (O.F. Müll.) T. Macbr., followed (in decreasing order of relative abundance) by L. epidendrum, Stemonitis axifera (Bull.) T. Macbr., Cribraria cancellata (Batsch) Nann.-Bremek., Stemonitopsis hyperopta (Meyl.) Nann.-Bremek., and Physarum viride (Bull.) Pers., all of which exhibited 10 colonies or more (Table 3). In autumn, two species-Lamproderma columbinum (Pers.) Rostaf. (Fig. 1C) and Cribraria macrocarpa Schrad. (Fig. 1D)-were prominent. The remaining species typically exhibited fewer than 10 colonies each.

Table 3 presents the average wood hardness and pH for the species. Most of the species were widely distributed on moderately decayed wood with a hardness depth range of 13–27 mm. However, two species [*Physarum atroviolaceum* G. Moreno, Y. Yamam. & A. Castillo and *P. viride* (Bull.) Pers.] showed preferences for hard wood with depth  $\geq$  29 mm. *Cribraria* and *Lindbladia* were the only species present on well-decayed wood (depth < 13 mm).

Figure 2 presents the pH and wood hardness of the 25 taxa exhibiting at least three colonies. Wood pH was 5.2–7.0 for most of the myxomycetes species; the autumn-fruiting myxomycetes species preferred wood with pH > 6.4, whereas summer-fruiting species preferred wood with pH < 6.4. In summer, *L. epidendrum* occurred at a hardness of 20.0 mm depth and pH of 5.9; in autumn, it occurred at 21.1 mm depth and pH of 6.7. In summary, myxomycetes substrate usage (and substrate chemistry) varied between the seasons.

Substrate condition, which depends on the type of wood rot, affected the myxomycete assemblage. Wood with white rot exhibited many colonies, hosting 31 taxa, whereas brown rot hosted 24 taxa. White rot exhibited higher species diversity (H' = 2.82, J' = 0.82) than brown rot (H' = 2.75, J' = 0.87) (Table 2). Thirteen taxa were present on both rot types (Table 3), 18 only on white rot, and 11 only on brown rot. Three species—*C. fruticulosa*, *L. epidendrum*, and *L. columbinum*—were significantly biased toward white rot, while *C. macrocarpa* was prominently biased toward brown rot.

Three species exhibited seasonal selectivity for the specific rot type with a particular substrate pH. *Lamproderma columbinum* exhibited a preference for wood with white rot in autumn (independent sample *t*-test, p < 0.05), with a hardness depth of 21.9 mm and pH of 6.7 (Table 3). Two species of *Cribraria* preferred wood

with brown rot (independent sample *t*-test, p < 0.01): *C. pyriformis* var. *notabilis* Rex occurred on average on wood with 14 mm depth and pH of 5.2 in summer, and *C. macrocarpa* on wood with 15 mm depth and pH of 6.0 in autumn. *Stemonitis pallida* Wingate and *Stemonitopsis amoena* (Nann.-Bremek.) Nann.-Bremek. exhibited a preference for brown rot (Table 3).

Brown rot accounted for 26% of the subalpine coniferous forest samples that we collected, although brown rot occurs on only a small proportion of pine logs in Japan (Fukasawa, 2015). Brown rot fungi, however, are more abundant than white rot fungi on Norway spruce logs (Rajala et al., 2011) and occur widely on well decayed *Picea abies* logs (Bütler et al., 2007). The present forest survey enabled us to examine myxomycete species diversity on brown rot in the subalpine coniferous forest.

The wood decay process involves two phases, the first involving white rot, and the second brown rot (Fukasawa et al., 2014). Thus, in this study, the wood with brown rot had decayed further and was softer than the wood with white rot. Under white rot, wood pH increases. In contrast, brown rot leads to acidification (Fukasawa et al., 2015), because the fungi responsible for brown rot produce oxalic acid as they degrade wood (Espejo & Agosin, 1991). Low substrate pH may limit the growth and development of several myxomycete species, limiting zygote formation (Shinnick et al., 1978), plasmodium formation (Collins & Tang, 1973), and sporulation (Gray, 1939, 1953). Our findings indicate that few species were suited to an acidic substrate and that species diversity was lower on wood with brown rot. Fukasawa et al. (2015) reported that lignin-rich logs with brown rot hosted Cribraria tenella and Cribraria intricata, which prefer a lower pH. Several Cribraria species have been reported on acidic decaying wood; these include Cribraria cancellata (mean pH 5.0) and Cribraria tenella (pH 5.1), as well as other species such as Arcyria cinerea (Bull.) Pers. (pH 5.2), L. epidendrum (pH 5.4), S. axifera (pH 5.4), and C. fruticulosa (pH 5.7) (Xavier de Lima & de Holanda Cavalcanti, 2015). Vlasenko et al. (2018) suggested that a preference for acidic substrates on decaying wood restricts the community composition to Cribraria and Comatricha species. These reports suggest that the myxomycete species assemblage is associated with wood rot type via effects of substrate pH.

The type of wood rot influences bacterial activity (Jurgensen et al., 1989) as well as the dominant bacterial taxa (Haq et al., 2022). This potentially affects the food source for the trophic stages of myxomycetes, indirectly enhancing the growth of their fruiting bodies. Acidity restricts the growth of most bacteria (Lauber et al., 2009). This is potentially a key driver of the reduced myxomycetes species diversity observed in this study under the lower pH of wood with brown rot.

This study described associations of myxomycetes community composition with wood hardness, acidity, and decay types. The

Table 2. Accuracy of assessments and myxomycete species diversity in rot types. The numbers in
brackets of taxa indicate the predicted species by Chao 1. Species diversity is indicated by the Shan-
non & Weaver index (H') and the numbers in brackets are the equitability index (J').

		White	Brown	Total
Summer				
	Taxa	18 (20)	19 (23)	27 (28)
	Accuracy (%)	90	82.6	96.4
Autumn				
	Taxa	14 (17)	5 (6)	16 (24)
	Accuracy (%)	82.3	83.3	66.7
Total	Taxa	31 (38)	24 (30)	42 (49)
	Accuracy (%)	81.5	80	85.7
	Species diversity	2.82 (0.82)	2.75 (0.87)	3.05 (0.82)

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$ \label{eq:product} product (C) (Aut: C) (Aut:$	$Taxa^a$	Abbreviation <sup>b</sup>	Colonies	Hardness	Hd	Colonies	Hardness	Hd	colonies
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Synool	Ceratiomyxa fruticulosa (O. F. Müell.) T. Macbr.	Cefu	34 *	18	6.1	9	21	5.5	40
Rement offer offer optimization (Construction)     Base and antipaction (Construction)     Base and antipaction     Base antipaction     Base antip	Lycogala epidendrum (L.) Fr.	Lyep-S	26 *	21	6.0	4	25	6.0	30
Strengt opport (Name)	Stemonitis axifera (Bull.) T. Macbr.	Stax	14	22	6.1	7	16	5.1	21
	Stemonitopsis hyperopta (Meyl.) NannBremek.	Sthy	14	20	6.1	З	23	5.1	17
	Cribraria cancellata (Batsch) NannBremek.	Crca	ю	16	5.9	$10^{**}$	15	5.3	13
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	Enerthenema papillatum (Pers.) Rostaf.	Enpa	7	25	6.4				7
Standard stands, mathematical stands, mathematis stands, mathamatical stands, mathematical stands, mathematical	Stemonitopsis aequalis (Peck) Y. Yamam.	Stae	5	24	6.5	1	25	5.0	9
	Stemonitis smithii T. Macbr.	Stsm	4	22	6.3	1	16	5.2	5
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Stemonitopsis gracilis (G. Lister) NannBremek.	Stgr	4	19	5.8				4
And the stand of a phylic	Cribraria piriformis var. notabilis Rex	Crpyn				**	14	5.2	4
	Arcyria cinerea (Bull.) Pers.	Arci	3	19	6.3				3
Controling balls (New Ed.).1 (Scholl.     Could     1     22     53     23     53     33       Strongling palls (New Ed.).1 (Scholl.     Stron     Strongling palls (New Ed.).1 (Scholl.     Strongling (New	Stemonitis virginiensis Rex	Stvi	ю	23	6.0				3
	Comatricha nigra (Pers. Ex J. F. Gmel.) J. Schröt	Coni	1	22	5.8	2	15	5.5	3
Symonholis Control     Sim	Stemonitis pallida Wingate	Stpa				3 *	21	5.4	3
$\label{eq:constraints} for four or for the form of t$	Stemonitopsis amoena (NannBremek.) NannBremek.	Stam				3 *	20	5.5	3
$\label{eq:constraints} \mbox{Lights} \mbox$	Arcyria obvelata (Oeder) Onsberg		2	26	6.5				2
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	Lindbladia tubulina Fr.		2	10	6.4				2
Arotia infectora (finate)     2     10     50     2       Semonitopics relations (in the finate)     2 </td <td>Fuligo septica (L.) Wiggers</td> <td></td> <td>1</td> <td>12</td> <td>6.2</td> <td>1</td> <td>27</td> <td>5.8</td> <td>2</td>	Fuligo septica (L.) Wiggers		1	12	6.2	1	27	5.8	2
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	Stemonitis flavogenita E. Jahn		1	32	6.5				1
	Comatricha pulchella (C. Bab.) Rostaf.					1	15	5.2	1
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$\label{eq:product} \mbox{Autum} Autu$	Cribraria vulgaris Schrad.					1	8	4.8	1
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Autumn								
$ \begin{array}{cccc} Cribratia macrocarpa Schad. \\ Cribratia macrocarpa Schad. \\ Trévia dezipeux (Pens.) T. Hat. \\ Coloderma cutam (Lippert) G. Lister \\ Coloderma cutam (Lippert) G. Lister \\ Coloderma cutam (Lippert) G. Lister \\ Diderma ungristeux (Taman. & A. Castilo \\ Phat. \\ Diderma ungristeux (Taman. & A. Castilo \\ Phat. \\ Trévia durgristeux (Taman. & A. Castilo \\ Phat. \\ Diderma ungristeux (Taman. & A. Castilo \\ Phat. \\ Trévia durgristeux (Taman. & A. Castilo \\ Phat. \\ Pha$	Lamproderma columbinum (Pers.) Rostaf.	Laco	33 *	22	6.8	б	21	6.1	36
Jycogale pidendrum (L.) Fr.   Lyep-A   8   21 $6.7$ 2   30   6.9   8     Trichia decipies (Pers.) T. Mach:   Tracha   Elec   6   24   6.9   6.7   2   30   6.9   6     Trichia decipies (Trichia ereta (Ras.) T. Rach:   Trichia ereta (Ras.) T. Guell   Trichia ereta (Ras.)   Trichia ereta (Ras.)   Trichia ereta (Ras.)   7   6.9   7   5     Trichia ereta (Ras.) J. F. Gmel.   Trichi   5   24   7.0   6.9   4   4   6   4   4   6   5   5   4   7   6   3   3   6   9   5   2   4   7   6   3   3   6   9   5   2   4   1	Cribraria macrocarpa Schrad.	Crma	13	15	6.7	18 **	15	6.4	31
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Blaeomyxa cerifera (G. Lister) Hagelst.Elce6246.96Trichia erecta RexTrel5247.05Trichia erecta RexTurie4166.95Tubifera fractaTurie3276.83Physarum acroviolaceum G. Moreno, Y. Yamam. & A. CastilloPhat3306.92Physarum acroviolaceum Hoffm.Cotoc3306.922Diderma ouglarow fractare Hoffm.1106.821Diderma aurantiacum Y. Yamam. & NamBremek. & Lado11176.91Diderma aurantiacum Y. Yamam. & NamBremek. & Lado11176.91Diderma aurantiacum Y. Yamam. & NamBremek. & Lado1176.91Diderma aurantiacum Y. Yamam. & NamBremek.1176.91Diderma aurantiacum Y. Yaman. & NamBremek.176.91Diderma aurantiacum Y. Yaman. & NamBremek.176.91Tribraria purpurea Schrad.176.911Tribraria purpurea Schrad.22 <td< td=""><td>Trichia decipiens (Pers.) T. Macbr.</td><td>Trde</td><td>9</td><td>19</td><td>6.7</td><td>2</td><td>30</td><td>6.9</td><td>8</td></td<>	Trichia decipiens (Pers.) T. Macbr.	Trde	9	19	6.7	2	30	6.9	8
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Trichia botytis (J.F. Gmei.) Pers.   1   25   7.0   1     Cribraria filijornis Nowotny & H. Neubert   1   4   6.7   1     Cribraria purpurea Schrad.   1   7   6.8   1     Number of colonies   222   80   302     Number of species   32   24   42	Lepidoderma tigrinum (Pers.) Rostaf.		1	17	6.9				1
Cribraria fuljornis Newolny & H. Neubert   1   4   6.7   1     Cribraria purpurea Schrad.   1   7   6.8   1     Number of colonies   222   80   302     Number of species   32   24   42	Trichia botrytis (J.F. Gmel.) Pers.		1	25	7.0				1
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growth of myxomycetes species on decaying wood is known to be affected by the decay state. Myxomycetes may play an important role in the decomposition of woody debris in forest ecosystems. Most myxomycetes exhibit spatial and temporal niche separation in woody debris, as do species of *Ceratiomyxa* (Rojas et al., 2008), potentially interacting with other organisms (Dudka & Romanenko, 2006) and slowing the decay of woody debris by preying on decomposers (Yoshida, 2015). The findings suggest potential directions for future research to clarify the mechanisms and ecological aspects of the functions of myxomycetes in forests.

# **Declarations of interest**

The authors declare no conflicts of interest. The survey undertaken complies with the current laws of Japan.

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