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Characteristics of the Symbiotic Flagellate Composition within the Termite Family Rhinotermitidae (Isoptera)

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Abstract

All lower termites contain a symbiotic flagellate community in their hindguts. Yamin (1979) listed the symbiotic protistan composition of 31 species that belong to the family Rhinotermitidae. In this study, the symbiont composition of additional 23 Rhinotermitid termite species were investigated. The flagellate genera *Spirotrichonympha, Pseudotrichonympha* and *Holomastigotoides* are prevalent among Rhinotermitid hosts. Members of the genus *Reticulitermes* lack the latter two flagellate genera, but possess the most diverse flagellate community of the Rhinotermitids, being the only genus to contain flagellates of the order Oxymonadida. A one to one host-species to symbiont-species relationship is found within the genera *Parrhinotermes* and *Termitogeton*.

Keywords: Termite, Rhinotermitidae, symbiotic flagellates

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1. Introduction

All lower termites possess flagellate symbionts in the swollen part of their hindguts (termed 'paunch'). These flagellates play an important part in the digestion of cellulose and may contribute up to 40% of the host wet weight (Honigberg, 1970; Breznak and Brune, 1994).

All flagellate symbionts found in lower termites belong to the three protistan orders Trichomonadida, Hypermastigida and Oxymonadida. Most host species possess more than one species of flagellate. Yamin (1979) summarized data from 229 papers and presented a list of symbiotic flagellates from hosts representing all 6 recognised families of lower termites. According to the list, 434 symbiont species were described from a total of 205 lower termite species. Individual termite taxa usually contain a specific flagellate species composition. However, some flagellate species are found in more than one host species. Some authors suggested that symbiont composition reflects phylogeny of the host termites (Kirby, 1937; Honigberg, 1970; Kitade and Matsumoto, 1993a,b). There have also been claims that symbionts can be transferred horizontally via interspecies aggression. Thorne (1990, 1991) presented evidence that this mode of transfer could explain the origin of flagellate symbionts in the wood-feeding cockroach genus Cryptocercus. Aggression between different termite species is a well documented phenomenon (Thorne and Haverty, 1991), however transfer of symbionts via necrophagy followed by hindgut colonization has yet to be demonstrated.

Worldwide, the lower termite family Rhinotermitidae contains 14 genera (Krishna 1970). Yamin (1979) listed the flagellate composition of 31 Rhinotermitid species belonging to 7 genera of this family. We investigated the symbiont composition of an additional 25 Rhinotermitid species, including members of one genus, *Parrhinotermes*, not included in Yamin's study.

2. Materials and Methods

Termite species investigated in this study and their localities are shown in Table 1. More than one colony of each species was examined wherever possible. Worker (or pseudergate) gut contents were placed directly on glass slides or fixed with Schaudin's fluid and impregnated with protalgol stain (Honigberg and Davenport, 1951) immediately after the termite collection. Symbionts were observed by Nomarski differential interference or normal light microscopy and identified to the generic level and species level where possible, by comparison with original flagellate descriptions (see Yamin, 1979 and references therein).

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Host species	Localities
Reticulitermes kanmonensis Takematsu (13)	Onoda, Yamaguchi Pref., Japan
R. amamianus (Morimoto) (11)	Amamioshima Is., Japan
R. miyatakei (Morimoto) (9)	Amamioshima Is., Japan
R. okinawanus (Morimoto) (9)	Okinawa Is., Japan
R. yaeyamanus (Morimoto) (19)	Ishigakijima Is., Japan
R. flaviceps (Morimoto) (5)	Taiwan Is., R.O.C.
R. guangzhouensis Ping (1)	Guangzhou, China
R. sp. 1 (1)	Beijing, China
R. sp. 2 (6)	Hongkong Is., Hongkong
R. sp. 3 (6)	Taipo, Hongkong
R. sp. 4 (2)	Taipo, Hongkong
Heterotermes tenuior (Haviland) (3)	Borneo Is., Malaysia
H. sp. 1 (2)	Altagracia, Venezuela
H. sp. 2 (1)	Queensland, Australia
H. sp. 3 (1)	Queensland, Australia
Coptotermes sp. 1 (2)	Altagracia, Venezuela
C. curvignathus Holmgren (1)	Borneo Is., Malaysia
C. sp. 2 (2)	Borneo Is., Malaysia
Schedorhinotermes sp. 1 (1)	Queensland, Australia
S. sarawakensis (Holmgren) (2)	Borneo Is., Malaysia
S. sp. 2 (3)	Borneo Is., Malaysia
Parrhinotermes queenslandicus Mjoberg (2)	Queensland, Australia
P. aequalis (Haviland) (3)	Borneo Is., Malaysia
Prorhinotermes japonicus (Holmgren) (1)	Ranyu Is., R.O.C.
Termitogeton planus (Haviland) (4)	Borneo Is., Malaysia

 Table 1.
 Termite species investigated in this study. Number of investigated termite colonies are shown in the brackets.

3. Results and Discussion

Table 2 shows the protistan genera found in each Rhinotermitid species investigated in this study. A total of 13 symbiont genera (including subgenera) were identified from 7 Rhinotermitid genera. It should be noted that within symbiont genera, species can differ between some host termite species. In individual termite species examined here, a specific flagellate composition (to the species level) was always found when more than one colony was examined. Similar symbiont genera are found within species of a genus. *Reticulitermes*

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Rhinotermitid species	Flagel Order Oxym	Flagellate genera (subgenera Order Oxymonadida Trichomona	ra (subg Order Tricho	a (subgenera) Order Trichomonadida	lida	Order Hypermastigida	permast	igida						
	Di	Py	St	He	Tr	Genus A*	Hoto	Mi	Sptr	Sp	Ho	Ps	Te	Trph
Reticulitermes kanmonensis	+	+	1	+	1		1	+	1	+	+	1	+	+
R. amamianus	+	+	I	+	1	I	ł	+	+	ł	+	1	+	+
R. miyatakei	+	+	1	+	I	1	ł	+	+	+	+	1	+	1
R. okinawanus	+	+	1	+	I	ł	1	+	+	ł	+	I	+	+
R. yaeyamanus	+	+	ł	+	I	1	I	+	I	+	+	E	+	+
flaviceps	+	+	I	+	I	1	I	+	1	+	+	I	+	+
R. guangzhouensis	+	+	I	+	1	ł	1	+	1	+	+	I	+	+
R. sp. 1	+	+	1	+	+	1	ł	+	+	1	+	ŀ	+	1
R. sp. 2	+	+	ł	+	t	I	I	+	T	+	+	ł	+	+
R. sp. 3	+	+	1	I	I	I	I	+	I	+	+	I	+	+
R. sp. 4	+	+	I	+	ł	I	I	+	1	+	+	I	+	+
Heterotermes tenuior	1	ı	I	I	I	I	+	ł	+	I	ł	+	I	i
H. sp. 1	I	ł	I	I	I	1	+	I	+	ł	I	+	ŀ	ł
sp. 2	1	I	ł	I	ł	ł	+	1	+	l	I	+	I	I
sp. 3	I	I	*	+	1	I	+	I	+	ł	I	+	I	I
Coptoternes sp.	1	ł	I	I	I	l	+	F	+	1	I	+	I	I
C. curvignathus	1	ł	I	1	ł	1	+	I	+	I	ł	+	ł	1
sp.	I	1	1	I	ł	ł	+	I	+	I	t	+	1	ı
Schedorhinotermes sp. 1	I	I	+	+	+	1	1	+	+	+	I	I	1	I
S. sarawakensis	I	I	I	+	+	i	1	+	+	+	I	ł	l	ł
sp. 2	I	I	I	+	+	I	t	+	+	+	I	ł	1	ł
Parrhinotermes queenslandicus	1	I	I	I	I	ł	T	I	ł	ł	I	+	1	ŧ
P. aequalis	ł	1	ł	ł	I	I	I	1	I	I	I	+	I	I
Prorhinotermes japonicus	I	ι	I	1	I	+	+	I	+	I	1	+	I	I
Termitogeton planus	ł	I	I	I	I	ı	I	1	I	ł	I	+	1	ŧ

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Flagellate genera	Rhinotermitid	-						
(subgenera)	Reticuli-	Hetero-	Copto-	Schedorhino-	Parrhino-	Psammo-	Prorhino-	Termito-
)	termes	termes	termes	termes	termes	termes	termes	geton
Order Oxymonadida								
inenympha	+	1	I	ł	I	1	I	I
Pyrsonympha	+	ı	ł	1	I	1	I	L
Order Trichomonadida								
Hexamastix	+	1	I	+	ŀ	1	I	i
Monocercomonas	+	١	1	I	I	I	I	I
Tricerocomitus	ł	1	+	ł	I	1	I	F
Stephanonympha	I	T	+	+	1	I	1	I
Trichomonas	+	+	I	+	I	I	1	I
Order Hypermastigida								
enus A	I	1	I	1	I	1	+	l
lolomastigotoides	1	+	+	+	ſ	+	+	I
licrojoenia	+	1	1	+	I	I	ł	ł
pirotrichonympha	+	+	+	+	ł	+	+	I
Spironympha	+	\$	I	+	1	Ľ	£.	١
lolomastigotes	+	Ι	I	ł	I	L	I	I
Pseudotrichonympha	1	+	+	+	+	+	+	+
Teranympha	+	1	ł	ţ	I	I	1	I
		-		1	I	+	I	I

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species examined always contained the flagellate genera Dinenympha, Pyrsonympha, Microjoenia, Holomastigotes and Trichonympha, and the majority of taxa contained Hexamastix, Spironympha and Teranympha. The latter symbiont genus has only been reported in Reticulitermes collected from Asian countries, and is not found in representatives from Europe and America (Yamin, 1979; Kitade and Matsumoto, 1993b). Reticulitermes and Heterotermes are thought to be phylogenetically close due to similarities in morphology, and comprise the subfamily Heterotermitinae (Krishna, 1979). However, symbiont composition does not agree with the apparent affinity between these groups. Rather, Heterotermes and Coptotermes (the sole genus of the subfamily Coptotermitinae) species share common symbiont compositions, both harbouring the protistan genera Holomastigotoides, Spirotrychonympha and Pseudotrichonympha. The three major symbiont genera found in Schedorhinotermes species are Microjoenia, Spironympha' and Pseudotrichonympha. Species of the Rhinotermitid genera Parrhinotermes and Termitogeton contained only one type of flagellate species of the genus Pseudotrichonympha. This phenomenon does not occur in any other lower termite genus examined thus far (Yamin, 1979).

A summary of symbiont composition in Rhinotermitidae incorporating data from this study and that of Yamin (1979) is given in Table 3. It is clear that symbiont species from the genera *Pseudotrichonympha*, *Holomastigotoides* and *Spirotrychonympha* are the most prevalent among Rhinotermitid taxa. In the genera *Heterotermes* and *Coptotermes*, reports of symbionts other than from the aforementioned genera are only fragmentary. In *Schedorhinotermes* species, *Holomastigotoides* has never been reported, but additional genera to *Pseudotrichonympha* and *Spirotrychonypha*, such as *Microjoenia* and *Spironympha* are commonly found in the hindgut fauna. *Reticulitermes* species generally have the most diverse symbiotic communities, harbouring 7–8 protistan genera in their hindguts, but interestingly lack *Holomastigotoides* and *Pseudotriconympha*, which are thought to be major components of the cellulose dissimilation process in *Coptotermes formosanus* (Yoshimura, 1993).

The presence of strikingly different protistan compositions within Rhinotermitid taxa, and particularly between genera which are apparently phylogenetically close (*Reticulitermes* and *Heterotermes*), raises obvious questions about how such a situation arose. Is it possible that these deviations in symbiont composition came about via interspecies aggression, perhaps between taxa of different lower termite families? Or, can these compositions be explained by occasional divergence and/or loss of symbiont species, occurring within hosts which are also undergoing speciation? We are currently undertaking an investigation of phylogenetic relationships between Rhinotermitid species using mitochondrial DNA sequence data. We hope this

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will help to identify patterns of symbiont gain and loss in the evolution of this lower termite family.

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