

Fungal Systematics and Evolution

VOLUME 1 JUNE 2018 PAGES 23–39

doi.org/10.3114/fuse.2018.01.03

Fusarium oligoseptatum sp. nov., a mycosymbiont of the ambrosia beetle Euwallacea validus in the Eastern U.S. and typification of F. ambrosium

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Key words:

Ailanthus altissima
Ambrosia Fusarium Clade
Camellia sinensis
phylogeny
shot-hole borer beetle

Abstract: Fusarium oligoseptatum sp. nov. was isolated from the invasive Asian ambrosia beetle Euwallacea validis (Coleoptera, Scolytinae, Xyleborini) and from the galleries that females had constructed in dying Ailanthus altissima (tree-of-heaven) symptomatic for Verticillium wilt in south-central Pennsylvania, USA. This ambrosia fungus was cultivated by Euwallacea validis as the primary source of nutrition together with a second symbiont, Raffaelea subfusca. Female beetles transport their fungal symbionts within and from their natal galleries in paired pre-oral mycangia. Fusarium oligoseptatum was distinguished phenotypically from the 11 other known members of the Ambrosia Fusarium Clade (AFC) by uniquely producing mostly 1–2 septate clavate sporodochial conidia that were swollen apically. Phylogenetic analysis of multilocus DNA sequence data resolved F. oligoseptatum as a genealogically exclusive species-level lineage but evolutionary relationships with other members of the AFC were unresolved. Published studies have shown that F. oligoseptatum can be identified via phylogenetic analysis of multilocus DNA sequence data or a PCR multiplex assay employing species-specific oligonucleotide primers. In addition, to provide nomenclatural stability, an epitype was prepared from an authentic strain of F. ambrosium that was originally isolated from a gallery constructed in Chinese tea (Camellia sinensis) by E. fornicatus in India, together with its lectotypification based on a published illustration.

Published online: 19 February 2018.

INTRODUCTION

Ambrosia beetles (Coleoptera, Curculionidae: Scolytinae and Platypodinae) are obligate mutualistic mycetophagous insects that cultivate ambrosia fungi as a source of nutrition typically in dead but occasionally in healthy woody hosts (Hulcr & Stelinski 2017). Most ambrosia beetles studied to date carry specific symbiotic ambrosia fungi within their mycangia, which are disseminated by females when they leave their natal galleries to establish new colonies (Hulcr & Cognato 2010, Hulcr & Dunn 2011). Genera in the tribe Xyleborini (Scolytinae) are considered to be the most ecologically successful ambrosia beetles (Hulcr & Stelinski 2017). Several well-studied fungusfarming beetles, including representatives of several tribes, have recently caused significant mortality of trees. Notable examples include the invasive Asian ambrosia beetle Xyleborus glabratus and its nutritional symbiont Raffaelea lauricola on redbay (Persea borbonia) in the southeastern United States (Fraedrich et al. 2008), and Platypus quercivorus and its symbiont Raffaelea quercivora on Japanese oak (Quercus serrata and Q. mongolica var. grosseserrata) in Japan (Kubono & Ito 2002, Seo et al. 2012).

Compared to their beetle partners, relatively few fungal symbionts have been formally described. Most of the ambrosia fungi described to date are ascomycetous fungi in the *Ophiostomatales*, including members of *Afroraffaelea*, *Ceratocystiopsis*, *Dryadomyces* and *Raffaelea* (von Arx & Hennebert 1965, Upadhyay & Kendrick 1975, Gebhardt *et al.* 2005, Harrington *et al.* 2008, 2010, Alamouti *et al.* 2009, Dreaden *et al.* 2014, Bateman *et al.* 2016, Hulcr & Stelinski 2017). The *Microascales* also include multiple groups of ambrosia fungi, some of which are important and widespread: *Ambrosiella*, *Meredithiella*, and *Phialophoropsis* (Mayers *et al.* 2015). Less common are symbionts belonging to the *Polyporales* (Li *et al.* 2015, Kasson *et al.* 2016, Simmons *et al.* 2016), *Hypocreales* (i.e., *Geosmithia*) (Kolařik & Hulcr 2009, Kolařik & Kirkendall 2010), and *Saccharomycetales* (van der Walt 1972, Hulcr & Stelinski 2017).

In addition to the symbionts mentioned above, Fusarium ambrosium (Hypocreales, Nectriaceae) is cultivated by Euwallacea fornicatus (formerly Xyleborus fornicatus) as a source of nutrition (Gadd & Loos 1947, Norris & Baker 1967, Brayford 1987, Nirenberg 1990). The taxonomic history of F. ambrosium, however, is complicated because the species was originally misclassified and established in Monacrosporium, as M. ambrosium. This fungus

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was isolated originally and described from galleries of the tea shot-hole borer, E. fornicatus, in Camellia sinensis (Chinese tea) and Ricinus communis (caster-oil tree) stems in Sri Lanka (Gadd & Loos 1947). Subsequently, F. bugnicourtii was described based on collections from galleries in Chinese tea in India, borer-damaged Hevea brasiliensis (rubber tree) and Theoborma cacao (cacao) in Sabah, Malaysia (Brayford 1987). Nirenberg (1990) synonymized F. bugnicourtii with M. ambrosium and recombined the latter as F. ambrosium based on nomenclatural priority. Brayford (1987) considered F. bugnicourtii to be conspecific with F. tumidum var. coeruleum (Bugnicourt 1939), but distinct from F. tumidum. Although the type of F. tumidum var. coeruleum based on a collection from H. brasiliensis appears to be phylogenetically distinct from F. bugnicourtii, the holotype of F. bugnicourtii selected by Brayford (IMI 296597 = NRRL 20438) is conspecific with F. ambrosium (Kasson et al. 2013).

Kasson et al. (2013) conducted an extensive multilocus molecular phylogenetic study on the ambrosial fusaria, based on isolates from beetles, their galleries, or from trees showing extensive borer damage and dieback. These included Camellia sinensis, Persea americana (avocado), Ailanthus altissima (treeof-heaven), Acer negundo (box elder), and Hevea brasiliensis from natural and cultivated ecosystems, and avocado in the United States, Israel and Australia. Seven different Fusarium species lineages were reported to be associated with Euwallacea ambrosia beetles within the Ambrosia Fusarium Clade (AFC) and one other species (i.e., Fusarium sp. AF-9) with Xyleborus ferrugineus in Costa Rica. The monophyletic AFC is nested within Clade 3 of the F. solani species complex (FSSC; O'Donnell 2000), which contains 60 plus phylogenetic species based on genealogical concordance phylogenetic species recognition (GCPSR; Taylor et al. 2000). The AFC comprises two strongly supported clades: the four species within Clade A typically produce curved fusiform septate macroconidia, which are typical of Fusarium, whereas nine of the 10 species within Clade B produce clavate macroconidia (Kasson et al. 2013, Aoki et al. unpubl.), described as 'dolphin-shaped' by Brayford (1987). O'Donnell et al. (2015) conducted a multilocus phylogenetic analysis of the AFC and Euwallacea and found evidence of repeated host shifts rather than strict co-evolution of this mutualism.

Freeman et al. (2013) described a new species, F. euwallaceae, based on isolates corresponding to the ambrosia species symbiotic with the Euwallacea sp. #1 sensu O'Donnell et al. (2015), which causes serious damage to avocado production in Israel and California, USA (Mendel et al. 2012, Eskalen et al. 2013). Fusarium euwallaceae is closely related morphologically to F. ambrosium, but it can be distinguished from the latter by the abundant production of bluish to brownish sporodochial conidia that form greenish masses on PDA after 1 mo in culture, together with hyaline conidia. To date only three of the 16 species within the AFC have been described formally (Kasson et al. 2013, O'Donnell et al. 2015, Na et al. 2018). Similar to E. validus, Euwallacea sp. #1 also carries additional symbiotic fungi, Graphium euwallaceae and Paracremonium pembeum (Freeman et al. 2016, Lynch et al. 2015). Recently, PCR multiplexes were developed to discriminate Fusarium symbionts of invasive Euwallacea ambrosia beetles that inflict damage on numerous tree species throughout the United States, including F. euwallaceae and F. kuroshium along with four unnamed AFC species-level lineages: AF-3, AF-4, AF-6 and AF-8 (Short et al. 2017). One of the undescribed species, which was informally referred to as Fusarium sp. AF-4, is cultivated by the ambrosia

beetle E. validus primarily in Verticillium wilt-stressed and dying stands of A. altissima, as well as from Verticillium wiltstressed Acer pensylvanicum (striped maple), Aralia spinosa (devils walkingstick) and Rhus typhina (staghorn sumac) in south-central Pennsylvania, USA (Schall & Davis 2009, Kasson et al. 2013, 2015). In the present study, this species is described as F. oligoseptatum sp. nov. based on a comparison with F. ambrosium (AF-1) and F. euwallaceae (AF-2) (Kasson et al. 2013, Freeman et al. 2013). In addition, because type material for F. ambrosium was not designated (Gadd & Loos 1947), or appears to have been lost (Nirenberg 1990), a line-drawing of a clavate conidium of the species from Gadd & Loos (1947) was selected as the lectotype. Furthermore, an epitype was prepared from an authentic strain of this species to stabilize its taxonomy, according to the International Code of Nomenclature for algae, fungi and plants (ICN, the Melbourne Code; McNeill et al. 2012).

MATERIALS AND METHODS

Fungal isolates and type specimens

Fusarium strains examined in this study (Table 1) are stored in the Agriculture Research Service Culture Collection (NRRL), National Center for Agricultural Utilization Research (NCAUR), U.S. Department of Agriculture in Peoria, Illinois, USA. These strains were originally isolated from Euwallacea ambrosia beetles and their galleries, or from host trees showing extensive borer damage (Kasson et al. 2013). The Pennsylvanian strains of F. oligoseptatum were isolated from E. validus ambrosia beetles that had colonized A. altissima. Beetles were surface disinfested for 15 s in 70 % ethanol and then washed three times in sterile deionized water. Whole beetles or their heads were macerated using sterile Tenbroek homogenizers (Pyrex, Corning, NY), or pellet pestles (Fisher Scientific, Hampton, NH), suspensions were diluted 1:10 and 1:100, and then spread evenly over half-strength Potato Dextrose Agar (PDA, BD-Difco™, Thermo Fisher Scientific, Waltham, MA) amended with 100 ppm streptomycin sulfate (Sigma-Aldrich, St. Louis, MO) as described in Kasson et al. (2013). Other related ambrosia fusaria or close relatives within the F. solani species complex (O'Donnell et al. 2008) were obtained from culture collections (Table 1). Isolates used in this study are available upon request from NRRL (http://nrrl.ncaur.usda. gov/cgi-bin/usda/), NARO Genebank, Microorganisms Section (MAFF), Genetic Resources Center, National Agriculture and Food Research Organization, Tsukuba, Ibaraki, Japan (http://www.gene. affrc.go.jp/about-micro en.php), and the Westerdijk Institute (formerly CBS-KNAW Fungal Biodiversity Center), Utrecht, the Netherlands (http://www.westerdijkinstitute.nl/). Isolates of four novel Taiwanese AFC species discovered very recently, i.e. AF-13 to AF-16 (Na et al. 2018), were not included in this study.

Holotype and epitype specimens newly prepared from the selected strains were deposited in BPI, US National Fungus Collection (https://nt.ars-grin.gov/fungaldatabases/specimens/specimens.cfm).

Incidence of Fusarium oligoseptatum and other fungi from Euwallacea validus mycangia across recently confirmed tree hosts

Mycangial fungal communities were characterized as previously described by Kasson et al. (2013) for adult female



Table 1. Strains of Ambrosia Fusarium Clade (AFC) species examined in present study. Bold text is used to identify ex-holotype of Fusarium oligoseptatum and ex-epitype of F. ambrosium.

Species	AFC clade #	NRRL strain #ª	equivalent nos. in other collections ^b	Host beetle °	Host plant	Origin	Country	Date	Collector	Remark
F. oligoseptatum	AF-4	62578	FRC S-2576	Euwallacea validus	Ailanthus altissima	Dauphin Co., Pennsylvania	USA	30-Jan-10	M. Kasson Bh17	Studied only phylogenetically
	AF-4	62579	FRC S-2581	Euwallacea validus	Ailanthus altissima	Dauphin Co.,	USA	30-Jan-10	M. Kasson Bh24	Ex-HOLOTYPE
			= MAFF 246283			Pennsylvania				
			= CBS 143241							
	AF-4	62580	FRC S-2594	Euwallacea validus	Ailanthus altissima	Franklin Co.,	USA	9-Mar-10	M. Kasson Ch19	
			= MAFF 246284			Pennsylvania				
			= CBS 143242							
	AF-4	62581	FRC S-2616	Euwallacea validus	Ailanthus altissima	Huntington Co.,	USA	27-Feb-10	M. Kasson Dh24	
			= MAFF 246285			Pennsylvania				
			= CBS 143243							
	AF-4	62582	FRC S-2627	Euwallacea validus	Ailanthus altissima	Mifflin Co., Pennsylvania	USA	1-Jul-09	M. Kasson Eh11	Degenerated
			= MAFF 246286							strain
			= CBS 143244							
F. ambrosium	AF-1	20438	IMI 296597	Euwallacea	Camellia sinensis	Chinchona,	India	17-Jul-85	anonymous	Ex-holotype of F.
			= MAFF 246291	fornicatus		Maharashtra				bugnicourtii
	AF-1	22345	BBA 65389	Euwallacea	Camellia sinensis	Upari Tea Institute	India	9-May-90	V. Agnihothrudu	
			= MAFF 246288	Jornicatus						
	AF-1	22346	BBA 65390	Euwallacea	Camellia sinensis	Upari Tea Institute	India	9-Мау-90	V. Agnihothrudu	Ex-EPITYPE
			= CBS 571.94	Jornicatus						
			= MAFF 246287							
	AF-1	36510	BBA 65390	Euwallacea	Camellia sinensis	Upari Tea Institute	India	9-Мау-90	V. Agnihothrudu	Duplicate of
			= MAFF 246289	fornicatus						NRRL 22346
	AF-1	46583	IMI 339338	Euwallacea	Camellia sinensis	High Forest Estate,	India	26-Mar-90	anonymous	Received as F.
			= MAFF 246290	Jornicatus		Anamaliais, Coimbatore District, Tamil Nadu				bugnicourtii



Table 1. (Continued).

Species	AFC clade #	NRRL strain #ª	Equivalent nos. in other collections ^b	Host beetle °	Host plant	Origin	Country	Date	Collector ^d	Remark
	AF-1	62605		Euwallacea fornicatus	Camellia sinensis	Tea Research Institute of Sri Lanka, St. Coombs, Talawakelle	Sri Lanka	Mar-12	S. Freeman 31.14	
F. euwallaceae	AF-2	54727	MAFF 243816 = CBS 135859	Euwallacea sp. #1	Persea americana	Volcani	Israel	17-Feb-10	S. Freeman 5-4	
	AF-2	62626		Euwallacea sp. #1	Persea americana	California	USA		A. Eskalen 1854	
Fusarium sp.	AF-3	62606		Euwallacea interjectus	Acer negundo	Gainesville, Florida	USA		J.A. Smith PL1499	
	AF-3	62628		Euwallacea interjectus	Acer negundo	Gainesville, Florida	USA		J.A. Smith 1190	
	AF-5	22231	IMI 110107	unknown	Hevea brasiliensis	Agriculture Research Centre Tuaran, Sabah, Borneo	Malaysia	19-Nov-64		Received as <i>F.</i> bugnicourtii
	AF-5	46518	FRC S-2075	unknown	Hevea brasiliensis		Malaysia			
	AF-6	62590		Euwallacea sp. #2	Persea americana	Miami, Florida	USA		R.C. Ploetz AF9	
	AF-6	62591		Euwallacea sp. #2	Persea americana	Miami, Florida	NSA		R.C. Ploetz AF10	
	AF-7	62610		Euwallacea sp. #3	Persea americana	Queensland	Australia		A.D.W. Geering 1	
	AF-7	62611		Euwallacea sp. #3	Persea americana	Queensland	Australia		A.D.W. Geering 2	
	AF-8	62584		Euwallacea sp. #2	Persea americana	Miami, Florida	NSA		R.C. Ploetz Amb2	
	AF-8	62585		Euwallacea sp. #2	Persea americana	Miami, Florida	NSA		R.C. Ploetz AF4	
	AF-9	22643	ATCC 44215	Xyleborus ferrigineus			Costa Rica		E.B. Smalley	
	AF-9	88099		unknown	Delonix regia	Florida	NSA			
	AF-10	62941	IMI 351954	unknown			Singapore			
	AF-11	62943		Euwallacea sp. #4	Camellia sinensis		Sri Lanka		P. Liyanage	
	AF-11	62944		Euwallacea sp. #4	Camellia sinensis		Sri Lanka		P. Liyanage	
F. kuroshium	AF-12	62945		Euwallacea sp. #5	Platanus racemosa	San Diego, CA	USA		A. Eskalen	
	AF-12	62946		Euwallacea sp. #5	Platanus racemosa	San Diego, CA	USA		A. Eskalen	

Table 1. (Continued)

			Equivalent							
Species	AFC clade #	AFC clade # NRRL strain # collections b	collections	Host beetle $^\circ$	Host plant	Origin	Country Date		Collector	Remark
a NRRL: ARS Cultu	re Collection, I	NCAUR-ARS-USDA	NRRL: ARS Culture Collection, NCAUR-ARS-USDA, Peoria, IL, USA.							
^b ATCC: American	Type Culture (Collection, Manas	sas, VA, USA; BBA	: Biologische Bundesa	ınstalt für Land- und	batc: American Type Culture Collection, Manassas, VA, USA; BBA: Biologische Bundesanstalt für Land- und Forstwirtschaft, Institut für Mikrobiologie (currently Julius-Kühn-Institut), Berlin, Germany:	ıt für Mikrobiologie	(currently Jul	lius-Kühn-Institut), E	serlin, Germany;

CBS: Westerdijk Institute (formerly CBS-KNAW Fungal Biodiversity Center), Utrecht, the Netherlands; FRC: Fusarium Research Center, The Pennsylvania State University, State College, PA, USA; IMI: CABI UK Centre, Egham, Surrey, UK; MAFF: NARO Genebank, Microorganisms Section, Genetic Resources Center, National Agriculture and Food Research Organization, Tsukuba, Ibaraki, Japan.

Weed Research, Agricultural Research Organization, The Volcani Center, Bet Dagan, Israel; A. Geering, University of Queensland, Brisbane, Australia; M. Kasson, Division of Plant and Soil Sciences, West d V. Agnihothrudu, Upari Tea Institute, India; A. Eskalen, Department of Plant Pathology and Microbiology, University of California, Riverside, CA, USA; S. Freeman, Department of Plant Pathology and Virginia University, Morgantown, WV, USA; R. C. Ploetz, University of Florida, Homestead, FL, USA; J. A. Smith, University of Florida, Gainesville, FL, USA The five unnamed Euwallacea spp. are distinguished by #1—#5 (O'Donnell et al. 2015). AF-12 was later described as F. kuroshium (Na et al. 2018).

beetles that harbor paired pre-oral mycangia. In addition to F. oligoseptatum, described in this paper, previous studies revealed that E. validus harbors a second less abundant symbiont, Raffaelea subfusca (Kasson et al. 2013). To determine if these trends held across other tree hosts and geographic locations, E. validus from Ailanthus and 16 other confirmed tree hosts across seven states were sampled. When available, a minimum of 10 adult females were included. Logextracted females were processed as previously described (Kasson et al. 2013). Following serial dilution plating of head macerates, individual fungal colony forming units (CFUs) were quantified by morphotype and representatives of each morphotype retained for molecular characterization using the ITS barcoding gene. Unlike the other fungal morphotypes, representative fusaria were subjected to an AF-3 / Fusarium oligoseptatum (AF-4) multiplex PCR recently developed by Short et al. (2017) to discriminate known Fusarium symbionts of E. interjectus and E. validus in the eastern U.S.

When comparing the CFUs recovered from individual beetle heads between the primary symbionts of E. validus, F. oligoseptatum and R. subfusca, a chi-squared test was performed across all tree species. To examine if there were differences in the relative amount of colony forming units (CFUs) recovered from individual beetle heads between the primary symbionts of E. validus, F. oligoseptatum and R. subfusca within individual species, a second chi-squared test was performed for each individual species. Results of the tests were deemed significant if p < 0.05.

Molecular systematics and biology

Methods for culturing mycelium, DNA extraction, PCR amplification, DNA sequencing and phylogenetic analyses followed published protocols (Kasson et al. 2013, O'Donnell et al. 2015). DNA sequence data included in this study were deposited in GenBank as JQ038007-JQ038034.

Phenotypic characterization

Strains were grown on PDA and synthetic low-nutrient agar (SNA; Nirenberg 1990, Nirenberg & O'Donnell 1998) in the dark, under continuous black light (Black light blue fluorescent tubes, FL8BL-B 8W/08, Panasonic, Osaka, Japan), or under an ambient daylight photoperiod. Strains were cultured on PDA in 9 cm Petri dishes at 20 °C in the dark to characterize colony color, odor and morphology. Kornerup & Wanscher (1978) was used as the color standard. PDA cultures were also used for determining mycelial growth rates in the dark at eight temperatures (5-40 °C) at 5 °C increments (Aoki et al. 2015). Culture plates were examined at 1 and 4 d post inoculation, and radial growth was calculated as arithmetic mean values per day by measuring 16 radii around the colony. Measurements of growth rate at different temperatures were replicated twice, and the data averaged for each strain. Cultures on SNA were used for examination of microscopic characters as described by Aoki et al. (2015). Conidia and conidiophores were examined in water mounts after culturing on SNA under continuous black light. Phenotypic characters were compared with data from the related AFC species, F. ambrosium (published as Monacrosporium ambrosium; Gadd & Loos 1947), F. bugnicourtii (synonymized as F. ambrosium; Brayford 1987, Nirenberg 1990), and F. euwallaceae (Freeman



et al. 2013). To compare the number of conidial septa in strains of *F. oligoseptatum* and *F. ambrosium*, they were incubated on SNA at 25 °C under continuous black light for one to two weeks and the number septa in the clavate sporodochial conidia were counted.

RESULTS

Incidence of Fusarium oligoseptatum and other fungi from Euwallacea validus mycangia across recently confirmed tree hosts

Mycangial communities were characterized from adult female beetles extracted from sixteen native host trees and Ailanthus (Fig. 1). Overall, F. oligoseptatum and Raffaelea subfusca comprised 84 % of all fungal CFUs from female heads across all plant hosts with F. oligoseptatum yielding significantly more CFU's (10 992) compared to R. subfusca (7 014; p < 0.0001). The remainder included miscellaneous yeasts and other fungi including Paracremonium sp. and a putatively novel Graphium sp. (Freeman et al. 2016, Lynch et al. 2016), as well as a variety of singleton taxa that were not further characterized. Incidence of the two primary symbionts from the heads of female E. validus was compared across and within plant hosts. Overall, significant differences were detected across hosts indicating that the relative proportion of the two symbionts varied across hosts with a majority of beetles from a majority of plant hosts yielding higher counts of F. oligoseptatum (Fig. 1). Of these, beetles from 11 of the 16 plant hosts, including Ailanthus, had significantly higher total CFU counts of F. oligoseptatum compared to R. subfusca. Only five species had a mean percent incidence of F. oligoseptatum below 50 %: Fraxinus americana (white ash), Pinus virginiana (Virginia pine), Populus grandidentata (bigtooth aspen), Quercus montana (chestnut oak), and Amelanchier arborea

(serviceberry) (Fig. 1). Of these, white ash, Virginia pine, and chestnut oak had significantly higher total CFU counts of *R. subfusca* compared to *F. oligoseptatum* (Fig. 1).

Molecular phylogenetics

A 4914 bp 31-taxon 4-locus dataset was constructed that included the internal transcribed spacer region and domains D1 and D2 of the nuclear ribosomal large subunit (ITS+LSU rDNA: 1004 bp alignment, 26 parsimony informative characters (PIC)), and portions of translation elongation factor 1- α (*TEF1*: 687 bp alignment, 53 PIC), DNA-directed RNA polymerase II largest (RPB1: 1588 bp alignment, 164 PIC) and second largest subunit (RPB2: 1635 bp alignment, 165 PIC) from 12 AFC species. Molecular phylogenetic analyses were conducted using maximum parsimony (MP) with PAUP v. 4.0b10 (Swofford 2003) and maximum likelihood (ML) with GARLI 2.01 (Zwickl 2006). Sequences of Fusarium neocosmosporiellum (≡ Neocosmospora vasinfecta; Geiser et al. 2013) NRRL 22468 and 43467 were used to root the phylogenies based on more inclusive analyses (O'Donnell et al. 2013). Fusarium oligoseptatum (AF-4) was poorly supported (MP and ML bootstrap = 63-56 %) as sister to a clade that included F. euwallaceae (AF-2) from Israel and California and Fusarium sp. (AF-3) from Florida. With the exception of F. ambrosium, the other AFC species represented by two or more strains received moderate to strong monophyly bootstrap support. The putative triparential hybrid strain F. ambrosium NRRL 62605 from Sri Lanka (Kasson et al. 2013), however, did not form a genealogically exclusive group with F. ambrosium from India. As reflected by poor bootstrap support along the backbone on the phylogeny, relationships among the species were generally unresolved (Fig. 2). The analyses did support monophyly of the AFC and the early diverging subclades designated Clade A and B (Fig. 2).

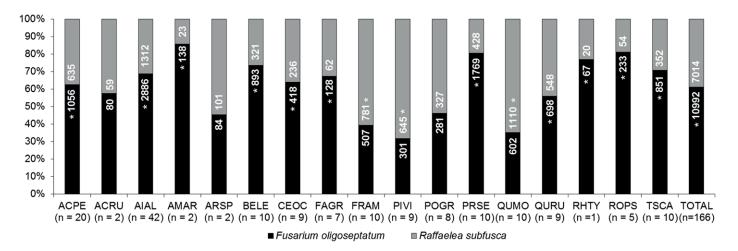


Fig. 1. Total number of *F. oligoseptatum* and *Raffaelea subfusca* CFUs recovered from macerated heads of adult female *E. validus* from 17 tree hosts. A significant difference between the two fungal CFUs within a specific host is indicated with an asterisk on the side corresponding to the higher count. Host plant IDs are based on USDA PLANTS Database (https://plants.usda.gov/java/) abbreviations, which are derived from the first two letters of the genus and species of the Latin binomial. Abbreviations are as follows: ACPE: *Acer pensylvanicum*, ACRU: *Acer rubrum*, AIAL: *Ailanthus altissima*, AMAR: *Amelanchier arborea*, ARSP: *Aralia spinosa*, BELE: *Betula lenta*, CEOC: *Celtis occidentalis*, FAGR: *Fagus grandifolia*, FRAM: *Fraxinus americana*, PIVI: *Pinus virginiana*, POGR: *Populus grandidentata*, PRSE: *Prunus serotina*, QUMO: *Quercus montana*, QURU: *Quercus rubra*, RHTY: *Rhus typhina*, ROPS: *Robinia pseudoacacia*, and TSCA: *Tsuga canadensis*.



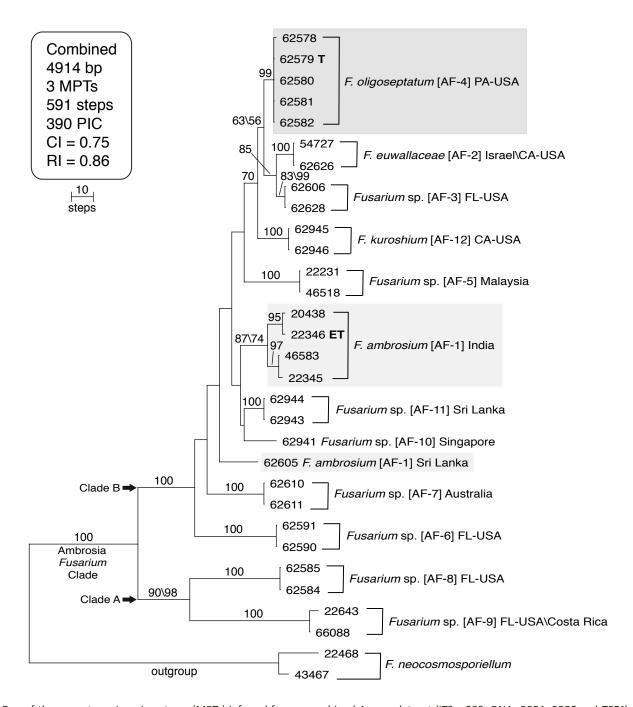


Fig. 2. One of three most-parsimonious trees (MPTs) inferred from a combined 4-gene dataset (ITS + 28S rDNA, *RPB1*, *RPB2* and *TEF1*) comprising 4914 bp of aligned DNA sequence data. The phylogram was rooted on sequences of *Fusarium neocosmosporiellum* NRRL 22468 and 43367 based on more inclusive analyses (O'Donnell *et al.* 2013). The 12 species within the Ambrosia *Fusarium* Clade (AFC) are identified as AF-1 through AF-12 using an ad hoc nomenclature (Kasson *et al.* 2013). Two early diverging monophyletic sister clades are identified as Clade A and B. Numbers above nodes represent maximum parsimony (MP) and maximum likelihood (ML) bootstrap based on 1000 pseudoreplicates of the data (MP-BS\ML-BS). The ML-BS value is only shown when it differed by ≥5 % of the MP-BS value. Note that 10 of the 11 AFC species with two or more strains received strong monophyly bootstrap support. However, *Fusarium ambrosium* highlighted in light gray was resolved as non-monophyletic because the interspecific hybrid strain NRRL 62605 from Sri Lanka did not group with the other strains of this species. The five strains of *F. oligoseptatum*, formally described herein, are identified using dark gray highlight. CI, consistency index; ET, ex-epitype; PIC, parsimony informative characters; RI, retention index; T, ex-holotype.

TAXONOMY

Fusarium oligoseptatum T. Aoki, M.T. Kasson, S. Freeman, Geiser & O'Donnell, *sp. nov.* MycoBank MB822305. Figs 3–5.

Etymology: oligo- + septatum, based on frequent production of sporodochial conidia with only 1–2 septa.

Diagnosis: Distinguished from F. ambrosium and F. euwallaceae by three times as many 0–2-septate sporodochial conidia (i.e., ≥75 % compared with values less than 25% in F. ambrosium and F. euwallaceae).

Type: **USA**, Pennsylvania: Dauphin Co., a dried specimen from a culture of NRRL 62579, isolated from a live female ambrosia beetle, Euwallacea validus, extracted from a gallery in a tree-



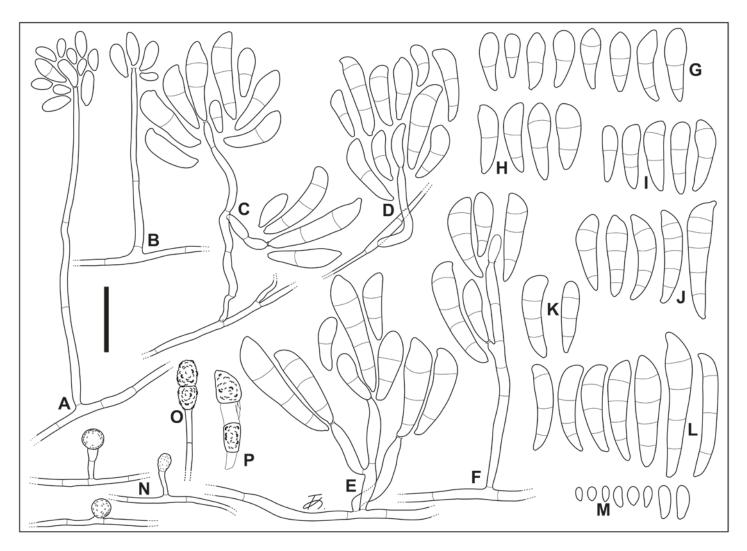


Fig. 3. Fusarium oligoseptatum cultured on SNA under black light. **A, B.** Tall and short aerial conidiophores forming mostly 0-septate conidia. **C, D.** Short aerial conidiophores forming conidia with relatively few septa. **E, F.** Sporodochial conidiophores forming clavate conidia with different septation. **G–L.** Septate conidia formed on sporodochial conidiophores, with some large clavate conidia (G–I: Short clavate conidia with 1–2 septa). **M.** 0-septate conidia formed on aerial conidiophores. **N–P.** Chlamydospores in hyphae (N, O) and in conidia (P). A–D, G, J, N from NRRL 62579 (exholotype); E, H, L, P from NRRL 62580; F, I, K, M, O from NRRL 62581. Bar = 25 μm.

of-heaven, *Ailanthus altissima*, 30 Jan. 2010, *Matthew T. Kasson (Kasson Bh24)* (BPI 910525 – holotype, designated in this study; NRRL 62579 = FRC S-2581 = MAFF 246283 = CBS 143241 – exholotype cultures).

Additional strains examined: **USA**: Pennsylvania: Franklin Co., isolated from a live E. validus female infesting an A. altissima tree, 9 Mar. 2010, Matthew T. Kasson (Kasson Ch19) (NRRL 62580 = FRC S-2594 = MAFF 246284 = CBS 143242); Pennsylvania: Huntingdon Co., isolated from a live E. validus female infesting an A. altissima tree, 27 Feb. 2010, Matthew T. Kasson (Kasson Dh24) (NRRL 62581 = FRC S-2616 = MAFF 246285 = CBS 143243; Pennsylvania: Mifflin Co., isolated from a live E. validus female infesting an A. altissima tree, 1 July 2009, Matthew T. Kasson (Kasson Eh11) (NRRL 62582 = FRC S-2627 = MAFF 246286 = CBS 143244; morphologically degenerated strain).

Description: Colonies on PDA showing radial mycelial growth rates of 2.2–3.6 mm/d at 20 °C and 3.3–4.6 mm/d at 25 °C in the dark. Colony color on PDA white (1A1) to yellowish-white (4A2) or orange white (5A2) in the dark, white (1A1) to yellowish-white (3–4A2) or pale yellow (3–4A3) under black light. Aerial mycelium white (1A1), sparsely formed or floccose in the dark,

more abundantly formed and covering entire surface of colonies under black light. Colony margin entire to undulate. Reverse pigmentation absent or yellowish-white (3-4A2) or pale yellow (3-4A3) in the dark and under black light. Exudates absent. Odor absent, or slightly moldy or sweet in some strains. Hyphae on SNA 1.5-7.5 µm wide. Chlamydospores present but formation delayed in hyphae and in septate sporodochial conidia, mostly subglobose to round ellipsoidal, intercalary or terminal, mostly single, sometimes in chains, ordinary hyaline to very pale-yellow, wall smooth or often minutely roughened, $6-23.5 \times 4.5-9 \mu m$. Sclerotia absent. Sporulation on SNA and PDA generally rapid and abundant under black light, delayed in the dark, sometimes less sporulation on PDA in the dark; light-colored on SNA and PDA under black light or under daylight; sporodochia formed sparsely on SNA, rare on PDA. Aerial conidiophores formed abundantly on SNA under black light, less frequently in the dark, erect, short or tall and narrow, mostly unbranched, rarely branched sparsely, up to 130 µm long, 3-5.5 µm wide at base, thin-walled, forming monophialides integrated in the apices. Phialides on aerial conidiophores simple, subcylindrical to subulate, tapering towards apex, often with a minute collarette at the tip, 10-62.5 × 2.5-5.5 µm. Aerial conidia mostly (1) elliptical, oblong-elliptical,



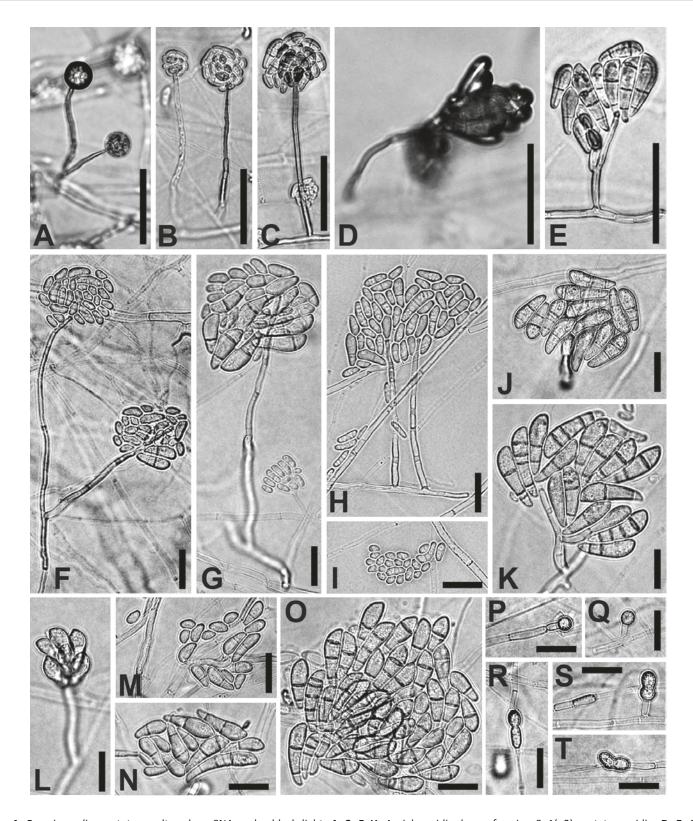


Fig. 4. *Fusarium oligoseptatum* cultured on SNA under black light. **A–C, F–H.** Aerial conidiophores forming 0–1(–2)-septate conidia. **D, E, J–L.** Sporodochial conidiophores forming clavate conidia, often swollen apically with 1–4 septa. **I.** Round to obovate 0-septate aerial conidia. **M–O.** Mostly clavate conidia formed on sporodochial conidiophores, including some 0-septate that are oblong or short-clavate. **P–T.** Chlamydospores formed in hyphae. A–I, K, M, N, Q–S from NRRL 62579 (ex-holotype); J, O, P, T from NRRL 62580; L from NRRL 62581. (A–E: Aerial view without a cover slip; F–T: Mounted in water with a cover slip). Bars: A–E = 50 μm, F–T = 20 μm.

fusiform-elliptical to short clavate, occasionally reniform, some obovate to subglobose, 0–1(–2)-septate; 0-septate on SNA in the dark: 3–13 \times 2–5.5 μm in total range, 5.3–8.5 \times 2.8–3.9 μm on average [ex type (NRRL 62579): 3.5–12 \times 2–4 μm in total range, 6.9±2.0 \times 2.8±0.5 μm on average ± S.D.]; 0-septate on SNA under black light: 3–17 \times 2–6.5 μm in total range, 6.0–9.0 \times 2.8–3.8

 μm on average [ex type (NRRL 62579): 4–17 \times 2.5–6.5 μm in total range, 9.0±2.8 \times 3.8±0.9 μm on average ± S.D.]; 1-septate on SNA under black light: 7.5–26 \times 2.5–8 μm in total range, 14.5–15.3 \times 4.6–4.9 μm on average [ex type: 10.5–21.5 \times 2.5–6 μm in total range, 15.1±2.7 \times 4.6±0.7 μm on average ± S.D.]; sometimes with (2) larger, falcate to clavate, or curved cylindrical,



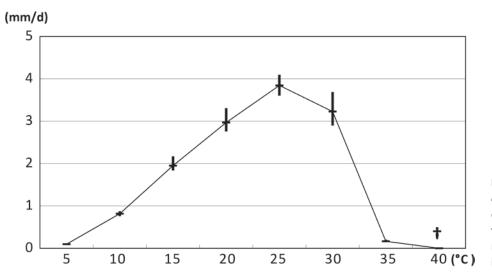


Fig. 5. Radial mycelial growth rate of *Fusarium oligoseptatum* per day on PDA cultured at eight different temperatures. Thick horizontal and vertical bars indicate means and total ranges, respectively, of the 4 isolates analyzed. All isolates failed to grow and died at 40 °C.

(1-)2(-3)-septate conidia, morphologically continuous with falcate sporodochial conidia. Sporodochial conidiophores generally shorter and thicker than aerial conidiophores, unbranched or sometimes sparsely branched, contorted, forming monophialides integrated apically, 20–145 \times 3–6 μ m, or sometimes adelophialides. Sporodochial phialides simple, subulate, lanceolate or subcylindrical, often with a conspicuous collarette at the tip, 9.5-44 × 2.5-5.5 µm. Sporodochial conidia hyaline, mostly falcate to long clavate, sometimes curved cylindrical, often swollen slightly or conspicuously in their upper part, tapering towards base, often with a rounded or papillate apical cell, and an indistinct foot-like or rounded basal cell, (0-)1-3(-5)-septate; swollen conidia sometimes 'dolphin-like' (Brayford 1987) or comma-shaped when 1- to 2-septate, formed on SNA frequently under black light, less frequently in the dark, very rarely formed on PDA under black light; 1-septate on SNA under black light: $11-32.5 \times 4-10 \mu m$ in total range, 18.2-20.1 \times 6.1–7.2 μ m on average [ex type: 13.5–30 \times 4.5–8.5 μ m in total range, $19.3\pm3.9 \times 6.2\pm1.0 \mu m$ on average \pm S.D.]; 2-septate on SNA under black light: 15.5-39.5 × 5.5-12 μm in total range, $24.8-26.5 \times 7.4-8.0 \ \mu m$ on average [ex type: $22-34 \times 5.5-12$ μ m in total range, 26.1 \pm 3.2 × 7.4 \pm 1.2 μ m on average \pm S.D.]; 3-septate on SNA under black light: $20-60 \times 5.5-12.5 \mu m$ in total range, $31.4-35.7 \times 8.3-8.7 \mu m$ on average [ex type: 23-60] \times 5.5–12.5 μ m in total range, 35.7 \pm 7.2 \times 8.7 \pm 1.3 μ m on average \pm S.D.]; 4-septate on SNA under black light: 28.5–67.5 \times 7–11 μm in total range, 38.7–47.9 × 8.8–8.9 μm on average [ex type: $28.5-67.5 \times 7-10 \,\mu\text{m}$ in total range, $47.9 \times 8.9 \,\mu\text{m}$ on average]. Together with multiseptate sporodochial conidia, often forming (0-)1(-2)-septate, oblong to naviculate or short-clavate, straight or curved conidia with a rounded apex and truncate base.

Clade-based diagnosis: Distinguished by phylogenetic analysis of multilocus DNA sequence data (Kasson *et al.* 2013, O'Donnell *et al.* 2015).

Substrates or hosts: All ex-holotype and authentic strains were isolated from *E. validus* in the galleries of *A. altissima* in Pennsylvania (PA), USA. *Fusarium oligoseptatum* has also been confirmed using multilocus sequence typing from Ohio (OH), Virginia (VA) and Maryland (MD) (O'Donnell *et al.* 2015), and from Tennessee (TN) and (West Virginia) WV using the AF-3 / *F. oligoseptatum* (AF-4) multiplex PCR assay (Short *et al.* 2017). Currently known from 16 additional plant hosts, all of which

have been confirmed molecularly as *F. oligoseptatum* using AF-3 / *F. oligoseptatum* (AF-4) multiplex PCR (Short et al. 2017): *Acer pensylvanicum* (PA, USA), *Acer rubrum* (PA, USA), *Amelanchier arborea* (VA, USA), *Aralia spinosa* (PA, USA), *Betula lenta* (PA, USA), *Celtis occidentalis* (WV, USA), *Fagus grandifolia* (OH, USA), *Fraxinus americana* (WV, USA), *Populus grandidentata* (PA, USA), *Prunus serotina* (GA, USA), *Quercus montana* (PA, USA), *Quercus rubra* (PA, USA), *Rhus typhina* (PA, USA), *Robinia pseudoacacia* (PA, USA), *Tsuga canadensis* (OH, USA), and *Pinus virginiana* (VA, USA).

Distribution: Presently confirmed from GA (Georgia), MD, OH, PA, TN, VA, and WV, USA.

Notes: Morphological data on sporodochial conidia was based mainly on NRRL 62579, 62580 and 62581. Strain NRRL 62582 appears degenerated and produced only 1-septate sporodochial conidia after 1 mo on SNA under continuous black light. Strains of this species were all isolated from female *E. validus* ambrosia beetles infesting *A. altissima* that were collected in different counties in Pennsylvania, USA. The most distinctive morphological feature of this fungus is the frequent production of sporodochial conidia with 1–2 septa (Table 2, Figs. 3E–I, 4E, J–O). This species formed sporodochial conidia with more than two septa, but the percentage of 0–2-septate conidia (76.5–81%) was much higher than observed in *F. ambrosium* (3.7–24.5%) and *F. euwallaceae* (Freeman *et al.* 2013), where more than 75% of the conidia were 3–5-septate. Cultures appear whitish to yellowish-white when aerial mycelium is sparse on PDA.

Fusarium ambrosium (Gadd & Loos) Agnihothr. & Nirenberg, Stud. Mycol. 32: 98. 1990. MycoBank MB130225. Figs 6–8. Basionym: Monacrosporium ambrosium Gadd & Loos, Trans. Brit. Mycol. Soc. 31(1 & 2): 13. 1947. MB288427. Synonyms: Dactylella ambrosia (Gadd & Loos) K.Q. Zhang, Xing

Z. Liu & L. Cao, *Mycosystema* **7**: 112. 1995. MB447506.

Neocosmospora ambrosia (Gadd & Loos) L. Lombard & Crous, Stud. Mycol. 80: 227. 2015. MB810957.

Fusarium bugnicourtii Brayford, Trans. Brit. Mycol. Soc. **89** (3): 350. 1987. MB133337.

Type: **India**, Upari Tea Institute, a dried specimen from culture of NRRL 22346, isolated from a gallery of *Euwallacea fornicatus* infesting a tea tree, *Camellia sinensis*, 9 May 1990, *V. Agnihothrudu*



Table 2. Percentage of clavate sporodochial conidia of *Fusarium oligoseptatum* and *F. ambrosium* with different numbers of septa cultured on SNA under black light at 25 °C.

		Per	centage of c	onidia with	different nu	mbers of sep	ota		
Species/strain	0-septate	1-septate	2-septate	3-septate	4-septate	5-septate	6-septate	7-septate	Total number of conidia counted
F. oligoseptatum ^a									
NRRL 62579 (ex-holotype)	9.5	41.0	30.5	15.2	1.9	1.9	0	0	105
NRRL 62580	12.1	46.8	20.6	16.8	3.7	0	0	0	107
NRRL 62581	3.8	47.1	25.5	18.9	3.8	0.9	0	0	106
F. ambrosium									
NRRL 20438	0	3.4	5.0	44.5	38.7	6.7	1.7	0	119
NRRL 22345	0	5.9	7.8	25.5	23.5	29.5	4.9	2.9	102
NRRL 22346 (ex-epitype)	0	0	3.7	40.2	46.8	8.4	0.9	0	107
NRRL 36510	0	0.9	12.3	30.2	43.4	13.2	0	0	106
NRRL 46583	0	4.4	5.3	28.1	44.6	12.3	5.3	0	114
NRRL 62605	1.0	4.9	18.6	52.0	18.6	4.9	0	0	102

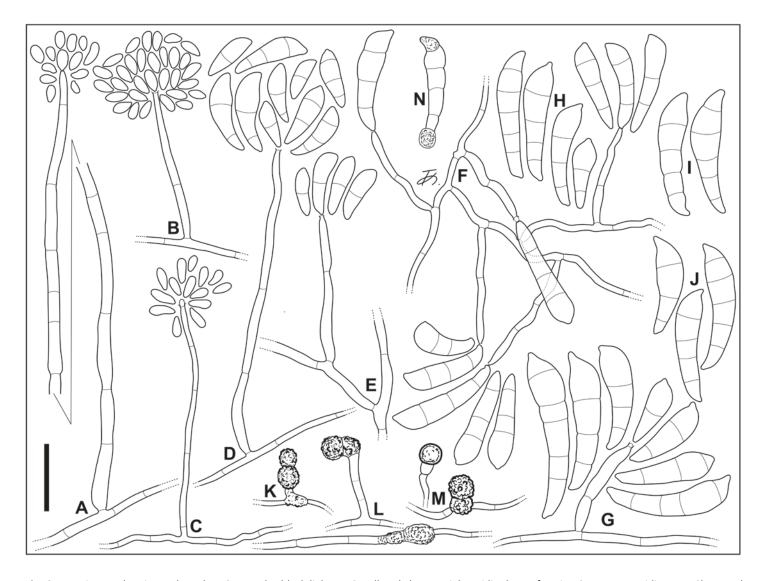


Fig. 6. Fusarium ambrosium cultured on SNA under black light. **A–C.** Tall and short aerial conidiophores forming 0-septate conidia. **D, E.** Short and tall aerial conidiophores forming septate conidia. **F, G.** Sporodochial conidiophores forming large clavate conidia. **H–J.** Clavate multiseptate septate conidia. **K–N.** Chlamydospores in hyphae (K–M) and conidium (N). A, E, F, L from NRRL 62605; B, D, H, K, N from NRRL 20438; C, G, M from NRRL 22346 (ex-epitype); I from NRRL 22345; J from NRRL 46583. Bar = 25 μm.



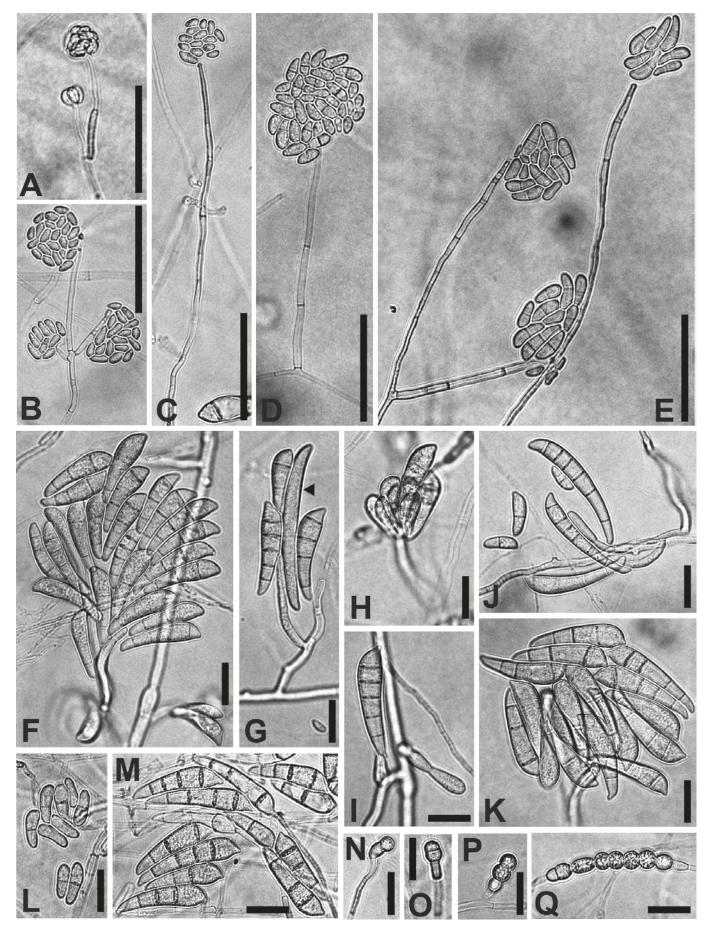


Fig. 7. Fusarium ambrosium cultured on SNA under black light. A–E. Aerial conidiophores forming 0–1(–2)-septate conidia. F–K. Sporodochial conidiophores forming mostly multiseptate clavate conidia, swollen apically with (2–)3–5(–6)-septa; (Arrowhead in G:) crescent-shaped conidium without septa. L. 0–1-septate conidia formed on aerial conidiophores. M. Clavate conidia formed on sporodochial conidiophores. N–Q. Chlamydospores formed in hyphae. A, L, M, O from NRRL 62605; B, C, E, I, K, N from NRRL 22346 (ex-epitype); D, F–H, J, P, Q from NRRL 20438. (A–E: Aerial view without a cover slip; F–Q: Mounted in water with a cover slip.) Bars: A–E = 50 μm, F–Q = 20 μm.

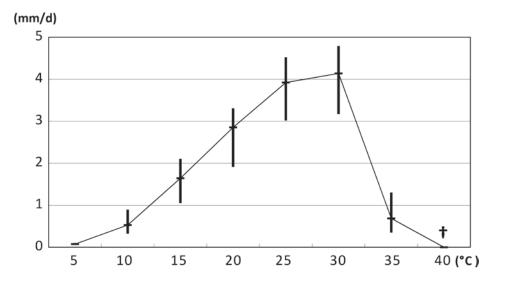


Fig. 8. Radial mycelial growth rate of *Fusarium ambrosium* per day on PDA cultured at eight different temperatures. Thick horizontal and vertical bars indicate means and total ranges, respectively, of the 6 isolates analyzed. All isolates failed to grow and died at 40 °C.

(BPI 910524 – epitype MBT 378232, designated in this study; NRRL 22346 = BBA 65390 = CBS 571.94 = MAFF 246287 – ex-epitype cultures). **Sri Lanka**, an illustration of a mature conidium of the fungus taken from a gallery of *Euwallacea fornicatus* infesting a tea tree, *Camellia sinensis*, C.H. Gadd & C.A. Loos (*Trans. Brit. Mycol. Soc.* **31** (1 & 2): 16, Text-fig. 5 (1987) – lectotype, MBT379562, designated in this study; the original description by Gadd & Loos (1947) did not designate type material of *M. ambrosium* and it was not found in the IMI, K, and BPI herbaria).

Additional strains examined: India, Maharashtra: Chinchona, isolated from a gallery of *E. fornicatus* infesting a *C. sinensis* tree, 17 Jul. 1985, (unknown collector) (NRRL 20438 = IMI 296597 = MAFF 246291, preserved as the ex-holotype strain of *F. bugnicourtii*); (State name of India not recorded): Upari Tea Institute, isolated from a gallery of *E. fornicatus* infesting *C. sinensis* tree, 9 May 1990, *V. Agnihothrudu* (NRRL 22345 = BBA 65389 = MAFF 246288; NRRL 36510 = BBA 65390 = MAFF 246289 as a duplicate of NRRL 22346); Tamil Nadu: Coimbatore District, Anamallais, High Forest Estate, isolated from a gallery of *E. fornicatus* infesting stem of *C. sinensis* tree, 26 Mar. 1990, (unknown collector) (NRRL 46583 = IMI 339338 = MAFF 246290, preserved as *F. bugnicourtii*). Sri Lanka, Talawakelle, St. Coombs, Tea Research Institute of Sri Lanka, isolated from a gallery of *E. fornicatus* infesting *C. sinensis* tree, Mar. 2012, *S. Freeman* (NRRL 62605).

Description: Colonies on PDA showing radial mycelial growth rates of 1.9–3.3 mm/d at 20 $^{\circ}$ C and 3.0–4.5 mm/d at 25 $^{\circ}$ C in the dark. Colony color on PDA white (1A1) to yellowish-white (4A2) in the dark, white (1A1) to yellowish-white (3-4A2) or pale yellow (4A3) under black light. Aerial mycelium white (1A1) sparse to floccose in the dark, more abundantly formed and often covering entire surface of colonies under black light. Colony margin entire to undulate. Reverse pigmentation absent or yellowish-white (3–4A2) or pale yellow (4A3) in the dark and under black light. Olive (3E–F5–8) to olive-brown (4E–F5–8) spots in some strains when sporodochia formed on PDA. Exudates absent. Odor absent, or slightly moldy or sweet in some strains. Hyphae on SNA 1.5-9.5 µm wide. Chlamydospores present in hyphae and in septate sporodochial conidia, mostly subglobose to round ellipsoidal, intercalary or terminal, single or often in chains, hyaline or slightly pale yellow, smooth to often minutely roughwalled, 5–31.3 \times 4.5–13 μ m. Sclerotia absent. Sporulation on SNA and PDA generally rapid and abundant under black light, less abundant in some strains, delayed or sometimes less production in the dark; generally light-colored on SNA and PDA under black light or under daylight; sporodochia formed sparsely on SNA, and rarely on PDA under daylight; olive (3E-F5-8) to olive brown (4E-F5-8) when produced in mass. Aerial conidiophores formed abundantly on SNA under black light, erect, short or tall, mostly narrow but rarely thicker, mostly unbranched, rarely branched sparsely, up to 320 μm long, 2.5–7 μm wide at base, thin-walled, forming monophialides integrated in the apices. Phialides on aerial conidiophores simple, subcylindrical to subulate, tapering towards apex, often with a minute collarette at the tip, 15–66 \times 2.5-4 µm. Aerial conidia typically (1) elliptical, oblong-elliptical, fusiform-elliptical to short clavate, occasionally reniform, some obovate, 0–1-septate; 0-septate on SNA in the dark: $3.5-14 \times$ 1.5–7 μ m in total range, 8.4–9.4 × 2.8–3.3 μ m on average [ex epitype (NRRL 22346): $3.5-14 \times 2-5.5 \mu m$ in total range, 9.4 ± 2.4 \times 3.2±0.6 μm on average ± S.D.]; 0-septate on SNA under black light: $3.5-22 \times 2-7.5 \mu m$ in total range, $7.8-10.8 \times 3.5-4.5 \mu m$ on average [ex epitype (NRRL 22346): $4.5-18.5 \times 2-7.5 \mu m$ in total range, 10.2±2.4 × 4.1±1.1 µm on average ± S.D.]; 1-septate on SNA in the dark: $8-26 \times 2.5-6.5 \mu m$ in total range, 13.6-16.3 \times 3.7–4.9 µm on average [ex epitype: 12–19 \times 3.5–5.5 µm in total range, $15.3\pm1.9 \times 4.8\pm0.4 \mu m$ on average \pm S.D.]; 1-septate on SNA under black light: $8.5-37 \times 2.5-10.5 \mu m$ in total range, $15.3-20.1 \times 5.4-6.6 \mu m$ on average [ex epitype: $10.5-36 \times 10^{-2}$ $2.5-10.5 \mu m$ in total range, $19.3\pm5.1 \times 6.2\pm1.5 \mu m$ on average ± S.D.]; sometimes with (2) larger, falcate to clavate, or curved clavate, (1–)2(–3)-septate conidia, morphologically continuous with falcate sporodochial conidia. Sporodochial conidiophores generally short, unbranched or rarely sparsely branched, contorted, monophialides integrated apically, $20-61.3 \times 3.5-5$ μm. Sporodochial phialides simple, subulate, lanceolate or subcylindrical, often with a conspicuous collarette at the tip, 13–60 × 3–6 μm. Sporodochial conidia hyaline, mostly falcate to long clavate, sometimes curved cylindrical, mostly swollen in the upper part, tapering towards base, often with a round or papillate apical cell, and a distinct or indistinct foot-like, or rounded basal cell, swollen conidia often appear "dolphin-like" (Brayford 1987), (0-)2-5(-7)-septate, formed on SNA under black light, less frequently in the dark, sometimes formed on PDA under black light, rarely in the dark; 2-septate on SNA under black light: $15-61.5 \times 3-12 \,\mu\text{m}$ in total range, $24.7-32.2 \times 6.9-9.1$ μm on average [ex-epitype: 15–61.5 × 4–10 μm in total range, $28.3\pm7.8 \times 7.8\pm1.2 \mu m$ on average \pm S.D.]; 3-septate on SNA under black light: $21-57.5 \times 3.5-13 \mu m$ in total range, 34.1-40.4



 \times 8.3–10.0 µm on average [ex epitype: 30–57 \times 7.5–12.5 µm in total range, 40.4±4.9 \times 8.8±1.1 µm on average ± S.D.]; 4-septate on SNA under black light: 25.5–78.5 \times 6–12.5 µm in total range, 40.7–45.6 \times 8.8–10.4 µm on average [ex epitype: 33–78.5 \times 7.5–11.5 µm in total range, 45.3±6.1 \times 8.8±0.9 µm on average ± S.D.]; 5-septate on SNA under black light: 30–64 \times 7–12.5 µm in total range, 42.9–52.1 \times 8.8–10.3 µm on average [ex epitype: 37–51.5 \times 7.5–12 µm in total range, 45.6±3.7 \times 8.8±1.1 µm on average ± S.D.]. Together with multiseptate sporodochial conidia, often forming (0-)1(-2)-septate, oblong to naviculate or short-clavate, straight or curved conidia, with a rounded apex and a truncate base.

Notes: Strain NRRL 62605 did not form conidia on SNA or PDA in the dark. Therefore, the description of conidia in the dark was based on the three other strains examined in this study. All of the strains studied were isolated from galleries of E. fornicatus infesting C. sinensis trees in India and Sri Lanka. Fusarium ambrosium was originally isolated from E. (Xyleborus) fornicatus galleries in stems of Chinese tea and caster-oil trees in Sri Lanka, and was described as a new species of Monacrosporium, i.e., M. ambrosium by Gadd & Loos (1947). Forty years later, it was redescribed by Brayford (1987) as F. bugnicourtii based on collections from beetle galleries in Chinese tea in India and borerdamaged Hevea brasiliensis and Theoborma cacao in Malaysia. Brayford (1987) considered F. bugnicourtii to be conspecific with F. tumidum var. coeruleum (Bugnicourt 1939) isolated from H. brasiliensis, but distinct from F. tumidum. Fusarium bugnicourtii was recognized as conspecific with Gadd and Loos' species, M. ambrosium from the shot-hole borer on tea, and synonymized under the new combination, F. ambrosium based on its nomenclatural priority (Nirenberg 1990). Because type material of *M. ambrosium* (\equiv *F. ambrosium*) was not designated in the original description by Gadd & Loos (1947) and not found in the IMI, K, and BPI herbaria, we selected a line-drawing (illustration) of a conidium from Gadd & Loos (1947) as the lectotype, according to Art. 9.2 and 9.3 of the ICNafp (McNeill et al. 2012). To supplement the lectotype, BPI 910524, a dried culture of NRRL 22346 (= BBA 65390 = CBS 571.94), isolated from a gallery of E. fornicatus infesting C. sinensis in India by V. Agnihothrudu, was selected as the epitype according to Art. 9.8 of the code. Because an authentic strain of F. ambrosium, IMI 296597 (= NRRL 20438), isolated from E. (Xyleborus) fornicatus galleries in tea tree from India in 1985, was designated as the holotype of F. bugnicourtii by Brayford (1987), this material was not selected for the epitype, per Art. 52.1 and 52.2 of the ICN (McNeill et al. 2012). Although NRRL 62605 was isolated from an E. fornicatus gallery in Sri Lanka from the same host and type locality, it was not selected as the epitype because it appears to be an interspecific hybrid that contains alleles from what appear to be two other AFC species (Kasson et al. 2013). The present epitypification was prepared to stabilize the taxonomy of this species.

Fusarium ambrosium is most similar morphologically to F. euwallaceae (Freeman et al. 2013, Kasson et al. 2013). Fusarium ambrosium and F. euwallaceae produce very similar falcate to clavate, septate sporodochial conidia that are swollen in their upper half, together with ovoid to ellipsoid, 0-septate aerial conidia. The conidial sizes and number of septa of these two fusaria are almost identical (Table 2, Figs 6, 7; Freeman et al. 2013). However, F. ambrosium and F. euwallaceae can be distinguished by the production of hyaline or olive to olive-

brown conidia when old in the former and bluish to greenish conidia in the latter when produced in mass on PDA after 1 mo (Freeman *et al.* 2013). Production of sporodochial conidia in *F. euwallaceae* is easily observed on SNA and PDA in the dark and under black light, but in *F. ambrosium* it is often delayed or limited without black light illumination, even if cultured on SNA.

A preliminary morphological comparison of the sporodochial conidia for 10 of the 12 AFC species has been conducted (T. Aoki et al. unpubl.). Ten of the species produced clavate sporodochial conidia that were swollen apically, and two, including AF-6 associated with Euwallacea sp. #2 in avocado in the Miami-Dade area of southern Florida, USA and AF-9 from Xyleborus ferrugineus in Costa Rica and Delonix regia (royal poinciana) in southern Florida only produced curved fusiform, septate sporodochial conidia in culture, as commonly observed in typical members of the F. solani species complex. Three novel AFC species that were reported recently (O'Donnell et al. 2015, Na et al. 2018), including AF-10 from Singapore, AF-11 from Sri Lanka, and AF-12 (= F. kuroshium) from San Diego County, California, produced clavate sporodochial conidia in culture.

The available data suggests that most of the AFC species might possess diagnostic phenotypic characters. For example, AF-3 ex Euwallacea interjectus infesting Acer negundo in Gainesville, Florida produced sporodochial conidia that were variable in size and shape; AF-5 from Malaysia ex Hevea brasiliensis produced the shortest sporodochial conidia when comparing those with the same number of septa produced by the members of the AFC; AF-7 from Euwallacea sp. #3 ex Persea americana in Australia produced sporodochial conidia that were densely and/or obliquely septate; AF-8 from Euwallacea sp. #2 ex avocado in the Miami-Dade area of southern Florida appeared to be unique in that it is the only AFC species that produced swollen clavate and curved fusiform sporodochial conidia (Kasson et al. 2013); AF-10 from Singapore produced clavate sporodochial conidia that were narrower than conidia of other AFC species with the same number of septa; AF-11 ex E. fornicatus in Chinese tea from Sri Lanka produced sporodochial conidia that were frequently pointed and curved or hooked to one side; F. kuroshium AF-12 ex Euwallacea sp. #5 from Platanus racemosa (California sycamore) and several other woody hosts in San Diego County, California was distinguished by the production of clavate sporodochial conidia together with crescent- or comma-shaped conidia (T. Aoki et al. unpubl.). By way of contrast, AF-6 ex Euwallacea sp. #2 in avocado in the Miami-Dade area of Florida and AF-9 from Costa Rica and the Miami-Dade area of Florida only produced curved cylindrical multiseptate conidia and elliptical to oblong aerial conidia (Kasson et al. 2013). However, the sporodochial conidia formed by AF-9 are atypical of the F. solani species complex because they were pointed at both ends, terminating in a short apical beak and a conspicuous foot-like basal cell. Detailed studies will be required to assess whether AF-6 and AF-9 possess morphological characters that distinguish them from other members of the F. solani species complex.

DISCUSSION

The AFC symbiont cultivated by the ambrosia beetle *Euwallacea* validus in Ailanthus altissima in eastern U.S. is formally described herein as *Fusarium oligoseptatum*. Sampling across 7 eastern U.S. states and 17 tree hosts confirmed that *F. oligoseptatum* is the



primary symbiont of E. validus and dominant, regardless of plant host with few exceptions. This species can be distinguished from the 11 other known AFC species by producing significantly more 0-2-septate clavate sporodochial conidia that are swollen apically and via multilocus molecular phylogenetics where it was strongly supported as a genealogically exclusive species-level lineage in the analyses reported here and in previous studies (Kasson et al. 2013, O'Donnell et al. 2015, Na et al. 2018). Fusarium oligoseptatum was strongly supported as a reciprocally monophyletic sister to F. euwallaceae + Fusarium sp. (AF-3) in Kasson et al. (2013), but the sister group relationship of F. oligoseptatum was unresolved in analyses that included the closely related F. kuroshium (AF-12) from San Diego, California (O'Donnell et al. 2015, and present study). Efforts to develop a robust hypothesis of evolutionary relationships among these four AFC species, which are estimated to have shared a most recent common ancestor approximately 1.6 Mya (O'Donnell et al. 2015), might benefit from the comparative phylogenomic analyses that are currently underway (Stajich et al., pers. comm.).

Herein, an epitype of F. ambrosium was designated based on material originally isolated from a gallery of E. fornicatus infesting Chinese tea in India to provide nomenclatural stability for this species. AFC species have been collected in eight different countries, including Sri Lanka (F. ambrosium AF-1 and Fusarium sp. AF-11), India (F. ambrosium AF-1), Malaysia (Fusarium sp. AF-5), Singapore (Fusarium sp. AF-10), Australia (Fusarium sp. AF-7), Israel (F. euwallaceae AF-2), Costa Rica (Fusarium sp. AF-9) and the United States (F. euwallaceae AF-2, Fusarium sp. AF-3, F. oligoseptatum AF-4, Fusarium spp. AF-6, AF-8, AF-9 and F. kuroshium AF-12)(Brayford 1987, Nirenberg 1990, Freeman et al. 2013, Kasson et al. 2013, O'Donnell et al. 2015, Short et al. 2017, Na et al. 2018). To date only three species within the AFC have been described formally, i.e., F. ambrosium (AF-1; Gadd & Loos 1947, Nirenberg 1990), F. euwallaceae (AF-2; Freeman et al. 2013) and F. oligoseptatum (AF-4; in this study). Although nine of the AFC species are currently unnamed, the prospects for naming them are excellent because most of them appear to possess unique phenotypic/morphological features. Delimitations of such features may, in time, help to uncover the mechanisms underlying the production of clavate conidia, a posited adaptation for the Euwallacea - Fusarium symbiosis (Kasson et al. 2013). Indeed, analogous adaptations in agaricalean fungi (i.e., gongylidia) farmed by higher and occasionally lower attine ants (Schultz & Brady 2008, Masiulionis et al. 2014) also appear to exhibit variation among closely related lineages. However, quality of the substrate, pH, and temperature have also been shown to affect the growth and size of gongylidia in some higher attine ant cultivars when cultivated under lab conditions (Powell & Stradling 1986).

It remains unclear whether *F. ambrosium, F. euwallaceae*, or *F. oligoseptatum* are each farmed by a single *Euwallacea* species, including within their native range, where evidence of hybridization and co-cultivation with other closely related AFC members have been reported (O'Donnell *et al.* 2015). However, it has been shown that *F. euwallaceae* from avocado is obligately required for the survival and development of *Euwallacea* sp. #1 sensu O'Donnell *et al.* (2015) currently occurring in Israel, whereas *F. ambrosium* does not support development of this beetle species (Freeman *et al.* 2012). Likewise, specificity exists for *F. ambrosium* and its beetle host. Future studies focused on vector specificity could help clarify the threats these beetle-fungus consortia pose to our

native ecosystems. This is especially important given that some AFC members such as *F. euwallaceae* have caused significant damage to orchard, landscape and forest trees and threaten avocado production worldwide (Mendel *et al.* 2012, Eskalen *et al.* 2013, Kasson *et al.* 2013), while other AFC members such as *F. oligoseptatum* appear to be quite innocuous when challenged against numerous plant species (Berger 2017).

The FSSC includes over 60 species (Zhang et al. 2006, O'Donnell et al. 2008, Short et al. 2013), a majority of which lack formal Latin binomials thus making it difficult to link specific plant diseases with specific phylogenetic species within the FSSC, including the AFC (Montecchio et al. 2015). The designation of formal Latin binomials for a majority of phylogenetic species within the FSSC coupled with recent abolishment of the dual system of fungal nomenclature will likely reduce confusion surrounding molecular identification of taxa within this large species complex. Nevertheless, the use of multilocus phylogenetic studies will remain the gold standard to discriminate closely related members in the FSSC.

Another avenue to further resolve these closely related phylogenetic species is to examine functional differences among closely related AFC. A recent study by Kasson et al. (2016) assessed the enzyme activity and wood degrading capacity of F. oligoseptatum and R. subfusca, the two known symbionts of E. validus in the eastern U.S. Polyphenol oxidase production was detected from F. oligoseptatum but not R. subfusca. An earlier study by Norris (1980) on AFC member Fusarium sp. AF-9 revealed this fungus was capable of degrading lignin. Further enzymatic studies among closely related AFC may compliment morphological and phylogenetic studies within the Euwallacea -Fusarium mutualism, revealing significant differences in enzyme activity. This is particularly important given recent studies by Aylward et al. (2015) that showed a diverse but consistent set of enzymes present in gongylidia, which are essential for initial degradation of plant substrates in the leaf-cutter ant-Leucoagaricus mutualism.

The results of this study suggest that many of the unnamed AFC species like *F. oligoseptatum* possess unique phenotypic/morphological features, which will facilitate formal description of these economically important pathogens. Phenotypic/morphological studies on the four additional AFC species from Tawain (Na *et al.* 2018) are, therefore, also fully expected. Our ongoing research is focused on advancing the systematics of the AFC to promote accurate communication within the global scientific community.

ACKNOWLEDGMENTS

We are pleased to acknowledge the skilled technical assistance of Gail Doehring, Stacy Sink and Nathane Orwig in generating the DNA sequence data reported in this study. Dr. Konstanze Bensch, MycoBank, is thanked for her helpful advice concerning typification of *F. ambrosium*.

DISCLAMER

Mention of trade names or commercial products in this publication is solely for the purpose of providing specific information and does not imply recommendation or endorsement by the U.S. Department of Agriculture. USDA is an equal opportunity provider and employer.

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