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Five new species of entomopathogenic fungi from the Amazon and evolution of neotropical *Ophiocordyceps*

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ABSTRACT

The neotropical biogeographic zone is a 'hot spot' of global biodiversity, especially for insects. Fungal pathogens of insects appear to track this diversity. However, the integration of this unique component of fungal diversity into molecular phylogenetic analyses remains sparse. The entomopathogenic fungal genus *Ophiocordyceps* is species rich in this region with the first descriptions dating to the early nineteenth century. In this study, material from various ecosystems throughout Colombia and Ecuador was examined. Molecular phylogenetic analyses of five nuclear loci including SSU, LSU, TEF, RPB1, and RPB2 were conducted alongside a morphological evaluation. Thirty-five specimens were examined representing fifteen different species of *Ophiocordyceps*, and five new species, *Ophiocordyceps blattariooides*, *Ophiocordyceps tiputini*, *Ophiocordyceps araracuarensis*, *Ophiocordyceps fulgoromorpha*, and *Ophiocordyceps evansii*, were described. An accurate identification of the host allowed us to conclude that host identity and host habitat are positively correlated with phylogenetic species of *Ophiocordyceps* and are probably strong drivers for speciation of neotropical entomopathogenic fungi.

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Introduction

The entomopathogenic genus *Ophiocordyceps* was established by Petch (1931) for fungi possessing clavate asci with gradually thickening apices and elongate, fusiform ascospores that do

not disarticulate into part-spores. The type species, *Ophiocordyceps blattae* Petch, from Sri Lanka, is rarely collected, but other more commonly collected fungi include *Ophiocordyceps unilateralis* (Tul.) Petch on Hymenoptera adults (worker ants) and *Ophiocordyceps sinensis* (Berk.) G.H. Sung, J.M. Sung,

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Hywel-Jones & Spatafora on Lepidoptera larvae. In other major taxonomic studies of the genus *Cordyceps* (L.) Link, Kobayasi (1941, 1982) and Mains (1958) both treated *Ophiocordyceps* as a subgenus of *Cordyceps*. With advances in the genetic characterization of fungi, a phylogenetic framework for *Cordyceps* (Ascomycota: Hypocreales) derived from molecular data has greatly clarified taxonomic systems previously based on morphology alone (Sung et al. 2007). *Ophiocordyceps* sensu Petch was resurrected as a genus, and expanded to include the majority of taxa with firm, darkly pigmented stromata based on the placement of *O. unilateralis*, but with a diversity of ascus and ascospore morphologies. Currently there are 184 species recognized in *Ophiocordyceps*, with many additional species described as asexual taxa (Quandt et al. 2014), making it the largest of the entomopathogenic genera in Hypocreales (Sung et al. 2007; Johnson et al. 2009). Molecular phylogenetic analyses have incorporated material from Asia, Europe, and North America, highlighting phylogenetic relationships for members of genus *Ophiocordyceps* in the Holarctic and Indo-Malaysian biogeographic zones, but relatively few neotropical specimens of the genus have been sampled (Sung et al. 2007). Based on work by mycologists such as Möller (1901), Henning (1904), and Spegazzini (reviewed in Kobayasi 1941), however, the neotropics are known to support a unique assemblage of *Ophiocordyceps*. More recently, Evans (1982) emphasized the importance of systematic collections of entomopathogenic fungi in the Amazon, a subregion within the neotropical zone. Producing a phylogeny of *Ophiocordyceps* inclusive of global diversity requires specimens from the neotropics, particularly considering the high diversity of entomofauna (Lopez & Miranda 2010).

In this study we sampled neotropical *Ophiocordyceps* species from Colombia and Ecuador, including the Imeri and Napo biogeographic provinces (Amazon subregion) and Magdalena province (Caribe subregion) proposed by Morrone (2006). Incorporation of these samples into existing molecular phylogenetic datasets, in combination with host identification to family and genus, allowed the description of new species and the development of hypotheses concerning the role of host association in the evolution and divergence of *Ophiocordyceps* species.

Material and methods

Material collected

Systematic collections were made in three sites of the Amazon region from Colombia with altitudes between 20 and 200 m above sea level, average temperatures of 28 °C, and relative humidity of 90 %. The first site was the Araracuara canyon of the Caquetá River (Puerto Santander, Amazonas) in May of 2009 and September of 2011. The second site was sampled in several expeditions to Uitoto San Francisco community along to the Igara Parana River (La Chorrera, Amazonas), in July of 2010 and March and September of 2011. The third site sampled was the scientific station El Zafire reserve in the Municipality of Leticia (Amazonas) in July of 2011 and January of 2012. Additionally, collections were made in Rio Claro (San Luis, Antioquia) in March of 2012 and

November of 2014, a region belonging to Magdalena biogeographic province. Some material collected in July 2004 in the Tiputini Research station (Napo, Ecuador) belonging to Napo biogeographic province was included in this study.

Collecting involved careful examination of leaf litter, downed wood, and elevated plant structures (e.g., leaves, twigs) to detect the emergence of stromata from insect cadavers. Dried specimens were placed in plastic bags with silica gel for transportation to the laboratory for identification. Collections were deposited in Antioquia University Herbarium (HUA), and the collection of Fungi of the Natural History Museum of Andes University (ANDES) in Colombia, and National Herbarium of Ecuador (QCNE). Additionally material provided by the National Herbarium of Colombia (COL) was revised. Fungal fruiting bodies were examined according to Sanjuan et al. (2014). Colour descriptions of stromata are according to Kornerup et al., 1984. The host insects were identified to the extent possible (e.g., family, genus, etc.) as allowed by the condition of the specimen.

DNA extraction, PCR amplification, and sequencing

In the field, small pieces of fresh tissue from stromata were placed in 50 µL CTAB extraction buffer (1.4 M NaCl; 100 mM Tris–HCl pH 8.0; 20 mM EDTA pH 8.0; 2 % CTAB w/v) and the DNA extraction process was performed following the method in Kepler et al. (2012). Attempts were made to amplify six nuclear loci including the nuclear ribosomal internal transcribed spacer region (ITS), small subunit ribosomal RNA (SSU) and large subunit ribosomal RNA (LSU), elongation factor-1 α (TEF), and the largest (RPB1) and second (RPB2) largest subunits of RNA polymerase II. The PCR reaction mixture consisted of 2.5 µL 10 × PCR buffer (Fermentas, Glen Burnie, MA), 2 µL MgCl₂ (25 mM), 0.5 µL each primer (10 µM), 0.5 µL dNTP (10 mM each), 1–2 µL template DNA, 0.2 µL Taq polymerase (5 U/µL) (Fermentas, Glen Burnie, MA) and dH₂O to a final volume of 25 µL.

Amplification of ITS was performed with primers ITS1f and ITS4 (White et al. 1990). SSU and LSU were amplified with NS1/SR7 and NS3/NS4 (White et al. 1990) and LROR/LR5 primers (Vilgalys & Sun 1994), respectively. Amplification of TEF was performed with the primers 983F and 2218R (Rehner & Buckley 2005). Amplification of RPB1 was performed with primers cRPB1-1 aF and cRPB1-CaR (Castlebury et al. 2004). Amplification of RPB2 was performed with primers fRPB2-5f2 and fRPB2-7 cR (Liu et al. 1999). The PCR reactions were performed in a Thermocycler 1000 (BIORAD, Hercules, CA) programmed as follows: 94 °C for 3 min; ten cycles of 94 °C for 30 s, 55 °C for 1 min, and 72 °C for 2 min; 35 cycles of 94 °C for 30 s, 50 °C for 1 min, and 72 °C for 2 min; one cycle of 72 °C for 3 min and 4 °C indefinitely (Kepler et al. 2011). Sequencing was performed with the amplification primers at the University of Washington (Seattle, USA) and Andes University (Bogotá, Colombia) sequencing centers. The ITS1 and LSU of the several specimens from this study were submitted to the IBOL initiative (<http://www.boldsystems.org/>).

Sequence alignment and phylogenetic analysis

Sequences from this study were edited using Geneious Pro version 4.8.5 (Drummond et al. 2009). A BLAST query of the

GenBank database was used to confirm that these sequences belonged in *Ophiocordycipitaceae*. New DNA sequences of genes generated in this study were submitted to GenBank and are listed in Table 1. A dataset of taxa in *Ophiocordycipitaceae* was assembled from previously published studies (Sung et al. 2007; Schoch et al. 2012; Kepler et al. 2013; Wen et al. 2013; Quandt et al. 2014) using SSU, LSU, TEF, RPB1, and RPB2 genes. Representatives from the genera *Metarhizium* and *Pochonia* (*Clavicipitaceae*) were used as the outgroup taxa. Alignment of SSU and LSU was performed using MUSCLE (Edgar 2004) with the default settings from Geneious Pro; alignment of protein coding genes TEF, RPB1 and RPB2 was performed with MAFFT (Katoh & Toh 2010) as part of the CIPRES gateway (Miller et al. 2010). Finally, each alignment was refined manually in BioEdit version 6.0.7. (Hall 1999) Alignments are deposited in TreeBase (accession number S15874).

Analyses for the best model of nucleotide substitution were performed using jModelTest version 2.1.3 (Posada 2008). The dataset was divided into 11 gene partitions, one each for SSU and LSU and one for each codon position for the three protein coding genes. We used the ML optimized base tree for likelihood calculations and used Bayesian information criterion (BIC) for model selection for each partition. Maximum Likelihood (ML) analyses were performed using RAXML-VI-HPC version 7.2 (Stamatakis 2006) with 1000 bootstrap replicates. All five genes were concatenated into a single dataset, and 11 data partitions were defined: one each for SSU and LSU, plus nine for each of the three codon positions for the protein coding genes TEF, RPB1, and RPB2 (Kepler et al. 2012). Nodes were considered supported by bootstrap values of greater than 70 %. Bayesian Inference (BI) was performed using MrBayes version 3.1.2 (Huelskenbeck et al. 2001). Substitution models and partitions were specified as in RAxML analyses. Ten million MCMCMC generations were performed, using a sample frequency of 1000 generations and a burn-in of 25 % of the total run. Two runs using four chains each (one cold and three heated chains) were performed, and each run was examined using Tracer version 1.5 (Drummond & Rambaut 2007) to verify burn-in parameters and convergence of individual chains. Nodes were considered supported by posterior probability values greater than or equal to 0.95.

Results

Phylogenetic analyses

We obtained 118 new sequences from 33 specimens. The combined dataset of 121 taxa and 5 genes consisted of 5427 bp (nrSSU 1496 bp, nrLSU 970 bp, TEF1 1073 bp, RPB2 1084 bp, RPB1 804 bp). Both ML and BI analyses displayed similar topologies resolving seven subclades of *Ophiocordycipitaceae* (Fig 1) including 1) 'gunnii' clade, 2) *Tolypocladium* (*Elaphocordyceps* sensu Sung et al. 2007), and the four subclades of *Ophiocordyceps*: 3)'ravenelii' (MLBS = 89; PP = 1.00), 4)'unilateralis' (MLBS = 62; PP = 1.00), 5) 'sobolifera' (MLBS = 100; PP = 1.00) and 6)'sphecocephala' (MLBS = 100; PP = 1.00). The genus *Polycephalomyces* sensu Kepler et al. (2013), recently accepted for inclusion in *Ophiocordycipitaceae* (Quandt et al. 2014), was also resolved in these analyses (MLBS = 88; PP = 0.94).

The 'ravenelii' clade included *Cordyceps gracillima* HUA 186132 (black circles in Fig 1) collected from Scarabaeidae (Coleoptera) larva in Puerto Santander, Amazonas (Colombia). For that reason *C. gracillima* was transferred to *Ophiocordyceps*. It was resolved as a sister species of *Ophiocordyceps melolonthae* which also infects Scarabaeidae larvae. In addition, in the 'gracilis' subclade the species *Ophiocordyceps heteropoda*, *Ophiocordyceps gracilis*, *Ophiocordyceps amazonica*, *Ophiocordyceps gracilioides*, and the new species described here, *Ophiocordyceps blattariooides*, were resolved as a well-supported clade (MLBS = 98; PP = 1.00). *Ophiocordyceps blattariooides* is pathogen of cockroaches, while the other species of the complex are pathogens of larvae of Lepidoptera (*O. gracilis*) and Coleoptera (*O. gracilioides*) and adult Orthoptera (*O. amazonica*).

The 'unilateralis' clade (MLBS = 62; PP = 1.00) is characterized by species typically possessing a *Hirsutella*-like asexual state and dark brown to black stromata that often produce an aperithecial tip. One new species described here, *Ophiocordyceps tiputini* (MLBS = 88; PP = 0.73), is a pathogen of Megaloptera larvae and is a member of this subclade, but represents a host affiliation that had not previously recorded for *Ophiocordyceps*. Although this clade contains numerous ant pathogens, they did not collectively form a monophyletic group. One group comprised *Ophiocordyceps pulvinata* and *Ophiocordyceps unilateralis*, which are pathogens of Camponotus ants (Formicinae). Meanwhile *Ophiocordyceps kniphofioides* var. *kniphofioides*, a pathogen of Cephalotes atratus (Myrmicinae tribe) and *O. kniphofioides* var. *ponerinarum*, a pathogen of Paraponera clavata (Ponerinae tribe) form a second ant clade that was inferred to share a more recent common ancestor with *O. tiputini*.

The 'sobolifera' clade (MLBS = 100; PP = 1.00) includes beetle pathogens, *Ophiocordyceps aphodii* and *Ophiocordyceps brunneipunctata*, and cicada (Hemiptera: Cicadidae) pathogens, including *Ophiocordyceps sobolifera*, *Ophiocordyceps yakusimensis*, *Ophiocordyceps longissima* and the new species described here, *Ophiocordyceps aracuanensis*.

The 'sphecocephala' clade is characterized by higher rates of nucleotide substitutions, which have complicated its inclusion in phylogenetic analyses (Sung et al. 2007, 2008). The 'sphecocephala' clade included 12 new specimens from the Amazon and displayed a topology concordant with previous studies (Sung et al. 2007, 2008; Kepler et al. 2013; Quandt et al. 2014). Although the internal nodes of the core were poorly resolved, species level clades (e.g., *Ophiocordyceps australis*, *Ophiocordyceps lloydii*, and *Ophiocordyceps dipterigena*) were strongly supported and two taxa *Ophiocordyceps fulgoromorpha* sp. nov., which attacks Fulgoridae (Hemiptera) (MLBS = 100; PP = 1.00), and *Ophiocordyceps evansii* sp. nov., which attacks Pachycondyla ants (MLBS = 100; PP = 1.00), are described as new species.

Taxonomy

Ophiocordyceps amazonica (Henn.) G.H. Sung, J.M. Sung, Hywel-Jones & Spatafora Stud. Mycol. 57: 40 (2007) (Fig 2A–C). MycoBank No.: MB504219.

Basionym: *Cordyceps amazonica* Henn, Hedwigia 43: 247. (1904)

Table 1 – Specimen voucher and Genbank accession numbers from Clavicipitaceae and Ophiocordycipitaceae species used in this study.

Taxon	Voucher#	Host/Substratum	ITS	nSSU	nLSU	TEF	RPB1	RPB2
<i>Ha. balanooides</i>	CBS 250.82	Nematoda	AJ292414	AF339588	AF339539	DQ522342	DQ522388	DQ522442
<i>Ha. sinense</i>	CBS 567.95	Nematoda	AJ292417	AF339594	AF339545	DQ522343	DQ522389	DQ522443
<i>Ha. zeosporum</i>	CBS 335.80	Nematoda	AJ292419	AF339589	AF339540	EF469062	EF469091	EF469109
<i>Hy. aurantiaca</i>	OSC 128578	Hymenoptera (adult ant)	JN049833	DQ522556	DQ518770	DQ522345	DQ522391	DQ522445
<i>Hy. dipterigena</i>	HUA 186134	Diptera (adult fly)			KJ917573			KC610712
<i>Hy. dipterigena</i>	NHJ 121702	Diptera (adult fly)	GU723771			GU797127		
<i>Hy. muscaria</i>	OSC 151902	Diptera (adult fly)		KJ878912	KJ878876	KJ878991	KJ878945	
<i>Hirsutella</i> sp.	OSC 128575	Hemiptera		EF469126	EF469079	EF469064	EF469093	EF469110
<i>Hirsutella</i> sp.	NHJ 12525	Hemiptera		EF469125	EF469078	EF469063	EF469092	EF469111
<i>M. anisoplae</i>	ARSEF 5714	Lepidoptera		AF543763	AF543787	AF543775	EF469092	EF469111
<i>M. anisoplae</i>	ARSEF 3145	Coleoptera		AF339579	AF339530	AF543774	DQ522399	DQ522453
<i>N. atypicola</i>	CBS 744.73	Arachnida		EF468987	EF468841	EF468786	EF468892	
<i>O. acicularis</i>	OSC 110987	Coleoptera (larva)		EF468950	EF468805	EF468744	EF468852	
<i>O. acicularis</i>	OSC 110988	Coleoptera (larva)		EF468951	EF468804	EF468745	EF468853	
<i>O. acicularis</i>	OSC 128580	Coleoptera (larva)	JN049820	DQ522543	DQ518757	DQ522326	DQ522371	DQ522423
<i>O. agriotidis</i>	ARSEF 5692	Coleoptera (larva)	JN049819	DQ522540	DQ518754	DQ522322	DQ522368	DQ522418
<i>O. amazonica</i>	HUA 186143	Orthoptera (Acrididae adult)		KJ917562	KJ917571	KM411989	KP212902	KM411982
<i>O. amazonica</i>	HUA 186107	Orthoptera (Acrididae imago)		KJ917561			KP212904	
<i>O. amazonica</i>	HUA 186113	Orthoptera (Acrididae imago)		KJ917566	KJ917571		KP212903	KM411980
<i>O. amazonica</i>	HUA 186136	Orthoptera (Acrididae imago)		KJ917560		KM411991	KP212905	
<i>O. amazonica</i>	HUA 186094	Orthoptera (Acrididae imago)		KJ917565		KM411986	KP212906	
<i>O. amazonica</i>	HUA 186100	Orthoptera (Acrididae imago)		KJ917564		KM411987	KP212907	
<i>O. amazonica</i>	HUA 186101	Orthoptera (Acrididae imago)				KM411988	KP212908	KM411981
<i>O. amazonica</i>	HUA 186106	Orthoptera (Acrididae imago)		KJ917563			KP212909	
<i>O. aphodii</i>	ARSEF 5498	Coleoptera		DQ522541	DQ518755	DQ522323		DQ522419
<i>O. appendiculata</i>	NBRC 100960	Coleoptera (larva)	JN943326	JN941728	JN941413		JN992462	
<i>O. araracuarensis</i>	HUA 186135	Hemiptera (cicada nymph)		KC610788	KC610769	KC610738	KF658665	KC610716
<i>O. australis</i>	HUA 186147	Hymenoptera (<i>Pachycondyla crassinoda</i> adult ant)	KF937351	KC610784	KC610764	KC610734	KF658678	
<i>O. australis</i>	HUA 186104	Hymenoptera (<i>Pachycondyla crassinoda</i> adult ant)	KC610783	KC610763	KC610733		KC610713	
<i>O. australis</i>	HUA 186097	Hymenoptera (<i>Pachycondyla crassinoda</i> adult ant)	KF937350	KC610786	KC610765	KC610735	KF658662	
<i>O. barnesi</i>	BCC 28560	Coleoptera (larva)		EU408776			EU408773	EU418599
<i>O. barnesi</i>	BCC 28561	Coleoptera (larva)		EU408775			EU408774	EU418572
<i>O. blattarioides</i>	HUA186093	Dyctioptera (imago <i>Blattodea</i> , imago)		KJ917559	KJ917570	KM411992	KP212910	
<i>O. blattarioides</i>	HUA 186108	Dyctioptera (<i>Blattodea</i> , imago)		KJ917558	KJ917569		KP212912	KM411984
<i>O. blattarioides</i>	HUA186093	Dyctioptera (<i>Blattodea</i> imago)		KJ917559	KJ917570	KM411992	KP212910	
<i>O. barnesi</i>	BCC 28560	Coleoptera (larva)		EU408776			EU408773	EU418599
<i>O. barnesi</i>	BCC 28561	Coleoptera (larva)		EU408775			EU408774	EU418572
<i>O. brunneipunctata</i>	OSC 128576	Coleoptera (Elateridae larva)		DQ522542	DQ518756	DQ522324	DQ522369	DQ522420
<i>O. communis</i>	NHJ 12581	Isoptera (adult termite)		EF468973	EF468831	EF468775		EF468930
<i>O. communis</i>	NHJ 12582	Isoptera (adult termite)		EF468975	EF468830	EF468771		EF468926
<i>O. curculionum</i>	OSC 151910	Coleoptera (adult Curculionidae)		KJ878918	KJ878885	KJ878999		
<i>O. dipterigena</i>	OSC 151911	Diptera (adult fly)		KJ878919	KJ878886	KJ878966	KJ879000	
<i>O. dipterigena</i>	OSC 151912	Diptera (adult fly)		KJ878920	KJ878887	KJ878967	KJ879001	
<i>O. dipterigena</i>	HUA 186102	Diptera (adult fly)		KC610787	KJ917568		KF658664	KC610715
<i>O. dipterigena</i>	MY 621	Diptera (adult fly)	GU723764			GU797126		
<i>O. elongata</i>	OSC 110989	Lepidoptera (larva)			EF468808	EF468748	EF468856	

<i>O. evansii</i>	HUA 186159	Hymenoptera (<i>Pachycondyla harpax</i> adult ant)	KP200889	KC610796	KC610770	KC610736	KP212916	
<i>O. evansii</i>	HUA 186163	Hymenoptera (<i>Pachycondyla impresa</i> adult ant)	KP200890	KC610797	KC610771	KC610737	KF658670	
<i>O. forquignonii</i>	OSC 151908	Diptera (adult fly)		KJ879003	KJ878947		KJ878922	KJ878889
<i>O. fulgoromorpha</i>	QCNE 186286	Hemiptera (<i>Fulgoridae</i> adult)		KC610793	KC610759			
<i>O. fulgoromorpha</i>	HUA 186139	Hemiptera (<i>Fulgoridae</i> adult)	KJ130991	KC610794	KC610760	KC610729	KF658676	KC610719
<i>O. fulgoromorpha</i>	HUA 186142	Hemiptera (<i>Fulgoridae</i> adult)		KC610795	KC610761	KC610730	KF658677	
<i>O. gunni</i>	OSC 76404	Lepidoptera (pupa)		AF339572	AF339522	AY489616	AY489650	DQ522426
<i>O. gracilis</i>	EFCC 3101	Lepidoptera (larva)		EF468955	EF468810	EF468750	EF468858	EF468913
<i>O. gracilis</i>	EFCC 8572	Lepidoptera (larva)	JN049851	EF468956	EF468811	EF468751	EF468859	EF468912
<i>O. gracillima</i>	HUA 186132	Coleoptera (<i>Scarabaeidae</i> larva)	KJ130990		KC610768	KC610744	KF658666	
<i>O. gracilioides</i>	HUA 186095	Coleoptera (<i>Elateridae</i> larva)		KJ917556			KP212914	
<i>O. gracilioides</i>	HUA 186092	Coleoptera (<i>Elateridae</i> larva)		KJ917555	KJ130992		KP212915	
<i>O. heteropoda</i>	EFCC 10125	Hemiptera (cicada nymph)	JN049852	EF468957	EF468812	EF468752	EP468860	EF468914
<i>O. heteropoda</i>	OSC 106404	Hemiptera (cicada nymph)		AY489690	AY489722	AY489617	AY489651	
<i>O. irangiensis</i>	OSC 128577	Hymenoptera (adult ant)	JN049823	DQ522546	DQ518760	DQ522329	DQ522374	DQ522427
<i>O. irangiensis</i>	OSC 128579	Hymenoptera (adult ant)		EF469123	EF469076	EF469060	EF469089	EF469107
<i>O. irangiensis</i>	NBRC 101400	Hymenoptera (adult ant)	JN943335	JN941715	JN941426		JN992449	
<i>O. kniphofiooides</i>	HUA 186148	Hymenoptera (<i>Cephalotes atratus</i> adult ant)		KC610790	KF658679	KC610739	KF658667	KC610717
<i>O. ponerinarum</i>	HUA 186140	Hymenoptera (<i>Paraponera clavata</i> adult ant)	KJ130987	KC610789	KC610767	KC610740	KF658668	
<i>O. poerinarum</i>	HUA 186133	Hymenoptera (<i>Paraponera clavata</i> adult ant)					KF658669	
<i>O. konnaana</i>	EFCC 7295	Coleoptera (larva)		EF468958			EF468862	EF468915
<i>O. konnaana</i>	EFCC 7315	Coleoptera (larva)		EF468959		EF468753	EF468861	EF468916
<i>O. lloydii</i>	HUA 186164	Hymenoptera (<i>Camponotus</i> adult ant)	KP200892	KC610805		KC610741		
<i>O. lloydii</i>	HUA 186130	Hymenoptera (<i>Camponotus</i> adult ant)	KP200893	KC610791				
<i>O. lloydii</i>	OSC 151913	Hymenoptera (<i>Camponotus</i> adult ant)		KJ878924	KJ878891	KJ878970	KJ879004	KJ878948
<i>O. longissima</i>	EFCC 6814	Hemiptera (cicada nymph)			EF468817	EF468757	EF468865	
<i>O. longissima</i>	TNS F18448	Hemiptera (cicada nymph)		KJ878925	KJ878892	KJ878971	KJ879005	
<i>O. longissima</i>	HMAS 199600	Hemiptera (cicada nymph)		KJ878926		KJ878972	KJ879006	KJ878949
<i>O. melolonthae</i>	OSC 110993	Coleoptera (<i>Scarabaeidae</i> larva)		DQ522548	DQ518762	DQ522331	DQ522376	
<i>O. myrmecophila</i>	MY 163	Hymenoptera (adult ant)	GU723759			GU797132		
<i>O. myrmecophila</i>	CEM 1710	Hymenoptera (adult ant)		KJ878927	KJ878893	KJ878973	KJ879007	
<i>O. myrmecophila</i>	HMAS 199620	Hymenoptera (adult ant)		KJ878929	KJ878895	KJ878975	KJ879009	
<i>O. nigrella</i>	EFCC 9247	Lepidoptera (larva)	JN049853	EF468963	EF468818	EF468758	EF468866	EF468920
<i>O. nutans</i>	OSC 110994	Hemiptera (<i>Pentatomidae</i> adult)		DQ522549	DQ518763	DQ522333	DQ522378	
<i>O. pulvinata</i>		Hymenoptera (<i>Camponotus</i> adult ant)		GU904208		GU904209	GU904210	
<i>O. ravenelli</i>	OSC 110995	Coleoptera (larva)		DQ522550	DQ518764	DQ522334	DQ522379	EF468923
<i>O. rhizoidea</i>	NHJ 12522	Isoptera (adult termite)	JN049857	EF468970	EF468825	EF468764	EF468873	EF468922
<i>O. rhizoidea</i>	NHJ 12529	Isoptera (adult termite)		EF468969	EF468824	EF468765	EF468872	
<i>O. robertsii</i>	KEW 27083	Lepidoptera (<i>Hepialidae</i> larva)			EF468826	EF468766		
<i>O. rubiginosiperitheciata</i>	NBRC 106966	Coleoptera (larva)	JN943344	JN941704	JN941437		JN992438	
<i>O. sinensis</i>	EFCC 7287	Lepidoptera (<i>Hepialidae</i> larva)	JN049854	EF468971	EF468827	EF468767	EF468874	EF468925
<i>O. sobolifera</i>	KEW 78842	Hemiptera (cicada nymph)	JN049855	EF468972	EF468828		EF468875	DQ522432
<i>O. sobolifera</i>		Hemiptera (cicada nymph)	AB027374	AB027328	AB027374			
<i>O. sobolifera</i>	TNS F18521	Hemiptera (cicada nymph)		KJ878933	KJ878898	KJ878979	KJ879013	
<i>O. sphecocephala</i>	OSC 110998	Hymenoptera (adult wasp)		DQ522551	DQ518765	DQ522336	DQ522381	
<i>O. sphecocephala</i>	NBRC 101416	Hymenoptera (adult wasp)	JN943348		JN941443		JN992432	EF468931
<i>O. stylophora</i>	OSC 111000	Coleoptera (<i>Elateridae</i> larva)	JN049828	DQ522552	DQ518766	DQ522337	DQ522382	
<i>O. tiputinii</i>	QCNE 186287	Megaloptera (larva)		KC610792	KC610773	KC610745	KF658671	

(continued on next page)

Table 1 – (continued)

Taxon	Voucher#	Host/Substratum	ITS	nSSU	nLSU	TEF	RPB1	RPB2
<i>O. tricentri</i>		Hemiptera (<i>Cercopidae</i>)		AB027330	AB027376			DQ522436
<i>O. unilateralis</i>	HUA 186161	Hymenoptera (<i>Camponotus</i> adult ant)		KC610799		KC610742	KF658674	KC610718
<i>O. unilateralis</i>	HUA 186126	Hymenoptera (<i>Camponotus</i> adult ant)		KC610798			KF658673	
<i>O. unilateralis</i>	OSC 128574	Hymenoptera (<i>Camponotus</i> adult ant)		DQ522554	DQ518768	DQ522339	DQ522385	
<i>O. variabilis</i>	OSC 111003	Diptera (larva)		EF468985	EF468839	EF468779	EF468885	EF468933
<i>O. variabilis</i>	ARSEF 5365	Diptera (larva)		DQ522555	DQ518769	DQ522340	DQ522386	DQ522437
<i>O. xuefengensis</i>	GZUHHN 13	Lepidoptera (<i>Hepialidae</i> , <i>Phassus nodus</i> , larva)	KC631804	KC631785	KC631790	KC631795		
<i>O. yakusimensis</i>	HMAS 199604	Hemiptera (cicada nymph)		KJ878938	KJ878938		KJ879018	KJ878953
<i>P. chlamydosporia</i>	CBS 504.66	Nematoda	AJ292398	AF339593	AF339544	EF469069	EF469098	EF469120
<i>P. chlamydosporia</i>	CBS 101244	Diplopoda		DQ522544	DQ518758	DQ522327	DQ522372	DQ522424
<i>Pol. cuboideus</i>	NBRC100941	Coleoptera (larva)	JN943329	JN941725	JN941416		JN992459	
<i>Pol. paracuboideus</i>	NBRC 101742	Coleopteran (larva)	JN943338	JN941710	JN941431		JN992444	
<i>Pol. prolifica</i>	NBRC 101750	Hemiptera (<i>Tana japonensis</i> , cicada nymph)	JN943340	JN941708	JN941433		JN992442	DQ522430
<i>Pol. formosus</i>	ARSEF 1424	Coleoptera	KF049661	KF049615	KF049634	KF049689	KF049651	KF049671
<i>Pol. nipponicus</i>	BCC 1881	Neuroptera		KF049618	KF049636	KF049692		KF049674
<i>Pol. ramosopulvinatus</i>	EFCC 5566	Hemiptera	KF049658		KF049627	KF049682	KF049645	
<i>Pol. ryogamiensis</i>	NBRC 103842	Coleoptera (larva)	JN943345	JN941701	JN941440		JN992435	EF468924
<i>Pu. lilacinum</i>	ARSEF 2181	Soil		AF339583	AF339534	EF468790	EF468896	
<i>Pu. lilacinum</i>	CBS 431.87	Nematoda	AY624188	AY624188	EF468844	EF468791	EF468897	EF468940
<i>Pu. lilacinum</i>	CBS 284.36	Soil	AY624189	AY624189	AY624227	EF468792	EF468898	EF468941
<i>T. capitatum</i>	OSC 71233	Fungi (<i>Elaphomycetes</i> sp.)			AY489689	AY489721	AY489615	AY489649
<i>T. capitatum</i>	NBRC 100997	Fungi (<i>Elaphomycetes</i> sp.)	JN943313	JN941740	JN941401		JN992474	
<i>T. fractum</i>	OSC 110990	Fungi (<i>Elaphomycetes</i> sp.)		DQ522545	DQ518759	DQ522328	DQ522373	DQ522425
<i>T. subsessilis</i>	OSC 71235	Coleoptera (<i>Scarabaeidae</i> larva)	JN049844	EF469124	EF469077	EF469061	EF469090	EF469108
<i>T. japonicum</i>	OSC 110991	Fungi (<i>Elaphomycetes</i> sp.)	JN049824	DQ522547	DQ518761	DQ522330	DQ522375	DQ522428
<i>T. ophioGLOSSOIDES</i>	OSC 106405	Fungi (<i>Elaphomycetes</i> sp.)		AY489691	AY489723	AY489618	AY489652	DQ522429
<i>T. ophoglossoides</i>	NBRC 8992	Fungi (<i>Elaphomycetes</i> sp.)	JN943316	JN941736	JN941405		JN992470	
<i>T. paradoxum</i>	NBRC 100945	Hemiptera (cicada nymph)	JN943323	JN941731	JN941410		JN992465	
<i>C. pleuricapitata</i>	NBRC 100745	Hemiptera (cicada nymph)		KF049606	KF049624	KF049679	KF049642	KF049667

Genus codes: Ha. *Haptocillium*; Hy. *Hymenostilbe*; M. *Metharizium*; N. *Nomuraea*; P. *Pochonia*; Po. *Polycephalomyces*; Pu *Purpureocillium*; O. *Ophiocordyceps*; T. *Tolypocladium*.

Herbarium codes: ARSEF, USDA-ARS Collection of Entomopathogenic, Ithaca, NY, USA; BCC, BIOTEC Culture Collection, Klong Luang, Thailand; CBS, Centraalbureau voor Schimmelcultures, Utrecht, the Netherlands; EFCC, Entomopathogenic Fungal Culture Collection, Chuncheon, Korea; GZUH, Herbarium of Guizhou University, Guiyang, Guizhou, China; HUA, Herbarium Antioquia University, Medellin, COL; KEW, mycology collection of Royal Botanical Gardens, KEW, Surrey, UK; NHJ, Nigel Hywel-Jones personal collection; OSC Oregon State University Herbarium, Corvallis, OR, USA; QCNE, National Herbarium of Ecuador, Quito, ECU. MY, Luangsard et al.(2011); NBRC, [Schoch et al. \(2012\)](#).

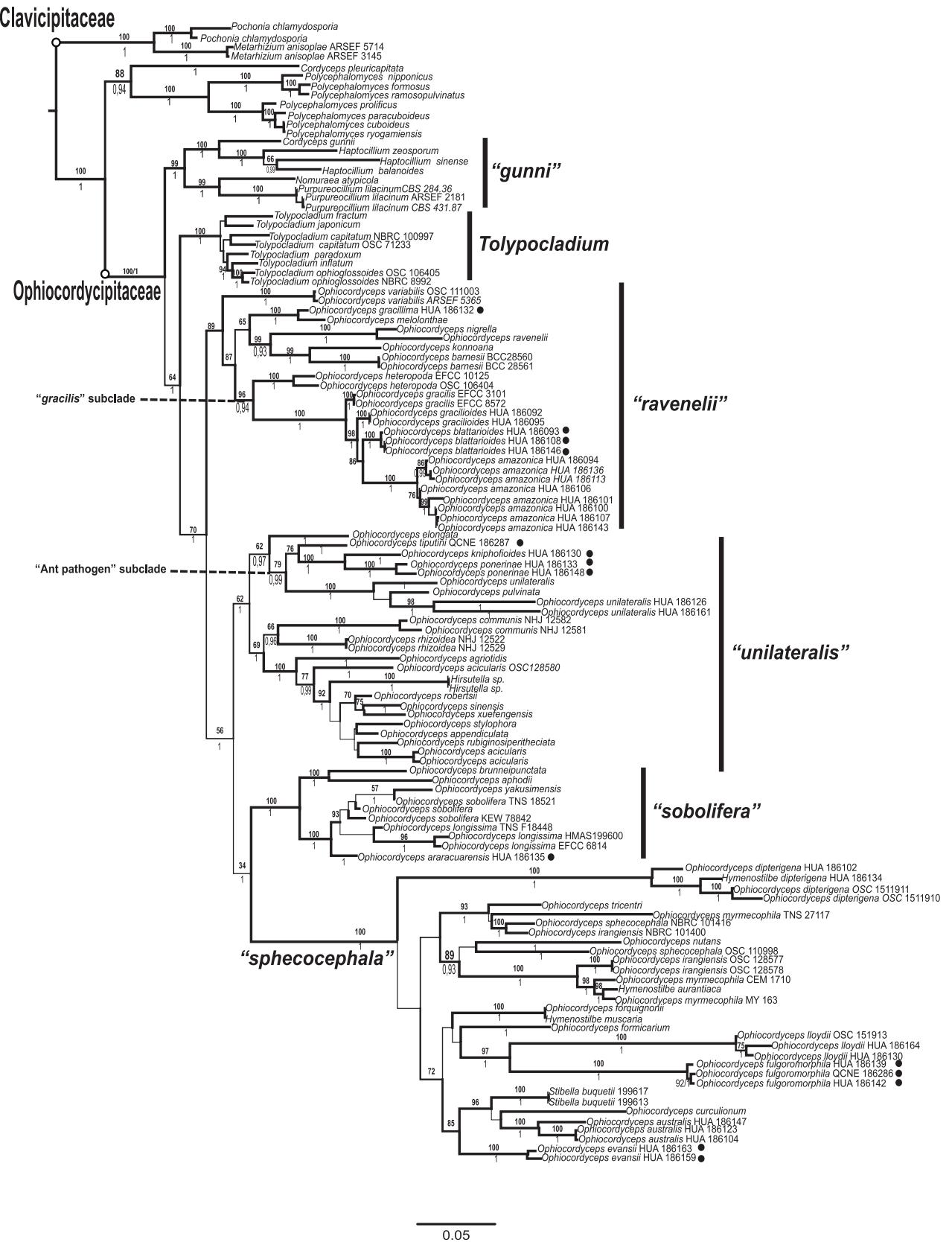


Fig 1 – The most likely tree from the maximum likelihood analysis of Ophiocordycipitaceae family including species collected in the Amazon region based on combined dataset of SSU rRNA, LSU rRNA, TEF, RPB1 and RPB2. Numbers above branches indicate bootstrap support from 1000 maximum likelihood replicates. Numbers below branches indicate posterior probability support in Bayesian analyses. Thinned branches show low resolution in one of the two analyses. The subclades of Ophiocordycipitaceae are named as follow: 1. ‘gunnii’ clade, 2. *Tolyphocladium* (*Elaphocordyceps* sensu Sung et al. 2007) and the four subclades of *Ophiocordyceps*: 3. ‘ravenelii’ 4. ‘unilateralis’, 5 ‘sobolifera’ and 6 ‘sphecocephala’. Black circles shown the new taxa proposed. The tree is rooted with *Metarhizium anisopliae* and *Pochonia chlamydosporia*.

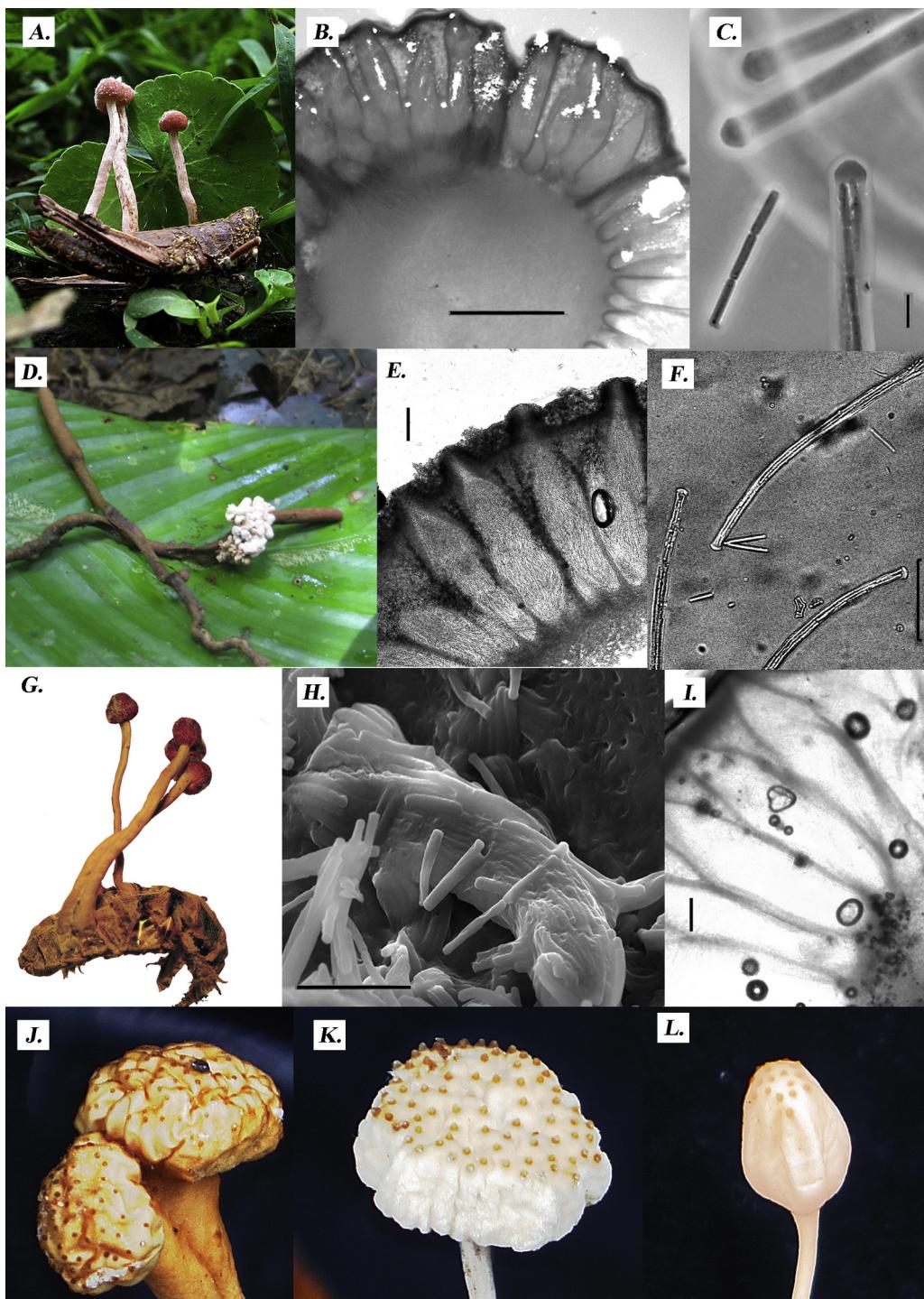


Fig 2 – Amazonian species of *Ophiocordyceps*. (A–C) Epitype of *Ophiocordyceps amazonica* HUA 186143 (A) Stromata on *Prionacris* sp. adult. (B) Perithecia. (C) Ascii and part-spores. (D–F) *Ophiocordyceps araracuarensis* sp. nov. HUA 186135 holotype. (D) Stromata and synnemata. (E) Perithecia on transversal arrangement. (F) Ascii and part-spores. (G–I) *Ophiocordyceps blattarioides* sp. nov. HUA 186108 holotype. (G) Stromata on cockroach adult host. (H) Part-spores on SEM. (I) Perithecia arrangement. (J) Close up of *Ophiocordyceps dipterigena* HUA 186102. (K) Close up of *Ophiocordyceps lloydii* HUA 186164. (L) Close up of *Ophiocordyceps evansii* sp. nov. HUA 186163. Bars = 200 μm 5 \times (B), 10 μm 100 \times (C, E–F), 10 μm 2500 \times (H), 500 μm 10 \times (I).

Description: Stromata solitary to single or gregarious, capitate 20–45 mm long. Fertile head spherical, 3–5.5 mm diam., pruinose when fresh, mammillate when dry, reddish brown (10D/8). Stipe rounded cylindrical to compressed, sometimes

twisted, 17–40 \times 1.0–2.0 mm, fleshy, glossy, pale red to dull red (9B4/9A3). Cortex composed of an ectal layer of longitudinal cells forming a pseudoparenchyma 25 μm thick, with perithecia arranged in a palisade. Perithecia ellipsoid, (760–

800–1100 × (220–) 250–400 µm (n = 20), wall 15–25 µm. Ascii cylindrical, 325–450 × 5 µm (n = 15), apex hemiglobose, 6–7 × 4–5 µm. Ascospores hyaline, filiform, multiseptate, breaking into regular cylindrical part-spores 9–15(–17) × 1.5–2 µm (n = 25). Not conidial state observed.

Lectotype: Brazil. Amazonas: Rio Juruá, Juruá, Miry, May 1901. Hennings, P. Fungi amazonici II. a cl. Ernesto Ule collecti. Tab IV f. 4a. (1904). 'lectotypus hic designatus'

Epitype: Colombia: Amazonas, Leticia, El Zafire research station, 4°0'21"S 69°53'55"W, alt 115 m, on adult of *Prionacris* sp. (Orthoptera: Romelidae) on the ground, 10 Jan 2012, A. Vasco-P. 2026 (HUA 186143).

Other specimens examined: Colombia: Amazonas, Leticia, El Zafire research station, 4°0'21"S 69°53'55"W, alt 115 m. 8 Mar 2012, A. Vasco-P. 1968 (HUA 186106), 18 Mar 2011, A. Vasco-P. 1856 (HUA 186100), 22 Mar 2011 A. Vasco-P. 1862 (HUA 186101), Peña Roja, 0°34'17.65"S 72° 8'24.55"W, alt 100 m, 09 Feb 2010, A. Vasco-P. 1781 (HUA 186084), Puerto Santander, Villa Azul, 0°34'17.65"S 72° 8'24.55"W, alt. 100 m, 9 May 2009, T. Sanjuan 697 (HUA 186136 – paratype); Antioquia: San Luis, Rio Claro 5°56'57.90"N 74°56'57.55"W, alt 250 m, 1 Sep 2012, T. Sanjuan 1001 (HUA 186113). Ecuador: Orellana, Yasuni National Park. Tiputini Research Station, 0°38'14.84"S 76°09'1.83"W, alt. 200 m. 15 Jul 2004. T. Sanjuan 496b (QCNE186285); Pichincha, Pedro Vicente Maldonado, Finca La Nueva Esperanza, 00°0.5'58.87"N 79°05'41.45" W, alt 800 m, 6 Nov 2005, T. Sanjuan 580 (QCNE186294).

Known distribution: Tropical lowlands forest in the Andes mountain of Colombia and Ecuador. Amazon region.

Notes about the host: On imagos and adults of grasshoppers from families Acrididae and Romelidae buried in the leaf litter. These families belong to the superfamily Acrido-morpha easily distinguishable by its large, well-developed hind legs. Localities for all collected specimens of *Ophiocordyceps amazonica* are consistent with the dispersal in the neotropics for Acridomorpha reported by Amedegnato & Des-camps (1982).

Ophiocordyceps araracuarensis T. Sanjuan & J.W. Spatafora, sp. nov. (Fig 2D–F).

MycoBank No.: MB805232.

Etymology: In reference to the Araracuara region in the locality where the type specimen was collected.

Holotype: Colombia: Amazonas, Puerto Santander, Villa Azul, Araracuara region 0°34'17.65"S 72° 8'24.55"W, alt 100 m, on nymph of Cicadomorpha buried in the ground, 8 May 2009, T. Sanjuan 696, (HUA 186135 – holotype).

Description: Stromata solitary, simple, cylindric, 83–97 mm long. Fertile head cylindrical, 25 × 4 mm, slightly punctate, dark brown (6E7) when fresh, dark brown (6F8) when dry. Stipe cylindrical, 62–78 × 2 mm, stiff and fibrous, dark brown (6F8), not straight, with scars where the conidial stage emerges. Perithecia distributed uniformly along of the fertile head, pyriform, 550–650 × 150–240 µm (n = 20), base with long neck of 8–10 µm wide, immersed in ordinal orientation. Cortex composed of indistinct branched, septate hyphae 3 µm diam. Ascii cylindric 300 × 5–6 µm, apex ovoid to lenticular 3 × 5–6 µm (n = 20). Ascospores filiform, hyaline, breaking irregularly into truncate part-spores 9–12 (–16) × 1 µm (n = 50). Asexual state formed by a coraliform growth or soboles (conidial lateral branches), pinkish to whitish, emerging from

the stipe of the stroma, composed by hyaline conidial cells with ovoid to ellipsoid conidia, 6–8 × 2–4 µm (n = 50).

Other specimen examined: Colombia; Amazonas, Puerto Santander, Villa Azul, Araracuara region, 0°34'17.65"S 72° 8'24.55"W, alt 100 m, on nymph of Cicadomorpha buried in the ground, 8 May 2009, T. Sanjuan 695 (HUA 186136 – isotype).

Known distribution: Amazon region of Colombia.

Notes about the host: The hosts were in poor condition because they were mature specimens buried in floodplain soil; however, the pronotum with scutellar suture and anterior legs suited for excavation were preserved and are diagnostic of the family Cicadidae. This family of plant sap-sucking insects is distributed worldwide, mostly in Southeast Asia, but more recently new species had been found in the tropics suggesting that more than 90 % of tropical cicada species remain undescribed (Dietrich 2005).

Ophiocordyceps blattarioides T. Sanjuan & J.W. Spatafora, sp. nov. (Fig 2G–I).

MycoBank No.: MB805784.

Etymology: In reference to the suborder of the host Blattaria (Dictyoptera).

Holotype: Colombia: Amazonas, Leticia, El Zafire research station, 4°0'21"S 69°53'55"W, alt 115 m, on adult of cockroach (Blattidae, Blattaria: Dictyoptera) buried in leaf litter, 16 Jan 2012, A. Vasco-P. 2098 (HUA 186108 – holotype).

Description: Stromata solitary to gregarious, capitate, 10–20 mm long. Fertile head ovoid to subgloboid, 2–3 × 1.5–2.5 mm, pruinose when fresh, mammillate when dry, reddish brown (8/9D/E6). Stipe rounded cylindrical to compressed, sometimes twisted, 10–14 × 0.5–0.8 mm, fleshy, glossy, grayish red to dull red (7/8B4). Cortex composed of an ectal layer of longitudinal cells forming a pseudoparenchyma of 15 µm thick, with perithecia arranged in a palisade. Perithecia ellipsoid, (650–) 760–800 × 220–300 µm (n = 20), wall 15–25 µm. Ascii cylindrical, 250–300 × 5 µm (n = 20), apex hemiglobose, 6–7 × 4–5 µm. Ascospores hyalines, filiform, multi-septate, breaking into regular cylindrical part-spores 6–12 (16) × 1.5 µm (n = 25). Not conidial state observed.

Other specimens examined: Colombia: Amazonas, La Chorrera. Viegas, 1°41'38.58"S 72°28'44.25"W, alt 160 m. 28 Jul 2010, T. Sanjuan 773 (HUA 186146 – paratype). Leticia, El Zafire research station, 4°0'21"S 69°53'55"W, alt 115 m. 16 Jan 2012, A. Vasco-P. 2099 (HUA 186109 – isotype); Puerto Santander, Villa Azul, 0°34'17.65"S 72° 8'24.55"W, alt 100 m. 16 Jun 2010, A. Vasco-P. 014 (HUA186093 – paratype).

Known distribution: Belize, tropical lowlands in the Andes mountain of Colombia and Ecuador. Amazon region.

Notes about the host: The triangular head, flattened-oval body and absence of ocelli are characteristics, which allow identification the host of specimens of HUA 186108–186109, 186146 as members of superorder Dictyoptera. Additionally the specimen HUA186093 has brown spots and short developed wings, which allow for identification as *Neostylopypga* (Blattaria: Blattinae).

Ophiocordyceps evansii T. Sanjuan, sp. nov. (Fig 2L; Fig 3A–B).

MycoBank No.: MB805413.

Etymology: This species is named in honour of Harold Charles Evans for his contributions to the knowledge of the entomogenous fungi parasitic on ants in the tropics.

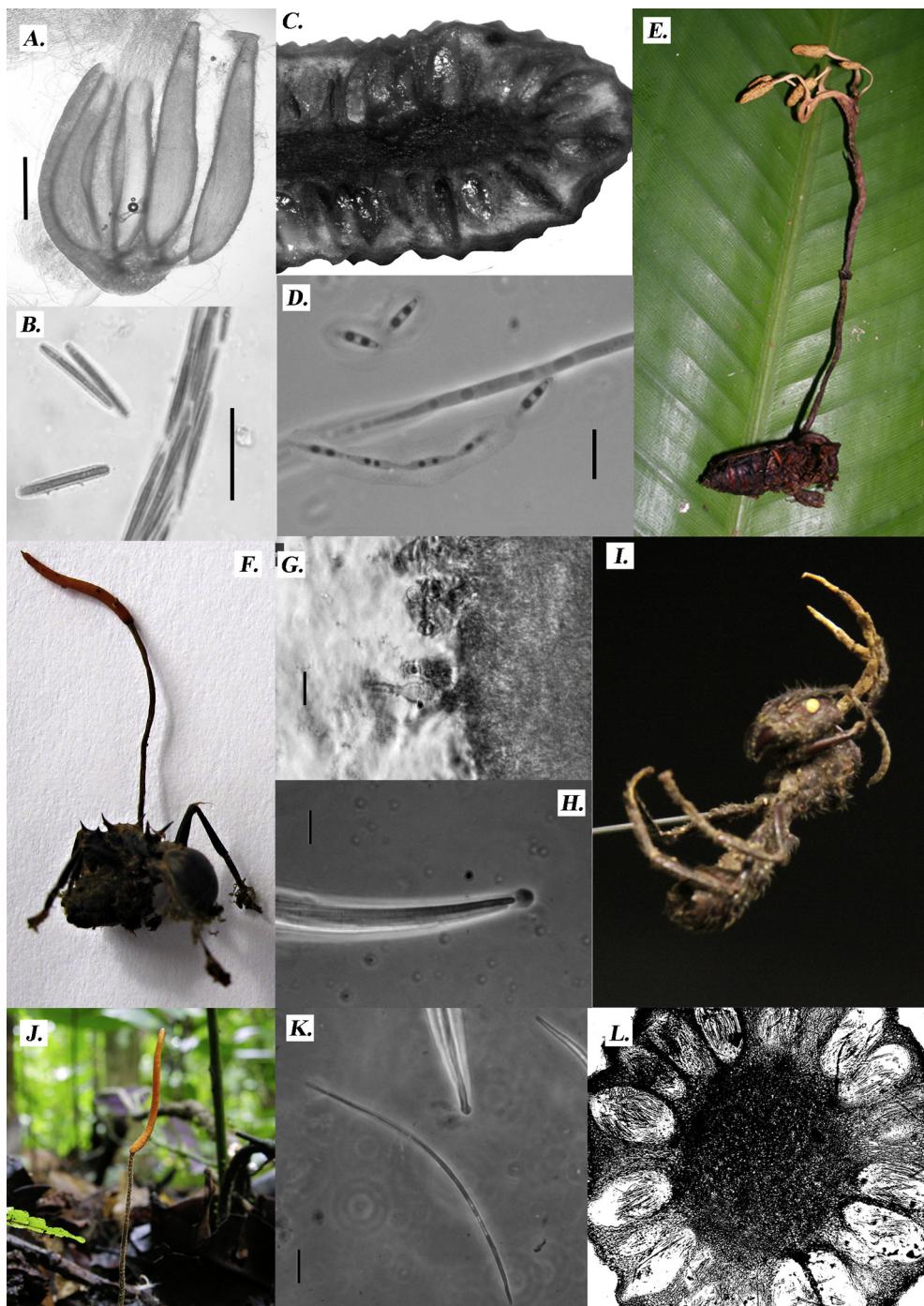


Fig 3 – Amazonian species of *Ophiocordyceps*. (A–B) *Ophiocordyceps evansii* sp. nov. HUA 186163 (A) Perithecia on longitudinal arrangement. (B) Part-spores. (C–E) *Ophiocordyceps fulgoromorphila* sp. nov. HUA 186139 holotype. (C) Perithecia on longitudinal arrangement. (D) Ascospores and part-spores. (E) Stromata on Fulgoridae adult. (F–H) *Ophiocordyceps kniphofioides* HUA 186130 (F) Stromata on *Cephalotes atratus* ants. (G) Conidial state Hirsutella-like shown bottle phialides. (H) Ascospores shown the tip. (I) Stromata of *Ophiocordyceps ponerinarum* HUA 186148 on *Paraponera clavata* ants. (J–L) *Ophiocordyceps tiputini* sp. nov. QCNE 186287 holotype. (J) Stromata. (K) Ascospores. (L) Ascospores. Bars = 300 µm 10× (A), 20 µm 100× (B), 10 µm 100× (D, G, H, K), 300 µm 10× (L).

Holotype: Colombia: Amazonas, La Chorrera, San Francisco, 1°21'40"S 72°51'48"W, alt 0 m, on the ant *Pachycondyla impressa* (Ponerinae) biting small roots in bark at 1 m from the ground, 18 May 2010, T. Sanjuan 858, (HUA 186159 – holotype).

Description: Stromata simple, capitellate, slender, 20–30 mm long, filamentous, bicolored. Fertile head flattened obovoid in vertical profile with erumpent ostioles, 1.5–2 × 0.5–1 mm, dull red to brownish red (8B6/9C7). Stipe slender, cylindric,

$18-28 \times 0.7$ mm glabrous, pliant, brownish black (7F8), concolorous with the head. Perithecia obclavate, $1600-1800 \times 260-300$ µm, with a prominent reddish brown ostiole, $60-80 \times 80-100$ µm, immersed vertically in clava consisting of very loosely interwoven hyphae with a very thick compact ectal layer, $40-50$ µm wide of oblong or globoid cells. Ascii cylindrical, $950-1200 \times 8$ µm when mature, apex hemispheric 8×10 µm. Ascospores arranged in parallel, filiform, easily breaking into part-spores, fusiform-ellipsoid, $17-20 \times 1.5-2$ µm. External mycelium brown dark, scarcely covering the host and attaching it sometimes to the substrate. Not conidial state observed.

Other specimens examined: Colombia: Amazonas: La Chorrera, San Francisco, primary forest, $1^{\circ}21'40"S\ 72^{\circ}51'48"W$, alt 250 m, on the ant *Pachycondyla harpax* (Ponerinae), 22 Mar 2011, T. Sanjuan 912, (HUA 186163 – paratype).

Known distribution: In the Amazon region of Colombia, Ecuador, and Brazil.

Notes about the host: *Pachycondyla* is the second most diverse genus in the Ponerinae tribe of ants. They are terrestrial hunters with some species making their nest on the ground while others are arboreal. *Pachycondyla* is characterized by triangular mandibles without a circular orifice. The adhesive pads of the tarsus are considered an adaptation to an arboreal ecology (Orivel et al. 2001).

Ophiocordyceps fulgoromorphila T. Sanjuan, sp. nov. (Fig 3C–E).

MycoBank No.: MB805227.

Etymology: In reference to the host family Fulgoridae (Hemiptera: Fulgoromorpha).

Holotype: Colombia: Amazonas, Puerto Santander, Araracuara Canyon, $0^{\circ}37'21"S\ 72^{\circ}24'26"W$, alt 115 m, on adult of Fulgoridae buried in leaf litter, 13 May 2009, T. Sanjuan 717, (HUA 186139 – holotype).

Description: Stromata solitary to gregarious, capitate, $87-110$ mm long, perpendicularly branched in secondary bi-colored stromata. Fertile head cylindrical (3.5–) $7-25 \times 2-3.5$ mm ($n = 4$), apex rounded, punctate, often rugose due to the ostioles, brownish yellow to brownish orange (4C5/7C4), ostioles yellow brown (5E6). Stipe slender, (57–) $80-120 \times 0.5-2.5$ mm, ligneous, grayish brown (5F3) becoming olivaceous buff (5C7). Secondary stipe glabrous, leathery, concolorous to the head above, $10-20 \times 0.5-1.5$ mm. The stipe extends to the fertile head as a columella composed by parallel, septate, brown, hyphae, 800 µm thick. Perithecia conical to ellipsoid, $780-1100 \times 280-380$ µm ($n = 20$), immersed in slightly oblique orientation in a cortex composed of brownish pseudoparenchyma with three layers of rounded cells and one of cylindrical base cells, 50 µm thick. Ascii cylindrical $300-600 \times 5-6$ µm, ($n = 20$) with a 4×5 µm thick rounded cap. Ascospores hyaline, filiform, multiseptate, breaking irregularly into fusiform part-spores $9-12$ (-16) $\times 1-2$ µm ($n = 30$) with two oil drops. Not conidial state observed.

Other specimens examined: Colombia: Amazonas, Puerto Santander, Araracuara Canyon, $0^{\circ}37'21"S\ 72^{\circ}24'26"W$, alt 115 m, 14 May 2009, T. Sanjuan 729, (HUA 186142 – isotype), Villa Azul, $0^{\circ}34'17.65"S\ 72^{\circ}8'24.55"W$, alt 100 m, 8 May 2009, T. Sanjuan 689 (ANDES paratype); Antioquia, San Luis, refugio del cañon de Rio Claro, $5^{\circ}56'57.90"N\ 74^{\circ}56'57.55"W$, alt

200 m, 9 Nov 2014, T. Sanjuan 1155 (ANDES). Ecuador: Orellana province, Yasuni National Park, Tiputini Research Station, $00^{\circ}38'14.84"S\ 76^{\circ}09'01.83"W$, alt 150 m, 13 Jul 2004. T. Læssøe 11468 (QCNE 186286 – paratype).

Known distribution: Tropical lowlands in the Magdalena River basin in Colombia and Amazon region of Bolivia, Ecuador and Colombia.

Notes about the host: Despite the conditions of the specimens TS 689, 717, and 729, they all possess the conserved insertion of antenna and carina covering the eyes' fossa, as well as specimen TL 11468 having a longer prolongation and spines in its abdomen. This unique morphology of the head is the most diagnostic character of the Fulgoridae. This family is one of the most diverse on the neotropics. Of the 500 described species of Fulgoridae, only 17 species are recorded from Holarctic (Urban & Cryan 2009).

Ophiocordyceps gracillima (Y. Kobayasi) T. Sanjuan & J.W. Spatafora, comb. nov.

Basionym: *Cordyceps gracillima* Y. Kobayasi Bull. Nat. Sci. Mus., Tokyo, Bot (7) 4: 126 (1981).

MycoBank No.: 811229.

Specimens examined: Colombia: Amazonas, Puerto Santander, Villa Azul, $0^{\circ}34'17.65"S\ 72^{\circ}8'24.55"W$, alt 100 m, 7 May 2009, on larva of Scarabaeidae, (Coleoptera) T. Sanjuan 679, (HUA 186132).

Known distribution: In the Amazon region of Colombia.

Ophiocordyceps kniphofioides sensu stricto H.C. Evans & Samson (1982). (Fig 3F–H).

MycoBank No.: 504288.

Synonym: *Cordyceps kniphofioides* var. *kniphofioides* H.C. Evans & Samson, Trans. Brit. Mycol. Soc. 79: 441. (1982).

Specimens examined: Colombia: Amazonas, La Chorrera, San Francisco, Chuquial forest, $1^{\circ}21'40"S\ 72^{\circ}51'48"W$, alt 50 m, on adults of ant *Cephalotes atratus* biting and hidden in moss on branches and trunks of trees, 30 July 2009, T. Sanjuan 790, (HUA 186148); 11 September 2011, T. Sanjuan 975, (HUA 186124).

Known distribution: In the Amazon region of Colombia, Ecuador, and Brazil.

Notes about the host: *Cephalotes atratus* nests in the canopy and forages in the lower forest where it feeds on plant sucking homopterans (Evans & Samson 1982). The Myrmicinae tribe is considered an army group with a higher level of social organization. However *C. atratus* is an opportunistic predator closely related phylogenetically to *Paraponera clavata* (Poneromorpha) (Keller 2011).

Ophiocordyceps ponerinarum (H.C. Evans & Samson) T. Sanjuan & R.M. Kepler, stat. nov. (Fig 3I).

MycoBank No.: MB810374.

Synonym: *Cordyceps kniphofioides* var. *ponerinarum* H.C. Evans & Samson, Trans. Brit. Mycol. Soc. 79: 441. (1982).

Ophiocordyceps kniphofioides var. *ponerinarum* (H.C. Evans & Samson) G.H. Sung et al., Stud. Mycol. 57: 37 (2007).

Specimens examined: Colombia: Amazonas, Puerto Santander, Araracuara Canyon, $0^{\circ}37'21"S\ 72^{\circ}24'26"W$, alt 115 m, on hunter ant *Paraponera clavata* and *Dinoponera longipes* on the ground, 13 May 2009, T. Sanjuan 727 (HUA 186140), Amazonas, La Chorrera, San Francisco, Chuquial forest, $1^{\circ}21'40"S\ 72^{\circ}51'48"W$, alt 50 m, 11 September 2011, T. Sanjuan 986–987, (HUA 186133).

Known distribution: In the Amazon region of Colombia, Ecuador, and Brazil.

Notes about the host: *Paraponera clavata* and *Dinoponera longipes* are the largest Amazon ants with lengths of 26 mm and 30 mm, respectively (Kempf 1971). They are hunters and make their nests in the bases of trees, palms, and lianas. *P. clavata* initially was included in Ponerinae tribe, but with the molecular phylogenetic approach this single species was placed in the tribe called Paraponerinae and is more related to Myrmicinae which includes *C. atratus*, the host of *Ophiocordyceps kniphofioides* (Keller 2011).

***Ophiocordyceps tiputini* T. Sanjuan & R.M. Kepler, sp. nov.** (Fig 3 J–L).

Mycobank No.: MB805228.

Etymology: Refers to the collection locality Tiputini River of the type.

Holotype Ecuador: Orellana province, Yasuni National Park, Tiputini Research Station, 00°38'14.84" S 76°09'01.83" W, alt 150 m, on larva of Megaloptera buried 10 cm in wet leaf litter, 13 Jul 2004. T. Læssøe 11465 (QCNE 186287 – holotype).

Description: Stromata single, simple or branched, capitate, slender, wiry, and stiff 150 mm long. Fertile head cylindric, dark orange to deep orange (5A8/6A8), 15–25 × 1.5–2.5 mm, having a central core and sterile acuminate tip. Stipe fibrillose, pliant, dark brown (7F8), 125–135 × 0.5 mm. Perithecia ovoid, 300 × 200–250 µm (n = 20), dark brown immersed, ordinal orientation. Cortex composed by a pseudoparenchyma with hexagonal cells. Ascii broadly fusoid, (100–)125–200 × 5 µm, narrowing to 2.5 µm at the apex, cap prominent 4.5 × 2.5 µm. Ascospores elongate-fusiform with rounded apex 120–180 × 1–1.5 µm, not breaking into part-spores. Not conidial state observed.

Known distribution: In the Amazon region from Ecuador and Bolivia.

Notes about the host: The larva is slightly flattened with sclerotized head and conspicuous mouthparts. The mandibles have the typical chewing morphology with prominent maxillae characteristic of Megaloptera. The size of the larva, more than 6 cm, suggests that it probably belongs to the Corydalidae family (e.g., dobsonfly). In Bolivia, similar entomogenous fungi on the same host that have been photographed in cloud forest of Coroico and the low rain forest of Mashaquipe show the typical coiled worm of the Megaloptera larvae (Winkler, 2014).

Discussion

This study represents the first extensive phylogenetic investigation of *Ophiocordyceps* and the family *Ophiocordycipitaceae* in the neotropics. *Ophiocordyceps* in the Amazon subregion displays high morphological diversity and a broad range of host affiliations. Two main hypotheses, the ‘host relatedness’ and ‘host habitat’ hypotheses (Brooks & McLennan 1991) are frequently invoked to explain host affiliations observed among *Cordyceps* s.l. In the ‘host relatedness’ hypothesis, host speciation functions as an isolating mechanism that results in cospeciation of the symbiont and congruent host-symbiont phylogenies. The ‘host habitat’ hypothesis involves host shifts among often distantly related hosts that occupy a common ecological niche. This second hypothesis has been invoked

to explain host switching among closely related species of *Cordyceps* s.l. that are found associated with distantly related host species between cicadas and truffles (Nikoh & Fukatsu 2000; Spatafora et al. 2007) and between scale insects, grasses, and fungi (Kepler et al. 2012). Here we conducted molecular phylogenetic analyses of *Ophiocordyceps* and discuss their phylogenetic diversity and the relative importance of these two hypotheses as explanatory mechanisms for speciation of Amazonian entomopathogenic fungi.

New Amazonian species in the ‘ravenelii’ clade

In the ‘ravenelii’ clade (Fig 1), the species *Ophiocordyceps heteropoda*, *Ophiocordyceps gracilis*, *Ophiocordyceps gracilioides*, *Ophiocordyceps amazonica*, and *Ophiocordyceps blattariooides* form the well-supported ‘gracilis’ clade that is characterized by the production of dark red to brown to tan capitate stromata with similar macroscopic and microscopic characteristics (Fig 1, Fig 2A,G; Table 2). *Ophiocordyceps heteropoda* attacks nymphs of Hemiptera (Cicadidae), *O. gracilis* attacks larval stages of Lepidoptera (mainly Hepialidae, Nymphalidae), *O. gracilioides* attacks larval stages of Coleoptera (Elateridae), *O. amazonica* attacks adult Orthoptera (Romaleidae, Acrididae) and *O. blattariooides* sp. nov. attacks adult Dictyoptera (Blattaria). Although Mains (1940) was unable to distinguish specimens collected on cockroaches in Belize from *O. amazonica* on grasshoppers based on morphological characteristics, our molecular phylogenetic analyses support the recognition of distinct phylogenetic species that are consistent with isolation by host within a common habitat, i.e., ecological niche. While *O. heteropoda* attacks cicada hosts buried in soil, the remainder of species in this clade infect hosts associated with the soil and leaf litter environments.

Despite the frequent collection of *O. amazonica* from the Amazon region, no type specimen exists owing to the destruction during WWII of the material on which the description published by Hennings (1904) was based. Here we conducted molecular and morphological analyses of 13 specimens of *O. amazonica*, and assigned an epitype from Amazonian material.

Ophiocordyceps gracillima is also located in the ‘ravenelii’ clade (Fig 1), in the ‘ravenelii – scarabaeidae’ subclade sensu Luangsa-ard et al. (2010), where most species parasitize beetle larvae. This species was described by Kobayasi (1981) from material collected in the Colombian Amazon region. It produces solitary cylindrical capitate stromata that are red-orange in colour and parasitize Scarabaeidae larvae buried in decaying, fallen tree trunks. In the phylogenetic analyses *O. gracillima* is a sibling species of an isolate of *Ophiocordyceps cf. melolonthae*. The specimen of *O. cf. melolonthae* analysed here was collected in the Appalachian Mountains of North America and occurred on a larva of Scarabaeidae buried in soil. This collection and others collected in the region (see <http://cordyceps.us>) were sulfur yellow in colour when fresh with a clavate stroma and irregular patches of perithecia. Differences in morphology and geography, along with nucleotide divergence are consistent with the recognition of *O. gracillima* as a distinct species, but the common host suggests a potential, larger ‘scarabaeidae’ clade of *Ophiocordyceps*. The close relationship between these two species and between their hosts, combined with their geographic separation, is consistent with the ‘host-

Table 2 – A morphological comparison of *Ophiocordyceps* species belongs to ‘gracilis’ subclade.

Species	Distribution	Host	Stromata	Fertile head	Ascomata	Ascii (μm)	Reference
<i>O. heteropoda</i>	Japan, Australia	Nymphs of Hemiptera (Cicadidae)	Simple, solitary, 12 cm long	Ovoid, cinnamon buff, 7–9 × 6–7 mm	Immersed, ampullaceous, 610–660 × 210 μm, neck 160 μm, peridium 5–10 μm	Cylindrical, 250–300 × 5.2–7 μm Part-spores truncate 6–7.7 × 0.9–1 μm	Kobayasi (1941)
<i>O. gracilis</i>	Asia, Europe, America, Africa and Oceania	Larvae of Lepidoptera (Hesperiidae, Nymphalidae)	Simple, usually solitary, 4–9 cm long	Globular ellipsoidal, red ochreous to pale orange, 4–9 × 4–7 mm	Immersed, elongate-oviform (320–)560–840 × 200–360 μm, peridium 20–40 μm	Cylindrical, (200–) 400–528 × 5–8 μm Part spores truncate 5–9 × 1.5–2 μm	Kobayasi (1941), Mains (1958), Samson & Brady (1983), Gorbunova et al. (2011)
<i>O. gracilioides</i>	Japan, China, Mexico, Colombia, Bolivia	Larvae of Coleoptera (Elateridae)	Simple, usually solitary, 2–5 (9) cm long	Spherical, pale rufous, 4–5.5 mm	Immersed, ellipsoid to naviform, (680–)830–900 × 200–280 μm, peridium 50 μm	Cylindrical, 450–700 × 5–6.5 μm Part spores truncate 7–10(–12) × 1–2 μm	Kobayasi (1941), Fan et al. (2001), and this study
<i>O. amazonica</i>	Argentina, Brazil, Colombia, Ecuador, Guyana, Peru	Imago and adults of Orthoptera (Romaleidae, Acriidae)	Simple, gregarious, 2–4.5 cm long	Subglobe, spherical, reddish brown, 2.5–5.5 mm	Immersed, ovoid-ellipsoid, (760–)800–1100 × (220–) 250–400 μm, peridium 25 μm	Cylindrical, 325–450 × 5 μm Part spores truncate 9–15(–17) × (0.5–) 1.5–2 μm	Henning (1904), Mains (1959), and this study
<i>O. blattarioridis</i>	Belize, Colombia, Ecuador	Adults of Blattaria (Dyctoptera)	Simple, gregarious, 1.4–2 long	Ovoid or subglobose, chestnut brown, 2–3 × 1.5–2.5 mm	Immersed, ellipsoid, (650–)750–800 × 220–300 μm, peridium 15–25 μm	Cylindrical, (180–250) (–300) × 4–5 μm Part spores truncate 6–12 (–16) × 1.5 μm	Mains (1940), and this study

related’ hypothesis of cospeciation with their hosts. However, sampling of additional isolates and species of both the fungi and hosts is necessary for a more definitive analysis. Further work is also necessary to determine the appropriate application of species names among members of this group, including examination of material collected from the type locality of *Ophiocordyceps melolonthae* by Tulasne in Boreal region of North America Boreal and both varieties mentioned by Mains (1958) *O. melolonthae* var. *melolonthae*, and *O. melolonthae* var. *rickii*.

A new species reveals a host-jumping event in the ‘unilateralis’ ant-pathogen subclade

In the ‘unilateralis’ clade (Fig 1), the pathogens of ants include species that are characterized by the production of filiform stromata with aperithecial tips, ascospores that are more fusiform than filiform and that do not disarticulate into part-spores. Each species parasitizes one kind of ant with *Ophiocordyceps pulvinata*, *Ophiocordyceps halabensis*, *Ophiocordyceps polyrhachis-furcata*, *Ophiocordyceps camponoti-leonardi*, *Ophiocordyceps camponoti-saundersi*, and *Ophiocordyceps unilateralis* parasitizing different species of *Camponotus* and *Polyrhachis* (Formicidae: Formicinae) (Kepler et al. 2011; Kobmoo et al. 2012). Our phylogenetic analyses support the use of host-identity as criteria for species discrimination within this group (Fig 1). Therefore, we propose recognition of *Ophiocordyceps kniphofiodes* varieties as an unique species; *O. kniphofiodes* is collected on adult *Cephalotes atratus* (Formicidae: Myrmicinae) and *Ophiocordyceps ponerinae* is collected on adults of *Paraponera clavata* and *Dinoponera longipes* (Formicidae: Ponerinae).

Despite its unique host affiliation, *Ophiocordyceps tiputini* is morphologically similar to species of the ‘unilateralis’ clade (e.g., size and shape of its stroma; Fig 2J–L) and represents a host-shift from ants (Formicidae) to dobsonflies (Megaloptera) again to ants (Fig 1), within the ant-pathogen subclade. The ant *C. atratus* is an opportunist predator that nests in the canopy but is parasitized in the low level of the forest. Meanwhile *P. clavata* and *D. longipes* ants are generalist hunters that make their nests at the base of trees (Lattke 2003). Megaloptera larvae are generalist predators of small invertebrates. They live in freshwater streams with pupation occurring in soils immediate to streams (Grimaldi & Engel 2005). The specimen studied here was found in leaf litter of the floodplain or ‘varzea’ (Wittman et al. 2011) of the Tiputini River. The ‘varzea’ with a dense layer of wet leaf litter is an ecological niche shared by predaceous Megaloptera larvae (Contreras-Ramos 2004) and Poneromorpha ants (Keller 2011). We suggest that the host shifts of the ‘unilateralis’ clade between Megaloptera and Poneromorpha occurred within a common habitat resulting in colonization of new host niches.

New cryptic species of Amazonian cicada pathogens in the “sobolifera” clade

The ‘sobolifera’ clade comprises pathogens of cicada, including the best known and most frequently collected of these species of cicada pathogens, *Ophiocordyceps sobolifera*, also known in China as ‘Chen chua’ or ‘flower of cicada’

(Kobayasi & Shimizu 1963). It produces a robust, fleshy, stipitate stroma with fertile clava. At the base of the fertile head are lateral, short branches, called soboles, which are the conidiogenous state of the fungus. *O. sobolifera* has been frequently recorded in China, Korea, Japan, and the Philippines as well as in the Caribbean islands, Florida, and Mexico (Petch 1942; Mains 1951). Also present in this clade is *Ophiocordyceps longissima*, another pathogen of cicadas; it is distinct from *O. sobolifera* in possessing a more elongated and cylindrical clava and typically in being attached to the host via a long tapering, rhizomorph-like stipe. A specimen identified as *O. sobolifera* was previously collected in Colombia (Kobayasi 1981) near the Caquetá River close to where our specimens (HUA 186135-136) were collected. The morphology of the Amazonian specimens studied here present some slight morphological differences with respect to *O. sobolifera*, including the shape of the fertile head, which is strictly cylindrical, and the measurement of the ascus, which is 100 µm smaller (Fig 3 A–C). Also, the perithecia of the Amazonian specimens have a pyriform shape with a short neck similar to *O. sobolifera*, rather than the ovoid perithecia of *O. longissima* (Fig 2 E).

Based upon these results, we describe *Ophiocordyceps aracuarensis* as a new cryptic species of cicada pathogen that was previously included in *O. sobolifera* based on the morphological species concept across a common host with a broad geographic distribution. Examples of cryptic species have been reported across fungi and their recognition has become common due to the increased use of molecular data and phylogenetic species concepts (e.g., Bickford et al. 2007; Le Gac et al. 2007; Bischoff et al. 2009; Rehner et al. 2011; Piątek et al. 2013; Sheedy et al. 2013; Sanjuan et al. 2014). Whether *O. sobolifera*, *O. longissima*, and *O. aracuarensis* occur on different subfamilies and genera of Cicadidae is difficult to ascertain owing to deterioration of host specimens. Dietrich (2005), however, affirmed that the diversity of cicadas in that region could be higher than in the rest of the world because of the diversity of arboreal plants on which they feed. An accurate assessment of the roles that biogeography and host association play in the diversification and speciation of this well identified hosts and a more intensive, global sampling.

New Amazonian species corroborate the adult host as an ecological characteristic of the 'sphecocephala' clade

The 'sphecocephala' clade is well supported (MLBS = 100; PP = 1.00) and is morphologically characterized by terminal fertile head with a peridium composed of well-defined pseudoparenchyma, perithecia that are mostly obliquely immersed, and filiform ascospores that disarticulate into fusiform part-spores. It is also ecologically unique in that the majority of species attack adult stages of insects from Coleoptera, Hemiptera, Diptera, and Hymenoptera. Numerous described species of Amazonian *Ophiocordyceps* were collected including *Ophiocordyceps dipterigena* on flies, and *Ophiocordyceps lloydii*, and *Ophiocordyceps australis* on ants. In addition to these taxa, two new species are described here: *Ophiocordyceps fulgoromorpha* is the first species reported as parasite of planthoppers in the Fulgoromorpha

(Hemiptera) (Fig 3 C–D) and is the first pathogen of Fulgoridae placed in the 'sphecocephala' clade. In addition to morphological characteristic of this clade, *O. fulgoromorpha* has the ability to produce secondary ramifications with new fertile heads after decay of the initial fertile head. *Ophiocordyceps fulgoromorpha* has yellow buff stromata and is resolved as a sibling species of *O. lloydii*. Combined morphological, ecological, and phylogenetic evidence support its recognition as a new species.

The other new species described in the 'sphecocephala' clade is *Ophiocordyceps evansii*. Initial observations of *O. evansii* led to its identification as an immature stage of *O. australis*, a parasite of the ant *Pachycondyla crassinoda* (Formicidae: Ponerinae). However, upon further examination of micromorphology and phylogenetic analyses, it is rejected as being conspecific with *O. australis*. With respect to some traits, *O. evansii* is morphologically more similar to *O. lloydii*, a parasite of *Camponotus* spp. ants (Formicidae: Formicinae), and *O. dipterigena*, a parasite of different dipteran families (Fig 2 J–L). These species have narrowly ovoid perithecia that are longer than 900 µm and immersed perpendicularly to the surface at the top of the fertile 'head' (Fig 3 I). But unlike the other species, the stroma of *O. evansii* has the appearance of a closed tulip flower and ellipsoid-fusiform part-spores greater than 10 µm (Fig 3B). In spite of the morphological similarities possessed by these species, they do not collectively form a monophyletic group within the 'sphecocephala' clade (Fig 1).

In conclusion, for *Ophiocordyceps*, as for many groups of fungi, morphological species concepts, when applied in absence of ecological and molecular data, are overly broad, resulting in the masking of evolutionarily unique taxa. Incorporation of new specimens of *Ophiocordyceps* from the neotropics within phylogenetic analyses revealed increased species diversity across all clades of *Ophiocordyceps*, supporting it as a diverse genus of neotropical entomopathogens. Moreover, the analysis revealed strong correlations between parasite and either host or habitat sufficient to distinguish morphologically cryptic phylogenetic species (e.g. new species in the 'unilateralis' subclade). Further investigation of this group will likely reveal many more undescribed species and parasite–host associations within the neotropics.

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REFERENCES

- Bickford D, Lohman DJ, Sodhi NS, Ng PKL, Meier R, Winker K, Ingram KK, Das I, 2007. Cryptic species as a window on diversity and conservation. *Trends Ecology & Evolution* 22: 148–155.
- Bischoff JF, Rehner SA, Humber R, 2009. A multilocus phylogeny of the *Metarhizium anisopliae* lineage. *Mycologia* 101: 512–513.
- Brooks DR, McLennan DA, 1991. *Phylogeny, Ecology and Behavior. A research program in comparative biology*. University of Chicago Press, Chicago, USA 434.
- Castlebury LA, Rossman AY, Sung G-H, Hyten AS, Spatafora JW, 2004. Multigene phylogeny reveals new lineage for *Stachybotrys chartarum*, the indoor air fungus. *Mycological Research* 108: 864–872.
- Contreras-Ramos A, 2004. Taxonomic and distributional notes on the dobsonflies of Ecuador (Megaloptera: Corydalidae). *Dugesiana* 11: 1–11.
- Dietrich CH, 2005. Keys to the families of Cicadomorpha and subfamilies and tribes of Cicadellidae (Hemiptera: Auchenorrhyncha). *Florida Entomologist* 88: 502–516.
- Drummond AJ, Rambaut A, 2007. BEAST: Bayesian evolutionary analysis by sampling trees. *BMC Evolutionary Biology* 7: 214.
- Drummond AJ, Rambaut A, Ashton B, Cheung M, Heled J, Kearse M, Moir R, Stones-Havas S, Thirer T, Wilson A, 2009. Geneious v4.8.5 Available from: <http://www.geneious.com>
- Edgar RC, 2004. MUSCLE: a multiple sequence alignment method with reduced time and space complexity. *BMC Bioinformatics* 5: 113.
- Evans HC, 1982. Entomogenous fungi in tropical forest ecosystems: an appraisal. *Ecological Entomology* 7: 47–60.
- Evans HC, Samson RA, 1982. Cordyceps species and their anamorphs pathogenic on ants (Formicidae) in tropical forest ecosystems. I. The Cephalotes (Myrmicinae) complex. *Transactions of the British Mycological Society* 79: 431–453.
- Fan MZ, Chun-Ru L, Li ZZ, 2001. *Cordyceps gracilioides* a new record for China. *Mycosistema* 20: 273–274.
- Gorbunova IA, Kryukov Y, Zibzeev EG, 2011. First records of the entomopathogenic fungus *Ophiocordyceps gracilis* (Ascomycota: Hypocreales) from Siberia. *Euroasian Entomological Journal* 92: 17–18.
- Grimaldi D, Engel M, 2005. *Evolution of the Insects*. Cambridge University Press, Cambridge, UK 755.
- Hennings P, 1904. Fungi amazonici II. a cl. Ernesto Ule collecti. *Hedwigia* 43: 246–249.
- Hall TA, 1999. BiEdit: a user-friendly biological sequences alignment editor analysis program for windows 95/98/NT. *Nucleic Acids Symposium Series* 41: 95–98.
- Huelsenbeck JP, Ronquist F, Nielsen R, Bollback JP, 2001. Bayesian inference of phylogeny and its impact on evolutionary biology. *Science* 294: 2310–2314.
- Johnson D, Sung GH, Hywel-Jones NL, Luangsa-ard JJ, Bischoff JF, Kepler RM, Spatafora JW, 2009. Systematics and evolution of the genus *Torrubia* (Hypocreales, Ascomycota). *Mycological Research* 113: 279–289.
- Katoh K, Toh H, 2010. Parallelization of the MAFFT multiple sequence alignment program. *Bioinformatics* 26: 1899–1900.
- Keller RA, 2011. A phylogenetic analysis of ant morphology (Hymenoptera: Formicidae) with special reference to the poneromorph subfamilies. *Bulletin of the American Museum of Natural History* 355: 1–90.
- Kempf W, 1971. A preliminary review of Ponerine ants genus *Dinoponera* Roger (Hymenoptera: Formicidae). *Studia Entomologica* 14: 369–394.
- Kepler RM, Kaitsu Y, Tanaka E, Shimano S, Spatafora JW, 2011. *Ophiocordyceps pulvinata* sp. nov., a pathogen with a reduced stroma. *Mycoscience* 52: 39–47.
- Kepler RM, Sung GH, Ban S, Nakagiri A, Chen A, Huang B, Li Z, Spatafora JW, 2012. New teleomorph combinations in the entomopathogenic genus *Metacordyceps*. *Mycologia* 104: 182–197.
- Kepler RM, Ban S, Nakagiri A, Bischoff J, Hywel-Jones NL, Owensby CA, Spatafora JW, 2013. The phylogenetic placement of hypocrealean insect pathogens in the genus *Polycephalomyces*: an application of one fungus one name. *Fungal Biology* 117: 611–622.
- Kobayasi Y, 1941. The genus *Cordyceps* and its allies. *Science Reports of the Tokyo Bunrika Daigaku Serie B* 84: 53–260.
- Kobayasi Y, 1981. Revision of the genus *Cordyceps* and its allies 2. *Bulletin of the National Science Museum Tokyo Serie B* 7: 123–129.
- Kobayasi Y, Shimizu D, 1963. Monographic Studies of *Cordyceps* 2. Group parasitic on Cicadidae. *Bulletin of the National Science Museum Tokyo* 6: 286–314.
- Kobmoo N, Mongkolsamrit S, Tasanathai K, Thanakitpattana D, Luansa-ard JJ, 2012. Molecular phylogenies reveal host-specific divergence of *Ophiocordyceps unilateralis* sensu lato following its host ants. *Molecular Ecology* 21: 3022–3031.
- Kornerup A, Wanscher JH, Pavée D, 1984. *Methuen Handbook of Colour*, 3rd edn. Hasting House, New York, USA 257.
- Lattke JE, 2003. Subfamilia Ponerinae. In: Fernández F (ed.), *Introducción a las hormigas de la región Neotropical*. Instituto de Investigación de Recursos Biológicos Alexander von Humboldt, Bogotá, pp. 261–270.
- Le Gac M, Hood ME, Fournier E, Giraud T, 2007. Phylogenetic evidence of host-specific cryptic species in the anther smut fungus. *Evolution* 61: 15–26.
- Lopez F, Miranda DR, 2010. A phylogenetic approach to conserving Amazonian biodiversity. *Conservation Biology* 24: 1359–1366.
- Liu YJ, Whelen S, Hall BD, 1999. Phylogenetic relationships among Ascomycetes: evidence from an RNA polymerase II subunit. *Molecular Biology and Evolution* 16: 1799–1808.
- Luangsa-ard JJ, Ridkaew R, Mongkolsamrit S, Tasanathai K, Hywel-Jones NL, 2010. *Ophiocordyceps barnesi* and its relationship to other melolonthid pathogens with dark stromata. *Fungal biology* 114: 739–745.
- Mains EB, 1940. Cordyceps species from British Honduras. *Mycologia* 32: 16–22.
- Mains EB, 1951. Notes concerning entomogenous fungi. *Bulletin of the Torrey Botanical Club* 78: 122–133.
- Mains EB, 1958. North American entomogenous species of *Cordyceps*. *Mycologia* 50: 169–222.
- Mains EB, 1959. Cordyceps species. *Bulletin of the Torrey Botanical Club* 86: 46–58.
- Miller MA, Pfeiffer W, Schwartz T, 2010. Creating the CIPRES science gateway for inference of large phylogenetic trees. In: *Proceedings of the Gateway Computing Environments Workshop (GCE)*, 11 April, 2015, New Orleans, LA, pp. 1–8.
- Möller A, 1901. Phycomyceten und ascomyceten, untersuchungen aus Brasilien. In: Schimper (ed.), *Botanische Mittheilungen aus den Tropen*, vol. 9, pp. 1–319.
- Morrone JJ, 2006. Biogeographic areas and transition zones of Latin America and the Caribbean Islands based on panbiogeographic and cladistic analyses of the entomofauna. *Annual Review of Entomology* 51: 467–494.
- Nikoh N, Fukatsu T, 2000. Interkingdom host jumping underground: phylogenetic analysis of entomopathogenic fungi of the genus *Cordyceps*. *Molecular Biology and Evolution* 17: 29–38.
- Orivel J, Malherbe MC, Dejean A, 2001. Relationships between pretarsus morphology and arboreal life in Ponerinae ants of the genus *Pachycondyla* (Formicidae: Ponerinae). *Annals of the Entomological Society of America* 94: 449–456.
- Petch T, 1931. Notes on entomogenous fungi. *Transactions of the British Mycological Society* 16: 55–75.
- Petch T, 1942. Notes on entomogenous fungi. *Transactions of the British Mycological Society* 25: 250–265.

- Piątek M, Lutz M, Chater A, 2013. Cryptic diversity in the *Antherospora vaillantii* complex on *Muscari* species. *IMA Fungus* **4**: 5–19.
- Posada D, 2008. jModelTest: phylogenetic model averaging. *Molecular Biology and Evolution* **25**: 1253–1256.
- Quandt CA, Kepler RM, Gams W, Araújo JP, Ban S, Evans HC, Hughes D, Humber R, Hywel-Jones NL, Li Z, Luangsa-ard JJ, Rehner S, Sanjuan T, Sato H, Shrestha B, Sung GH, Yao Y, Zare R, Spatafora JW, 2014. Phylogenetic-based nomenclatural proposals for Ophiocordycepsitaceae (Hypoocreales) with new combinations in *Tolypocladium*. *IMA fungus* **5**: 121–134.
- Rehner SA, Buckley EP, 2005. A *Beauveria* phylogeny inferred from nuclear ITS and EF1- α sequences: evidence for cryptic diversification and links to *Cordyceps* teleomorphs. *Mycologia* **97**: 84–98.
- Rehner SA, Minnis AM, Sung GH, Luangsa-ard JJ, Devotto L, Humber RA, 2011. Phylogeny and systematics of the anamorphic, entomopathogenic genus *Beauveria*. *Mycologia* **103**: 1055–1073.
- Sanjuan T, Tabima J, Restrepo S, Læssøe T, Spatafora JW, Franco-Molano AE, 2014. Entomopathogens of Amazonian stick insects and locusts are members of the *Beauveria* species complex (*Cordyceps* sensu stricto). *Mycologia* **106**: 260–275.
- Schoch CL, Seifert KA, Huhndorf S, Robert V, Spouge JL, Levesque CA, Griffith GW, 2012. Nuclear ribosomal internal transcribed spacer (ITS) region as a universal DNA barcode marker for Fungi. *Proceedings of the National Academy of Sciences* **109**: 6241–6246.
- Sheedy EM, Van de Wouw AP, Howlett BJ, May TW, 2013. Multi-gene sequence data reveal morphologically cryptic phylogenetic species within the genus *Laccaria* in southern Australia. *Mycologia* **105**: 547–563.
- Samson RA, Brady BL, 1983. *Paraisaria*, a new genus for *Isaria dubia*, the anamorph of *Cordyceps gracilis*. *Transactions of the British Mycological Society* **81**: 285–290.
- Spatafora JW, Sung GH, Sung JM, Hywel-Jones HJ, White JR, 2007. Phylogenetic evidence for an animal pathogen origin for ergot and the grass endophytes. *Molecular Ecology* **16**: 1701–1711.
- Stamatakis A, 2006. RAxML-VI-HPC: maximum likelihood-based phylogenetic analyses with thousands of taxa and mixed models. *Bioinformatics* **22**: 2688–2690.
- Sung G-H, Hywel NL, Sung J, Luangsa JJ, Shrestha B, Spatafora JW, 2007. Phylogenetic classification of *Cordyceps* and the clavicipitaceous fungi. *Studies in Mycology* **57**: 5–59.
- Sung G-H, Poinar Jr, George O, Spatafora JW, 2008. The oldest fossil evidence of animal parasitism by fungi supports a Cretaceous diversification of fungal – arthropod symbioses. *Molecular Phylogenetics and Evolution* **49**: 495–502.
- Urban JM, Cryan JR, 2009. Entomologically famous, evolutionarily unexplored: the first phylogeny of the lanternfly family Fulgoridae (Insecta: Hemiptera: Fulgoroidea). *Molecular Phylogenetics and Evolution* **50**: 471–484.
- Vilgalys R, Sun BL, 1994. Ancient and recent patterns of geographic speciation in the oyster mushroom *Pleurotus* revealed by phylogenetic analysis of ribosomal DNA sequences. *The Proceedings of the National Academy of Sciences of the United States of America* **91**: 4599–4603.
- Wen TC, Zhu RC, Kang JC, Huang MH, Tan DB, Ariyawansha H, Liu H, 2013. *Ophiocordyceps xuefengensis* sp. nov. from larvae of *Phassus nodus* (Hepialidae) in Hunan Province, southern China. *Phytotaxa* **123**: 41–50.
- White TJ, Bruns T, Lee S, Taylor JW, 1990. Amplification and direct sequencing of fungal ribosomal RNA genes for phylogenetics. In: *PCR Protocols a Guide to Methods and Applications*. Academic Press, pp. 315–322.
- Winkler D, 2014. <http://www.ophiocordyceps.com/> December 9, 2014.
- Wittmann F, Schöngart J, Junk W, 2011. Phytogeography, species diversity, community structure and dynamics of central Amazonian floodplain forests. In: Wolfgang J, Junk W, Piedade M (eds), *Amazonian Floodplain Forests Ecophysiology, Biodiversity and Sustainable Management*. Ecological Studies, vol. 210. Springer, Netherlands, pp. 61–102.