

# Hysterangium bonobo: A newly described truffle species that is eaten by bonobos in the Democratic Republic of Congo

Elliott, Todd; Georgiev, Alexander V.; Lotana Lokasola, Albert; Smith, Matthew

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1	Short title: Bonobo mycophagy
2	Hysterangium bonobo: a newly described truffle species that is eaten by bonobos in the
3	Democratic Republic of Congo
4	
5	Todd F. Elliott <sup>a</sup> , Alexander V. Georgiev <sup>b</sup> , Albert Lotana Lokasola <sup>c</sup> Matthew E. Smith <sup>d*</sup>
6	<sup>a</sup> Ecosystem Management, University of New England, Armidale, New South Wales 2351,
7	Australia; <sup>b</sup> School of Natural Sciences, Bangor University, Bangor, UK; <sup>c</sup> Vie Sauvage,
8	Kokolopori Bonobo Reserve, Province Equateur, Democratic Republic of Congo; <sup>d</sup> Department
9	of Plant Pathology, University of Florida, Gainesville, Florida 32611, USA
10	Corresponding author email: <u>*trufflesmith@ufl.edu</u>
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# 24 ABSTRACT

Many animals have been shown to eat fungi, and most truffle-like fungi depend on animals for 25 spore dispersal via mycophagy. Although these interactions are widespread, they are 26 understudied in many habitats. In this study, we show that bonobos (Pan paniscus) forage and 27 feed on an undescribed truffle species in the rainforests of the Democratic Republic of Congo. 28 29 Based on morphological and molecular assessment of collections, we show that the species eaten by bonobos is a previously undescribed taxon described here as Hysterangium bonobo. This 30 31 species is known in the local Bantu language (Bongando) as simbokilo and is used for baiting 32 traps to catch several species of small mammals. Our findings highlight the need for further research into mycophagy and systematics of sequestrate fungi in Africa. 33 **KEY WORDS:** African fungi, Hysterangiales, Phallomycetidae, primate mycophagy, truffle 34

35 taxonomy, 1 new taxon

#### **36 INTRODUCTION**

37 Fungi play a significant role in the diets and nutrition of diverse vertebrates. Many fungi, particularly truffle-like taxa, have evolved close associations with animals that help to disperse 38 spores via mycophagy (Fogel and Trappe 1978; Elliott et al. 2019a, b). These animal-fungus 39 40 associations are frequently overlooked, but they are an important part of functional ecosystems and imperative for the dispersal of fungi through these systems. There is a growing interest in 41 42 better understanding the ecological significance of these associations. Mycophagy is nutritionally 43 important for animals and simultaneously aids dispersal of mycorrhizal fungi (Cázares and Trappe 1994; Hussain and Al-Ruqaie 1999; Colgan and Claridge 2002; Kalač 2009; Wallis et al. 44 2012). These interconnected associations have been studied in various parts of the world but 45 46 have been mostly overlooked in Africa.

The specialization of these associations varies depending on the fungal taxa and types of 47 animals that consume and disperse them. Most studies of vertebrate mycophagy have focused on 48 small mammals in regions outside of Africa (Fogel and Trappe 1978; Claridge and May 1994). 49 In the Middle East and arid North Africa, some bird species are known to feed on truffle-like 50 fungi and may play important roles in the health of desert ecosystems (Elliott et al. 2019b), but in 51 52 Sub-Saharan Africa only a few vertebrate species have been reported as mycophagous. There are a handful of reports of African primates eating fungi (Kano and Mulavwa 1984; Hanson et al. 53 2003 and references therein; Isbell and Young 2007; Georgiev et al. 2010; Georgiev et al. 2011; 54 55 Buyck et al. 2016). Among African vertebrates other than primates, four mammal species (Elliott et al. 2019c) and eleven terrestrial turtle species (Elliott et al. 2019a) have been reported to eat 56 fungi. Buyck et al. (2016) noted that animals in the Central African Republic, including different 57 species of duikers, primates, and wild pigs, all consumed *Elaphomyces*. However, this study did 58 not provide the genera and species of the African mycophagous mammals that were observed. 59 As has been shown in other regions, this paucity of mycophagy reports is most likely due 60 to insufficient sampling or the use of methods that do not detect fungal spores. The low number 61 of reports does not necessarily mean that Africa has a low diversity of mycophagous animals 62 63 (Elliott and Vernes 2019; Elliott et al. 2019b). For many vertebrates it can be difficult to determine whether or not fungi are regularly ingested solely by observing their feeding behavior. 64 Microscopic and/or DNA analysis of feces are useful techniques to determine the presence and 65 66 diversity of fungi in the diet. It is clear that these methods need to be more thoroughly utilized to study the diets of vertebrates in Africa. 67

68 Schmitt and Mueller (2007) documented 2,250 described species of macrofungi from
69 Africa (Schmit and Mueller 2007). Mammals on other continents have been shown to eat a wide

variety of morphologically diverse fungi (Fogel and Trappe 1978; Claridge and May 1994; 70 Nuske et al. 2017). Fungi that produce sequestrate (enclosed) and/or hypogeous (below ground) 71 fruiting bodies are presumably more dependent on animal vectors than non-sequestrate species 72 because sequestrate and/or hypogeous taxa typically lack the ability to forcibly discharge spores 73 (Trappe et al. 2009). There are more than a dozen species of sequestrate fungi reported from 74 75 Sub-Saharan Africa (Dissing and Lange 1962; Dring and Pegler 1978; Castellano et al. 2000; Eberhardt and Verbeken 2004; Beenken et al. 2016; Castellano et al. 2016a, b; Orihara and 76 Smith 2017). Several species have also been reported from the arid Kalahari desert and the 77 78 surrounding region of southern Africa (Taylor et al. 1995; Trappe et al. 2008; Trappe et al. 2014). Many of these taxa have been recently described and occur in habitats where primates are 79 known to forage, but no studies have carefully examined the mycophagy of sequestrate fungi by 80 primates or many other African vertebrates. Here we provide new observations of bonobo 81 mycophagy and show that the fungus consumed by these apes is a previously undescribed 82 83 species of truffle named here as *Hysterangium bonobo* sp. nov..

#### 84 MATERIALS AND METHODS

Animal observations. — The feeding ecology of the Hali-Hali bonobos (Georgiev et al. 2011), a
wild unprovisioned community, was studied at the Nsondo Camp (0°12'N, 22°51'E),

87 Kokolopori Bonobo Reserve, Province Equateur, Democratic Republic of Congo from Oct 2006

through Jul 2007 with experienced field assistants familiar with the bonobos and the ecology of

the forest (Georgiev et al. 2010; Georgiev et al. 2011). The study site is located approx. 30 km

90 east of a long-term bonobo study site at Wamba and shared many of its ecological features

91 (Hashimoto et al. 1998; Kano and Mulavwa 1984). Three main habitat types are present: 1) dry

92 primary forest with portions dominated by trees in Fabaceae subfam. Detarioideae, including

ectomycorrhizal (ECM) species of *Gilbertiodendron* and *Brachystegia*, 2) seasonally inundated,
riparian swamp forest where *Guibourtia demeusei* (Detarioideae) and ECM *Uapaca* spp.
(Phyllanthaceae) are common, and 3) secondary disturbed forest heavily influenced by slashand-burn agriculture expanding from nearby villages. The seasonality of precipitation at
Kokolopori is similar to Wamba, which typically has up to 2900 mm of rainfall per year during
one light rainy season from Mar – May and a heavier rainy season from Sep – Nov (Hashimoto
et al. 1998; Mulavwa et al. 2008).

The Hali-Hali bonobos have been habituated since 2000 by local conservation NGO Vie 100 101 Sauvage with support from the Bonobo Conservation Initiative (BCI). By Oct 2006, the apes were sufficiently accustomed to human presence to allow behavioral observations on a regular 102 basis. Although the apes allowed detailed monitoring when feeding in the canopy, ground 103 104 observations were less frequent because they did not always tolerate close human proximity during ground foraging. Observational conditions on the forest floor were also limited to 15-20 105 m or less by dense understory vegetation. The bonobo truffle foraging data presented here are 106 thus considered a conservative estimate of truffle consumption. 107

The diet of the bonobos was scored at 15-min intervals by recording food species and 108 109 plant parts (if any) eaten by the majority of individuals in view on the sampling point. A 'running food list' was also recoded on a daily basis to note all foods the bonobos were seen to ingest, 110 whether they happened on the 15-min sample point or not (Georgiev et al. 2010; Georgiev et al. 111 112 2011). We presented data on truffle consumption via a simple dietary score to document the frequency of truffle-eating over the study period, calculated as the monthly proportion of days on 113 which at least one episode of truffle-eating was seen from the total number of days on which 114 115 bonobos were observed and feeding data were recorded. In some cases it was possible to directly

116	observe truffle consumption by bonobos. At other times, truffle feeding was inferred because the
117	bonobos were feeding on the ground and then observers moved into the area after the bonobos
118	began to move away and were able to observe digging and discarded pieces of truffle
119	basidiomata. In Aug 2007 we were able to directly view bonobos consuming basidiomata and
120	then to collect fresh specimens when they were done feeding (see below).
121	Morphological studies. — Four basidiomata were collected from the exact location where
122	bonobos were observed to be feeding on truffles. Specimens were preserved in 99% ethanol. The
123	collection has been accessioned at the Fungal Herbarium of the Florida Museum of Natural
124	History as FLAS-F-64335. Field collected truffles from the Kokolopori bonobo site (FIG. 1)
125	were also directly compared with herbarium specimens from the Oregon State University
126	Mycological Collection (OSC) of Aroramyces radiatus (Lloyd) Castellano, Verbeken &
127	Walleyn, one of the only related truffle species known from tropical Africa (see below).
128	Descriptions of macromorphological characters were based on fresh material and photos.
129	Colors were described in general terms. Microscopic characters were examined based on hand
130	sectioned dried tissues rehydrated in Melzer's reagent, 3% KOH, and water. Photomicrographs
130 131	sectioned dried tissues rehydrated in Melzer's reagent, 3% KOH, and water. Photomicrographs were taken in water. Basidiospore measurements are based on 20 randomly selected
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131 132 133 134 135	<ul> <li>were taken in water. Basidiospore measurements are based on 20 randomly selected</li> <li>basidiospores. Given the remote nature of the region, logistical difficulties of returning to the</li> <li>site, and the infrequence of hypogeous fungal collections in the region, we were unfortunately</li> <li>forced to base the description of this new species on a single collection.</li> <li><i>Molecular methods and phylogenetic analyses.</i> — Clean fungal tissues were taken from inside</li> </ul>
131 132 133 134 135 136	were taken in water. Basidiospore measurements are based on 20 randomly selected basidiospores. Given the remote nature of the region, logistical difficulties of returning to the site, and the infrequence of hypogeous fungal collections in the region, we were unfortunately forced to base the description of this new species on a single collection. <i>Molecular methods and phylogenetic analyses.</i> — Clean fungal tissues were taken from inside the dried specimens that had been previously preserved in ethanol. DNA was extracted using a

139	(White et al. 1990) and the Phusion Hot Start Flex DNA Polymerase standard protocol (New
140	England BioLabs Inc., Ipswich, Massachusetts). Amplification of a portion of nuc 28S rDNA
141	(28S) was performed using the same protocol with forward primer LROR and reverse primer
142	LR3 (Hopple and Vilgalys 1994). PCR products were visualized on 1.5% agarose gels stained
143	with SYBR Green I (Molecular Probes, Eugene, Oregon). Amplicons were cleaned with EXO
144	(Exonuclease I) and SAP (shrimp alkaline phosphatase) enzymes (Werle et al. 1994) and
145	sequenced by GENEWIZ (South Plainfield, New Jersey). Sequences were then edited with
146	SEQUENCHER 5.0.1 (Gene Codes Inc., Ann Arbor, Michigan). The ITS and 28S sequences
147	from our Hysterangium collection were compared with those in the NCBI database using the
148	BLASTn tool (Altschul et al. 1990).
149	For phylogenetic analysis, the 28S sequences of FLAS-F-64335 were placed in an
150	alignment of 28S and mitochondrial ATP synthase membrane subunit 6 (ATP6) DNA sequences
151	of Hysterangiales and allied fungi in Phallomycetidae previously generated by Hosaka et al.
152	(2006, 2008). We were unable to obtain ATP6 sequences from FLAS-F-64335. Sequences were
153	downloaded directly from GenBank and aligned in MESQUITE 3.2 (Maddison and Maddison
154	2018) with the aid of MUSCLE 3.8.31 (Edgar 2004). Independent analyses of 28S and ATP6
155	showed no conflicting phylogenetic signal (data not shown), so the two loci were combined into
156	a single concatenated analysis. The alignment was edited manually to exclude gaps and
157	ambiguously aligned regions.
158	The concatenated alignment was analyzed with maximum likelihood (ML) and Bayesian
159	Inference (BI) as performed in the Cyberinfrastructure for Phylogenetic Research Science
160	Gateway (CIPRES) 3.1 (Miller et al. 2010). ML was run via RAXML 8.2.10 (Stamatakis 2014)
161	with 1000 bootstrap iterations and a GTRGAMMA model under the default paramaters

(Stamatakis 2015). BI was performed in MRBAYES 3.2.7a (Ronquist et al. 2012) using the 162 GTR+I+G model following Hosaka et al. (2008). BI analysis was run on two separate chains 163 using a chain length of one million generations, sampling frequency of 1000, and discarding the 164 first 25% of the samples as the burn-in. The multilocus ML tree was visualized and rooted in 165 FIGTREE 1.4.3 (Rambaut 2016) and Bayesian posterior probability (PP) values were added in 166 ADOBE ILLUSTRATOR CS5.1 (San Jose, California). Nodes were considered strongly 167 supported if ML bootstrap values were  $\geq$ 75% and PP $\geq$ 0.95. 168 169 RESULTS 170 Truffle consumption by bonobos. — Between Nov 2006 and Jul 2007 bonobos were observed on 155 days (range: 12–26 days per month). Truffles were eaten on 38 observation days in 7 out 171 of the 9 study months on a mean of 4.2 days per month (range: 0–10 days), or for a mean of 172 173 23.1% of monthly observation days (range: 0.0–52.9% of monthly observation days). Truffles were, however, a minor element of the diet in terms of their overall contribution to feeding 174 observations (less than 3% of all 15-min, group-level feeding scans for all months). 175 BLASTn and phylogenetic analysis. — BLASTn analysis of both the ITS and 28S sequences 176 generated from FLAS-F-64335 showed clear affinities with other species of ECM Hysterangiales 177 178 and related Phallomycetidae. The ITS from FLAS-F-64335 had the highest homology with 179 uncultured Hysterangiales sequences from ECM roots (e.g., KT461360 from an unknown 180 miombo woodland tree from the DRC; KM402914 from Pseudotsuga menziesii from British 181 Colombia) as well as specimens of Hysterangium (e.g., DQ974810) and Ramaria (e.g., FJ627035). However, the highest hits were only 83–84% similar to the ITS sequence from 182 183 FLAS-F-64335. The 28S also exhibited obvious affinities with ECM Hysterangiales, including 184 92–93% similarity to a wide range of Phallomycetidae such as species of *Hysterangium*,

185	Ramaria, and Austrogautieria (e.g., AF336259, JQ408235, KP191776), as well as ECM root tips
186	of Hysterangiales (e.g., JX316465 from a root of Nothofagus pumilio from Argentina).
187	Our alignment included 770 nucleotides of aligned 28S sequences and 691 nucleotides of
188	aligned ATP6 sequences from 171 taxa. Of these, 76 nucleotides were excluded from the 28S
189	portion of the alignment and 31 nucleotides were excluded from the ATP6 alignment. The ML
190	phylogeny (FIG. 2) depicts FLAS-F-64335 nested among ECM Hysterangium and resolved in a
191	clade separate from Aroramyces radiatus, the only other described ECM Hysterangiales from
192	tropical Africa. The analyses also revealed that Aroramyces is a strongly supported monophyletic
193	group nested within Hysterangium, making the latter paraphyletic. This result was previously
194	found by Hosaka et al. (2006, 2008).
195	TAXONOMY
196	<i>Hysterangium bonobo</i> M.E. Sm. & T.F. Elliott, sp. nov. FIG. 2
197	MycoBank MB834363
198	Typification: DEMOCRATIC REPUBLIC OF THE CONGO (DRC). TSHUAPA
199	PROVINCE: Djolu Territory, Kokolopori Bonobo Reserve, in mixed rainforest with
200	ectomycorrhizal trees in the genera Uapaca (Phyllanthaceae), Brachystegia (Fabaceae), and
201	Gilbertiodendron (Fabaceae), Aug 2007, A. Georgiev MES-127 (holotype FLAS-F-64335).
202	GenBank: $ITS+28S = MT111903$ .
203	Etymology: bonobo, in reference to the common name of the primate Pan paniscus,
204	which digs and eats this fungus.
205	Description: Basidiomata hypogeous to partially emergent, up to 50 mm broad, more or
206	less globose to irregularly globose. Peridium up to 2 mm thick, light to dull brown, apparently
207	bruising brown when damaged or handled, sometimes cracked, smooth, not easily separable

208	from the gleba, with a somewhat hairy appearance in patches. Gleba dark brown, solid, with
209	narrow meandering hollow veins and small open pockets that are not gel-filled, radiating from
210	indistinct off-white columella less than 1 mm wide at the base but becoming indistinct as it
211	radiates upward through the center of the gleba.
212	Peridium 70–150 $\mu$ m thick, comprised of two layers; outer layer 10–65 $\mu$ m thick,
213	composed of interwoven hyphae $3-5 \ \mu m$ wide, hyphae in the outer layer notably darker in color
214	than the inner layer and prominently encrusted with irregular warts $2\times4~\mu m$ or larger, peridial
215	cystidia not observed, clamp connections rare or absent, debris sometimes adhering to the outer
216	layer. Inner peridial layer up to 130 $\mu$ m thick, composed of thin-walled, tightly packed, highly
217	interwoven hyphae 1–3(–5) $\mu$ m wide. Gleba trama of tightly interwoven and unorganized,
218	hyaline, gelatinized hyphae 0.5–1.5 $\mu$ m wide, forming a layer mostly 20–40(–55) $\mu$ m broad.
219	Columella composed of hyaline interwoven hyphae mostly 1–2 $\mu$ m wide, up to 100 $\mu$ m thick at
220	the base but rapidly decreasing in width as it radiates out into the gleba tissues.
221	Basidiospores 14–16 $\mu$ m × 8.5–10 $\mu$ m (mean 15 × 9.5 $\mu$ m), basidiospore walls 1–2 (–3)
222	$\mu$ m thick, Q = 1.40–1.68, mean Q = 1.58, brown in mass, spore wall thickness often irregular but
223	in most spores the walls notably thicker toward the apical end of the spores near the attachment
224	to the sterigmata, ornamentation of very short and somewhat indistinct spines or warts (less than
225	$0.5 \ \mu m$ tall), apical attachments notable and sometimes with a piece of the sterigma broken off
226	from the basidium and still attached to the spore at maturity, faint oil droplets apparent in some
227	spores, spores somewhat dextrinoid in Melzer's reagent. Basidia mostly 2-sterigmate, difficult to
228	view, deflated and irregularly shaped, apparently collapsing after spore dehiscence; sterigmata
229	irregular in shape, visibly running through the spore wall to attach to the spore apex, $2-5 \ \mu m$
230	long and $1-2 \ \mu m$ wide.

231	Notes: One species of Hysterangium and one species of Aroramyces (Hysterangiales)
232	have been previously described from Sub-Saharan Africa, H. niger Lloyd and A. radiatus.
233	Hysterangium niger was originally described from South Africa but is morphologically divergent
234	from all other species of Hysterangium. It was transferred by Zeller and Dodge (1929) to
235	Rhizopogon. We suspect that it is a Rhizopogon species introduced to South Africa with pine
236	seedlings, but the type has not been recently studied. However, based on microscopic and
237	macroscopic morphology it is clearly not conspecific with <i>H. bonobo</i> .
238	Aroramyces radiatus was originally described as Hymenogaster radiatus Loyd but has
239	also been placed in <i>Dendrogaster</i> Buchholtz and <i>Gymnoglossum</i> Massee (Castellano et al. 2000).
240	The genus Dendrogaster, however, is currently considered a synonym of Hymenogaster
241	(Castellano et al. 2000), and the identity of Gymnoglossum remains unverified by sequence data
242	(viz, there are no publicly available sequences in GenBank from the Australian type species
243	Gymnoglossum stipitatum Massee). Accordingly, Castellano et al. (2000) established the genus
244	Aroramyces to accommodate two species, Aroramyces gelatinosporus (J.W. Cribb) Castellano
245	and A. radiatus. The genus was morphologically characterized by brown ornamented spores and
246	brownish gleba, the presence of a columella, a multi-layered peridium, and gelatinized tissues.
247	Castellano et al. (2000) reported A. radiatus with spores that are strongly truncate at the base of
248	the sterigmata attachment and covered by a wrinkled utricle that conceals the spore ornaments
249	when viewed with SEM. Although Castellano et al. (2000) considered A. radiatus in the
250	Cortinariaceae, subsequent phylogenetic studies indicate that A. radiatus is a member of
251	Hysterangiales, and that Aroramyces is nested within Hysterangium (Hosaka et al. 2008).
252	Accordingly, Hysterangium is currently a paraphyletic genus and additional taxonomic revisions
253	are needed but are beyond the scope of this study (Hosaka et al. 2008).

The type of *H. bonobo* (FLAS-F-64335) from a bonobo foraging site at the Kokolopori 254 Bonobo Reserve is superficially similar to the descriptions of A. radiatus in Castellano et al. 255 (2000) and also our direct observations of specimens cited by Castellano et al. (2000) from 256 Zimbabwe. Both species are from tropical Africa and also have a brown gleba, broadly ellipsoid 257 spores and irregular basidia that collapse at maturity. However, the spores of A. radiatus are 258 much smaller than those of *H. bonobo* (mean of  $10.8 \times 6.9 \,\mu\text{m}$  in *A. radiatus* versus  $15 \times 9.5 \,\mu\text{m}$ 259 260 in *H. bonobo*), and the spore ornaments of *A. radiatus* are much larger and more notable than in 261 *H. bonobo*. Furthermore, the spore walls are notably thicker near the attachment to the sterigmata 262 in *H. bonobo*, whereas in *A. radiatus* the spores usually taper the opposite direction and are most narrow near the attachment to the sterigmata. The two species also differ in the peridium 263 morphology. Although both species have encrusted hyphae on the outer peridial layer, A. 264 radiatus has a 3-layered peridium up to 400 µm thick, whereas H. bonobo has a 2-layered 265 peridium typically around 100 µm thick. Phylogenetic analysis also clearly separates these taxa 266 (FIG. 2). 267

We do not know which ECM trees are the symbiotic hosts for *H. bonobo*. However, 268 Hysterangium spp. are known to be obligate ECM fungi (Hosaka et al. 2008) and H. bonobo was 269 found in forests with several confirmed ECM host plants, including species of Uapaca, 270 Brachystegia, and Gilbertiodendron. Species of Uapaca and Gilbertiodendron are also known to 271 272 form ECM associations with several other African truffles (Castellano et al. 2016a, b; Orihara 273 and Smith 2017). Bermejo et al. (1994) also noted the presence of Uapaca and Gilbertiodendron species at sites where they reported bonobos successfully foraging for unidentified truffles. 274 275 DISCUSSION

Many mammals rely on aromas released by mature hypogeous fungi to detect where they 276 are fruiting (Mills 1978; Maser et al. 2008; Stephens et al. 2020). Birds and animals with more 277 limited olfactory abilities may rely on visual cues, including mimicking the color or shape of 278 fruits (Beever and Lebel 2014; Elliott and Marshall 2016; Elliott et al. 2019b; Elliott and Elliott 279 2019; Elliott and Vernes 2019). Some of the truffles described from Africa are brightly colored, 280 281 possibly indicating that their dispersers may rely on their visibility rather than odor (Castellano et al. 2000). It is difficult to determine what alerts bonobos to the presence of subterranean fungi, 282 but it is likely a combination of visual and olfactory cues. Bermejo et al. (1994) described a 283 bonobo that used olfactory cues to locate a truffle: "...standing quadrupedally, digs up the earth, 284 first with one hand, then with the other, in search of subterranean truffles. She puts her face 285 closer to the hole that she has dug and looks closely. Then she carefully puts one hand into the 286 287 hole and withdraws it immediately, putting her fingers to her nose to detect the scent of truffles." Similarly, our observations of the Hali-Hali bonobos foraging on the ground are consistent with 288 the idea that bonobos rely on olfactory cues to detect hypogeous fungi. 289 The Hali-Hali bonobos consumed truffles on a regular basis (23% of all sampling days), 290 indicating that truffles are a component—but not a staple—in the diet of this community of 291 292 bonobos. Hysterangium bonobo was collected directly after we observed bonobos feeding on truffles, therefore leading to the conclusion that bonobos consume *H. bonobo* as a food source. 293 294 Although we expect that bonobos may consume other truffle species in the region, further studies 295 are needed to confirm this hypothesis. Previous studies on bonobo diets in the DRC have also reported that bonobos eat truffles (Yalosidi: Kano 1983; Wamba: Kano & Mulawa 1984; 296 297 Lilungu: Bermejo et al. 1994), but truffles have always been considered a minor component in

their diet. The most detailed report on bonobo truffle-eating comes from the Ikela study site,

where the apes consumed truffles on 18 days over a 605-day-long study, totaling 686 hrs and 47 299 min of direct observations (Bermejo et al. 1994). We saw bonobos consume truffles on 38 days 300 (from a total of 155 observation days), so it is likely that the Ikela bonobos eat fewer truffles than 301 the bonobos at Kokolopori. The identity of the truffles from previous studies have remained 302 unknown, although Kano (1983) suggested that the puffball Langermannia fenzlii (Reichardt) 303 Kreisel might be a food source for bonobos. However, we can find no evidence or specimens to 304 support this hypothesis. Despite being consumed infrequently and in low quantities, the 305 nutritional value of fungi can still constitute an important part of an animal's diet (Wallis et al. 306 307 2012).

Because the truffles at Ikela (Bermejo et al. 1994) were never collected or identified we 308 cannot determine whether the bonobos observed in that study consumed H. bonobo, A. radiatus, 309 or other truffle species. It is also possible that the truffle species at the Ikela site were less 310 appealing to bonobos or that the Ikela study occurred during seasons or years with less fungal 311 fruiting than our studies at Kokolopori. Near Kokolopori the local villagers use the Bongando 312 word "simbokilo" to refer to truffles (Takemoto 2017). There is some evidence of wider 313 consumption of simbokilo by other mammals because the truffles are used to bait traps during 314 315 forest hunting expeditions (Kimura et al. 2015). Simbokilo is specifically useful for trapping *Cricetomys emini* (Emin's pouched rat) but has also been documented by local people to attract 316 317 Atherurus africanus (African brush-tailed porcupine), at least three species of duiker 318 (Cephalophus monticola, C. callipygus, C. nigrifrons), and several species of squirrels (A. Lokasolac, pers. obs.). The word "simbokilo" is derived from 'simba' (don't go away) and 319 320 'bokilo' (brother-in-law) and derived from a longer phrase "do not allow your brother-in-law to 321 go away because there will be plenty of food coming from traps using simbokilo as a bait." This etymology of simbokilo is indicative of the regular use of this truffle by local people (A.
Lokasolac, pers. obs.). However, as far as we know, our collection of *H. bonobo* is the only
collection of simbokilo that has been examined microscopically or molecularly. It therefore
remains unclear whether this word refers to *H. bonobo* specifically or to a suite of truffles. More
direct observations and collections are needed to determine whether simbokilo is one truffle
species or several truffle species.

There have been scattered reports of mycophagy among primates in Africa and other 328 parts of the world (see Hanson et al. 2003). Most studies are based on visual observations of 329 330 feeding, but the fungal taxa are rarely or never identified. Most reports of mycophagy give vague descriptions of the macroscopic morphology of the fungi that provide little assistance to 331 taxonomists, e.g. "bracket fungi consumed by gorillas" (Fossey 1983). We urge zoologists 332 working with animal diets in the future to collect, photograph, and preserve voucher specimens 333 of the fungal taxa eaten in order to allow for more in-depth taxonomic studies by fungal 334 335 biologists.

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# 527 FIGURE LEGENDS and FOOTNOTES

Figure 1. Morphological features of the holotype collection of *Hysterangium bonobo* collected 528 in a bonobo foraging site after bonobos had recently been feeding. A. Fresh fruiting body of H. 529 bonobo in hand, note the slight brownish discoloration from handling. B. Broken H. bonobo 530 fruiting body revealing brown gleba and columella at the center. C. Peridial hyphae with brown 531 encrusted warts. D. Overview of the gleba showing locules of hymenial tissue lined with basidia 532 and basidiospores interleaved by tramal plates composed of densely interwoven sterile hyphae. 533 Interwoven peridial hyphae on the inner surface of the peridium is visible in the upper left-hand 534 corner of the image. E. Thick-walled basidiospores that are still attached to narrow basidia are 535 536 visible at the far left and right of the image. Note that the sterigma appear to grow through the thick cell walls. The basidiospore in the center of the image shows the fine ornamentation that is 537 538 present on mature basidiospores at high magnification. Bars: A-B = 2 cm; C,  $E = 10 \mu$ m, D = 30539 μm.

Figure 2. ML phylogenetic tree of Hysterangiales and other fungi in the Phallomycetidae based
on analysis of 28S and *ATP6* showing phylogenetic placement of *Hysterangium bonobo* sp. nov.

542 within the genus *Hysterangium*. Note that *H. bonobo* is resolved in a clade that is

- 543 phylogenetically distant from *Aroramyces radiatus*, the only other described species of
- 544 Hysterangiales from tropical Africa. Several taxa in Agaricales, Thelephorales, and Boletales
- served as outgroups. Support values are shown above the nodes using the following format: ML
- 546 bootstrap values  $\geq 75\%$  / posterior probabilities  $\geq 0.95$ .