

1 Which web to invade? Argyrodine kleptoparasites
2 differentiate among architecturally different host webs
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30 **Abstract**

31 Kleptoparasitism, the theft of resources from another organism, is a survival strategy
32 found across the animal kingdom. Many argyrodoxinae cobweb spiders (Theridiidae) are
33 obligatory kleptoparasites that have largely abandoned web building, relying instead on
34 webs of larger hosts. Theory predicts, and limited prior research indicates, that
35 kleptoparasites are not randomly distributed among host webs, and prior studies indicate
36 that web architecture (size and tenure) and prey availability impact kleptoparasite
37 abundance and host choice. We investigate the distribution of multiple argyrodine
38 kleptoparasite species among three contrasting host types in Madagascar's
39 Analamazaotra National Park: *Nephilingis* (Nephilidae, large nocturnal orb weavers),
40 *Caripetella* (Pisauridae, large nocturnal sheet web builders), and *Anelosimus* (Theridiidae,
41 small cathemeral social spiders constructing three-dimensional tangle webs). Although
42 small in scope, this study is the first to take a real-time snapshot of multi-species endemic
43 communities of spider kleptoparasites and their in-situ distribution across contrasting host
44 webs. We found an unexpectedly high diversity of ten kleptoparasite species that,
45 remarkably, are not conspecific with the five that were previously known from all of
46 Madagascar. Kleptoparasite species composition and abundance varied across the three
47 hosts: some appeared host-specific while others seemed versatile. In general, argyrodine
48 kleptoparasites evidently discriminate among hosts but choose among them with varying
49 degrees of astucity. At the community level, very limited data exist, and based on our
50 study we can only speculate that species assembly into host webs involves a complex
51 interplay between host preference and species competitive hierarchy. Future field
52 research should investigate this system as a community of multiple interacting species to
53 gain a more comprehensive understanding of the rules that may govern the assembly of
54 diverse kleptoparasites into equally variable host webs. Synergistically, laboratory
55 experiments are needed to elucidate key cues facilitating kleptoparasite web detection
56 and choice; likely involving eavesdropping on pheromones intended for the hosts male.

57

58 **1. Introduction**

59 Species interactions are diverse, ranging from indirect to direct and from mutually
60 beneficial to hostile. Leung and Poulin [1] define a symbiotic relationship as “an intimate

61 interaction between different organisms, where at least one of the parties is obligatorily
62 dependent on the association as a part of its life history". Symbiosis can be mutualistic,
63 where both parties benefit, commensal, where one species benefits and the other is
64 unaffected, or parasitic, where the parasite benefits and the host is harmed [1-4].
65 Predation and parasitism are prominent forms of antagonistic animal interactions,
66 foraging modes involving the exploitation of innate resources—bodily tissue, fluid,
67 behavior—of the victim. Kleptoparasitism is another exploitative foraging strategy
68 observed among various animals including birds, mammals, insects, and arachnids [5-6].
69 It involves the theft of extrinsic resources that are of potential value to the host that has
70 secured or produced them. Kleptoparasitism is an intriguing adaptive strategy that
71 requires balancing, among other things, the energy costs of searching for food against
72 the benefits and costs of stealing resources from others [7]. While facultative
73 kleptoparasitism is common [e.g., 6], natural selection has engrained this strategy in
74 some groups, where host-selection may be a more critical foraging decision than the
75 exact nature of the stolen resource. Precious little, however, is known about host choice
76 in many kleptoparasitic organisms like spiders.

77 Obligate kleptoparasitic spiders have largely abandoned web building, relying
78 instead on symbiosis in the webs of larger hosts to forage in [8-11]. This strategy
79 represents a complex interplay of evolutionary adaptations, ecological dynamics,
80 behavioral decision making, and doubtless, a variety of largely unknown sensory cues [5,
81 12-13]. The majority of known spider kleptoparasites belong to the subfamily Argyrodoxinae
82 (Theridiidae, cobweb spiders), a versatile group employing diverse strategies to exploit
83 various host species. Among their hosts are 1) large nephilid orb weavers such as *Nephila*,
84 *Trichonephila*, and *Nephilingis*, that seem to be particularly important/preferred hosts in
85 most subtropical and tropical environments [12], 2) large araneid orb weavers such as
86 *Caerostris*, *Cyrtophora*, and *Gasteracantha*, 3) large sheet weavers including various
87 agelenids, the pisaurid *Caripetella*, the tengellid *Tengella*, and others, 4) social and
88 colonial spiders (mostly Araneidae, Eresidae, and Theridiidae) that build complex silken
89 networks, and 5) some smaller species (notably Theridiidae and Linyphiidae) that are
90 locally abundant [14-21].

91 Argyrodinae spiders are speciose [13, 22-24] and vary in size, morphology,
92 coloration, and tactics to obtain resources from host webs [5, 11]. These tactics include
93 pilfering small insects ignored by the host, stealing wrapped prey, feeding on digested
94 prey alongside the host, opportunistic predation on the host, its offspring, and eggs, and
95 even the consumption of host web silk [5, 12, 25]. Their hosts are also highly diverse,
96 differing in size, innate aggression (solitary versus communal or social spiders), web
97 architecture, and prey choice. Given this diversity of kleptoparasites and hosts,
98 fundamental ecological foraging and game theories predict that kleptoparasitic species
99 do not distribute randomly among potential host webs [6, 26-28]. Field observations
100 corroborate these fundamental predictions: large, long-lived webs tend to contain
101 kleptoparasites, while they are rarely found in webs of spiders that prefer small prey, and
102 those that have short duration in space and time [5, 12, 28-31]. The few available field
103 studies broadly agree that 1) distribution of kleptoparasites is non-random among hosts,
104 2) large and predictable webs are favored by kleptoparasites, 3) larger webs have more
105 kleptoparasite individuals and species, and 4) kleptoparasite species range from versatile
106 generalists to relative host specialists [5, 9-12, 28, 32-38]. However, most prior studies
107 focus on a single kleptoparasite or host species [32, 37-39], and none have investigated
108 the in-situ distribution of a species-rich kleptoparasite community among architecturally
109 diverse host webs.

110 Here, we investigate the real-time distribution of a highly endemic spider
111 community: multiple argyrodine species among three hosts in the small Analamazaotra
112 National Park in Madagascar. The three host species are among the most abundant web
113 builders in the reserve but contrast starkly in web architecture and behavior (Fig. 1, see
114 Methods): We surveyed multiple webs of all three hosts and conducted translocation
115 experiments between *Nephilingis* and *Caripetella* webs to explore kleptoparasite diversity,
116 the assembly of kleptoparasite communities into host webs, and evaluate the evidence
117 for host choice.

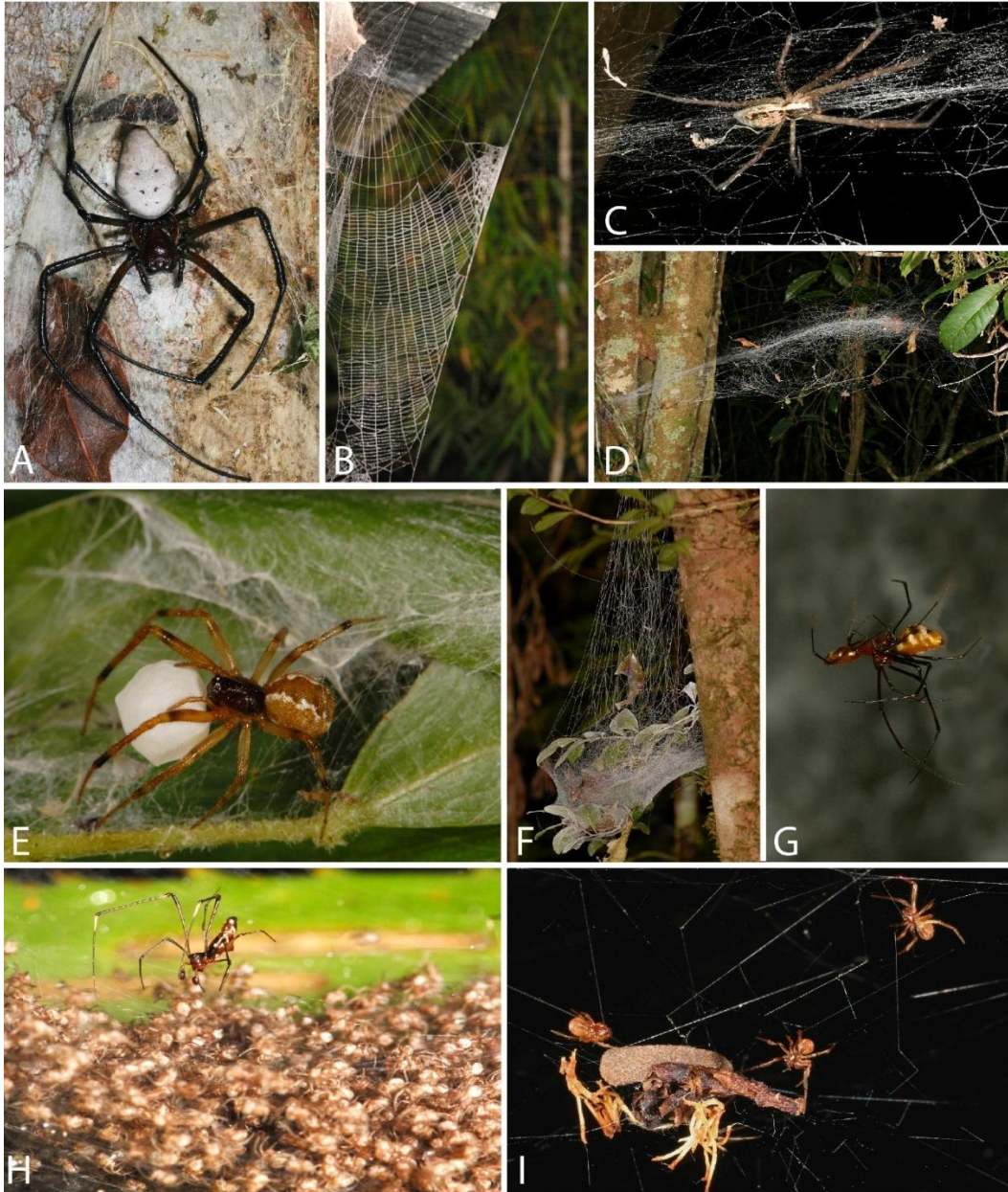
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119 **2. Materials and Methods**

120 2.1. Study system

121 The host species belong to three distantly related families and while all build long lasting
122 webs, they differ in web architecture, size, activity patterns, and innate aggression (Fig.
123 1). *Nephilingis livida* [29] is a large mostly nocturnal nephilid building vertical orb webs
124 that contain sticky silk. The pisaurid *Caripetella madagascariensis* [40] is a large, strictly
125 nocturnal, solitary spider in horizontally domed sheet webs made of dry silk. Spiders of
126 the genus *Anelosimus* are subsocial, cathemeral, and build three-dimensional tangles of
127 dry silk that contain numerous individual spiders. Analamazaotra is home to about 10
128 highly similar species of *Anelosimus* that we sampled indiscriminately. A few webs of the
129 subsocial pisaurid *Dendrolycosa* were also sampled, highly similar in web structure and
130 content to *Anelosimus*. All three web types have extensive networks of non-sticky
131 structural threads separated from the capture area of the web, where the kleptoparasites
132 accumulate when not foraging.

133



134
 135 Figure 1: Study system. A-B, *Nephilingis livida* host. A, female. B, vertical orb web. C-D,
 136 *Caripetella madagascariensis* host. C, female in web. D, horizontal domed sheet web. E-
 137 F, *Anelosimus vondrona* Agnarsson & Kuntner, 2005 host. E, female with egg sac in web.
 138 F, social 3D tangle web. G-I, kleptoparasites in action. G, *Argyrodes* sp. 1 from this study,
 139 pair in copula in *N. livida* web. H, *Argyrodes* sp. 3 from Montagne d'Ambre, male
 140 approaching freshly hatched *N. livida* spiderlings. I, *Argyrodes* kleptoparasites stealing
 141 prey from *Nephilingis borbonica* [29] in Réunion. Photographs are not to scale, total length
 142 of females (front of carapace to spinnerets) *N. livida* ~20 mm, *C. madagascariensis* ~18-

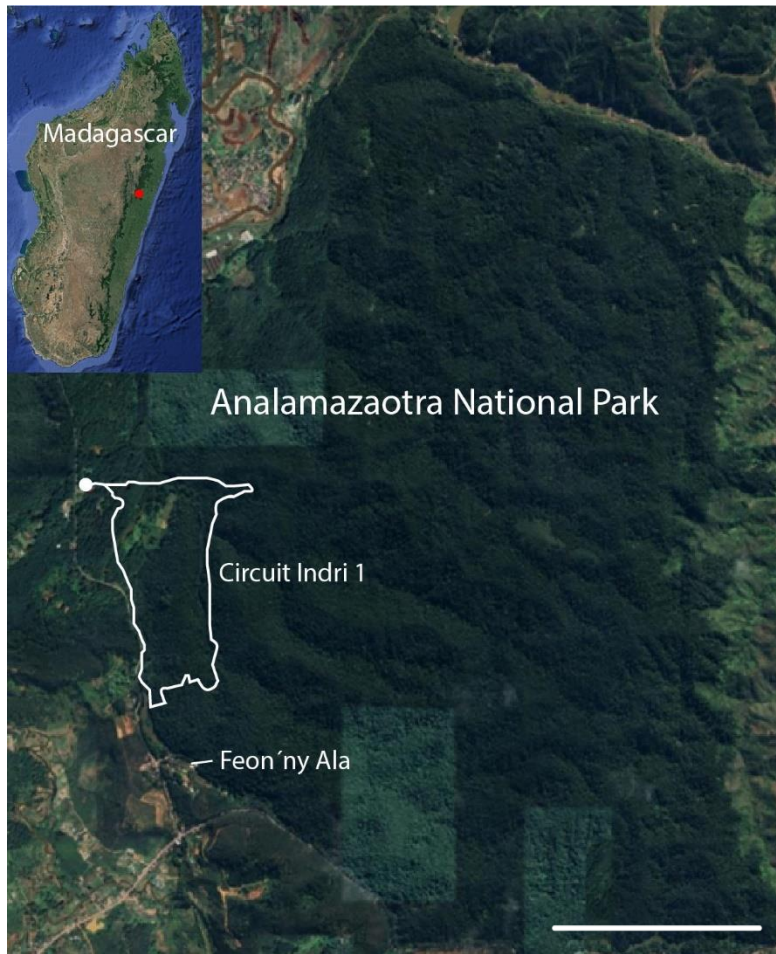
143 25 mm, *A. vondrona* ~5 mm, argyrodines ~3-6 mm. Web size (longest axis), *N. livida* orb
144 ~40-80 cm high, *C. madagascariensis* sheet ~30-60 cm wide, *A. vondrona* basket ~10-
145 25 cm width, up to ~30 cm tangle above. Photos A, C, D, I by M. Kuntner, B, E-H by I.
146 Agnarsson.

147

148 2.2. Survey of kleptoparasites in host webs

149 This study was conducted in the small Analamazaotra National Park, a montane
150 rainforest reserve in eastern Madagascar (25.5 km², 900-1050 m alt., center ca.
151 18°56'10"S, 48°25'42"E, Fig. 2), during 21-28 May 2024. Samples were taken of
152 kleptoparasitic argyrodine (Theridiidae) spiders, all presumably endemic to Madagascar,
153 found in three types of host webs: *Nephilingis livida*, *Caripetella madagascariensis*, and
154 subsocial webs, most belonging to *Anelosimus vondrona*, *A. nazariani*, and *A. may*, and
155 a few *Dendrolycosa* sp. (Fig. 1, Table 1). The host species are likewise endemic to
156 Madagascar, except *N. livida* that also occurs on small islands in the northern Indian
157 Ocean [41]. *Caripetella* and various highly similar *Anelosimus* species [42-44] are
158 common in the forest and were mainly collected along the 2.5 km long Indri 1 trail loop
159 that spans the western portion of the reserve (Fig. 2). While *Nephilingis* webs were also
160 sampled within the forest, most samples were taken at the forest edge from the grounds
161 of the Feon'ny Ala cottages (Fig. 2).

162 The kleptoparasitic spiders were readily visible in webs of *Nephilingis* and
163 *Caripetella* and were captured with manual aspirators (pooters) from ~20 webs of each
164 host and placed in live collection vials. In contrast, in the dense 3D tangle webs of
165 subsocial spiders, kleptoparasites were challenging to detect among the many inhabitants,
166 and the host species cannot be reliably identified in the field. The entirety of 87 subsocial
167 colonies (75 *Anelosimus* and 9 *Dendrolycosa*) were therefore collected in plastic bags
168 and later dissected, with all the web inhabitants preserved in ethanol. Kleptoparasites
169 from *Nephilingis* and *Caripetella* webs were kept alive and used in the translocation
170 experiments described below, before being preserved in ethanol. Taxonomic
171 identification and molecular studies were subsequently performed at the University of
172 Iceland.



173

174 Figure 2: The small Analamazaotra National Park (25.5 km²), located in eastern
175 Madagascar, see inset map (note that for visibility, the red dot covers a vastly larger area
176 of Madagascar than the reserve). The edges of the park are clearly visible by changes in
177 vegetation, evident by dark green forest cover, surrounded by lighter green vegetation
178 and brownish human developments. The research was done along the approximately 2.5
179 km Indri trail loop (circuit Indri 1), and on the grounds of the Feon'ny Ala cottages. Scale
180 bar in the lower right corner is 1 km.

181

182 2.3. Translocation experiments

183 To test whether the kleptoparasitic spiders show host-specific or generalist tendencies, a
184 simple field experiment was conducted. Kleptoparasitic spiders from two of the host
185 species, *N. livida* and *C. madagascariensis*, were collected and kept in live vials until they
186 were placed onto the web of the other (*N. livida* kleptoparasites into *C. madagascariensis*

187 webs and vice versa). Kleptoparasitic spiders were placed onto webs that had been
188 cleared of native kleptoparasites, who were used in the reverse experiment. In a total of
189 17 translocation experiments, four to seven foreign kleptoparasitic spiders were
190 successfully placed onto each cleared host web: 54 native *Caripetella* kleptoparasitic
191 spiders on 10 *Nephilingis* webs and 34 native *Nephilingis* kleptoparasitic spiders on seven
192 *Caripetella* webs. All spiders were placed using their lifeline, inasmuch as possible and
193 depending on access to the web, feet-first to prevent entanglement, onto anchor threads,
194 frame threads, spiral or radius threads, near the outer edges of the web. In the first
195 experiment, the behavior of both the introduced kleptoparasitic spiders and hosts were
196 observed for a period of 45-60 minutes. Our observations were focused on kleptoparasite
197 movement within webs, the nature of which can possibly reflect a kleptoparasitic spider's
198 evaluation of risk [45] and, to an extent, outside the web. Additional behaviors, such as
199 silk deposition, were also noted. Movements of the kleptoparasitic spiders were cataloged
200 into the following categories: 1) stayed on the threads of the host web 2) stayed in the
201 vicinity of host web, either on its own lines connected to the host web or sitting within 5
202 cm of host web support lines, 3) left host web completely. The translocation experiments
203 were performed during daytime.

204

205 2.4. Statistical analyses

206 To investigate the association between kleptoparasitic species and host web types we
207 used Fisher's Exact Test, appropriate for small sample sizes and low expected values.
208 Given the size of our contingency table, we used a simulation-based approach with one
209 million replicates to ensure accurate p-value calculations.

210 The contingency table, detailing the distribution of kleptoparasitic species across
211 host web types, was first constructed in Excel (Microsoft, USA) and then imported into
212 RStudio [46] for analysis. To investigate the association between kleptoparasitic species
213 and host web types, we performed statistical tests in RStudio using this table as the
214 dataset. The RStudio code for these analyses is available as supplementary material.

215

216 3. Results

217 3.1. Survey of kleptoparasites in host webs

218 A total of 316 argyrodine individuals were sampled (Table 1), belonging to a total of 10
 219 putative argyrodine species based on morphology and DNA barcodes. These species
 220 were preliminarily assigned to the groups *Argyrodus* (Arg sp. 1-4), *Argyrodus minax*-
 221 group (Ami sp. 1-4—tentatively, denoting a group of species similar to ‘*Argyrodus*’ *minax*,
 222 clearly distinct from other *Argyrodus*, but not well placed in existing genera), *Argyrodella*
 223 (Ard sp.1—tentatively, this monotypic genus is only known from the Seychelles), and an
 224 additional unknown genus (‘Unk’).

225

226 *Table 1.* The distribution and abundance of the ten kleptoparasite species found among
 227 the studied three host webs (Arg, *Argyrodus*; Ard, *Argyrodella*; Ami, *Argyrodus minax*
 228 group; Unk, unknown). Cell-shading marks the relative abundance of kleptoparasite
 229 species for each host web.

Host	Arg. 1	Arg. 2	Arg. 3	Arg. 4	Ard. 1	Ami. 1	Ami. 2	Ami. 3	Ami.4	Unk	Total
<i>Nephilingis</i>	72 66.1 %	2 1.8 %	0 0 %	0 0 %	1 0.9 %	23 21.1 %	5 4.6 %	5 4.6 %	0 0 %	1 0.9 %	109
<i>Caripetella</i>	0 0 %	0 0 %	13 9.3 %	1 0.7 %	8 5.7 %	78 55.7 %	5 3.6 %	21 15.0 %	13 9.3 %	1 0.7 %	140
<i>Anelosimus</i>	0 0 %	0 0 %	4 6.0 %	0 0 %	0 0 %	32 47.8 %	12 17.9 %	0 0 %	4 6.0 %	15 22.4 %	67

230

231 3.2. Translocation experiments

232 Of the total 54 *Caripetella* native kleptoparasitic spiders introduced onto 10
 233 *Nephilingis* webs, 15 left the web entirely, while 36 stayed in the host web vicinity, less
 234 than 5 cm from the support lines, sometimes connected to them with the kleptoparasite
 235 own lines, but none stayed on the host capture area. The *Nephilingis* host responded
 236 twice towards the introduced kleptoparasitic spiders by attacking and killing them.

237 Of the total 34 *Nephilingis* native kleptoparasites introduced onto seven *Caripetella*
 238 webs, three left the web entirely, 24 stayed on the host capture area, and seven stayed
 239 in the vicinity of the host web. The three spiders that left immediately were all large and
 240 presumed to belong to *Argyrodus* sp. 1. Many introduced kleptoparasitic spiders were
 241 observed moving onto the numerous vertical support threads of the *Caripetella* domed
 242 sheet webs. It was not always clear whether the spiders that migrated towards these
 243 upper vertical support threads were staying on them or on their own attached silk threads.

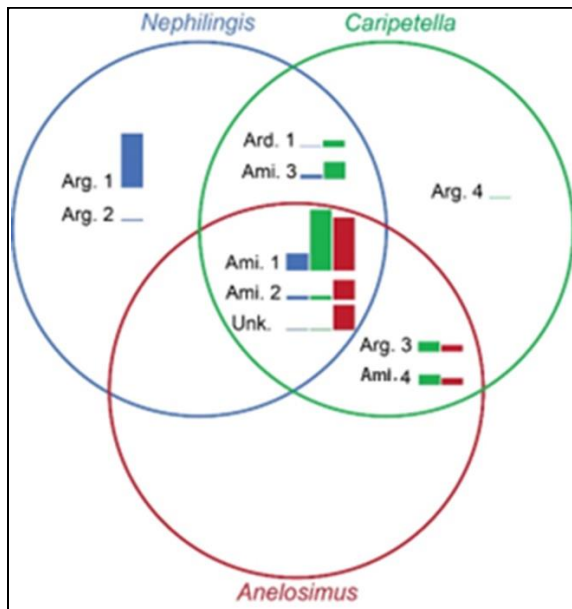
244 No response was recorded towards introduced kleptoparasitic spiders by *Caripetella*
 245 hosts.

246 While few details can be concluded from the translocation experiments, the
 247 behavior of introduced kleptoparasitic spiders clearly differs between the different host
 248 webs. None of the *Caripetella* native kleptoparasitic spiders introduced onto *Nephilingis*
 249 webs stayed on the web itself, while approximately 70% of the *Nephilingis* natives
 250 stayed on the *Caripetella* web. In addition, approximately 28% of native *Caripetella* left
 251 *Nephilingis* webs entirely, while approximately 9% of native *Nephilingis* left *Caripetella*
 252 webs.

253

254 3.3. Distribution of kleptoparasitic spiders among host webs.

255 Fisher's Exact Test with simulated p-values based on one million replicates
 256 revealed a significant difference in kleptoparasite species association with the three
 257 host web types ($p = 1 \times 10^{-6}$). These results suggest 1) webs of the three host species
 258 represent three different habitats, and 2) at least some kleptoparasite species show
 259 strong host-preferences.



260
 261

262 Figure 3. Distribution and overlap of kleptoparasitic species found on different host
 263 webs. Arg, *Argyrodes*; Ard, *Argyrodella*; Ami, *Argyrodes minax* group; Unk, unknown.

264

265 **4. Discussion**

266 4.1. Kleptoparasite diversity and distribution among host webs

267 This study, conducted in the small Analamazaotra National Park in Madagascar,
268 investigates the distribution of kleptoparasitic argyrodine species among the webs of
269 three host species. As a labor intense but brief student-driven field-course project, the
270 study provides a unique, real-time snapshot of the distribution of an impressively diverse
271 and endemic community of kleptoparasites, across contrasting webs of equally endemic
272 hosts. The hosts include *Nephilingis livida*, a large, mostly nocturnal spider constructing
273 a huge and sticky vertical orb web [47], *Caripetella madagascariensis*, an exclusively
274 nocturnal giant nursery web spider that makes one of the world's largest horizontal domed
275 sheet webs, and the cathemeral *Anelosimus* and *Dendrolycosa* form subsocial spider
276 communities and construct three-dimensional tangle webs, whose numerous
277 inhabitants—a mother and her offspring—are barely larger than the kleptoparasites [41].
278 Through direct sampling, and simple translocation experiments, we discovered a high
279 diversity of kleptoparasite species and show that while generally versatile, kleptoparasites
280 do distinguish among the different host webs.

281 Remarkably, at least ten argyrodine species were sampled in the webs of three
282 host species (among a large number of potential local host species), along a 2.5 km linear
283 transect. The study area represents less than 1/200.000th of this vast island, yet we
284 sampled double the diversity of kleptoparasitic argyrodine spiders hitherto known from all
285 of Madagascar. Furthermore, it appears that none of our ten species are among the
286 Madagascar's currently reported five. This observation underscores both the astounding
287 biodiversity of Madagascar and the high conservation value of protected forests, even
288 tiny patches. While remarkable, this discovery aligns with the reputation of Analamazaotra
289 and the neighboring Mantadia National Parks as rich in spider diversity, e.g., as a world
290 hotspot of the cosmopolitan *Anelosimus* spiders [43-44] and old world *Caerostris* spiders
291 [48].

292 Kleptoparasite species composition and abundance differed significantly among
293 host webs. Some kleptoparasites appear to be highly selective, for example, the most
294 common kleptoparasite in *Nephilingis* webs, *Argyrodes* sp. 1, was absent in other host
295 webs, and most other species were biased towards a particular host species (Fig. 3). One

296 kleptoparasite species, *Amin* sp. 1, was common in all host webs (though most abundant
297 in *Caripetella* webs), and some occurred in webs of two out of three hosts. While
298 *Caripetella* and *Nephilingis* webs thus represent distinct habitats for kleptoparasites, each
299 occupied by 7-8 argyrodine species, they also support a shared argyrodine community of
300 five species.

301 Currently, it is impossible to know whether all kleptoparasite species found in a
302 particular host web actively use these as resource sites. Given that some kleptoparasite
303 species occur in low abundance it seems likely that, in search of preferred webs, they
304 may occasionally wander into webs they do not use for resource stealth. This caveat
305 aside, our results suggest that most kleptoparasite species are, to some degree, versatile
306 and capable of utilizing a variety of (perhaps) suboptimal hosts. Notably, relatively few
307 species seem to make use of *Anelosimus* webs. Only five of the ten detected species
308 were found there, and all occurred in low abundance, averaging less than one individual
309 per web. Yet one species, marked 'Unknown', is almost exclusively found in subsocial
310 webs of *Anelosimus* and *Dendrolycosa*. The relatively small size of *Anelosimus* webs may
311 restrict kleptoparasite abundance, though a comparison between the size of 2D and 3D
312 webs is not straightforward. Beyond that, parasitizing social webs inhabited by numerous,
313 ever active, juvenile spiders not much larger than the kleptoparasites themselves, may
314 require distinct behavioral adaptations compared to living in a web with a solitary giant
315 nocturnal host.

316 Our translocation experiments provide further support for these patterns. For
317 example, many individuals collected in one host web and introduced into a kleptoparasite-
318 free web of another host, chose to leave. Notably, three *Argyrodes* sp. 1 individuals we
319 could identify with some confidence in the field, were the only ones to immediately leave
320 a *Caripetella* web when introduced. As *Argyrodes* sp. 1 was never found in *Caripetella*
321 webs in nature, indicating the ability of this species to rapidly detect non-preferred host
322 webs. Furthermore, the introduced kleptoparasitic spiders were generally more accepting
323 of *Caripetella* webs than *Nephilingis* webs, reinforcing the prediction that kleptoparasites
324 differentiate among hosts. Most *Caripetella* native kleptoparasitic spiders introduced onto
325 *Nephilingis* webs stayed in their vicinity, although some were not found in *Nephilingis*
326 webs in natural conditions, indicating that this nephilid might serve at least as a facultative

327 host for most of kleptoparasitic species. However, these experiments were preliminary
328 and must be interpreted with caution: they might have been impacted by external
329 variables of an unpredictable rainforest habitat (e.g., wind and rain) and limited by our
330 inability to monitor individual kleptoparasite species in the field. Furthermore, due to time
331 constraints, we were unable to conduct control experiments by reintroducing *Caripetella*
332 kleptoparasites to *Caripetella* webs and vice versa.

333 Taken together, our evidence supports that argyrodine kleptoparasites are
334 generally versatile, but not randomly distributed or indifferent to host species. While these
335 results may appear somewhat inconclusive, this likely simply reflects the complexity of
336 the system—kleptoparasite species do not neatly fall into square categories such as
337 ‘specialist’ and ‘generalist’. The current results align well with prior research and allow
338 some general conclusions and promising avenues for future research. Attempting to fit
339 argyrodines into categorical bins, e.g., host generalist vs. specialist, is tempting, but even
340 for relatively choosy kleptoparasites [e.g., 8, 49-50], such boxes have proven to be ill-
341 fitting [11-12, 38, 51-53]. Instead, argyrodines may better be characterized as generally
342 capable of discriminating among host webs, but with varying degrees of astucity or
343 concern. Prior research, focusing mostly on single kleptoparasitic species in isolation, has
344 established that web architecture and prey availability are important factors in individual’s
345 host choice [14, 32-33, 38, 49, 51, 54]. However, in diverse communities of kleptoparasite
346 and host species, underlying mechanisms of host choice are, no doubt, more complex
347 and interactive. Therefore, to understand the distribution of kleptoparasitic species in
348 ecosystems, we need to address host choice at the level of community assembly.

349 Based on our findings and prior research, we here offer some speculative
350 hypotheses that may serve to guide future research. First, some kleptoparasite species
351 show minimal preference (e.g., *Argyrodes minax* group sp. 1 in this study), perhaps being
352 species that are highly versatile and/or stealthy. Other species astutely choose hosts: For
353 example, *Argyrodes* sp. 1 is the largest kleptoparasite in this study and may specialize
354 on hosts that are the most likely to catch relatively large prey, such as nephilid spiders
355 [55]. An alternative hypothesis, though not mutually exclusive, is that the observed
356 distribution of kleptoparasites results from competition rather than microhabitat
357 specialization. *Argyrodes* sp. 1 may, for example, have competitive advantage due to size

358 and occupy the best habitat patches; Nephilidae appears to be the preferred argyrodine
359 host globally. In a similar lane, the small and delicate 'Unknown' species may prefer social
360 nests or may be competitively inferior to other argyrodines and thus restricted to smaller
361 and less favorable habitats.

362 Second, while '*Argyrodes* sp. 1' and '*Argyrodes minax*-group sp. 1' approximately
363 exemplify opposite ends of the host-choice spectrum, being host-specialists and -
364 generalists, respectively, we hypothesize that other species in this study, and perhaps
365 most argyrodines, likely fall somewhere in between. Thus, deciphering the role of
366 specialization vs competition in shaping kleptoparasite distribution among hosts is an
367 urgent priority. Determining how argyrodine kleptoparasites locate host webs will be a
368 critical segue into testing these hypotheses, as we first need an understanding of the
369 mechanisms of web discovery and kleptoparasite determination of habitat suitability. The
370 extent to which host choice is guided by host specific cues, such as airborne pheromones
371 (that may operate both locally and at a distance, see supplementary data), should offer
372 evidence on specific host choice. Insights into the role of intra-guild competition could, in
373 turn, be gained by examining how kleptoparasites already present in an encountered web,
374 impact further settlement into host webs.

375 In conclusion, our study shows that argyrodine kleptoparasites are apt at locating
376 host webs and choosing among them. This choice, or more generally, the community
377 assembly of kleptoparasite species into host webs, likely results from an interplay of
378 host-specialization and interspecific competition. We predict that host-specific chemicals
379 (likely pheromones in silk, M. Elgar pers. comm.) inform the kleptoparasites on host
380 choice and speculate that sensory adaptations stemming from an existing system
381 guiding male spiders to conspecific female webs, have played a role in the evolution of
382 web kleptoparasites. This fascinating and accessible study system of giant hosts and
383 tiny thieves occurring in highly distinct habitats—large spider webs—remains poorly
384 investigated while offering a wealth of opportunities for topical research including
385 foraging/game theory, coevolution, diversification, and the role of assembly rules in
386 community structuring. To facilitate such studies, fundamental field research is needed
387 on the diversity and degree of specialization of web kleptoparasites (both in host choice

388 and foraging mode), as are basic laboratory experiments on pheromones and other
389 cues that play a role in kleptoparasite web detection and choice.

390

391 **Data availability**

392 COI barcoding sequences will be submitted to GenBank (in progress). Other data are
393 available from the authors.

394

395 **Conflicts of Interest**

396 The authors declare no conflict of interest regarding the publication of this paper.

397

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408

409 **References**

410

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