

1 Article

2 Trade in Zambian edible orchids – DNA barcoding 3 reveals use of unexpected orchid taxa for *chikanda*.

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31 **Abstract:** In Zambia wild edible terrestrial orchids are used to produce a local delicacy called
32 *chikanda*, which has become increasingly popular throughout the country. Commercialization puts
33 orchid populations in Zambia and neighbouring countries at risk of overharvesting. Hitherto, no
34 study has documented which orchid species are traded on local markets, as orchid tubers are
35 difficult to identify morphologically. In this study, the core land-plant DNA barcoding markers rbcL
36 and matK were used in combination with nrITS to determine which species were sold on Zambian
37 markets. Eighty-two interviews were conducted to determine harvesting areas, as well as possible
38 sustainability concerns. By using nrITS DNA barcoding, a total of 16 orchid species in six different
39 genera could be identified. Both rbcL and matK proved suitable to identify the tubers up to genus-
40 or family level. *Disa robusta*, *Platycoryne crocea* and *Satyrium buchmanii* were identified most
41 frequently and three previously undocumented species were encountered on the market. Few
42 orchid species are currently listed on the global IUCN Red List. Local orchid populations and
43 endemic species could be at risk of overharvesting due to the intensive and indiscriminate
44 harvesting of *chikanda* orchids and we therefore encourage increased conservation assessment of
45 terrestrial African orchids.

46 **Keywords:** CITES; Chikanda; Conservation; DNA barcoding; Orchids; Species delimitation;
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48 1. Introduction

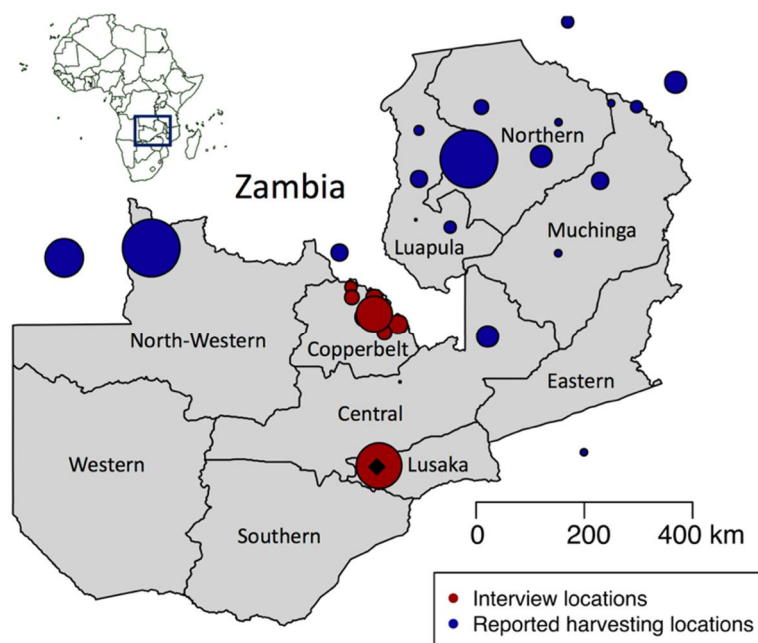
49 Terrestrial orchids have been used for medicinal and culinary purposes for centuries [1], with
50 the most notable example being the use of orchid tubers to make *salep*, a traditional Turkish creamy
51 drink or ice cream, consumed in Asia Minor and several countries on the Balkan peninsula [2–5]. In
52 south-eastern Africa terrestrial orchid tubers are mixed with peanut flour, salt, baking soda and
53 chili powder to make a traditional Zambian meat-like cake known as *chikanda* or African polony [6–
54 8]. Although initially not highly regarded [9], *chikanda* has more recently become popular
55 throughout the country. It is sold as a snack along the streets, on markets, in supermarkets and on
56 the menu of high-end restaurants [10] and recipes as well as cooking tutorial videos can be found
57 online [11]. Orchids used for *chikanda* are harvested exclusively from the wild, and although it is
58 unlikely that traditional village consumption poses a serious threat to orchid populations, the
59 increased popularity and subsequent commercialization of *chikanda* has led to the exhaustion of
60 Zambian orchid resources [8]. Collecting tubers means the end of a perennial and generally long-
61 lived orchid, since the entire plant is removed in the harvesting process.

62 Soweto market in the Zambian capital Lusaka is the hub of the *chikanda* trade. Surveys
63 performed on this market have shown that a large part of the *chikanda* tubers sold are sourced from
64 Tanzania and that Zambian *chikanda* orchids are collected from the Luwingu, Mporokoso and
65 Kasama districts in the Northern Province and Serenje in the Central province [7,8]. According to
66 local *chikanda* vendors, another region with a flourishing *chikanda* trade is the Kitwe region in the
67 Copperbelt Province, but so far no surveys have been performed there. Despite international
68 legislation (CITES) banning prohibiting cross-border trade, an estimated 2–4 million orchid tubers
69 are transported annually from Tanzania to Zambia [8,12]. Import from the surrounding countries of
70 Angola, DRC, Malawi and Mozambique is also documented [8,10]. Orchid species originally
71 reported as ingredients for *chikanda* are *Disa robusta* N.E.Br. and *Satyrium buchananii* Schltr. [12,13],
72 whereas recently at least 32 species belonging to the genera *Brachycorythis*, *Disa*, *Eulophia*, *Habenaria*,
73 *Roeperocharis* and *Satyrium* were suggested to be used for *chikanda* production based on collections
74 in the field [12,14–19] and one metabarcoding study of ready-made *chikanda* cakes [19]. To date,
75 however, no study identified the orchids traded at the local markets, since the tubers lack
76 sufficient morphological characters for taxonomic identification to species level [8,19]. Local
77 classification systems categorize the tubers based on texture, harvesting locality, soil colour and
78 phenology, but these are not likely to be congruent with scientific classifications [6,15].

79 Knowing which orchid species are currently being collected for the expanding *chikanda* trade
80 enables the identification of species susceptible to overharvesting and can inform conservation
81 planning. Molecular methods such as DNA barcoding can be applied to identify samples when
82 morphological diagnostic characters for identification are lacking [20]. DNA barcoding and
83 metabarcoding has proven to be effective in the authentication of commercial wood species (Jiao,
84 2018), medicinal plants [21] and *salep*-producing orchids on Iranian markets [5]. The analysis of
85 ingredients in Tanzanian *chikanda* cake with DNA metabarcoding revealed the presence of 21
86 different orchid species [19], but a DNA barcoding approach has not yet been applied to individual
87 orchid tubers used to make this product. The aim of this study was to test to what extent species
88 delimitation using standard molecular markers yields robust identification of *chikanda* orchid tubers
89 traded on Zambian markets. Molecular identification can enable mapping of harvesting and trade
90 of specific Zambian orchid species and facilitate identification and implementation of targeted
91 conservation strategies. Within that framework this study also aimed to identify conservation issues
92 associated with the *chikanda* trade, and addresses the following questions: 1) Which species are used
93 for *chikanda* production in the Lusaka and Kitwe districts of Zambia and what is their geographic
94 origin? 2) Can *chikanda* tubers be identified up to species level using DNA barcoding? 3) How do
95 local classification systems relate to scientific species concepts? 4) What are the main conservation
96 issues associated with *chikanda* trade in the Lusaka and Kitwe districts?

98 **2. Materials and Methods**99 **2.1. Interviews and sample collection**

100 Fieldwork in Zambia was conducted in 2016 in the Kitwe, Kalulushi, Luanshya, Ndola,
 101 Mufulira, Chingola and Chililabomwe districts of the Copperbelt Province; the Kapiri Mposhi and
 102 Serenje districts in the Central Province and in the capital Lusaka (Figure 1). Semi-structured
 103 interviews were conducted with harvesters, middlemen and vendors to obtain insight in *chikanda*
 104 commercialization, harvesting times, preferences and availability. The questionnaires consisted of
 105 three sections, one on informant and interview characteristics, one with general questions about
 106 *chikanda* posed to all informants, and a third section with questions more specifically designed for
 107 each interviewee category: harvester, middleman and vendor. All research was conducted in
 108 accordance with the International Society of Ethnobiology Code of Ethics [22]. Ethical clearance was
 109 obtained from the Humanities and Social Science Research Ethics Committee of the University of
 110 Zambia. The interviews were performed in English or Bemba, with a translator affiliated with the
 111 University of Zambia and the Copperbelt University. Informants were selected using the snowball
 112 technique [23], by asking people whether they could direct us to people harvesting or selling
 113 *chikanda*. All informants were provided with information about the study and signed a prior
 114 informed consent sheet. Fieldwork took place during June and July, the peak season for *chikanda* [8],
 115 to ensure collection of both fresh and dried *chikanda* tubers on the market and in the field. A
 116 collection was made each time a specific vernacular *chikanda* type was bought from a specific
 117 vendor, and assigned a collection number (SJK1, SJK2, etc.). Each individual tuber within the
 118 collection received a subsample number within that collection (SJK1.1, SJK1.2, SJK 1.16, etc.). The
 119 fresh tubers were sliced and stored with silica gel in plastic bags. All *chikanda* samples were brought
 120 to Sweden under a CITES inter-institutional exchange agreement between the University of Zambia
 121 (ZM001) and the Botany Section of the Evolutionary Museum in the Evolutionary Biology Center in
 122 Uppsala (SE009). Export permission was obtained from the Zambian CBD and Nagoya Protocol
 123 focal point at the Ministry of Natural Resources and Environmental Protection. Upon arrival to
 124 Sweden some of the *chikanda* tubers had sprouted. Those were transferred to the Uppsala Botanical
 125 Garden for cultivation and subsequent sampling of fresh leaf tissue for DNA barcoding as well as
 126 morphological identification.



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Figure 1. Overview of interview localities and reported provenance of the *chikanda* tubers. Dot size corresponds to the number of informants.

131 2.2. Reference taxon sampling

132 Herbarium specimens were collected with associated silica-dried material for DNA extraction
133 and spirit collections during fieldwork in Zambia in January and February 2017. In addition,
134 putative orchid mycorrhizal fungi were sampled from roots and tubers for isolation, culture and
135 identification at RBG Kew (results not reported here). All material was collected and exchanged in
136 accordance with national and international legislation. The collections are deposited at the Division
137 of Forest Research (Kitwe, Zambia) and RBG Kew (UK) and field identifications verified at the
138 Bews Herbarium (South Africa). A total of 94 novel Orchidaceae reference vouchers were collected
139 for this study, representing 4 *Brachycorythis*, 9 *Disa*, 16 *Habenaria*, 6 *Satyrium* species and 26 species
140 in other orchid genera. Voucher specimens of all taxa sampled are listed in Table S1
141 (Supplementary Material). In addition, 88 ITS, 71 *matK* and 45 *rbcL* *Habenaria* sequences generated
142 for a forthcoming phylogenetic study [24], and 510 ITS, 522 *matK* and 213 *rbcL* sequences
143 corresponding to 311, 325 and 100 taxa in the previously mentioned orchid genera downloaded
144 from GenBank were included in the reference database.

145 2.3. From sample to sequence

146 Out of the 1284 individual tubers in 48 different sample collections, 304 samples were selected
147 for DNA extraction. A few tubers per sample (2-8) were extracted if the sample was
148 morphologically homogenous, whereas more (6-33) were selected if the sample was diverse. DNA
149 was extracted using a CTAB protocol [25] modified with 3 to 5 extra washing steps with STE buffer
150 (0.25 M sucrose, 0.03 M Tris, 0.05 M EDTA) [26,27], to reduce the gelatinization effect of the large
151 amount of polysaccharides in the starch-rich orchid tubers. Total DNA was stored in 70-100 µl
152 10mM Tris-HCl buffer, pH 8.0. DNA concentration was measured with a Qubit 3.0 fluorometer
153 (Thermo Fisher Scientific, Oakwood, USA). The core land plant barcoding markers *rbcL* and *matK*
154 were amplified using the primers and protocols described in Kress et al. [28] and Dunning and
155 Savolainen [29] respectively. The reactions were performed in a total reaction volume of 25µl with
156 14.725µl ddH₂O, 2.5µl DreamTaq Buffer (Thermo Fisher Scientific, Oakwood, USA), 0.5µl 25mM
157 dNTP, 0.65µl 2% Bovine Serum Album (BSA), 0.125µl DreamTaq Polymerase, 2.5µl 5pmol forward
158 and reverse primer and 1.5µl template DNA. Nuclear ribosomal nrITS was amplified using the Sun
159 et al. [30] primers and protocol in a total reaction volume of 25µl containing 15.25µl ddH₂O, 2.5µl
160 DreamTaq Buffer (Thermo Fisher Scientific, Oakwood, USA), 0.5µl 25mM dNTP, 0.125µl 2% BSA,
161 0.125µl DreamTaq Polymerase (Thermo Fisher Scientific, Oakwood, USA), 2.5µl 5pmol forward and
162 reverse primer and 1.5µl template DNA. For ITS an additional protocol was used with Q5 high-
163 fidelity polymerase: reactions were performed in a total reaction volume of 23.5µl including
164 10.875µl ddH₂O, 5µl Q5 reaction buffer, 0.5µl 25mM dNTP, 5µl Q5 GC enhancer, 0.125µl Q5 high-
165 fidelity polymerase, 1.5µl 5pmol forward and reverse primer and 0.5µl template DNA. The PCR
166 program for the ITS primers in combination with the Q5 polymerase was an initial heating step of
167 30s at 98°C; 35 cycles of 10s 98°C, 30s 56°C, 30s 72°C; and a final elongation of 2min at 72°C. PCR
168 products were cleaned using eight times diluted ExoSAP (Thermo Fisher Scientific, Oakwood,
169 USA) and analysed on an ABI3730XL Sanger sequencer by Macrogen Europe (Amsterdam, The
170 Netherlands). The obtained trace files were assembled using Pregap4 and Gap4 [31] as
171 implemented in the Staden package [32]. Sequences shorter than 200 bp were discarded from the
172 analysis and all sequences have been deposited in NCBI GenBank (Table S2, Supplementary
173 Material). The NCBI BLAST algorithm was used to assess the identification of all the obtained
174 query sequences using the Python BLAST tabular parser script
175 (https://github.com/SLAment/Genomics/blob/master/BLAST/BLAST_tabularparser.py). Similarity
176 scores, query coverage, expect value (E-value), and max identity percentage were calculated, and all
177 the information of the top 5 hits were automatically mined and tabulated per marker (Table S3-S5,
178 Supplementary Material), and summarized per individual tuber sample (Table S6, Supplementary
179 Material).

180 2.4. Phylogenetic analysis and species delimitation

181 Alignments for nrITS, *matK* and *rbcL* were made using AliView [33], combing the query

182 sequences and local reference databases consisting of sequences from NCBI GenBank and reference
183 collections from fieldwork and unpublished data from collaborators [33]. Species delimitation was
184 performed using the Poisson Tree Processes (PTP) model, as it has been shown to outperform the
185 Generalized Mixed Yule Coalescent (GMYC) approach as well as OTU-picking methods when
186 evolutionary distances between species are short enough [34]. For all alignments a maximum
187 likelihood (ML) search for the best-scoring tree was performed using the RAxML web server [35] to
188 generate input trees for species delimitation analysis using bPTP [34]. GTRCATI was used to
189 implement the CAT approximation, and the final tree was evaluated using the traditional GTR
190 model. The bPTP.py script settings was 100,000 MCMC chain iterations for all trees; sampling
191 interval thinning value of 100; burn-in of 25%; and a random seed of 1234. No outgroup was
192 defined. The generated convergence curve was visually confirmed. The description of each voucher
193 is built up as following: 'sample #_BLAST search result_identification %_reported city/region of the
194 origin_the country,,'; for example: 'SJK04.09_S.buchananii_97.893_Mwinilunga_Zambia' (Figure S1-
195 3, Supplementary Material). For the reference sequences, accession number and species names are
196 described for *Disa* and *Satyrium*. The reference *Habenaria* species and others are described as species
197 name and voucher (sample) number.

198 3. Results

199 3.1. Market surveys and interviews

200 We visited 25 markets in ten Zambian cities: seven cities in the Copperbelt Province, two in the
201 Central Province, and one in Lusaka (Figure 1). Eighty-two persons involved in *chikanda* trade were
202 interviewed of which 8 were harvesters, 44 middlemen, 29 vendors and one informant was both
203 harvester and middleman. The term 'harvesters' refers to people collecting *chikanda* in the field,
204 'middlemen' to people selling either dried or fresh *chikanda* orchid tubers on the market, and
205 'vendors' to informal street vendors selling ready-made *chikanda* cakes contained in a basket carried
206 on the head. Since one of the harvesters also acted as middleman, in total 83 individual interviews
207 were conducted: 72 participants were female and 10 male. The ages of the respondents varied
208 between 18 and 78, with an average of 41 years. The majority of the respondents belonged to the
209 Bemba tribe (62%), while the other the respondents (38%) belonged to smaller ethnic groups. Most
210 interviews were conducted in the Copperbelt Province (55), eight in the Central Province and 19 in
211 Lusaka.

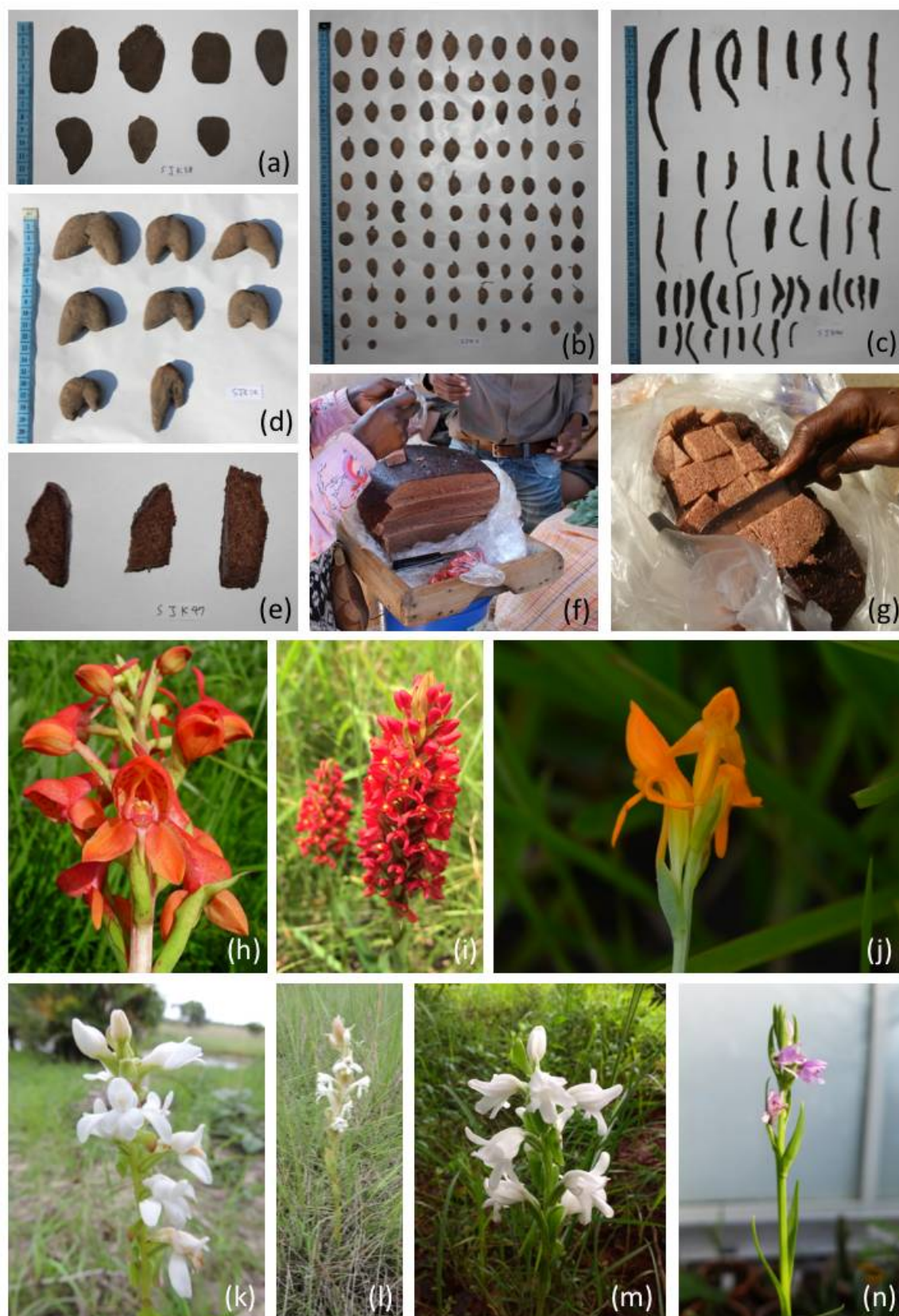
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213 3.2. Local classification system

214 Fifty-three different vernacular names for the various *chikanda* types were recorded during the
215 interviews. The most common way to distinguish between *chikanda* tubers was by using the terms
216 original (*myala*) and fake (*mbwelenge* or *msekelele*). In some cases it was the shape of the tuber that was
217 used to differentiate between the different tubers: *mshilamshila* means root-like in Bemba and referred
218 to the elongated, root-shaped tubers whereas *mampanda* referred to the heart-shaped tubers. It also
219 appeared common to use the origin of the tuber as a trade name: *mwinilunga*, *chozi*, *luwingu* and
220 *kasama* are for example all Zambian city names, *sumbawanga* and *iringe* are referring to Tanzanian
221 cities (Sumbawanga and Iringa) and *angola* refers to one of the countries bordering Zambia. In some
222 cases the *chikanda* tubers were sold pre-mixed, whereas other vendors marketed the different types
223 of tubers separately. The morphology and size of the tubers varied both within and between
224 collections of a certain *chikanda* type. Tubers could be heart-shaped, rounded, egg-shaped to
225 elongated and almost root-like. The largest tubers were the heart-shaped ones, which could be up to
226 5 cm long and 6 cm wide. The elongated tubers were up to 9 cm long and maximally 2 cm wide.
227 Harvesters, middlemen and vendors themselves indicated that they distinguish the tubers based on
228 the size of the granules inside the tubers, which can be large or small and in some tubers a concentric
229 ring was said to be present. Figure 2 illustrates the different tuber types and ready-made *chikanda* that
230 were encountered on the local markets, as well as some of the orchid species producing these tubers.

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Figure 2. Chikanda tubers, cake and orchids. (a) Myala – real chikanda; (b) Mbwelenge – fake chikanda; (c) Mshilamshila – supposedly *Brachycorytis* sp.; (d) Mampanda. (e-g) Chikanda cake; (h) *Disa robusta*; (i) *Disa welwitschii*; (j) *Platycoryne crocea*; (k) *Satyrium carsonii*; (l) *Satyrium buchananii*; (m) *Satyrium kitimboense*; (n) *Brachycorytis* cf. *friesii*; Photographs (a-g) by Seol-Jong Kim, (h) by Robert v. Blittersdorff, (i,k and l) by Nicholas Wightman, (j) by Warren McClelland, (m) by Ruth E. Bone and (n) by Sarina Veldman.

240 3.3. *Chikanda* trade and availability

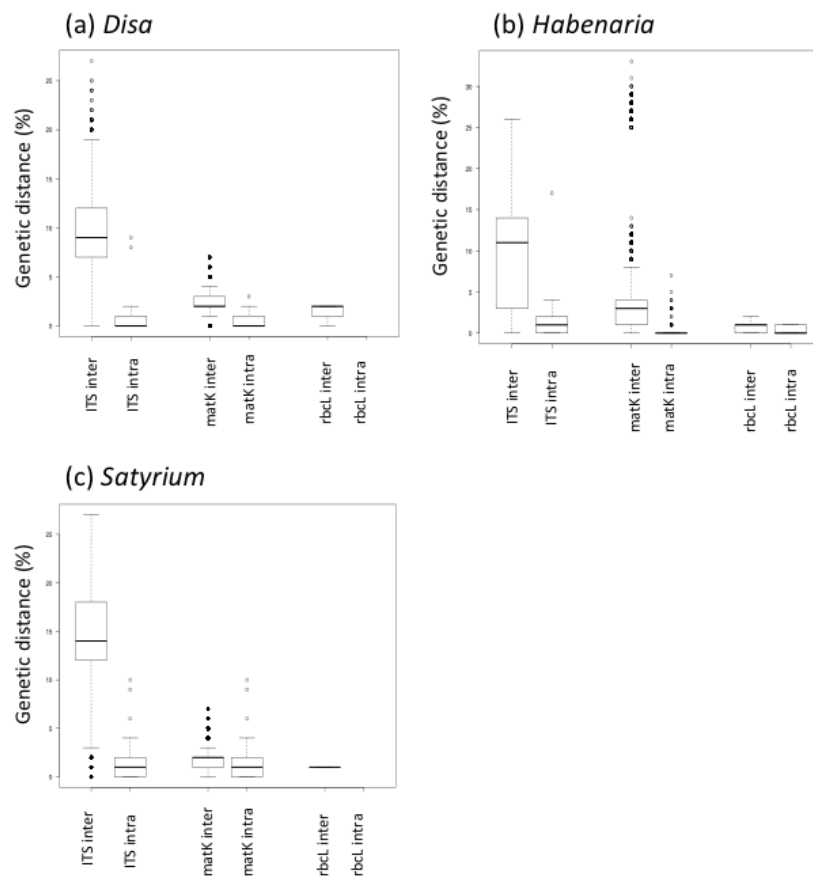
241 Many participants had a relatively long experience (average 13.5 year) in the *chikanda* trade. They
242 indicated that they were asked by family, friends or neighbors to get involved in the *chikanda*
243 business, or simply because it seemed a profitable industry. Some of the participants, especially
244 harvesters, stated that they started collecting *chikanda* because the plants were easily accessible and
245 were found growing close to their areas of residence. In general, people involved in *chikanda* trade
246 indicated that this work alone did not suffice to fully support themselves and their families, and these
247 traders therefore supplemented their income by trading additional natural products such as fruits,
248 maize, groundnuts, beans, mushrooms, snacks, herbs, and *kapenta* (a dried fish from Lake
249 Tanganyika). In Lusaka, however, income generated with *chikanda* trade seemed to be sufficient for
250 subsistence, and this suggests that there are significant local differences in profits generated with
251 *chikanda* business.

252 The *chikanda* trade was structured in several different ways. Some of the middlemen loaded the
253 orchid tubers in trucks directly from the harvesting areas and brought them to larger cities such as
254 Lusaka and Kitwe, whereas others relied on agents to gather a certain amount of *chikanda* tubers,
255 which they paid for through the East-African mobile payment system M-Pesa and received as cargo
256 from one of the local buses. In addition to the Tunduma and Nakonde markets that are the trade-
257 hubs between Tanzania and Zambia, other centers of trade were identified on the border with the
258 Democratic Republic of Congo (Chililabombwe and Kasumbalesa) and Angola (Mwinilunga), where
259 both *chikanda* tubers and ready-made *chikanda* cake was sold. An overview of all interview localities
260 and the reported provenance of the *chikanda* tubers is given in Figure 1. Most of the participants
261 indicated that *chikanda* plants are becoming locally rare. Middlemen emphasized the decrease of
262 quantity, whereas harvesters were concerned about the decline in both quantity and quality (size and
263 preferred *chikanda* type), which may also depend on the species. The participants from urban areas
264 stated that access to *chikanda* tubers was managed by the chief of each tribe, who seasonally
265 designated the harvestable *dambo* (wet meadow) within the chiefdom so that the collectors could
266 maintain the quality of the harvests. Nevertheless, interviewed harvesters encountered in the *dambo*
267 areas of Serenje (Central Province) claimed to be free to harvest tubers whenever available.

269 3.4. Molecular identification of traded orchids

270 During the DNA extraction many samples formed a thick jelly-like layer in the extraction tubes,
271 despite repeated washing steps with STE buffer. This resulted in a very small water phase and likely
272 negatively influenced the downstream steps of the extraction process. The average DNA
273 concentration of the *chikanda* tuber samples was 4.96 ng/ μ l, while the average DNA concentration
274 from leaf samples from *chikanda* orchids was 28.5 ng/ μ l. Out of the 304 samples selected for DNA
275 extraction, 232 samples produced detectable DNA. Amplification was attempted for each barcoding
276 marker for all of the samples. A nrITS sequence was obtained for 159 samples, *rbcL* for 117 samples
277 and *matK* only for 45 samples. Sequences from all three markers were obtained for 40 samples,
278 sequences from two markers for 58 samples, and 55 samples only yielded sequences for a single
279 marker. Analysis of the inter- and intraspecific variation of nrITS was performed for 141 sequences
280 representing 124 *Disa* species, 73 sequences representing 59 *Habenaria* species and 110 sequences
281 representing 67 *Satyrium* species. In the case of *matK* 135 sequences belonging to 122 *Disa* species were
282 included, 507 *Habenaria* sequences belonging to 239 species and 116 sequences belonging to 60
283 *Satyrium* species. For *rbcL* the available reference material was quite limited: 45 sequences for 37
284 *Habenaria* species, eight *Satyrium* sequences for two species and four *Disa* sequences for four species.
285 A graphical overview of the inter- and intraspecific variation per genus and per marker can be found
286 in Figure 3. The interspecific variation for nrITS was significantly higher as compared with *matK* and
287 *rbcL* in all three genera: on average 10.2% in *Disa*, 9.51% in *Satyrium* and 8.75% in *Habenaria*. For *matK*
288 the interspecific variation was 2.39% for *Disa*, 2.79% for *Habenaria* and 1.39% for *Satyrium*. The
289 intraspecific distances for nrITS and *matK* were respectively 1.37% and 0.61% for *Disa*, 1.98% and
290 0.36% for *Habenaria* and 1.45% and 0.48% for *Satyrium*. The limited reference sequences that were
291 available for *rbcL* showed little pairwise interspecific distances (1.44% for *Disa*, 0.82% for *Habenaria*
292 and 1.13% for *Satyrium*) and only allowed for the intraspecific distance calculation of *Habenaria*

293 (0.38%), indicating that *rbcL* is unsuitable as barcode for species level identification of *chikanda*
 294 orchids. The calculated thresholds were subsequently used to evaluate the identifications made with
 295 blastn.



296 **Figure 3.** Boxplots showing the inter- and intraspecific variation for *Disa* (a), *Habenaria* (b) and
 297 *Satyrium* (c) based on genetic diversity.
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 300 In total 15 orchid species were identified using DNA barcoding: *Brachycorythis* sp. SJK7, *Disa caffra*
 301 Bolus, *Disa celata* Summerh., *Disa robusta* N.E.Br., *Disa satyriopsis* Kraenzl., *Disa welwitschii* Rchb. f.,
 302 *Disa* sp. SJK4.1., *Habenaria* sp. SJK31.15, *Habenaria* aff. *helicoplectrum* Summerh., *Habenaria* cf. sp.
 303 DO112, *Platycoryne crocea* Rolfe, *Satyrium buchananii* Schltr., *Satyrium carsonii* Rolfe, *Satyrium*
 304 *kitimboense* Kraenzl. as well as one species in an unidentified genus, which seems to be closely related
 305 to *Habenaria* based on the similarity-based BLAST identification. Additionally, one orchid that
 306 flowered from a sprouting tuber was identified based on morphology as *Brachycorythis* cf. *friesii*
 307 (Schltr.) Summerh.

308 bPTP analysis for *matK* and *rbcL* showed lumping of several supposedly different species within
 309 one clade on several occasions (Figure S1 and S2, Supplementary Material). In case of nrITS the bPTP
 310 outcome tree often reflected the expected species boundaries, although the posterior probabilities on
 311 the nodes were often too low to determine with confidence if species delimitation had been
 312 performed correctly (Figure S3, Supplementary Material). Something that could be observed from
 313 both the nrITS BLAST identification, as well as the bPTP analysis, was that some of the identifications
 314 showed ambiguity: in collection SJK41, SJK44 and SJK46 some samples were identified
 315 unambiguously as *Platycoryne crocea*, whereas others showed a mix of *Habenaria* and *Platycoryne* top
 316 hits. The *matK* BLAST identification showed an unambiguous *Platycoryne crocea* identification for
 317 these samples, despite the presence of several *Habenaria* species in the reference database and despite
 318 the fact that *matK* shows a lower level of interspecific variation. A similar observation could be made
 319 for several *Disa* samples, but here a geographic pattern could be observed. The samples that could

320 unambiguously be identified as *Disa robusta* were all collected from Tanzania, where the reference
321 sequence originated from. Some of the Zambian samples also showed *Disa robusta* as the closest
322 relative, but not with a high enough percentage identity match to confirm this identification. The
323 bPTP result in this case shows a lumped and mixed clade with several *Disa* species and several other
324 samples included, which confirms that sequence divergence is too limited to resolve the relationship.
325 These samples were identified as *Disa* sp. 1, to reflect the fact that they all grouped together in the
326 same clade. In other cases, such as for certain *Habenaria* species, we named the closely related
327 sequences in the identification, whereas in this case the clade was too large to allow for this, since it
328 contained the following sequences: *Disa engleriana* Kraenzl., *D. erubescens* Rendle., *D. miniate*
329 Summerh., *D. ochrostachya* Rchb.f., *D. satyriopsis*, *D. ukingensis* Schltr., *D. verdickii* De Wild., *D.*
330 *welwitschii*, *D. zombiaca* N.E.Br., an unidentified *Disa* species and several samples that showed a
331 highest percentage identity match with *D. robusta*, *D. satyriopsis* and *D. welwitschii*. The posterior
332 probability for this clade is only 0.25, and combined with the posterior probabilities present on the
333 within-species nodes, indicates a lot of uncertainty for this identification.

334 Overall, the most frequently encountered species were *Satyrium buchananii*, with 41 samples in
335 eleven different collections, *Platycoryne crocea* with 19 samples in three different collections and *Disa*
336 *robusta* with 4-16 samples in two to seven different collections. *Myala* or original *chikanda* seems to
337 contain *Disa robusta*, *D. welwitschii* and *Satyrium buchananii*. *Mbwelenge* or fake *chikanda* seems to
338 correspond only to *Satyrium buchananii* and *mshilamshila* to one or several *Brachycorythis* species.
339 *Kasebelele* and *kapapa* referred to *Habenaria* spp., *Platycoryne crocea*, *Satyrium carsonii* and *Satyrium*
340 *kitimboense*. The mixed collections contained all of the above-mentioned species as well as an
341 unidentified *Habenaria* species. An overview of the local *chikanda* classification types, their collection
342 numbers, identifications and number of samples can be found in Table 1.

343 4. Discussion

344 4.1. Species used for *chikanda*

345 Using DNA barcoding as an identification tool for *chikanda* tubers sold on local Zambian markets
346 has allowed us to determine for the first time what orchid species are sold on local markets. Previous
347 studies identifying orchids used for *chikanda* relied on voucher collections made with collectors in the
348 field and their morphological identification, which requires a qualified orchid taxonomist
349 [7,12,14,17,36]. Additionally, relying on local harvesters for details on *chikanda* collection might not
350 always lead to collection in the areas where actual intensive harvest is taking place, since some
351 harvesters do not like to divulge where the best places to harvest are and in some initial harvesting
352 areas such as the Kitulo Plateau in Tanzania's Southern Highlands, collection is now prohibited [37].
353 The current study identified 16 orchid species present on the markets, including at least three
354 previously undocumented ones: *Brachycorythis* cf. *friesii*, *Platycoryne crocea* and an unidentified species
355 in a genus, which appears to be closely related to *Habenaria*, but is not present in our reference
356 database.

357 Orchids used for *chikanda* seem to be harvested from several provinces in Zambia, as well as at
358 least two regions in Tanzania. Moreover, three international *chikanda* trade-hubs in towns on or close
359 to the border with the DRC (Chililabombwe and Kasumbalesa) and Angola (Mwinilunga) were
360 identified, in addition to the already known trade-hub Tunduma-Nakonde on the Tanzanian border
361 [8,12]. However, unlike reported in other studies, no harvest from Malawi was mentioned by the
362 people interviewed in this study, which could mean that trade from this country is currently not
363 taking place. Another explanation is that this information is lost on the way and that only Tanzania,
364 being geographically closer and thus more easily accessible for the Bemba people, is mentioned as a
365 region of origin for *chikanda* traded in Zambia.

366 4.2 DNA barcoding performance

367 Since the term DNA barcoding was coined in 2003 [20], a plethora of studies applying DNA
368 (meta)barcoding has been performed ranging from retrieving orchids from paleoenvironments [38],
369 preserved in mammoth dung [39] to the identification of Iranian orchid tubers used for *salep* [5]. In

371 this study a combined use of the core plant markers *matK* and *rbcL* and the nuclear ribosomal ITS
372 region was used to attempt species level identification of tubers traded on Zambian markets.
373 Although genetic distance calculations showed limited interspecific distances between closely related
374 species for all three barcoding markers, DNA barcoding allowed for species-level identification for
375 several of the frequently sold *chikanda* species. The data shows that the core land-plant DNA
376 barcoding markers *rbcL* and *matK* were not suitable because of limited variability between species
377 (*matK* and *rbcL*), amplification problems (*matK*), and/or a limited sequence reference database (*rbcL*).
378 nrITS was shown to be more suitable as a barcode marker to distinguish between different *chikanda*
379 species, but is not discriminative enough to enable reliable species level identification in certain
380 orchid clades, such as the clade with *Platycoryne crocea*, *P. buchananiana* and *Habenaria buchananii*; the
381 clade with *H. schimperiana*, *H. kyimbilae* and *H. microsaccos* and the clade containing *Disa* sp. 1. Another
382 drawback of nrITS are the multiple ITS paralogs present in the ribosomal genome. Usually these
383 copies would show high similarities due to concerted evolution [40,41], but this is not always the case
384 [42–45]. In our case potentially different nrITS ribotypes became fixed in different orchid populations
385 and having only one of them in our reference database could lead to the unresolved identifications
386 observed. Our bPTP results show that even nrITS has too little resolution to reliably delimit species
387 with high posterior probability support using this method. It does demonstrate, however, how
388 valuable the use of tree-based methods can be, since it shows the relations between the sequences
389 and can be used to determine if some of the unidentified samples are likely to belong to the same
390 species or different species within the same genus. Even if no species-level identification can be made
391 for these samples, it is possible to use the clustering to determine the diversity of species used.
392 Although several samples can only be reliably identified as *Habenaria* sp., we find that they are likely
393 to belong to at least three different species (*Habenaria* aff. *helicoplectrum*, *Habenaria* cf. sp. DO122 in the
394 clade with *H. schimperiana*, *H. kyimbilae* and *H. microsaccos* and lastly the *Platycoryne* sp./*Habenaria* sp.,
395 which group together with *H. buchananii*, *P. buchananiana* and *P. crocea*. Expansion of the reference
396 database, by including at least one individual per species, and preferably multiple individuals per
397 species from different populations and countries, could ultimately solve remaining challenges, and
398 this seems the way forward in identification of the traded *chikanda* tubers, as well as other species
399 unidentifiable based on morphology. Similar studies using DNA barcoding for the identification of
400 unknown samples show comparable results for the employed barcoding markers. In a study on the
401 identification of orchids used for *salep*, nrITS showed an over three-time higher sequencing success
402 than *matK* as well as a two-times higher species-level identification success [5]. Moreover, the
403 similarity-based approach seemed to outperform the tree-based identification method (ML) in this
404 study as well with 57% and 39% species-level identifications, respectively. nrITS also shows the
405 highest identification performance in studies on the identification of medicinal plants [46,47], which
406 supports the idea that it is recommendable to add a more discriminative marker to the two core land-
407 plant barcodes in studies where it is needed to distinguish between closely-related species. Moreover,
408 our results stress the need for a phylogenetically underpinned taxonomic framework, which is
409 currently available for *Disa* [48] and *Satyrium* [49], but not yet for *Habenaria* and related genera.

410

411 4.3. Local versus scientific classification of *chikanda*

412 The results of our identifications of *chikanda* orchids traded in the Zambia show that the local
413 classification systems for *chikanda* are not in congruence with the botanical classification of orchid
414 species. The orchids sold on the markets were grouped according to area of origin, tuber consistency
415 preference, or shape of the tuber, but often the tubers offered for sale were mixtures. The different
416 local types of *chikanda* sometimes show a variation in orchid species that are identified within these
417 local grouping. When we look at the *chikanda* type known as *kapapa* for example, SJK44 contains
418 *Platycoryne crocea*, whereas SJK11, which is supposed to be a mix of *kasebelela* and *kapapa*, only seems
419 to contain *Satyrium* species. In case of other *chikanda* types there seems to be more consistency: *myala*
420 or real *chikanda* referred to *Disa robusta*, *D. welwitschii* and *Satyrium buchananii*, *mshilamshila* samples

Table 1. Overview of the different local *chikanda* classification types, their collections and the identified scientific species.

Vernacular name	Collections	Reported origin	barcoding IDs	# samples
Fungulwe	SJK16	unknown	<i>Disa robusta</i>	1
Iringe	SJK17	Tanzania	<i>Satyrium buchananii</i>	1
			<i>Satyrium carsonii</i>	1
John White	SJK39	Mporokoso, Zambia	<i>Satyrium buchananii</i>	1
Kabula seke	SJK46	Serenje, Zambia	<i>Habenaria</i> sp. (Clade <i>H. schimperiana</i> , <i>H. kyimbilae</i> , <i>H. microsaccos</i>)	1
			<i>Platycoryne crocea</i> Rolfe	7
Kapapa	SJK44	Mporokoso, Zambia	<i>Platycoryne crocea</i> Rolfe	6
Kasebelela, John White and Myala	SJK41	Chinsali and Mporokoso, Zambia and Tanzania	<i>Habenaria</i> cf sp. DO122 (Clade <i>H. schimperiana</i> , <i>H. kyimbilae</i> , <i>H. microsaccos</i>)	4
			<i>Platycoryne crocea</i> Rolfe	4
			<i>Platycoryne</i> sp./ <i>Habenaria</i> sp.	2
Kasebulela and Kapapa	SJK11	Luwingu, Zambia	<i>Satyrium kitimboense</i>	6
			<i>Satyrium carsonii</i>	5
Mbwelenge	SJK5	Luwingu, Zambia	<i>Satyrium buchananii</i>	11
			<i>Satyrium</i> sp.	1
	SJK32	Serenje, Zambia	<i>Satyrium buchananii</i>	6
Mshilamshila	SJK7	Luwingu, Zambia	<i>Brachycorythis</i> sp.	1
			<i>Brachycorythis</i> sp.	1
			<i>Brachycorythis</i> cf. <i>friesii</i>	1
Myala	SJK4	Mwinilunga, Zambia;	<i>Disa robusta</i>	4
			<i>Disa welwitschii</i>	1
	SJK18	Sumbawanga, Tanzania	<i>Satyrium buchananii</i>	4
			<i>Disa robusta</i>	4
			<i>Satyrium buchananii</i>	1

Myala	SJK37	Kawambwa, Zambia	<i>Satyrium buchananii</i>	1
Myala and nampana	SJK21	Luapula, Zambia	<i>Disa welwitschii</i>	2
			<i>Satyrium buchananii</i>	1
Ntonkonshi	SJK25	Democratic Republic of Congo	<i>Disa robusta</i>	1
Sumbawanga	SJK20	Sumbawanga, Tanzania	<i>Disa satyriopsis</i>	1
mixed	SJK31	Serenje, Zambia	<i>Disa caffra</i>	1
			<i>Disa robusta</i>	1
			<i>Habenaria</i> cf sp. DO122 (Clade <i>H. schimperiana</i> , <i>H. kyimbilae</i> , <i>H. microsaccos</i>)	1
			<i>Satyrium buchananii</i>	6
unknown-mixed	SJK19	Luwingu, Zambia	<i>Satyrium carsonii</i>	1
			<i>Habenaria</i> aff. <i>helicoplectrum</i> (BB3151)	1
unknown	SJK8	Mwinilunga, Zambia;	<i>Disa miniata</i>	1
			<i>Disa robusta</i>	2
			<i>Disa welwitschii</i>	1
			<i>Satyrium buchananii</i>	2
	SJK9	Luwingu, Zambia	<i>Satyrium carsonii</i>	1
	SJK13	Kawamba, Zambia	<i>Disa celata</i>	1
<i>Disa welwitschii</i>			1	
			<i>Satyrium buchananii</i>	3

422 were identified as *Brachycorythis* species and *mbwelenge* or fake *chikanda* was made of *Satyrium*
423 *buchananii*. However, our previous study on the analysis of Tanzanian *chikanda* cakes showed that the
424 cake made with fake *chikanda* tubers also contained *Disa miniata*, *Satyrium anomalum*, *S. comptum*, *S.*
425 *elongatum*, *S. riparium*, *S. shirense* and *S. volkensii*, indicating that there might be some differences
426 between fake *chikanda* samples as well [19]. It is well-known from literature that local species concepts
427 are not necessarily congruent with scientific classifications and that species might be subject to over-
428 or underdifferentiation [50,51]. In this case, the grouping of the orchids according to the area of origin,
429 shape or consistency preference or plainly under the general term *chikanda* is clearly a case of
430 underdifferentiation as a much higher diversity was retrieved when employing DNA barcoding. In
431 order to more reliably identify the orchid species used for a particular *chikanda* type more samples
432 per local classification need to be analyzed.

433

434 4.4. Orchid availability and conservation

435 Throughout recent decades *chikanda* has made a remarkable leap in popularity. The first record of
436 *chikanda* use made by Audrey Richards [9], described the relish as a poor man's food, eaten in times
437 of famine. Recent studies, from 2002 onwards, show that *chikanda* has emerged as a Zambian snack
438 popular throughout the country. Studies on *chikanda* report that with this rise in popularity, the
439 orchid harvest has escalated and is pressuring local Zambian orchid populations as well as those in
440 neighboring countries [52,12,10,53]. Many people involved in *chikanda* trade indicated that *chikanda*
441 plants were becoming scarce, and many were concerned about both the quality as well as the quantity
442 of the orchids available. Our study also confirmed a significant international trade network for
443 *chikanda* sources in several regions in Tanzania, as well as in the Democratic Republic of Congo and
444 Angola. In our current study, we found at least 16 different orchid species sold as *chikanda* on the
445 Zambian markets and an overview of previous studies contains 46 species reported to be used for
446 *chikanda* (Table S7). This brings us from the use of an initial two orchid species reported for *chikanda*
447 [13] to a total of 49 species in eight different genera. The increased harvesting pressure in combination
448 with the indiscriminate harvesting and use of many more species than earlier assumed pose a threat
449 to nearly half of the terrestrial orchids occurring in these regions. Despite the establishment of Kitulo
450 National Park in Tanzania, with orchid conservation as a prime concern, it seems that harvesting
451 continues even there, since *iringe* tubers found in this study come specifically from this region [12,53].
452 Currently there are only seven *Disa* species from Zambia and surrounding countries registered on
453 the global IUCN Red List and no species from other genera used for *chikanda* [54]. Most of the orchid
454 species used for *chikanda* seem to have a widespread distribution, but local populations as well as
455 endemic species could be at risk of overharvesting, and we urge to add the most frequently traded
456 *chikanda* species, such as *Disa robusta* and *Satyrium buchananii* to the IUCN Red List. Although there
457 seems no stopping to commerce, people involved in *chikanda* trade seem genuinely concerned about
458 welfare of local orchids and interested in exploring other options. Since especially the *chikanda*
459 harvesters seem to be in a vulnerable position, where they have to rely on surrounding natural
460 resources to secure their livelihoods [16], it is essential that when trying to protect orchids used for
461 *chikanda*, the situation of the people dependent on the trade is taken into account as well. Currently
462 the development of sustainable cultivation or *chikanda* orchids is attempted in collaboration with the
463 Cape Institute of Micropropagation (Barrydale, South Africa) and possible alternative sources of
464 income for the people involved in *chikanda* trade, such as honey production, are being explored [55].
465 Alternatively, since the purpose of the *chikanda* orchids mainly is to bind and create an elastic
466 structure to the cake, it might be possible to encourage the use of an alternative source of starch to
467 replace the tubers.

468

469 5. Conclusions

470

471 DNA barcoding using the nuclear ribosomal ITS marker proved to be useful in identifying
472 terrestrial orchid species traded as *chikanda* on local Zambian markets and outperformed
473 identification using the core land-plant barcoding markers *matK* and *rbcl*. Sixteen orchid species, of
474 which three previously undocumented, were identified from marketed *chikanda* tubers, bringing the

475 total number of orchid species used for *chikanda* to at least 49. The species most frequently found on
476 the markets were *Disa robusta*, *Satyrium buchananii* and *Platycoryne crocea*. However, the results are
477 only as good as the reference material is and an increased reference database in combination with an
478 underpinned phylogenetic framework for *Habenaria* and related genera would likely ameliorate the
479 reliability of the identifications. Tubers are harvested from various regions in Zambia and Tanzania,
480 and additional international *chikanda* trade-hubs have been identified on the border with the
481 Democratic Republic of Congo and Angola. People involved in *chikanda* trade indicate that both
482 orchid quality as well as quantity are decreasing and are willing to consider alternatives to *chikanda*
483 trade to secure their income.

484 Authors should discuss the results and how they can be interpreted in perspective of previous
485 studies and of the working hypotheses. The findings and their implications should be discussed in
486 the broadest context possible. Future research directions may also be highlighted.

487 **Supplementary Materials:** The following supplementary figures and tables are available online at
488 www.mdpi.com/xxx/s1

489 **Figure S1.** bPTP analysis of all *chikanda* matK query and reference samples.

490 **Figure S2.** bPTP analysis of all *chikanda* rbcL query and reference samples.

491 **Figure S3.** bPTP analysis of all *chikanda* nrITS query and reference samples.

492 **Table S1.** Brahm's RDE file of novel voucher specimens of orchid taxa samples in this study.

493 **Table S2.** Genbank accession numbers of the *chikanda* tuber sequences.

494 **Table S3.** Hit table with the first 5 BLAST top hits per sample for nrITS.

495 **Table S4.** Hit table with the first 5 BLAST top hits per sample for *matK*.

496 **Table S5.** Hit table with the first 5 BLAST top hits per sample for *rbcL*.

497 **Table S6.** Successfully sequenced samples with their vernacular name, reported origin, identification
498 based on sequence-similarity for nrITS and *matK*, identification based on the tree-based bPTP analysis
499 and the consensus ID.

500 **Table S7.** Orchid species used for *chikanda* according to literature.

501 **Author Contributions:** S.V., R.B. and H.d.B. conceived and designed the study in collaboration with
502 S.J.K., D.C., R.V. and N.W.; S.J.K. performed the *chikanda* collection and interviews in collaboration
503 with D.C., G.M., N.W. and R.V.; R.B., N.W. and K.Y. collected reference vouchers in the field, which
504 were identified in collaboration with B.B.; B.B., G.N. and F.M. provided the reference sequence
505 database for *Habenaria* species and some species in closely related genera; S.J.K. performed the
506 labwork for the *chikanda* collections and M.B.F. for the reference collections; S.J.K. performed the
507 data analysis under supervision of S.V. and H.d.B.; S.V. and S.J.K. wrote the manuscript in
508 consultation with the other authors and the manuscript was edited and reviewed in detail by B.G.,
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518

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