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5 **The destructive sampling conundrum and**  
6 **guidelines for effective and ethical sampling of herbaria**  
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32

33 **Abstract**

34

35 **The use of herbaria for science and conservation is revolutionizing the discovery,**  
36 **exploration, and protection of biodiversity at unprecedented scopes and scales. The**  
37 **Global Metaherbarium—a digitally interlinked, open-access resource—is**  
38 **stimulating these efforts and helping to facilitate massive investigations that utilize**  
39 **aggregated digital derivatives of physical herbarium specimens. Simultaneously, the**  
40 **growing use of this virtual resource is expanding the use of physical collections by**  
41 **researchers from many scholarly domains who increasingly are sampling specimens**  
42 **for multiomic investigations (e.g., genomics, transcriptomics, metabolomics,**  
43 **proteomics, and microbiomics). These investigations are leading to new scientific**  
44 **insights and supporting the development of conservation actions, but they come with**  
45 **a substantial cost: the (partial) destruction of priceless and often irreplaceable**  
46 **specimens, which constitute a global heritage that should be permanently**  
47 **safeguarded for future reference. The absence of a comprehensive set of “best**  
48 **practices” for destructively sampling herbarium specimens leads to confusion and**  
49 **uncertainty from researchers and institutions alike and risks over-exploitation of**  
50 **precious collections when the research is executed. Here, we provide a set of best**  
51 **practices aimed at reducing these uncertainties and creating a framework for**  
52 **sustainably and ethically sampling herbarium specimens. Our recommendations are**  
53 **intended for two complementary but overlapping audiences—users and stewards—**  
54 **who together build, use, and protect herbarium collections.**

55

56

57 **Keywords:** biodiversity, herbaria, genomics, metabolomics, multiomics, museum ethics,  
58 natural history collections

59

60 The world's herbaria house approximately 400 million specimens<sup>1,2</sup>, including extinct  
61 species or populations, and represent the diversity required to understand the origin, past  
62 and future evolution, and ecology of plants and fungi. Herbaria are one of the most  
63 important places in which to study nature in the face of continued threats to biodiversity  
64 and its importance to humankind for food, medicine, spiritual comfort, and shelter<sup>3</sup>. They  
65 also are being used increasingly by educators and artists to draw attention to broader  
66 societal issues<sup>3</sup>, and to develop plans for conservation, restoration, and sustainable use of  
67 natural resources.

68

69 Traditional uses of herbaria mainly have included species descriptions and associated  
70 floristic investigations and monographic treatments; determining species' geographical  
71 ranges; and historical perspectives on collectors and collections<sup>3,4</sup>. Since the 1920s,  
72 additional uses of collections have expanded tremendously, continue to grow and, in  
73 combination with 21<sup>st</sup>-century digitization and next-generation integrative multiomic  
74 approaches, are ushering in a new era of herbarium and biodiversity science<sup>3,5</sup>. In short,  
75 the 'Herbarium of the Future' is propelling basic and applied science in ways that were  
76 never imagined when these institutions and their collections were established centuries  
77 ago<sup>3</sup>.

78

79 We applaud the ongoing renaissance of herbaria, but over many years in our professional  
80 roles we have become increasingly aware of what we could call the "destructive sampling  
81 conundrum:" how to foster innovative research while permitting destructive sampling of  
82 specimens meant to be protected permanently. However, best practices for destructively  
83 sampling herbarium specimens have not yet been explicitly developed or agreed. The lack  
84 of common guidelines and best practices results in inconsistent practices by stewards of  
85 these collections, especially when newcomers to the field want to destructively sample  
86 specimens but are uncertain how to do so effectively and responsibly. Conversely, the pool  
87 of herbarium users has expanded to include students, researchers, and other professionals  
88 from a wide range of disciplines, many of whom are not educated in collection stewardship  
89 and do not understand the range of costs of their requests. In this **Perspective**, we propose  
90 clear guidelines for the effective and ethical sampling of herbaria. A central premise of

91 these guidelines is that herbarium specimens are a limited and priceless resource that must  
92 be protected for current and future generations.

93

94 These guidelines imply best practices, and we outline below best practices for two groups  
95 of stakeholders: i) herbarium *users*, and ii) herbarium *stewards*. Herbarium stewards  
96 include a broad swath of backgrounds and expertise, including digitizers, curatorial  
97 assistants, collections managers, keepers, curators, and directors. We recognize that  
98 individuals may be both users and stewards, and although some of our specific guidelines  
99 may appear to treat them separately, recommendations for users and stewards intersect  
100 and inform one another. Whereas institutional priorities may vary, we hope that our  
101 recommended best practices will stimulate discussions that lead to establishing clear  
102 guidelines for destructively sampling herbaria. Finally, although our focus is on herbaria,  
103 many of our guidelines and best practices apply equally to any natural history collection.

104

### 105 **Background and rationale**

106

107 The online mobilization of digitized herbaria is sparking a renaissance in the use of natural  
108 history collections<sup>6</sup>. Inherent to digitizing and mobilizing herbarium collections is the  
109 creation of the ‘Global Metaherbarium,’ a ‘common, digitally interlinked and open-access  
110 resource that will stimulate large-scale and novel science’<sup>3</sup>. The Global Metaherbarium is  
111 more than the sum of its parts: it is an ideal platform for connecting essential biodiversity  
112 data (often physically scattered) in a virtual, searchable framework, and represents far  
113 more information content than is available from physical specimens alone. ‘Extended’  
114 specimen data may include not only digital derivatives of the physical specimen (i.e., a  
115 digital photograph and label transcription) but also its genomic sequence, field images, and  
116 inferred species distribution models (SDMs), literature citations, among other data types  
117 that can be linked to it<sup>7,8</sup>. Although the Global Metaherbarium is virtual, it makes specimens

118 and their extended data  
 119 interconnected and  
 120 discoverable, which in turn  
 121 stimulates increased visitation  
 122 of herbaria by diverse scholars  
 123 who study and sample  
 124 physical collections<sup>6,9</sup> (Fig. 1).

125  
 126 Alongside these exciting  
 127 advancements, we have  
 128 witnessed a dramatic increase  
 129 in the destructive sampling of

130 herbarium specimens for  
 131 scientific and scholarly pursuit  
 132 (Fig. 2). Research topics that  
 133 necessitate destructive  
 134 sampling include: expanded

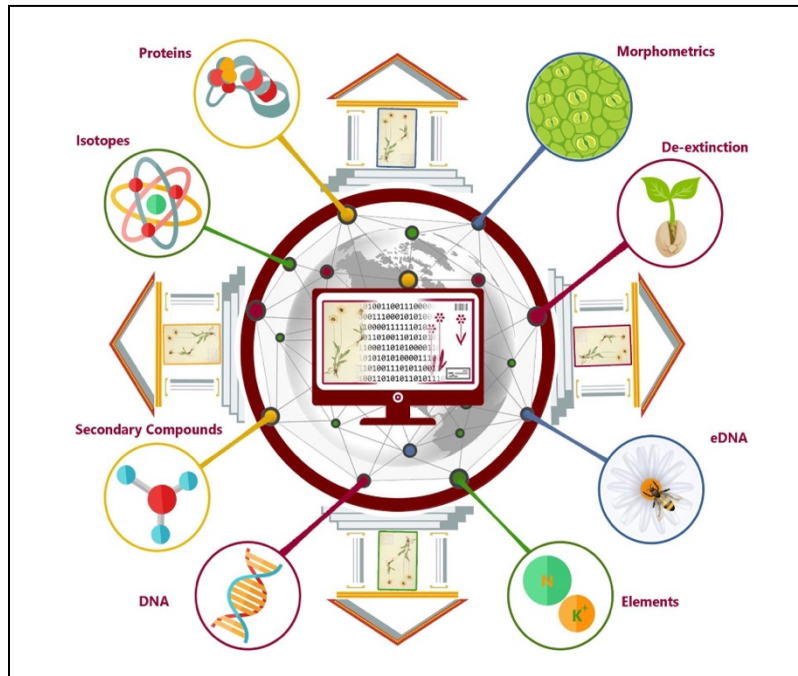
135 DNA barcoding efforts; high  
 136 resolution phylogenomics;  
 137 tracking the genetic history of  
 138 domestication; illuminating  
 139 human migration;

140 characterizing anthropogenic  
 141 change during the Quaternary;  
 142 mapping the origin and spread  
 143 of crop pathogens; exploring  
 144 the genetic basis of species

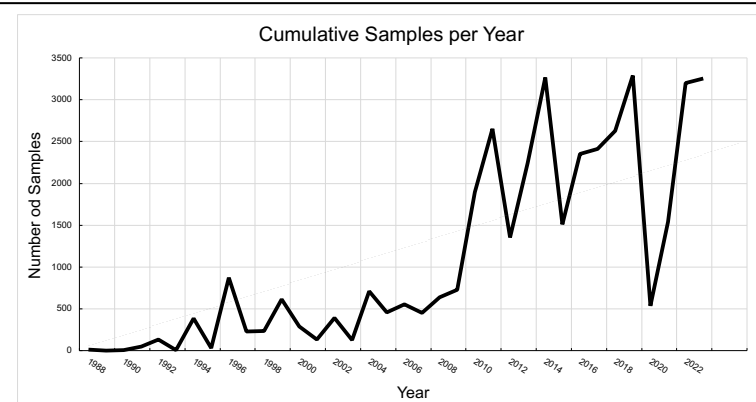
145 invasions and extinctions;

146 investigating microbial associations across time and space; resolving species boundaries;  
 147 and incorporating historical variation to investigate biodiversity change through time<sup>3,5,10-</sup>

148 <sup>16</sup>. Relatedly, we have observed growth in destructive sampling of herbaria for isotopic



**Figure 1.** The Global Metaherbarium seamlessly accommodates the extended specimen and greatly facilitates searchability. Using this virtual framework, key extensions of the physical specimen, such as multiomic data, can be recorded easily and integrated for findability at a global scale, facilitating more effective destructive sampling. For example, if a duplicate at one herbarium were sampled for genomic sequencing, similarly sampling a duplicate from the same collection at another herbarium should be avoided or receive lower priority. Open access icons by macrovector/FreePik.



**Figure 2.** Annual cumulative growth in the number of destructively sampled specimens for DNA sequencing, from 1988–2023. Data tabulated from records from the Harvard University Herbaria (HUH), the Missouri Botanical Garden herbarium (MO), the New York Botanical Garden herbarium (NY), and the Royal Botanic Gardens herbarium (K).

149 assessments to characterize temporal changes in atmospheric CO<sub>2</sub> and other pollutants<sup>17</sup>,  
150 and to extract and characterize small metabolites of potential medicinal value<sup>18-20</sup>.

151  
152 Destructive sampling of herbarium collections is not new. It has occurred continually since  
153 the earliest herbaria were established in the 1500s and 1600s (e.g., boiling flowers to  
154 measure the living size of their floral organs; harvesting anthers and pollen grains for  
155 detailed microscopic investigation). Since the late 1980s, the discoverability of specimens  
156 facilitated by digitization combined with cutting-edge multiomic approaches have  
157 bolstered macroevolutionary investigations<sup>21,22</sup> and fueled explosive growth of molecular  
158 systematics<sup>23</sup>. However, destructively sampling specimens for multiomic investigations and  
159 other research has a serious cost: the (partial) destruction of the specimen. This cost will  
160 continue to grow as the Global Metaherbarium becomes fully realized, new questions arise,  
161 and novel tools are developed. The convenience and relative ease of sampling herbarium  
162 specimens, as opposed to generating new collections from fieldwork, also may have  
163 undesirable and unintended consequences. We must work collectively to ensure that  
164 collections are preserved and research using their finite resources can be pursued  
165 vigorously and fairly. Our proposed set of best practices are designed to facilitate and  
166 encourage research while minimizing the destructive cost to priceless collections.

167

### 168 **Best practices for herbarium users**

169

170 *1. Always consult other available resources before destructively sampling herbarium*  
171 *specimens.* There are a variety of tissue resources distinct from conventional pressed and  
172 dried herbarium specimens that can be destructively sampled and are especially useful for  
173 multiomic investigations. Such resources include DNA cryofacilities held at institutions  
174 (such as at K and NY; herbarium acronyms follow Thiers<sup>1</sup>) and extensive silica-dried leaf  
175 collections (such as at K and MO). These other resources should always be consulted *before*  
176 resorting to destructive sampling of conventional herbarium specimens. Silica-dried  
177 samples, albeit finite, are an excellent resource for initial exploration because they tend to  
178 produce higher quality DNA with greater yields than conventionally dried herbarium  
179 specimens<sup>24,25</sup>. This option also can be used to establish positive lab controls for

180 downstream investigations with more degraded tissues. Another viable option is to contact  
181 and ask researchers who have access to living plants or have destructively sampled  
182 herbaria previously for multiomic investigations to share tissue or aliquots. Finally, the  
183 [Plant Search](#) database allows searching for taxa held within a network of > 1000 living  
184 collections. Any use of shared resources normally should lead to a formal collaboration  
185 (including co-authorship).

186

187 *2. Herbaria are not a substitute for fieldwork.* Herbaria increasingly are viewed as the first  
188 target of genetic sampling<sup>3,26</sup>. However, continuing to build these collections increases their  
189 value and sustains their utility for future research<sup>27</sup>. Thus, whenever possible, herbaria  
190 should not be the first resort for sampling. Rather, new specimens collected during  
191 fieldwork—including generating new herbarium collections as vouchers for future  
192 generations—should be sampled first; herbarium specimens should be used only as needed  
193 to fill gaps or if sampling from herbaria is the only practical means to conduct a study (e.g.,  
194 in situations of wars, political strife, funding restrictions).

195

196 This ‘field-first’ approach is how herbaria have been used historically for multiomic  
197 investigations, in part because of the inability to obtain reliable ‘omic data from so-called  
198 ‘antique samples.’ Although next-generation approaches are removing some obstacles and  
199 further unlocking these collections, these advances should not lead to less field sampling or  
200 collecting whenever and wherever possible. Fieldwork inspires curiosity and question-  
201 oriented science and provides researchers and students with a first-hand understanding of  
202 the threats and ecological dependencies that species face. Targeted field work, even done  
203 on a small scale, also can help build collections and generate goodwill when making later  
204 requests for destructive sampling. In-country institutions should be the primary partners  
205 in fieldwork and joint participation from study planning through collection and publication  
206 are important aspects of Access and Benefit Sharing under the [Convention on Biological  
207 Diversity](#).

208

209 *3. Ensure that institutional permission to destructively sample specimens is granted.* Herbaria  
210 typically require explicit permission to destructively sample their collections; users must

211 obtain appropriate permission from institutions and those who steward these collections.  
212 If users are uncertain how to do this, simply ask. Institutional guidelines vary among  
213 herbaria and, in some cases, among collections within an institution, and users should not  
214 assume that sampling protocols and permits are the same in every herbarium.  
215 Furthermore, permission to destructively sample usually are granted only on a specimen-  
216 by-specimen basis; permission to sample one specimen normally does not imply or grant  
217 permission to sample other materials. Destructive sampling of type specimens is strictly  
218 forbidden by some institutions, but others may consider such requests while subjecting  
219 them to additional evaluation and scrutiny, which may involve input from multiple  
220 stakeholders (e.g., curatorial boards at local institutions or colleagues and stakeholders  
221 spanning multiple institutions). Such scrutiny and evaluation can take months, so requests  
222 for permits should be made well in advance. In some cases, institutions also may have  
223 formal understandings with governmental agencies or other stakeholders that go beyond  
224 international treaties like the Nagoya Protocol that restrict particular kinds of multiomic  
225 sampling<sup>28,29</sup>. Many countries have developed legislation governing access to genetic  
226 resources, their use, and related benefit sharing. Collections will enter into written  
227 agreement with institutions or government agencies in the provider country and in some  
228 cases these agreements will prohibit external transfer to third parties without prior  
229 permission<sup>30,31</sup>. To prove that institutional permission to sample was granted, publishers  
230 and aggregators of genomic data (e.g., GenBank) normally require copies of permits before  
231 a paper can be published or sequences deposited, respectively. Proof of permissions also  
232 may be required by federal granting agencies.

233

234 *4. Prioritize destructive sampling of more recent collections.* Newer specimens generally  
235 yield higher quality and more abundant DNA<sup>18,24,32</sup>. Whenever possible, older specimens  
236 should not be used and historic vouchers should not be destroyed. In some cases,  
237 sequencing the type specimen of the species—which in many cases also is the oldest  
238 specimen—may be thought to be crucial<sup>33,34</sup>. However, some institutions may not allow  
239 destructive sampling of type specimens (see #3, above).

240



241 *5. Destructively sample only the amount of tissue required for the analysis.* A comprehensive  
242 sampling plan for a project should be determined before initiating any destructive  
243 sampling and the plan should be submitted along with an application for a permit or visit to  
244 the herbarium. Because, many biodiversity investigations require only a single placeholder  
245 taxon, e.g., a single species representing a much larger clade (genus) may suffice. In such  
246 cases, rare or threatened species should be sampled only as a last resort, only after  
247 protocols have been well established, and only for science deemed most impactful.

248

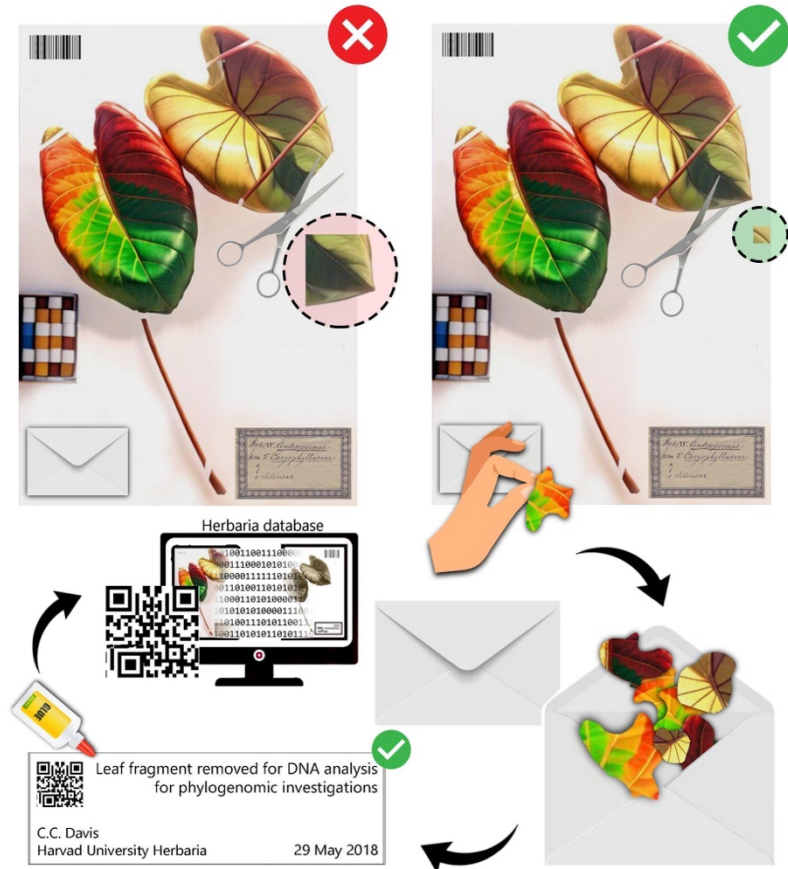
249 Data acquired through destructive sampling often improve the value of the physical (and  
250 extended) specimen, but destructive sampling should be done skillfully and minimally (i.e.,  
251 measure twice, cut once). In general, a good rule of thumb is to sample in such a way that it  
252 is impossible to identify that the collection has been sampled, except of course for the  
253 required addition of an annotation label indicating that the specimen was sampled (**Fig. 3**).  
254 Ideally, the annotation would be both physical and digital to enhance local and more global  
255 efforts to track sampling. We strongly advocate using QR codes to allow rapid access to  
256 specimen metadata and annotation histories with readily available devices, such as phones  
257 or tablets. Documenting and tracking this information with globally unique identifiers<sup>35</sup> is a  
258 hallmark of good science and stewardship of these collections.

259

260 Any destructively sampling must  
 261 maintain the future utility of a  
 262 specimen. Thus, from any  
 263 specimen, take only what is  
 264 required for the planned  
 265 research—i.e., do not take ‘rainy  
 266 day’ surplus samples for potential  
 267 or undefined future uses.  
 268 Sampling should always be  
 269 prioritized from collections with  
 270 abundant materials (**Fig. 3**).

271  
 272 It is also imperative to ensure  
 273 that crucial developmental stages  
 274 or morphological features are not  
 275 destroyed completely. For  
 276 example, the upper and lower  
 277 surfaces of leaves may have  
 278 distinct morphological differences  
 279 that are important for species  
 280 identification. Thus, if only a single  
 281 abaxial (lower) surface of a leaf is  
 282 showing on an herbarium sheet,  
 283 leaves in which the adaxial (upper) surface is exposed should be prioritized for sampling.  
 284 This recommended approach also preserves, for example, key climate-change response  
 285 traits such as stomatal size and density, which tend to be abundant on the lower leaf  
 286 surface<sup>17,36</sup>.

287  
 288 *6. Apply effective and proven methods.* It is important to apply state-of-the-art protocols and  
 289 experimental designs whose success has been demonstrated. For example, in the last three  
 290 decades, numerous DNA extraction techniques have been developed and evaluated,



**Figure 3.** Recommended guidelines for destructive sampling of herbaria. Remove only very small tissue fragments and prioritize specimens with abundant material; start first with packeted fragments when they are available. Upon sampling, it is essential to immediately add an annotation label documenting the destructive sampling event. This can be accomplished with a conventional physical label as shown or with a digital QR barcode for more easily viewing the entire history of specimen metadata, including a detailed annotation timeline.

291 including bulk cesium chloride extractions, cetyltrimethylammonium bromide (CTAB)-  
292 based extractions, silica columns, and magnetic beads<sup>37-39</sup>. A key to successful experimental  
293 design and implementation is to continually refine and perfect multiomic methodologies to  
294 require only miniscule amounts of tissue. For some requests for DNA extraction may be for  
295  $\geq 4 \text{ cm}^2$  of leaf tissue (D. White, pers. comm.) even though  $1 \text{ cm}^2$  of leaf may be more than  
296 adequate and is a rule of thumb dimension. We advocate more sophisticated explorations  
297 of these parameters—see below.

298

299 Good models for such efforts comes from the burgeoning field of ancient DNA (aDNA)<sup>40</sup> and  
300 DNA-based forensics<sup>11</sup>, where novel laboratory and bioinformatics techniques have  
301 sparked methodological sea changes in multiple fields of scholarship. New methods and  
302 best practices for extracting tissues that were thought to be intractable, including formalin-  
303 <sup>41</sup> or ethanol-preserved tissues, also have been developed<sup>42</sup>. As they become available and  
304 more widespread, these advanced methods also should be applied to destructively  
305 sampling herbarium specimens<sup>43-45</sup>.

306

307 Research limitations resulting from taxon or sample specificity also should be explored and  
308 identified by researchers before beginning any larger destructive sampling initiative. In  
309 addition to age-related degradation of DNA, DNA extraction from conventionally prepared  
310 herbarium specimens from wet tropical environments may be poorer for multiomic  
311 investigation than species from drier tropical environments<sup>24</sup>. Similarly, there is  
312 substantial variation in DNA extraction success across clades (e.g., Cyperaceae and  
313 Sapindaceae have higher yields than Fabaceae and Melastomataceae; <sup>24</sup>).

314

315 If multiomic extractions are to be treated as loans to be returned (see Best Practice #3 for  
316 Herbarium Stewards), the information on extraction method and concentration and quality  
317 should also be provided and catalogued. Sequences should be accessed simultaneously into  
318 local institutional databases and in NCBI/Genbank or similar publicly accessible database  
319 so that others can use the data and avoid unnecessary sampling of the same or related  
320 specimens.

321

322 Finally, if proposed methodologies are based on data-free folklore or rules of thumb, or are  
323 new and unproven, it is important to provide proof-of-concept alongside institutional  
324 requests to sample herbaria. For example, 'rule-of-thumb' tissue sizes often overlook its  
325 thickness, density, or the reliability of extraction success within a particular taxon or from a  
326 particular habitat or biome. The amount of tissue yields can vary greatly by tissue density,  
327 age, taxon, habitat, or preservation mode<sup>24</sup>, and more careful and systematic investigations  
328 of these factors considering specific protocols to be used should be done before  
329 destructively sampling specimens. In such cases, 'test' herbarium samples can be sampled  
330 destructively to demonstrate feasibility (e.g., harvesting spectral signature data for species  
331 identification and functional trait assessments<sup>46</sup>).

332

333 *7. Confirm specimen determinations prior to sampling.* Many herbarium users assume that  
334 previous determinations of the identity of specimens are always correct. However,  
335 specimens are often misidentified or are labeled with outdated names (e.g.,<sup>47</sup>). Confusing or  
336 nonsensical research results can occur simply because sampled specimens were mis-  
337 identified or mis-labeled<sup>48</sup>. Thus, users should verify the determinations of all specimens  
338 prior to sampling. For systematics and phylogenetic research, this issue can be addressed  
339 by deliberately factoring in some revisionary taxonomic work as part of the project. As  
340 intuitive as such a proposition sounds, it is more common than not for users to request  
341 destructive samples from an herbarium while declining to see and verify the physical  
342 vouchers.

343

344 For broader-scale phylogenetic or non-systematics projects, where researchers lack the  
345 experience necessary to identify requested specimens, collaboration with a taxonomist,  
346 systematist, or phylogeneticist should be the norm. Similarly, if expert identifications are  
347 already available for a group of specimens because of the recent work of a taxonomist who  
348 is alive and active, those determinations should be recognized for the genuine intellectual  
349 contributions they are and collaborations involving co-authorship should be seriously  
350 explored (see also Best Practice #9 for Herbarium Users, below).

351

352 *8. Annotate specimens after sampling.* Clear annotation, both physical and digital, and  
353 associated record-keeping are essential for documenting destructive sampling of  
354 herbarium specimens. An annotation should be made immediately upon sampling and  
355 should minimally include: the name of the sampler or research team, contact information of  
356 sampler, date of collection when the sample was taken, what kind of material was removed,  
357 and the title and goal of the project (**Fig. 3**). This annotation should be digitized  
358 simultaneously as part of the extended specimen both in local institutional databases and  
359 in large data aggregators such as GBIF and iDigBio. If herbarium staffing is minimal, the  
360 user should extend support to help complete this important step.

361

362 *9. Make data from destructive sampling publicly available immediately.* Any data derived  
363 from destructive sampling should be made publicly available as soon as it is produced.  
364 Larger federal funding agencies, such as the United States National Science Foundation,  
365 have timeline requirements for data delivery and serve as good models. Similar examples  
366 of such policies include the [Darwin Tree of Life](#) project and the [Fern Tree of Life](#).

367

368 *10. Ensure credit to herbaria in publications and grants.* Natural history collections,  
369 including herbaria, are under-funded and under-appreciated<sup>49,50</sup>. To remedy this, herbaria  
370 should be formally recognized for supplying tissues for destructive sampling. Minimally,  
371 herbarium vouchers utilized for multiomic initiatives should be prominently cited in  
372 publications ensuing from these initiatives. Such recognition should be prominent and  
373 extend beyond simple acknowledgement statements that conclude a scientific paper<sup>26</sup>.  
374 Specifically, more formal attributions of herbaria and natural history collections are  
375 required, including possibly co-authorship of institutions that is tracked by citation  
376 aggregators such as Google Scholar<sup>51</sup>. Minimally, authors should formally recognize the  
377 herbaria that supply tissue in some format that is trackable and citable, such as working  
378 with herbaria to ensure that clear records to track sampled specimens are available via  
379 GBIF, which can provide a citable DOI for relevant datasets. Such metrics are essential for  
380 justifying the use and importance of herbaria for stewardship and funding. FAIR standard  
381 conventions should apply here<sup>52</sup>.

382

383 Similarly, curators, support staff, and especially collectors of sampled herbarium specimens  
384 should be invited to participate in research as co-authors whenever possible. Contributions  
385 should be indicated through the Contributor Roles Taxonomy (CRediT) system, especially  
386 for larger, more substantial requests<sup>26</sup>. We have witnessed researchers systematically  
387 harvest in a single short visit hundreds of samples that encompass the entire career of  
388 collectors and researchers at those institutions. In most cases, these individuals were not  
389 consulted or extended an invitation to be involved in the research produced by the  
390 destructive sampling or the ensuing large grants and publications derived from them.  
391 These approaches are predatory and should be avoided at all costs. An appropriate  
392 alternative to such sampling is to canvas more broadly the community of herbarium staff  
393 who supported these collection and identification efforts and offer formal collaborations or  
394 co-authorship.

395  
396 Finally, destructive sampling requests make enormous demands on staff and their  
397 institutions, whose budgets are already thin. Research grants that use herbarium  
398 specimens should explicitly include funds to herbaria to offset staff costs and the associated  
399 curation and stewardship of such collections, which in many cases have been ongoing for  
400 centuries. Indeed, funding for participating herbaria that have contributed sampling  
401 specimens for multiomics should be formally included in all budgets, just as is done for  
402 other field and lab expenses and fees. Specifically, we advocate that users explicitly include  
403 herbaria as funded subcontractors/subawardees in grants that are written in good faith  
404 and with careful consultation with partner herbaria. Moreover, the funding allocation of  
405 any request should scale with the age and rarity of material requested. Appropriate and fair  
406 cost models need to be established for such efforts. To this point, NY has created a  
407 sophisticated calculator for its curators that can be used to calculate all costs associated  
408 with accessioning new specimens into the herbarium. Such approaches can be used to  
409 estimate the long-term and substantial costs of stewarding any individual specimen.

410  
411 **Best practices for herbarium stewards**

412

413 *1. Evaluate individually each request for destructive sampling.* Destructive sampling requests  
414 should be considered on a specimen-by-specimen basis, ideally with each sampled  
415 specimen recorded in a Material Transfer Agreement (MTA). This transparency ensures  
416 that no material is sampled without the explicit permission of the institution; stewards of  
417 collections can check to ensure institutional obligations have been met (e.g., Nagoya  
418 Protocol requirements, CITES, Memoranda of Understanding (MOUs), permits or other  
419 agreements with other countries or institutions, bioculturally sensitive materials, rare or  
420 threatened species restrictions)<sup>53</sup>; and users of those collections can be confident that they  
421 have documented permission to conduct their proposed project. Best practice dictates that  
422 institutions should always track transfers of their specimens and derivative samples and  
423 for countries and material falling under the scope of the Nagoya Protocol, which is a legal  
424 commitment for signatories<sup>30</sup>.

425  
426 *2. Destructive sampling of bioculturally sensitive plants requires special considerations.*  
427 Certain specimens, plant species or populations, and research topics necessitate an  
428 additional level of care and attention when it comes to destructive sampling. Many plants  
429 have spiritual and ceremonial importance and are essential to Indigenous groups who  
430 steward or govern nearly 40 million km<sup>2</sup> of land and inland waterways in 82 countries,  
431 including 25% of earth's terrestrial lands<sup>54,55</sup>. The colonial legacy of herbaria is well  
432 documented, and most herbarium collections are housed in the Global North<sup>56</sup>. This creates  
433 additional responsibilities for most stewards of herbaria to ensure these specimens are  
434 treated with care and are not used as a means of circumventing the sovereignty of nations  
435 and Indigenous people as pertains to their biodiversity and associated knowledge.  
436 Established guidelines for respectful, responsible, and ethical partnership with Indigenous  
437 communities in the context of field biology<sup>57</sup> should be extended to herbarium sampling. If  
438 an activity would be considered unethical in the context of newly collected specimens, it  
439 should not be done using herbarium specimens.

440  
441 Multiomic sampling of extinct, rare, or bioculturally sensitive plants (whether alive or  
442 preserved in herbaria) should proceed only after careful user consideration and  
443 stewardship deliberations<sup>20</sup>. Users should be made aware that requests to sample such

444 materials make unique demands that potentially impinge on many groups and individuals.  
445 In these cases, provenance, acknowledgement, and collaboration are essential to advancing  
446 research with such collections<sup>58</sup>, whose use also may be regulated by international treaties  
447 and domestic laws (e.g., CITES). Stewards must make users aware of these added  
448 requirements and sensitivities during planning and permitting discussions.

449

450 *3. Treat destructive multiomic samples from collections as loans.* The active loan and  
451 exchange of physical specimens among herbaria has been ongoing for more than a century.  
452 They have been an essential and effective means to more comprehensively study  
453 biodiversity and obviate extensive, unnecessary, and expensive travel. Multiomic  
454 derivatives resulting from destructive sampling should be considered a part of the physical  
455 collections and should be treated in a similar way as loans of physical specimens. Unused  
456 materials and aliquots of extracted DNA should be returned to the institutions who steward  
457 these collections, or at least repatriated to a related herbarium where duplicates of  
458 sampled specimens are held (see Best Practice #6 for Herbarium Users, above). We  
459 recommend that multiomic loans derived from specimens be returned within two years of  
460 sampling regardless of publication status and following established MTAs and Data Use  
461 Agreements (DUAs). Such loan returns can be accomplished using infrastructure within an  
462 institution (e.g., cryofacility) or by aggregated stock facilities that supply samples, such as  
463 the algal facility at Bigelow Labs (<https://ncma.bigelow.org/>).

464

465 Relatedly, just as digitization has created a revolution in the biodiversity sciences and has  
466 increasingly become a standard curatorial practice in herbaria<sup>6,9</sup>, the community should  
467 embrace curating multiomic samples in botanical collections. For example, genomic and  
468 biochemical resources in herbaria are invaluable elements of the extended specimen and  
469 should be stewarded accordingly. Such stewardship will vary with institutional resources,  
470 e.g., providing extraction in on-site labs (at K, for example), cryogenic facilities for storing  
471 extracted DNA aliquots, or partnerships with existing facilities to safeguard these samples  
472 on behalf of the herbaria which house their associated voucher specimens. Uncountable  
473 aliquots of DNA from finite and irreplaceable specimens have been discarded or lost when  
474 researchers move, retire, or die, and because of the lack of a standardized process for



475 depositing extracted samples. Curating a collection of extracted DNA (or other related  
476 multiomic sample) is costly and time-consuming, but will extend the research value of  
477 herbarium specimens immeasurably into the future<sup>30</sup>.

478

479 *4. Develop a succession plan for derivative (e.g., multiomic) collections from destructive*  
480 *sampling.* Academics frequently move as their career trajectories develop, and eventually  
481 retire or pass on. It is imperative that a clear succession plan be in place for multiomic  
482 samples after their initial use has been completed. This can be clarified in an MTA.  
483 Minimally, the sample should be searchable online and available to the community.  
484 Succession plans are of particular importance for large research labs where faculty and  
485 staff careers may span decades, during which time a very large number of potentially useful  
486 samples may accumulate. In this case, users will assume a stewardship role and work with  
487 herbarium leadership to ensure safe transfer of derivative collections.

488

489 *5. Protect against hoarding.* Any serious inquiries involving exceptionally large requests  
490 should be vetted carefully by internal and, where appropriate, external stakeholders.  
491 Moreover, material requests involving larger sampling efforts should be initiated as part of  
492 a formal collaboration with associated MOUs, MTAs, and DUAs negotiated prior to the  
493 onset of such efforts. These agreements also should include a clear statement of when the  
494 data and metadata will be released and how aliquots will be archived and shared. As noted  
495 earlier, herbaria should guard against being seen as a 'back door' to access the genetic  
496 diversity of another country (see Best Practice #3 for Herbarium Users). Large requests for  
497 sampling from a particular country should be directed first to its national institutions so  
498 appropriate collaborations can be organized by the user. Whenever possible, all reasonable  
499 efforts should be made to accommodate requests from in-country representatives to  
500 sample their country's biodiversity. There may be additional rules restricting use of  
501 collections from particular countries, and in all cases for country-focused studies,  
502 collaboration with in-country partners must be the norm<sup>56,57,59</sup>.

503

504 Finally, the primary voucher set of any such derivative collections (e.g., DNA aliquots)  
505 should reside at the herbarium providing the samples. If adequate curation of such

506 derivative collections is not possible, a third-party institution should be designated, ideally  
507 another herbarium with appropriate domain knowledge and expertise. Similarly, following  
508 long-standing practice among herbaria to deal with physical specimens, large derivative  
509 sample sets should be dispersed to all collaborating institutions and institutional partners  
510 rather than to single individuals or labs to mitigate against loss.

511  
512 *6. Ensure proper institutional permitting is up to date, held, and available for inspection.* This  
513 includes national level permits for CITES-listed plants, relevant import/export permits,  
514 permits for plants deemed illicit, and any additional permits required by sub-national  
515 jurisdictions. Sampling should be restricted wherever possible to specimens that can be  
516 clearly documented with all relevant permits. We recognize that this high standard is not  
517 always possible with older material, but it should be standard practice with newer  
518 collections, which are also preferable in terms of metabolites and DNA quality (see Best  
519 Practice #4 for Herbarium Users;<sup>30</sup>)

520  
521 *7. Destructive sampling should coincide with barcoding and specimen digitization.* Every  
522 specimen should be barcoded, imaged, and transcribed, ideally *prior* to destructive  
523 sampling. Users should be asked to support financially or, if staffing is unavailable, help  
524 input these data following institutional guidance and standards at the time of sampling. A  
525 visit is not considered complete until this essential bookkeeping is finished. Stewards  
526 should also expect that users will verify and update the taxonomic accuracy of each  
527 specimen sampled, which for some groups and regions of the world may be highly  
528 inaccurate<sup>47</sup>. Ensuring that researchers inform herbaria of newer, better determinations is  
529 critical to the vitality and continued relevance of herbaria.

530  
531 Relatedly, stewards should coordinate with organizations, networks, and initiatives to  
532 provide greater access to herbarium resources (e.g., GBIF, iDigBio, DiSSCo, and BCoN),  
533 further extend specimen data, and grow the Global Metaherbarium<sup>2,49</sup>. Annotations of  
534 collections for multiomic sampling should be added to the digital specimen record with the  
535 associated researcher who collected the sample and the focal project (**Fig. 3**). These efforts  
536 should also be further coordinated with ongoing large efforts to sequence genomes across

537 the Tree of Life, such as the [Earth Biogenome Project](#)<sup>60</sup>, the [African Biogenome Project](#)<sup>61</sup>,  
538 and the [10,000 Plants Genome Project](#)<sup>62</sup>. Multiomics data arising from the conducted  
539 research always should be made available in publicly accessible databases within a  
540 reasonable period regardless of its publication status. Herbaria should consider restricting  
541 sampling of their material if this is not the case.

542

543 *8. Maintain thorough records and hold users accountable.* Ideally, an institution should be  
544 able to link any specimen cited in a publication resulting from destructive sampling directly  
545 to an MTA or DUA certifying the researcher had permission to take that material and use it  
546 according to the reported methods. When researchers fail to honor the conditions of these  
547 agreements (e.g., deliberately sampling specimens not included in the MTA, using materials  
548 for different projects than the ones proposed, withholding data resulting from samples),  
549 they should be prevented from future access. Repeat offenders should be reported to their  
550 institutions and to the herbarium community so other collections can protect their  
551 specimens accordingly. Using material without permission is theft and makes the task of  
552 responsibly safeguarding collections impossible.

553

554 *9. Destructive sampling requests should balance the importance of current discovery against*  
555 *future needs.* This is complicated and potentially more subjective, but nevertheless  
556 important. Certain research questions and potential discoveries ensuing from them should  
557 be prioritized for destructive sampling, especially when rare or sensitive specimens are  
558 under consideration. This includes extinct species and populations, and species central to  
559 humankind, including not only crops and their wild relatives but also plants of medicinal  
560 and spiritual importance. At present, crops and their wild relatives are not well preserved  
561 in seed banks and germplasm resources<sup>63</sup>. However, when available, such material allows  
562 researchers to grow plants for study, thus obviating the need to sample herbaria and  
563 expanding their ability to conduct broader investigations. The same likely applies to other  
564 plants of human use. Herbaria offer another important and underused path to preservation  
565 of these groups and should be treated with great care and caution. Permission to sample  
566 these collections should be granted only in cases where research promises to add

567 significant additional value to the extended specimen and where we should do our very  
568 best to facilitate these advancements for the benefit of humanity.

569

570 *10. Establish transparent policies for destructive sampling and treat them as 'living*  
571 *documents.'* Biodiversity will continue to change. Similarly, new science will develop and  
572 lead to new technologies. It is important that documentation on how specimens may be  
573 sampled in herbaria also evolve. Herbarium steering committees, leadership teams,  
574 curators, staff, researchers, and other stakeholders should work to keep these documents  
575 current and relevant to facilitate responsible stewardship and support impactful science.  
576 Although some amount of judgment by collections stewards will always be necessary,  
577 guidelines for researchers about sampling policies should be as clear and consistently  
578 applied as possible. Ideally, these policies should form part of the broader strategic plan for  
579 any herbarium that explicitly balances preservation with use. Although institutions will  
580 vary in their needs and resources and it is impossible to establish a one-size-fits-all policy,  
581 there is value to both the herbarium community and its various users in having consistent  
582 guidelines across institutions.

583

## 584 **Summary and conclusions**

585 New technologies have revolutionized the uses of herbarium specimens, yielding  
586 invaluable insights into evolution, ecology, taxonomy, conservation, ethnobotany, history,  
587 and the social sciences. Although these developments have been exciting and herbaria have  
588 certainly benefited from a renewed global interest in their resources, a lack of clear  
589 community guidelines and best practices for destructive sampling of irreplaceable  
590 specimens risks the long-term sustainability of the only verifiable record of plant and  
591 fungal life on Earth. The lack of best practices also places undue burdens on the stewards  
592 and users of these collections, resulting in needless confusion and consternation.

593 We have outlined here what we hope will be a starting point for much-needed  
594 conversations within and among herbaria and their associated research communities. The  
595 more than 3,500 herbaria across the globe <sup>1</sup> are nearly as diverse as the specimens they

596 contain, so it is nearly impossible to write one-size-fits-all prescriptions. We hope the  
597 recommendations presented here will be a useful framework for the community to begin to  
598 clarify and create a path towards greater common understanding about how herbarium  
599 specimens can be sampled sustainably and responsibly while continuing to expand our  
600 knowledge and understanding of botanical diversity and preserve their use for generations  
601 to come.

602 Although we have suggested separate recommendations for users and stewards, we feel  
603 strongly that progress requires that both users and stewards better understand and  
604 appreciate one another's needs and concerns. Taken collectively, our proposed best-  
605 practices will help ensure that: i) physical collections will be preserved indefinitely and will  
606 continue to grow for the benefit of future generations; ii) the expertise required to build  
607 and maintain collections will be valued, recognized, and cultivated; iii) the rights of nations  
608 and Indigenous communities for their biodiversity and knowledge will be respected; iv)  
609 access to samples for research will be fair and equitable; and v) knowledge generated from  
610 destructive sampling will build upon and complement the expanded specimen network,  
611 creating a truly powerful, accessible, and comprehensive Global Metaherbarium.

612

### 613 **Author Contributions**

614

615 CCD, JT, AP, AA, and ES jointly conceived the premise of this paper. CCD wrote the paper  
616 with key initial editorial input on the stewards and conclusions sections from JT. All  
617 authors revised and approved the final text. Figures were coordinated and/or compiled by  
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619

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621

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636

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