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## Selection and characterization of botanical natural products for research studies: a NaPDI center recommended approach†

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Dietary supplements, which include botanical (plant-based) natural products, constitute a multi-billion-dollar industry in the US. Regulation and quality control for this industry is an ongoing challenge. While there is general agreement that rigorous scientific studies are needed to evaluate the safety and efficacy of botanical natural products used by consumers, researchers conducting such studies face a unique set of challenges. Botanical natural products are inherently complex mixtures, with composition that differs depending on myriad factors including variability in genetics, cultivation conditions, and processing methods. Unfortunately, many studies of botanical natural products are carried out with poorly characterized study material, such that the results are irreproducible and difficult to interpret. This review provides recommended approaches for addressing the critical questions that researchers must address prior to *in vitro* or *in vivo* (including clinical) evaluation of botanical natural products. We describe selection and authentication of botanical material and identification of key biologically active compounds, and compare state-of-the-art methodologies such as untargeted metabolomics with more traditional targeted methods of characterization. The topics are chosen to be of maximal relevance to researchers, and are reviewed critically with commentary as to which approaches are most practical and useful and what common pitfalls should be avoided.

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1. **Introduction**
2. **Terminology relevant to botanical natural products**
3. **Journal and government agency guidelines for characterization of natural product study material**
4. **Literature research**
5. **Obtaining botanical natural product study material**
  - 5.1 **Voucher specimens**
  - 5.2 **Contracting an independent laboratory to prepare study material**
  - 5.3 **Purchasing study material from a commercial supplier**
  - 5.4 **Optimal number of samples to analyze**
  - 5.5 **Purchasing authenticated botanical reference materials**
  - 5.6 **Collecting cultivated or wild plant material**
6. **Authentication**
  - 6.1 **Authentication by microscopic and morphological characteristics**
  - 6.2 **Authentication by DNA barcoding**
  - 6.3 **Authentication by targeted analysis of “marker compounds”**
  - 6.4 **Authentication by “chemical fingerprints”**
  - 6.5 **Authentication by untargeted metabolomics**
  - 6.5.1 **Metabolomics data analysis**
  7. **Selecting a “representative” sample**
    - 7.1 **Selecting a formulation**
    - 7.2 **Standardized products**
    - 7.3 **Selection of extraction solvent**
  8. **Quality control and comparison of botanical product composition**
    - 8.1 **Quantitative metrics for comparing similarity of natural product mixture**
    - 8.2 **Adulteration of botanical natural products**
    - 8.3 **Consideration of contamination**
  9. **Identification, structure elucidation, and quantitative Analysis of bioactive compounds in the selected study material**

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- 9.1 Determining which constituents of a botanical natural product are biologically active
- 9.1.1 Advantages and limitations of *in vitro* assays
- 9.1.2 Approaches for integrating *in vitro* data with chemical composition
- 9.2 Obtaining “pure” standards
- 9.3 Structure elucidation of constituents of botanical natural products
- 9.4 Quantitative analysis
- 10. Conclusions

- 12. Conflicts of interest
- 13. Acknowledgements
- 14. References

## 1. Introduction

More than 50% of Americans acknowledge taking dietary supplements,<sup>1,2</sup> with sales more than tripling since passage of the Dietary Supplement Health and Education Act in 1994.<sup>3,4</sup> In many cases, these dietary supplements are prepared from



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*rose through the ranks of RTI and eventually directed the Natural Products Laboratory. In 2009, he moved his group to the Department of Chemistry & Biochemistry at the University of North Carolina at Greensboro, where he helped to grow their Ph.D. program in Medicinal Biochemistry. His lab largely focuses on the characterization and development of chemical entities from natural sources. This includes efforts that seek new drug leads, as well as, projects, like this, that focus on insuring the safety and quality of herbal preparations.*

botanical (plant based) material, and as such are referred to as 'botanical natural products.' Countless studies have been devoted to the scientific evaluation of the safety and/or efficacy of botanical natural products. Investigators involved in such studies face a unique set of challenges (Table 1). The research methodology for evaluating safety and efficacy of conventional (pharmaceutical) drugs operates with the assumption that the product being tested is a single compound of known purity, identity, and concentration. Natural products differ from their pharmaceutical counterparts in that they are typically complex mixtures, for which the identities and quantities of components present are not fully known. The composition of these natural products can vary depending on the method of preparation or source material used.<sup>5</sup> Such variability can impact the interpretation of *in vitro*, non-clinical *in vivo*, and/or clinical studies.

To enable studies to be as robust and as widely applicable as possible, selecting, characterizing, and ensuring continued quality and consistency of a botanical natural product is an essential task. The ideal characteristics of a botanical natural product used for research studies include that it is authenticated (of known identity), well-characterized in terms of potentially active constituents, and stable (Table 1). At present, many studies are conducted with botanical natural products that lack one or more of these characteristics. Herein, we seek to provide guidance for effective selection of botanical natural products prior to *in vitro*, non-clinical *in vivo*, or clinical evaluation. We discuss the relative merits of potential analytical approaches, including state-of-the-art metabolomics techniques, and recommend specific, effective and practical guidelines (Fig. 1). The recommendations we provide are informed by the coauthors' experience for several decades researching botanical natural products, culminating in our role as investigators participating in the Center of Excellence for Natural Product Drug Interaction Research (NaPDI Center). The National

**Table 1** Characteristics of the ideal botanical natural product for use in research studies

- (1) Representative of what is commonly used by consumers
- (2) Authentic (species verified)
- (3) Well-characterized
  - (3.1) Active constituents known
  - (3.2) Concentrations of active or marker metabolites have been characterized and comply with monograph
- (4) Free of contamination and adulteration
- (5) Sufficient material available to conduct *in vitro* and/or clinical studies
- (6) Material consistent for duration of the trial
  - (6.1) Shelf life (*i.e.* stability)
  - (6.2) Batch-to-batch reproducibility

Center for Complementary and Integrative Health established the NaPDI Center to address the need for robust methods to study the potential for natural products to precipitate clinically significant pharmacokinetic interactions with conventional medications (natural product–drug interactions).<sup>6</sup> While the focus of the NaPDI Center is specifically on the evaluation of interactions between conventional drugs and natural products, the recommendations provided herein are more broadly applicable to the selection of botanical natural products for any *in vitro* or *in vivo* investigation (including studies to evaluate efficacy or toxicity).

## 2. Terminology relevant to botanical natural products

"Natural products" are a broad range of substances that can come from a variety of sources, including bacteria, fungi, marine organisms, and plants. The term can be used to describe both complex mixtures and single isolated compounds that come from these mixtures. For this publication, we focus specifically on botanical natural products, *i.e.* natural products derived from plants. Such botanical natural products are available for use by consumers in a variety of forms, including capsules containing raw or extracted material, extracts, teas (typically raw plant material that is extracted in hot water prior to use), tinctures (ethanolic extracts), and traditional formulations such as powders used in Traditional Chinese Medicine or Ayurvedic practices. In this manuscript, we use "botanical natural products" as an umbrella term to describe complex plant-based preparations. Terms used elsewhere include: supplements, herbal medicines, herbal drugs, herbs, botanical preparations, nutraceuticals, phyto-medicines, and botanical medicines. Additional definitions for terms used in this manuscript are provided in Table 1S (ESI†).

From a regulatory perspective, distinctions are often made between various botanical natural products depending on how they are used by consumers. The regulatory approaches and classifications of botanical natural products are complex and differ from country to country, and a detailed comparison is beyond the scope of the current work.<sup>7</sup> The majority of botanical natural products consumed in the US (including the examples discussed in this manuscript) are regulated either as foods or dietary supplements. From a scientific perspective, the way in which the products are classified and regulated is less



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*group at UNCG, for which a major focus in the development of metabolomics as a tool to understand synergy and complexity in biologically active botanical natural products. This work has been continuously funded by the National Institutes of Health for more than 15 years, and was awarded the Jack L. Beal Award from the Journal of Natural Products in 2011. Dr Cech is a member of the Center of Excellence for Natural Product Drug Interaction Research, and Co-Director of the Medicinal Chemistry Collaborative (<https://mcsquared.uncg.edu/>).*



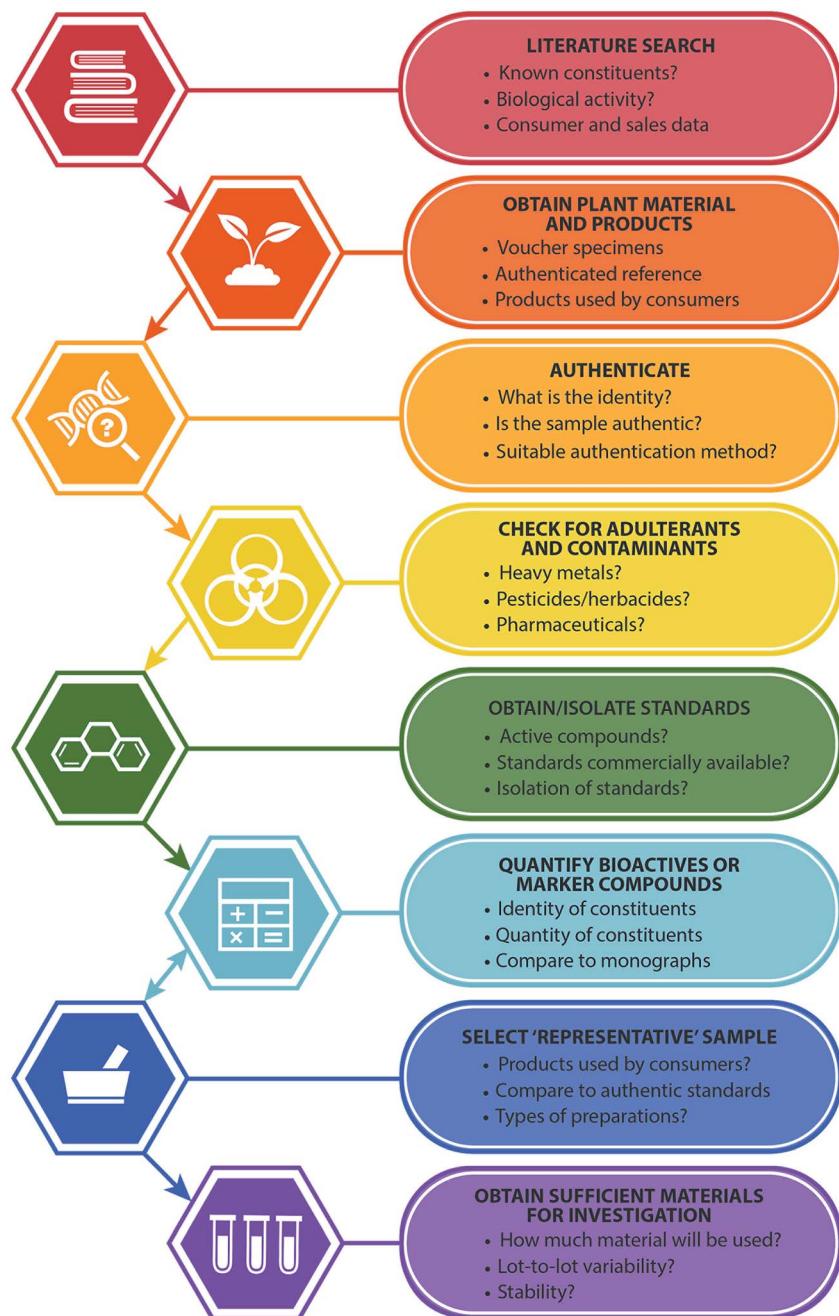


Fig. 1 Recommended steps in the selection of botanical natural products for research purposes (*in vitro* and/or *in vivo* studies), and important questions for consideration with each step.

important than the material from which they are prepared, and much of the information presented herein is relevant to botanical natural products broadly defined, regardless of their regulatory classification.

### 3. Journal and government agency guidelines for characterization of natural product study material

Several agencies provide guidance for the type of information that should be obtained about a natural product prior to its use

for research purposes. The United States Food and Drug Administration (FDA) recommendation regarding botanical drug clinical trials specifies that the investigational new drug application contains “a chemical identification for the active constituents or characteristic markers in the drug substance, if possible”.<sup>8</sup> The National Center for Complementary and Integrative Health (NCCIH), a component of the National Institutes of Health, funds a large portion of the research conducted on botanical natural products in the United States. NCCIH has established a “Natural Product Integrity Policy”, which requires that researchers provide information about the identity,



extraction solvent, characterization (*i.e.*, metabolite profile), stability, standardization, and storage of all natural products that will be used in NCCIH funded studies.<sup>9</sup> Some individual journals also provide specifications for how complex botanical natural products should be characterized before data on these products can be published.<sup>10–13</sup> Journal editors have a great deal of influence in this regard, since their publishing requirements cross geopolitical borders.<sup>14</sup>

## 4. Literature research

Before beginning botanical natural product studies, it is useful to obtain information from the peer-reviewed literature regarding: (1) identities of major metabolites, especially those that may be biologically active; (2) potential *in vitro* or *in vivo* targets and activities; and (3) current knowledge gaps. A recent publication by the NaPDI Center addresses useful strategies for critically evaluating botanical natural product literature to identify gaps in the knowledge base.<sup>6</sup> Due to the complexity of natural product constituents, it is important to note that reported chemical structures may contain errors or inconsistencies. Thus, the literature review by an organic or natural product chemist trained in structure elucidation is recommended.

Another important topic to research during sourcing of botanical natural products is information regarding traditional and consumer usage of the product to be studied. The investigator will wish to answer questions such as: what are the most common species used medicinally? Which plant parts (for example, roots, leaves, berries) are typically used? What types of preparations (capsule, tincture, tea, compress, topical *versus* internal) are employed? Some information of this nature may be available in the peer-reviewed literature, but it is often necessary to seek other sources. Common products used by consumers can be evaluated *via* national surveys, such as the National Institute of Health's Office of Dietary Supplement's Dietary Supplement Label Database,<sup>15</sup> National Health and Nutrition Examination Survey (NHANES),<sup>16</sup> consumer quality reports<sup>17,18</sup> as well as industry or retail sales reports.<sup>4,19,20</sup>

### Box 1 Summary of recommendations: literature review for botanical natural products

- Examine references to identify known constituents and biological targets.
- Investigate botanical usage information, including common species used, physiological portions harvested, and preparation.
- Utilize both peer-reviewed resources, government reports, and industry-based information.

## 5. Obtaining botanical natural product study material

Prior to conducting *in vitro*, non-clinical *in vivo*, and/or clinical studies, it is important that researchers identify an authentic

natural product (*i.e.* correct assignment of genus and species) that is available in sufficient quantity to conduct the studies. In the case of clinical studies, this product should be selected to closely resemble (or be identical to) the commercial product(s) used by the target population. Where is such a product obtained? One option is to purchase this material from a commercial distributor. Alternately, researchers may opt to obtain raw material and prepare a natural product themselves or to employ a contract laboratory for formulation.<sup>21</sup> Finally, several suppliers provide high quality reference materials of botanicals for research purposes. Here we discuss the importance of voucher specimens and compare the relative merits of various sources of botanical material.

### 5.1 Voucher specimens

Under optimal circumstances, a voucher specimen of any botanical natural product to be studied is collected at the same time and from the same lot that the study material is obtained. The voucher specimen consists of an intact, dried sample of the plant material, including the flower when possible, and as many parts as can reasonably be collected, *i.e.* roots, stem, leaves, flowers and/or seeds.<sup>22</sup> The voucher specimen is used for taxonomic identification of the study material by a trained botanist or otherwise qualified individual. Herbaria have existed for centuries,<sup>23</sup> and these vouchers serve as historical records of plant specimens, and modern uses of these go beyond taxonomy, including areas like conservation biology.<sup>24</sup> Thus, a pressed and dried sample of the voucher is deposited in a regional or national herbarium, where it is catalogued and stored for future reference. Herbarium vouchers are essential for preserving a record of the original sample tested and provide lasting, public access to that material in perpetuity. Authentication and retention of voucher specimens is required by major natural product journals.<sup>10–12,25</sup> For example, for the *Journal of Natural Products*, authors who “purchase dried ‘herbal remedies’ or other materials from companies must make provision for their proper deposit in a herbarium or other permanent repository, for access by future workers”.<sup>10</sup> Another example is *Phytochemistry*, which mandates that samples, “must also include a reference to voucher specimen(s) and voucher number(s) of the plants or other material examined” as well as “the name and address of the authority who identified each non-cultivated plant investigated”. Similar guidelines are in place for other major natural product journals.<sup>11,12,25</sup>

### 5.2 Contracting an independent laboratory to prepare study material

One way to obtain quality study material is to contract a laboratory to prepare the botanical natural product study material from raw materials to pre-determined specifications. This approach has been employed by other research groups.<sup>21,26,27</sup> A major advantage of employing a contract laboratory to produce study materials is that the researchers can be responsible for ensuring the quality, consistency, and rigor with which the material is prepared. However, contracting a laboratory to prepare a formulation can be expensive, and the production of



such formulations is time-consuming and requires access to specialized facilities and expertise. Even more importantly, a botanical natural product prepared by a contract laboratory may not reflect commercial products being consumed by the general public. Thus, many researchers opt instead to conduct studies with commercial products (the same products used by consumers). This was the approach that we took with the NaPDI Center's studies of both green tea [*Camellia sinensis* (L.) Kuntze (Theaceae)]<sup>28</sup> and goldenseal [*Hydrastis canadensis* L. (Ranunculaceae)].<sup>29</sup>

### 5.3 Purchasing study material from a commercial supplier

Many botanical natural products can be purchased from online distributors, health food stores, grocery stores. The advantage of purchasing commercial botanical natural products for study is that it is possible to select material that directly reflects what is being used by consumers (with the caveat that there may be a great deal of variability among commercial products). In addition, for clinical studies, commercial products have the advantage of being "food grade" meaning that they can be administered to human subjects. On the other hand, it is difficult to control quality and ensure correct identity of botanical natural products obtained from commercial sources. Commercial botanical natural products typically do not come with associated voucher specimens. Additionally, adulteration or mis-identification of commercial natural products (both intentional and unintentional) is common.<sup>30-32</sup> For example, in the process of selecting commercial products of goldenseal for clinical studies by the NaPDI Center, we identified several preparations sold as goldenseal (*Hydrastis canadensis*) that were a different botanical, Chinese goldenthread (*Coptis chinensis*).<sup>29</sup>

### 5.4 Optimal number of samples to analyze

When a study will be conducted with commercial products, the question arises as to how many samples should be evaluated before one is selected for study material. The goal is to select a sample set that is representative of the variability in products used by the target population. How can this be accomplished? Purchasing every potential commercial botanical natural product would theoretically be an ideal strategy. However, this is not always feasible due to constraints of time and cost. If a botanical natural product has a relatively small commercial footprint, it is possible to acquire all available products at a given time. As an illustrative example, the 35 commercial goldenseal samples selected for the second NaPDI Center study represented an estimated >90% of the readily available commercial products.<sup>29</sup> However, when a botanical natural product is found in hundreds or potentially thousands of products, the logistical and potentially financial hurdles of sampling a great number of samples must be taken into account during this planning stage.

For analyses of commercial samples, there has not been a conventional minimum value set on the number of samples needed to yield a robust analysis. Previous studies on botanical natural products have sampled as few as five products,<sup>33</sup> with many sampling 10–18 samples;<sup>34-37</sup> while some procure over 75

commercial products.<sup>38</sup> We elected to select a minimum of 30 commercial samples for each of the NaPDI Center's studies on green tea (34 products)<sup>28</sup> and goldenseal (35 products).<sup>29</sup> Importantly, these samples were selected based on sales data to represent the products most widely purchased for use by US consumers (our target population). The result was a sample set that captured variability among the products most widely consumed and could be realistically handled within constraints of processing and analysis time.

### 5.5 Purchasing authenticated botanical reference materials

Several sources provide authenticated botanical reference materials. The National Institute of Standards and Technology (NIST, <https://www.nist.gov>) offers a number of botanical natural products as verified reference materials available for testing and authentication purposes, as does the U.S. Pharmacopeia (USP, <https://www.usp.org>). Additionally, phytochemical reference suppliers like ChromaDex (Irvine, CA, <https://chromadex.com>) provide different grades of botanical reference material. A major advantage of these reference materials is that they have often been characterized already, and information is available about known genetic markers and the presence and identity of known metabolites. Some sources, such as ChromaDex, may also provide information regarding their internal voucher specimens. Even if the intention is to use a commercial product for the final studies, it is advisable, where possible, to obtain an authenticated reference material. The composition of this reference material can then be compared to that of commercial botanical natural products to verify identity and compare quality.<sup>28,29</sup>

### 5.6 Collecting cultivated or wild plant material

Certain botanicals are unavailable as authenticated references, and references must be obtained by other means. In some cases, it is possible to contract third-party vendors to obtain vouchered botanical samples. Another option is to purchase raw or minimally processed plant material from a nursery or farm, or, in the case of a botanical that grows wild, to harvest it from its native environment (sometimes termed "wild crafting"). In these cases, a voucher specimen should be collected at the time of harvest. For wild-harvested plants, permission of the land-owner or overseeing agency should be sought prior to collection.<sup>39</sup> Researchers seeking to collect botanical material for research purposes should also be aware of regulations that govern intellectual property related to that material. In 1992, the United Nations, through the U.N. Convention on Biodiversity, enacted recognition of traditional knowledge and the natural resources that go with that knowledge. The Convention on Biological Diversity codified enforcement of ecologically sound harvesting practices and sharing of any intellectual or economic benefits that may arise as a result of the utilization of that traditional knowledge.<sup>40-42</sup> Even though the United States did not ratify this treaty, many funding agencies, particularly those of the National Institutes of Health, have chosen to adopt the principles of this convention.<sup>43</sup> In addition, during the 10th conference of parties of the Convention on Biological



Diversity,<sup>43</sup> the Nagoya Protocol was adopted, which is a legally binding protocol for access to genetic resources and benefit sharing.<sup>44</sup> Again, since the United States is not a party to the Convention, this creates a gray area. However, this is another example where journal editors have a great degree of influence, as some journals (for example, *Planta Medica*) now require verification that study materials were acquired in accordance with the Nagoya Protocol, as it is interpreted by the country from where the sample was collected. This latter point is important, as individual countries have interpreted how to handle the Nagoya Protocol differently. Ultimately, collecting a plant specimen is not a trivial matter, especially when done outside the United States, and we recommend working with local authorities and seeking guidance and consultation from individuals with relevant experience before doing so.

#### Box 2 Summary of recommendations: obtaining study material

- When harvesting cultivated or wild botanical specimens, collect additional material for identification and accession to an herbarium.
- Procure sufficient number of commercial products to be representative of the overall market population. In our studies we evaluated a minimum of 30 different products.
- For commercial samples that are dried, powdered, or pre-extracted, obtain additional authenticated reference material for identification.
- Either a commercial source or a contract laboratory can be used to provide botanical natural product study material. The advantage of a commercial source is that the product is representative of what is used by consumers, but care must be taken to ensure that the material is authentic, consistent, and stable.
- Obtain proper permissions before harvesting; collection of samples could be restricted due to local, state, federal, or international laws.

## 6. Authentication

Once a botanical sample (or series of samples) has been collected for analysis, an important step is “authentication,” *i.e.* the definitive determination of the species represented by the sample.<sup>13,45,46</sup> Botanical samples can be authenticated by multiple means, including examination of voucher samples, DNA barcoding, and both untargeted and targeted characterization of chemical constituents. Whenever possible, more than one technique should be applied to confirm the identity of a botanical sample. However, the type of processing that a sample has been subjected to will determine which techniques are effective for authentication, as described in the following section.

### 6.1 Authentication by microscopic and morphological characteristics

Traditional methods used to authenticate botanical samples rely on examining the morphological and microscopic characteristics of representative voucher specimens. For example, macroscopic observations surrounding the arrangement (*e.g.*, alternate, opposite, whorled) and shape (*e.g.*, elliptic, ovate,

cordate) of the leaves, the shape of inflorescence (*e.g.*, cruciform, head, spurred) as well as its type (*e.g.*, raceme, umbel, spike), the classification of the root system (*e.g.*, taproot, fibrous root) can provide insight into the genus and species of the plant material. Similarly, microscopic examination of the histological characteristics of plant parts (*e.g.*, stems, roots and rhizomes, bark, leaves, flowers, seeds, wood) can also aid in positive identification. These techniques, whether independently or in combination, have formed a baseline of approaches for botanical natural product quality control, and they continue to serve as essential methods used in pharmacopeia worldwide.<sup>47</sup> However, accurate microscopic and morphological analysis requires a trained expert, and the robustness of these procedures is also reliant on the presence (and expert knowledge) of diagnostic characteristics that enable one natural product to be separated from another. Challenges arise for closely related species and plant parts that share similar morphological structures yet could have differing biological activities.

Studies conducted using natural product extracts, botanical supplements, and commercial material are subject to several obstacles to authentication. When using commercial suppliers as material sources, voucher specimens are not always readily available. Identification by morphology or microscopy is not suitable for extracted natural product materials, and it may be difficult to determine the identity to the species level. Thus, to address the need for additional, more robust methods, researchers developed complementary techniques of differentiating botanical natural products.

### 6.2 Authentication by DNA barcoding

DNA barcoding is a genetic technique used to identify species using specific differences in small regions of DNA.<sup>48</sup> Barcoding is an effective tool in authentication of botanical natural products that are comprised of fresh, dried, or powdered material, where intact DNA is still present.<sup>46,49,50</sup> However, the DNA barcoding approach is more difficult when applied to botanical extracts. The manufacturing process for botanical natural products may involve heat treatment, oxidation, extraction, distillation, filtration, pressing, encapsulation, spray drying and/or UV light exposure, and these often lead to removal or degradation of DNA. DNA barcoding is not feasible for processed botanical products where the DNA is either not present or potentially highly degraded, or where there are two or more species present (either mixtures of active botanicals or the use of botanical “filler” such as rice).<sup>50</sup> Unfortunately, DNA barcoding is occasionally used to characterize processed botanical natural products without recognition of these limitations. For example, in 2015, the New York State Attorney General investigated potential adulteration of commercial products. This investigation employed DNA barcoding methodology to suggest that only five of 24 samples actually contained the purported botanical natural product,<sup>51</sup> and as a result many of the products were subjected to a forced recall. However, the methods and results of the study were suspect. Among other concerns, there were reports of cross-contamination of the samples, and the investigation focused on botanical extracts, not whole plant



material, such that intact DNA may not have been present even if the correct botanical natural products were used.<sup>52</sup> This controversy may have been avoided with the use of complementary analytical approaches suitable for characterizing extracts (such as described in 6.3 and 6.4).

Although beyond the scope of this review, there are other factors that make DNA barcoding of plants challenging, such as hybridization and polyploidy, slow mutation rates compared with speciation rates, as well as lineages that show rapid and recent divergence rates. In some species of plants, these factors can influence the discriminatory power of standard plant barcoding markers.<sup>53–58</sup> Despite these limitations, the use of two core plant DNA barcode regions *rbcL* (ribulose-bisphosphate carboxylase) and *matK* (maturase K) as well as two supplementary regions, *trnH–psbA* (chloroplast intergeneric spacers) and internal transcribed spacer (ITS) have been used for species discrimination in identifying plants *via* barcoding.<sup>58</sup> Given all these complexities and controversies, it is recommended that DNA barcoding of botanical materials be conducted in consultation with experts.

### 6.3 Authentication by targeted analysis of “marker compounds”

Information about the identities of known small molecule constituents (metabolites) present in a botanical extract (known as “chemotaxonomy”) can be extremely useful in the authentication process. For most botanicals sold commercially as dietary supplements, there is a broad literature base that describes known metabolites and quantities in which they are likely to be present. When these constituents are unique to the botanical of interest, they are often referred to as “marker compounds”. Information about the chemical composition of a given plant is available in the primary peer reviewed literature, and in the form of “monographs” that summarize the chemical content of a particular botanical. Monographs are published by several sources, including the German Commission E,<sup>59</sup> US Pharmacopeia,<sup>60</sup> and *Tyler's Herbs of Choice*.<sup>61</sup> To make use of these monographs, the sample under question is analyzed by a method appropriate for detecting and quantifying the key marker compounds (see Section 9, Identification, structure elucidation, and quantitative analysis of bioactive compounds in the selected study material), and their identities and concentrations are compared with literature values. The most common analytical techniques for analyzing botanical constituents are ultraviolet/visible (UV/VIS) spectrophotometry, mass spectrometry (MS), and Nuclear Magnetic Resonance (NMR) spectroscopy. Because of the complicated nature of botanical samples, it is also typical to perform chromatographic separation prior to analysis by spectrometric approaches. These chromatographic separations are most commonly carried out using gas chromatography (GC), high-performance thin-layer chromatography (HPTLC), high-performance liquid chromatography (HPLC), or a modern advancement on HPLC, termed ultrahigh performance liquid chromatography (UHPLC) prior to analysis by spectrometric approaches. The relative merits of different separation and analysis approaches for botanical natural products have been

previously described.<sup>62–65</sup> It is advised to use multiple approaches to characterize a single sample such that orthogonal data can be collected and compared.

### 6.4 Authentication by “chemical fingerprints”

In addition to specifically analyzing individual marker compounds, researchers seeking to authenticate botanical natural products often collect a “chemical fingerprint” that represents a more comprehensive profile of metabolites produced by the plant. This chemical fingerprint can be compared between two or more samples to assess similarity or difference (Fig. 2).<sup>66</sup> The most common techniques used to measure chemical fingerprints are gas or liquid chromatography, coupled to either spectrophotometric or mass spectrometric detection.<sup>62,65</sup> The resulting data can be viewed as a chromatogram, which displays detector response as a function of time (Fig. 2), such that each “peak” in the chromatogram corresponds to a chemical constituent (or multiple constituents) from the sample. Quantitative NMR (qNMR) is another technique which provides chemical fingerprinting capable of distinguishing species and authenticating botanical origins.<sup>67</sup>

The relative intensities, retention times, and spectroscopic data obtained by chromatographic analysis can be used to putatively identify a sample. Small molecules are more stable than DNA, and they typically survive the processing involved with preparing botanical natural products. Thus, it is possible to authenticate many samples based on their metabolite profiles that may not be characterizable by DNA barcoding. However, it is important to note that the “fingerprint” of which metabolites are present differs depending on the extraction procedure as well as analytical technique used to collect the data and the parameters of the specific analysis.<sup>68</sup> Thus, it is necessary to include authenticated reference material for comparison in the same analysis with the sample under question and processed by the same extraction procedure and solvent system (see Purchasing authenticated reference materials, Section 5.5) (Fig. 2).

### 6.5 Authentication by untargeted metabolomics

Often, it is of interest to compare more than two botanical natural product samples to each other and assess their relative differences and similarities. The technique, currently referred to as “untargeted metabolomics,” can be very useful for this purpose. The “metabolome” is defined as the complete set of small molecules produced by a biological sample. Metabolomics techniques rely on measuring as many of these small molecules as possible (although it is never possible to measure the entire metabolome, due to limitations in analytical methods). Metabolomics has been employed to characterize the relationships between the metabolome of a given natural product and corresponding genotype, origin, quality, or other biotic or abiotic attributes.<sup>28,69–71</sup> Two main analytical techniques are currently employed for metabolomic studies – MS and NMR. The relative advantages of these approaches have been discussed elsewhere.<sup>28,72–74</sup> The ability of untargeted metabolomics analyses to simultaneously profile a large



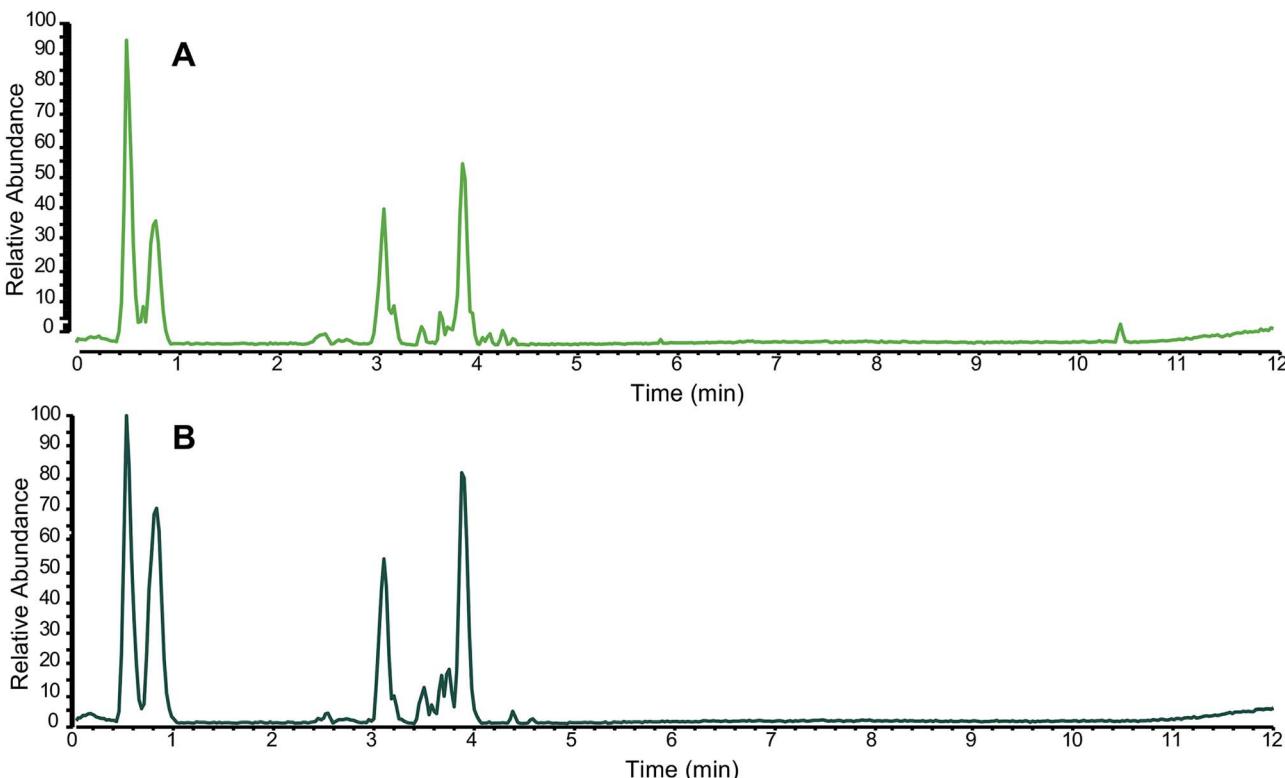


Fig. 2 Representative liquid chromatography-mass spectrometry (LC-MS) fingerprinting of a green tea sample (A) compared against an authentic green tea (*Camellia sinensis*) standard from the National Institute of Standards and Technology (NIST 3254) (B). Both chromatograms were collected in the negative ion mode using ultrahigh performance liquid chromatography (UHPLC) coupled to electrospray ionization mass spectrometry; (reproduced with permission from ref. 28). Note that on balance, the majority of the peaks, both in retention time (x axis) and relative abundance (y axis), match between the two samples.

fraction of the phytochemical make-up of the botanical, and offer detailed characterization of the samples in question has led metabolomics to be a very useful tool in the authentication of botanical natural products.

The process of data collection for a metabolomics analysis is essentially the same as that described for chemical fingerprinting, except that it is typical to analyze scores to hundreds of samples in a single analysis. This need to collect large datasets requires careful attention to experimental design, because runtimes are often long, and drift in instrument response can cause artifacts in the data. Quality control checks throughout the run, as well as attention to sample order, blanks, and replication, are critically important.<sup>75,76</sup>

**6.5.1 Metabolomics data analysis.** The major challenge in metabolomics studies is not in data collection, but in interpreting the very large datasets that are generated. The process known as “chemometrics” refers to the statistical approaches used to analyze metabolomics datasets. Various approaches are used to simplify and compare the data (reviewed in Kellogg *et al.* (2016)).<sup>77</sup> Most common among these is Principal Component Analysis (PCA), in which a dataset is transformed into a set of orthogonal variables (components) that account for the greatest degree of variability in the data. An example of how PCA can be employed to compare botanical samples is shown in Fig. 3. Visual analysis of an untargeted metabolomic PCA scores plot of

various commercial green tea preparations (Fig. 3A) yielded distinct clusters of loose-leaf green tea samples (green symbols), green tea supplements (blue symbols), and a non-green tea sample (orange). Targeted metabolomics, using only “marker compounds” found in green teas, produced less distinct separations (Fig. 3B).<sup>28</sup>

#### Box 3 Summary of recommendations: authentication of natural products

- Authenticate botanical materials to ensure the validity of *in vitro*, non-clinical *in vivo*, and/or clinical studies.
- Select an authentication method that is suitable for the type of specimens being analyzed. For example, extracted botanical supplements cannot readily be identified using DNA barcoding techniques.
- Authentication using only a single technique may be insufficient to ensure accurate identification of the botanical material, and the application of orthogonal methods (*i.e.* targeted analysis of marker compounds plus untargeted metabolomics or DNA barcoding) is recommended.

## 7. Selecting a “representative” sample

Faced with a range of commercial samples purported to have been prepared from a given botanical natural product, how does the researcher select a representative product to advance to *in*



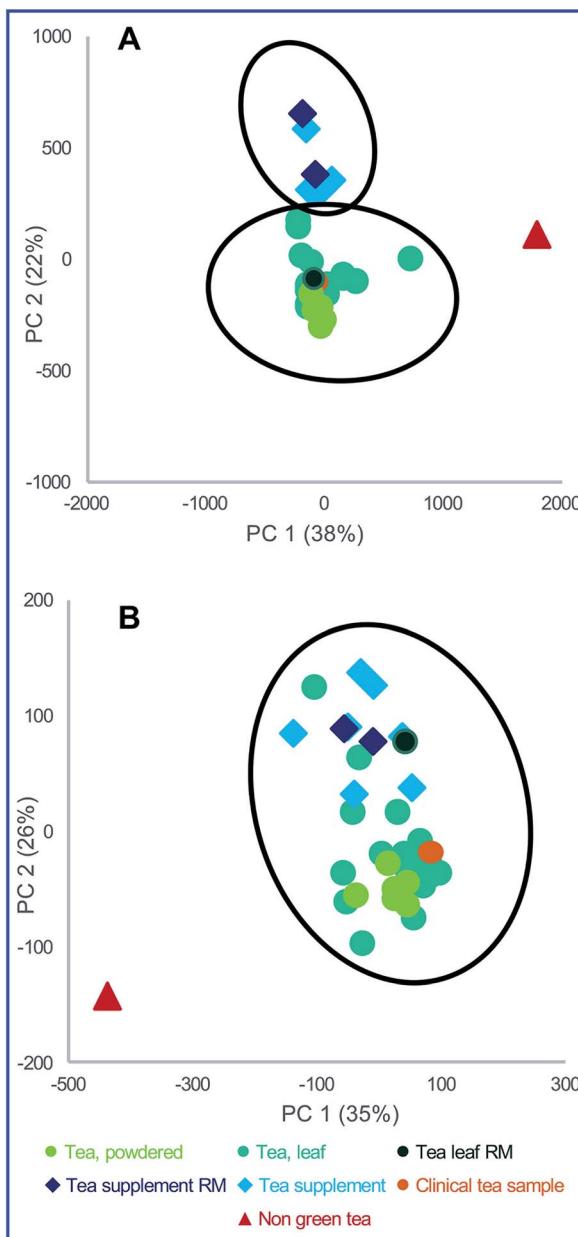


Fig. 3 An example of principal component analysis (PCA) scores plots for green tea samples. Data points represent averaged triplicate extractions and plots are drawn with Hotelling's 95% confidence ellipses. "RM" represents green tea standard reference material from the National Institute of Standards and Technology (NIST). (A) Untargeted metabolomics analysis yielded clusters of distinct sample types (green tea supplements, green teas, and the negative control (indicated as "non-green tea")). (B) Targeted mass spectrometry analysis, using 15 quantified standards, was employed to differentiate between green tea samples and the negative control. However, distinction between green tea supplements and green teas was not as clearly defined, and the RM samples were clustered together, irrespective of their origin. Reproduced with permission from Kellogg *et al.*<sup>28</sup>

*in vitro*, non-clinical *in vivo* or clinical studies? Numerous factors influence the composition of a botanical natural product. For example, with green tea, cultivar, geography, elevation, sun exposure, climate, harvest time, harvest location (where on the

plant the leaves were removed), as well as processing, fermentation, and manufacturing methodologies all led to discernable differences in final phytochemical make-up of the product.<sup>78–83</sup> These details are not readily available to the consumer (or post-harvest researcher); thus, choosing one product out of a multitude of available options constitutes a major challenge. Researchers apply a wide variety of criteria for choosing natural products for further study, including convenience,<sup>84</sup> consumer preference,<sup>28</sup> or a custom prepared extract formulation contracted by the research team.<sup>85</sup> As a result, the natural product interventions that are evaluated may or may not have a resemblance to the products that are currently on the market or used by the greatest numbers of consumers. While it is typically not possible to select a single product that represents all of the products used by consumers, it is worthwhile to capture a "snapshot" of the variability among the pool of products available, such that a desired product can be selected rationally. To conduct such a comparison, it is first necessary to obtain multiple products that represent those commercially available. For example, when the NaPDI Center prepared to conduct studies with green tea (*Camellia sinensis*) and goldenseal (*Hydrastis canadensis*), we purchased 34 and 35 commercially available samples of each botanical, respectively. We obtained fully characterized green tea samples from NIST, and several botanical reference standards of goldenseal from ChromaDex and laboratory collections.<sup>28,29</sup> These products were then all extracted in the same fashion, and their metabolite profiles were compared to those of the reference materials to select a representative product for *in vitro* and clinical studies. Here we describe our selection process in more detail, using green tea as an example. A similar approach has been employed by other laboratories for other studies of botanical natural products.<sup>30,86,87</sup>

### 7.1 Selecting a formulation

Botanical natural products are available to the public in a variety of formulations. Commercially available formulations include tinctures (ethanolic or glycerol extractions of raw plant material), tea bags for making hot water extracts, raw plant material for human consumption, capsules, and tablets. A different constituent profile may be obtained by extracting in alcohol *versus* hot water and by using different plant material (for example, roots *versus* leaves) as starting material. Not all capsules are prepared in the same fashion. Some are made of raw plant material encapsulated in some coating (*i.e.* gelatin), some have a liquid extract directly encapsulated, whereas others are prepared by spray drying an extract on some solid support, such as cellulose. The formulation used can alter the chemical composition of the product. Spray-dried extract capsules usually are more concentrated than capsules prepared from raw plant material and may also be chemically simpler or modified by the extraction process. Because of these differences, we recommend that multiple representative formulations and, if applicable, multiple plant parts, of a given natural product be subjected to chemical evaluation. For example, for studies our group conducted with green tea, both spray-dried and raw plant material

supplements and tea formulations in the 34 products were included in the initial evaluation. Metabolomics analysis demonstrated differences in profiles of the loose-leaf tea *versus* the encapsulated green tea supplements. The PCA loadings plot revealed several metabolites present in higher concentrations in the green tea supplements than the leaf teas: myricetin, kaempferol, and quercetin aglycones, as well as theaflavin 3-O-(3-O-methyl) gallate and the dimer epicatechin (4 $\beta$ →8)-epigallocatechin-3-O-gallate (Fig. 4).<sup>28</sup>

Another consideration when selecting natural product formulation is that the dissolution and absorption of the

bioactive constituents, and by extension human systemic exposure, can differ depending on formulation.<sup>88,89</sup> These factors should be considered when designing *in vitro*, non-clinical *in vivo*, or clinical studies involving a botanical natural product.

Selection of product formulation should be guided by consumer usage. For example, we elected to conduct a clinical study with green tea as a beverage (hot tea) rather than a supplement based on popular use.<sup>16</sup> Similarly, a goldenseal capsule containing dried botanical material was selected because this formulation is most commonly used as a botanical dietary supplement.

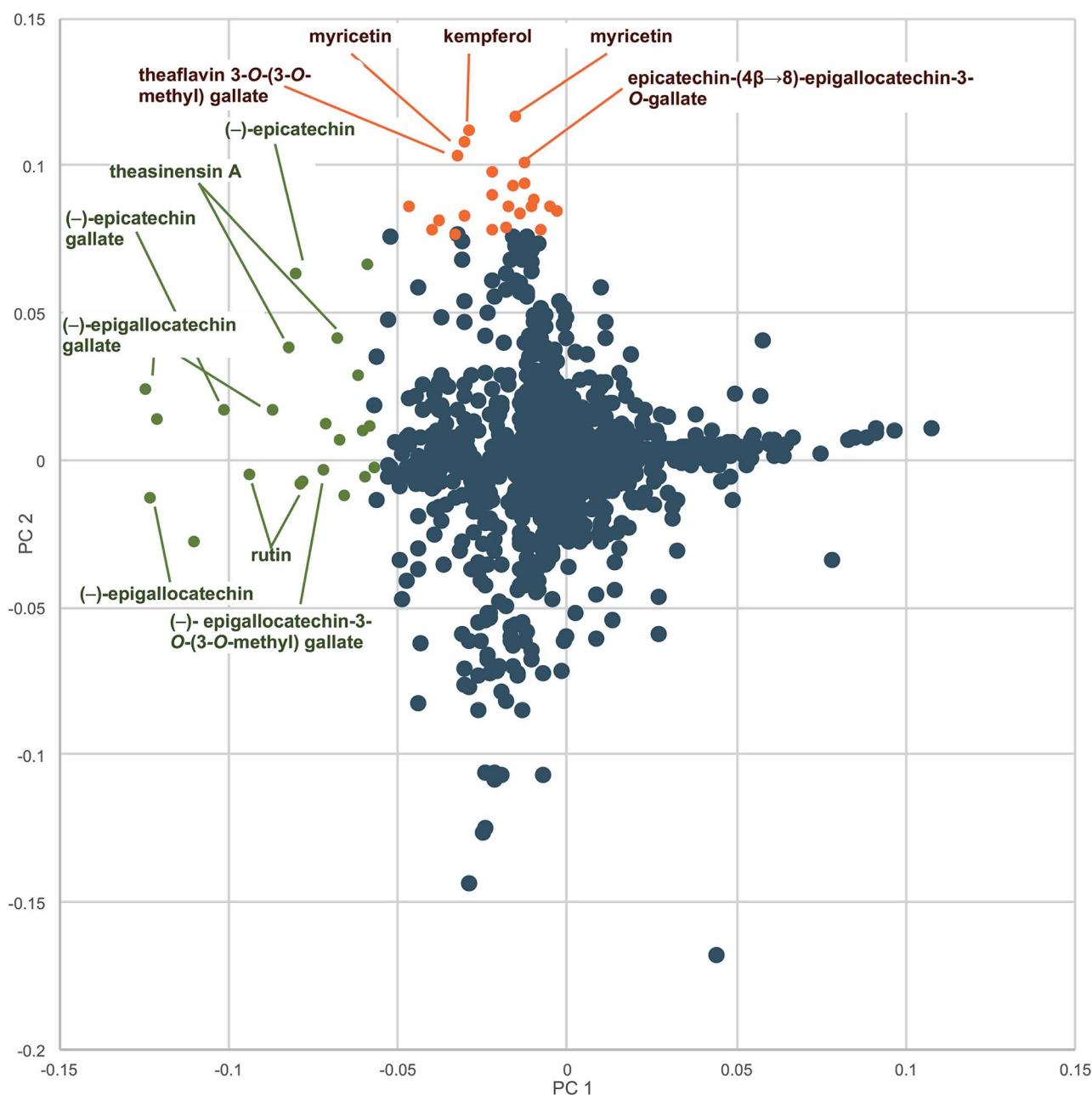


Fig. 4 Loadings plot to identify constituents responsible for differences among commercial preparations of green tea. Metabolites with more negative values along the x-axis (PC1, green labels) were present in higher concentrations in the green tea samples *versus* the non-green tea (negative) control. Labeled metabolites with greater positive values along the y-axis (PC2, brown labels) were more heavily represented in green tea supplement samples compared to green tea leaf and powder samples (reproduced with permission from Kellogg *et al.*)<sup>28</sup>



## 7.2 Standardized products

Some commercially available botanical natural products are sold with no information about chemical makeup, while others are “standardized” to (reportedly) contain a specific amount of a known constituent or constituents.<sup>27</sup> An advantage of standardization is that researchers, health care practitioners, and consumers are provided information that may be useful for selecting and adjusting dosage.<sup>26</sup> The standardization process may also provide consistency across products, with the caveat that methods used for standardization must be reproducible across multiple labs.<sup>90,91</sup> Standardization can involve multiple analytical methods, including qNMR,<sup>92</sup> UV-VIS absorbance patterns,<sup>90,93</sup> and MS profiling.<sup>94</sup> Even for standardized products, label claims about content and identity of a natural product should be confirmed in-house or by a contract laboratory prior to conducting *in vitro*, non-clinical *in vivo*, and clinical studies.

A disadvantage of standardization is that the focus shifts from the quality of the entire mixture to a single or limited set of constituents. Standardized extracts may be chemically simpler than non-standardized counterparts, because the process of enhancing the concentration of constituents may involve extraction procedures that reduce extract complexity. Reduced extract complexity can be a disadvantage when the true “bioactive” constituent is not known, such that standardizing to one constituent may result in a lower dosage (or complete lack) of a constituent responsible for the presumed biological effect. How does a researcher decide whether to use a standardized product for an *in vitro*, non-clinical *in vivo*, or clinical study? One consideration is whether the most popular product(s) used by consumers are standardized. It is also advisable to compare metabolite profiles of standardized products to those of unstandardized counterparts to determine if components contributing to biological activity might have been lost during the standardization process (see Section 9.1).

## 7.3 Selection of extraction solvent

Once a series of representative products and authenticated standards has been selected, the first step in chemical analysis is to prepare extracts from these products. Typical methods of extracting botanical natural products for human consumption include the use of aqueous ethanol or aqueous glycerin (to prepare tinctures), or hot water (to prepare a tea). Applying these same extraction procedures when preparing samples for chemical analysis may seem prudent. However, water, ethanol and glycerin extracts are difficult to manipulate in a laboratory setting. Drying botanical natural product extracts (removing solvents) for long term storage is ideal because extracts are most stable and least likely to degrade when solvent is not present.<sup>95,96</sup> Removing ethanol is a challenging process, removing water is more so, and removing glycerin is most arduous. Additionally, water extracts are unstable and contain polar constituents that confound chemical analysis and result in sticky, tar-like samples.

Unlike the aforementioned solvents, methanol is an outstanding solvent for preparing botanical natural product

extracts prior to analysis. Methanol has an intermediate polarity that effectively solubilizes a wide range of structurally diverse small molecules, is inexpensive, and can be rapidly and efficiently removed under vacuum or a nitrogen stream. As such, many natural product extracts are prepared in methanol or similar organic solvents.<sup>77,97,98</sup> A disadvantage of these solvents is that they are unsafe for human consumption, raising concern that the resulting extracts are not representative of consumer use. While characterizing various green tea products prior to clinical evaluation, we addressed this concern by comparing hot water extraction (an obvious choice for tea) to methanol extraction. Methanol and hot water extracts contained similar quantities of catechins<sup>28</sup> and showed qualitatively similar biological activity *in vitro*. Based on these results, methanol extraction was selected for metabolomics analysis of green tea products. Because methanol enabled rapid and reproducible extraction of samples, the extraction process did not confound comparisons of the chemical composition of the products themselves. A follow up study showed that accelerated solvent extraction improved extraction efficiency of green tea components in methanol.<sup>99</sup> An alternative to methanol extraction includes “comprehensive” extraction techniques, with a gradient of solvents producing a variety of crude extracts with differing polarities.<sup>100</sup> Ultimately, the ideal extraction solvent will depend on the type of material being extracted, the methods used to prepare preparations for consumer use, and the study question. Extraction solvent choice should be carefully considered and justified based both on experimental data and practical considerations. As the extracts prepared for research purposes often are not intended for human use, practical considerations of component solvation, ease of evaporation, and other physiochemical properties of the solvent may outweigh concerns about whether the extraction solvent is identical to that used to produce food grade products.

### Box 4 Summary of recommendations: selecting a botanical natural product sample

- Acquire multiple representative examples of different preparations and/or plant parts.
- Use industry and consumer data to guide selection of products that are representative of the consumption patterns of the population of interest.
- Consider whether a standardized product is appropriate for a given application; there are advantages and disadvantages of standardization.
- Select an extraction solvent based on the research question, being mindful that there are many advantages to methanol extraction for stability, ease, and repeatability of laboratory work.

## 8. Quality control and comparison of botanical product composition

Targeted chemical analysis of known markers and untargeted metabolomics to authenticate botanical natural product samples can be used to characterize and compare a series of candidate botanical natural products prior to selecting a product for *in vitro*, non-clinical *in vivo*, and/or clinical study.



The advantage of untargeted chemical analysis for this purpose is that the analysis is relatively simple, can be highly sensitive and selective (leading to low limits of detection) and can be quantitative (when relevant reference standards are available; see Section 9.2). The disadvantage of targeted analysis for comparing samples is that the differences between samples may not be due to known “marker compounds”. For example, in green tea studies conducted by the NaPDI Center, samples of tea leaves or powder were shown to differ chemically from green tea supplements when untargeted metabolomics was applied, as indicated by the distinct grouping in a PCA plot (Fig. 3). However, when a PCA plot was generated using data from targeted analysis of 15 known green tea constituents, the differences among samples were less clear. As evidence of this lack of clear distinction, the points in the PCA plot in Fig. 3B representing supplement and tea samples overlap, suggesting (incorrectly) that the chemical makeup of the different samples is indistinguishable.

### 8.1 Quantitative metrics for comparing similarity of natural product mixture

Untargeted metabolomics approaches comparing botanical natural products yield a tremendous quantity of data. Multivariate statistical techniques (e.g., PCA and partial least squares discriminate analysis (PLS-DA)) have become routine analytical

methods used to analyze these data<sup>101</sup> and serve as the backbone of metabolomic statistical analyses. PCA enables visualization of similarity and differences among samples by plotting them as points in 3D or 2D space (Fig. 3).<sup>102</sup> With appropriate validation, PCA serves as an effective approach to compare qualitative differences among botanical samples.<sup>103,104</sup> Other approaches, such as cosine calculations, measures based on vector distance, and measures based upon probabilistic difference, are used to generate quantitative metrics to compare chemometric or metabolomic spectral datasets.<sup>105–108</sup> Recently, to facilitate selection of a green tea product for *in vitro* and clinical studies, we developed a new approach for quantitative comparison among samples in a metabolomics dataset, termed the composite score (CS).<sup>28</sup> The CS is based on the scores and loadings of the totality of a multivariate PCA model, rather than pairwise comparisons of components (as in traditional PCA scores plots). The scores and loadings are integrated into a single correlation matrix, which can be used to compare similarity between multiple samples (Fig. 5). The CS ranges from  $-1.00$  to  $1.00$ . Values close to  $1.00$  indicate similarity among two samples (Fig. 5B), whereas values further from  $1.00$  indicate more dissimilar samples (Fig. 5C). The CS approach was used to evaluate which commercial sample was most similar to a given standard reference material.<sup>28,29,99</sup> The PCA plot (Fig. 3A) from the green tea study yielded distinct

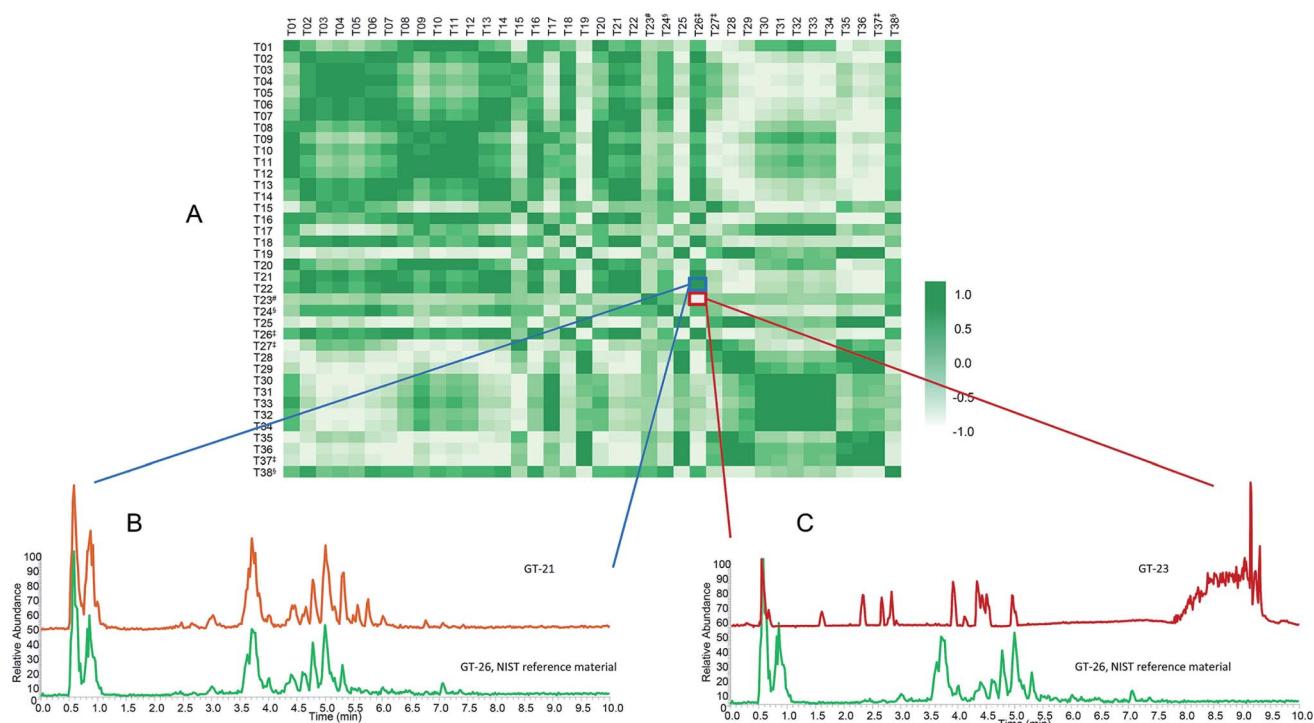


Fig. 5 Composite score (CS) analysis for green tea samples. (A) Heat map correlation matrix of all green tea samples (indicated by codes on the horizontal and vertical axes). CS was calculated from the reproduced correlation coefficient matrix comprised of a four-component PCA model. In the heat map, darker shades (values closer to 1.00) represent stronger correlation between samples, while lighter shades (values approaching  $-1.00$ ) represent decreased correlation between samples. The correlations' relationship to the metabolome are represented in two examples. (B) High correlation between two samples (CS = 0.98, dark color) indicates a strong similarity of the metabolomic profiles. (C) Low correlation (CS =  $-0.85$  light color) indicates two samples with diverging metabolomic profiles. #Negative control (T23); †NIST standard reference materials (T26, T27, and T37); ‡green teas with botanical additives (T24 and T38).



clusterings of green tea supplements *versus* loose leaf green tea products (teas and powders). However, resolution was lacking to discern which specific tea was most similar to the chosen standard, NIST T26; indeed, a different answer regarding which teas are most similar would be obtained depending on which two principal components were plotted against each other (PC1 *versus* PC2, PC2 *versus* PC3, *etc.*). CS analysis (Fig. 5) was used to evaluate which products were most similar to the reference, as well as to select a commercial product (T21) for *in vitro* and clinical studies.<sup>28</sup>

## 8.2 Adulteration of botanical natural products

Several reports have described suppliers of natural products selling products that are either unintentionally misidentified or intentionally adulterated (contaminated with botanical or other material different from that listed on the label).<sup>64,87,109</sup> Adulteration can occur in a variety of ways, including spiking synthetic compounds into the plant material or adding or substituting a different (less expensive) species, representing cost-savings to a dishonest supplier.<sup>110</sup> When studies are unknowingly conducted with adulterated or misidentified plant material, results may be irreproducible in future studies or irrelevant to the botanical supposedly under evaluation. Thus, testing botanical natural products for adulteration is critical.

Several research groups have used targeted<sup>64,87,90,111–113</sup> or untargeted methods<sup>114–116</sup> to identify adulterants in botanical natural products. For example, we studied 35 commercial products listed as “goldenseal”. PCA of untargeted metabolomics revealed a distinct grouping of goldenseal material, including both root/rhizome- and leaf-based samples (Fig. 6), yet three samples (GS-07, GS-20, and GS-33) were distinctly separated from the other clusters.<sup>29</sup> Separation of these samples from the others in the PCA scores plot raised suspicion that they were misidentified or adulterated with other botanical products.

To tentatively identify the species present in the outlier materials, reference material for three non-goldenseal species that serve as common adulterants were included in the dataset: *Coptis chinensis* rhizome (GS-39) and root (GS-40), *Mahonia aquifolium* leaf (GS-41) and root (GS-42), and *Berberis vulgaris* root (GS-43).<sup>110,117,118</sup> The resulting PCA scores plot (Fig. 6B) showed close clustering of GS-33 with *C. chinensis*, GS-20 with *M. aquifolium*, and GS-07 in the middle, possibly as a blend of multiple species. Examining specific marker compounds for each of the four species (berberine,  $m/z$  336.1229 [M]<sup>+</sup>; hydrastine,  $m/z$  384.1440 [M + H]<sup>+</sup>; canadine,  $m/z$  340.1545 [M + H]<sup>+</sup>; palmatine,  $m/z$  352.1543 [M]<sup>+</sup>; coptisine  $m/z$  320.0917 [M]<sup>+</sup>; and dihydrocoptisine,  $m/z$  322.1074 [M]<sup>+</sup>) revealed specific concentration patterns (Fig. 7) for the putative adulterants that supported the hypotheses generated by the metabolomic analysis.<sup>29</sup> This example illustrates the usefulness of both untargeted and targeted methods for identifying adulterated samples. PCA plots generated from untargeted metabolomics can be used to give a broad sweep analysis of the data and identify potential outliers, and follow up targeted analyses can help verify that adulteration has occurred and, if sufficient literature precedent exists, determine the identity of adulterants.<sup>29</sup>

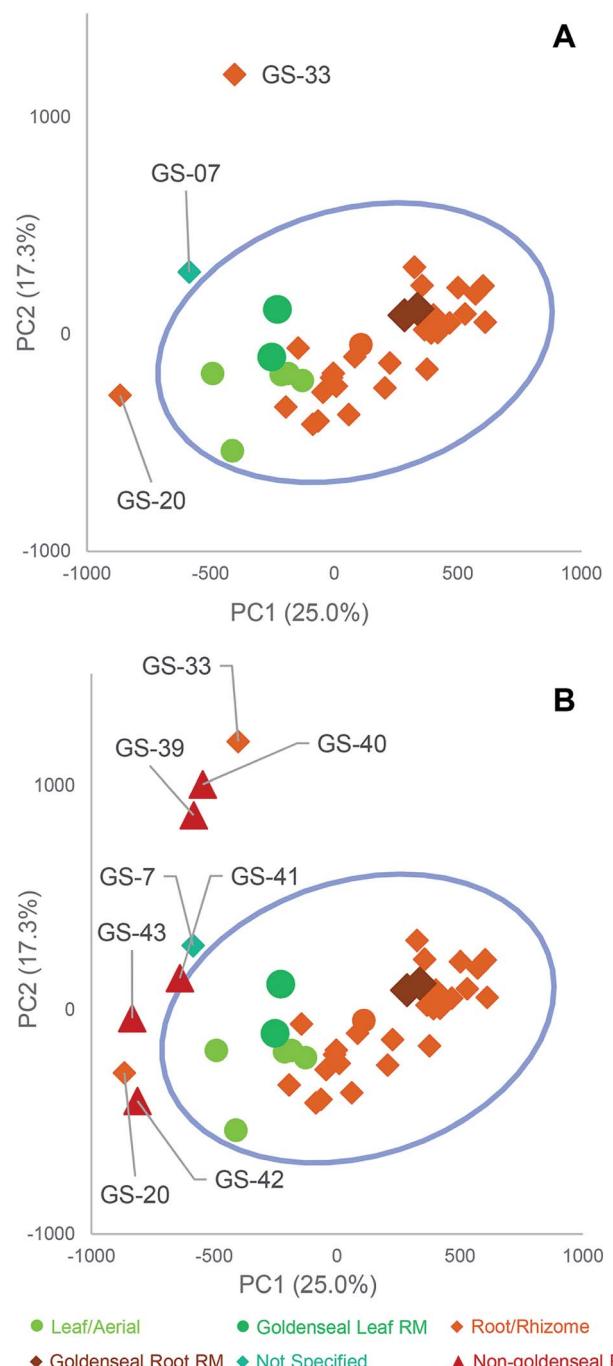


Fig. 6 Principal component analysis (PCA) scores plots from untargeted mass spectrometry metabolomics analysis of commercial goldenseal product samples, plotted with Hotelling's 95% confidence interval (as shown by blue circle). (A) PC1 versus PC2 (25.0% and 17.3% explained variance, respectively) allowed for visualization of the samples (aerial portions and root/rhizome portions), with corresponding reference material (RM) located within the main cluster. The three samples labeled GS-07, GS-20, and GS-33 are located distinctly from other goldenseal samples and were considered outliers. (B) Follow-up metabolomics profiling and PCA scores plot including non-goldenseal reference materials (*Coptis chinensis*, GS-39 and GS-40; *Mahonia aquifolium* GS-41 and GS-42; *Berberis vulgaris*, GS-43) highlighted the potential sources of adulteration in the three outlier samples. Reproduced with permission from Wallace *et al.* (2018).<sup>29</sup>



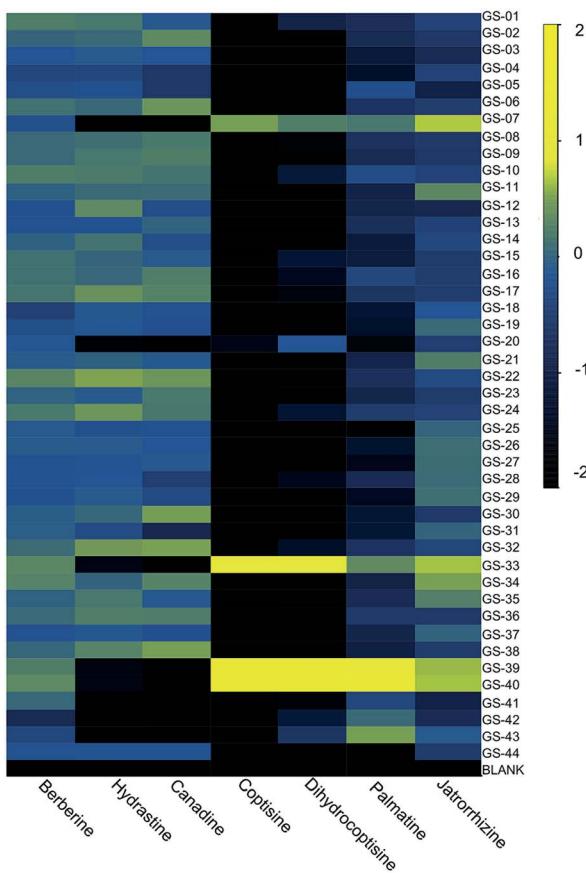


Fig. 7 Heatmap of purported adulterants: six primary alkaloids present in *H. canadensis*, *C. chinensis*, *M. aquifolium*, and *B. vulgaris*. Heatmap was generated using the  $\log_{10}$  of the peak area for the relevant ion detected by high resolution LC-MS. Reproduced with permission from Wallace *et al.*<sup>29</sup>

### 8.3 Consideration of contamination

A potential risk with botanical natural products is contamination by heavy metals,<sup>119</sup> pesticides and herbicides,<sup>120</sup> residual solvents,<sup>121</sup> fungal spores and mycotoxins,<sup>122</sup> or microbes.<sup>123</sup> These concerns have received considerable attention, and recent reviews of the risks, policies, and analyses of contaminants in botanical natural products are available.<sup>27,31,112,124</sup> In 2010, the FDA instituted the requirement that botanical natural products must be prepared using good manufacturing practices (GMP) that provide guidance for safety, consistency, and reproducibility of these products.<sup>21,125</sup> GMP producers certify that their products are consistently produced and controlled according to quality standards. Whether the product is custom formulated<sup>21</sup> or obtained from commercial sources, testing is required for contaminants, including “heavy metals, microbial limits, residual pesticides, adventitious toxins (e.g., aflatoxins), [and] endogenous toxins (e.g., pyrrolizidine alkaloids)” according to 21 CFR 111.70(b)(3). Product batches that do not meet the limits of contamination must be rejected.<sup>125,126</sup> FDA guidance also specifies that at least one test must be used to verify botanical ingredients using a validated method.<sup>112</sup> However, because the manufacturer is responsible for developing GMP

procedures for each botanical natural product, there is no standardized procedure across the industry. In addition, compliance enforcement is limited and does not require evaluation or verification of authenticity or screening for contaminants by a third party.<sup>124</sup> Given these limitations, testing botanical natural product study materials for contamination by pesticides, heavy metals, microorganisms, and residual solvents is advised. Such tests can be accomplished in house or through a contract laboratory. Screening is done in a targeted fashion, *i.e.* the analyst specifies a list of common or likely contaminants *a priori* and analyzes the sample to determine levels of these contaminants. The challenge with this approach is selecting which contaminants to test for among the almost unlimited number of potentials. Selection of contaminants can be guided by consultation with the literature or based upon knowledge regarding the environment in which the sample was grown and the processing it was subjected to post-harvest. Nonetheless, it is easy to imagine that some contaminants may be unexpected and, therefore, overlooked. Untargeted metabolomics may serve as a useful tool for identifying unexpected contaminants.<sup>29,111,115,127</sup> However, it should be noted that untargeted methods are likely to have higher limits of detection than targeted methods; thus, the former may fail to detect low-level contaminants. Furthermore, some contaminants are detectable only with specialized analytical methods. For example, GC-MS is appropriate for the analysis of certain pesticides and residual solvents, but does not respond to many non-volatile contaminants, and a specialized system such as an inductively coupled plasma mass spectrometer (ICP-MS) is needed to detect heavy metal contaminants. With these challenges in mind, the analyst should carefully consider what the most likely contaminants are in a given sample and apply multiple analytical techniques and a combination of both targeted and untargeted methods to maximize the likelihood of detecting as many contaminants as possible.

#### Box 5 Summary of recommendations: quality control and comparison of botanical material

- PCA is an effective method to compare qualitative differences among botanical samples.
- The composite score (CS) represents a useful statistical approach for comparing the similarity or dissimilarity of multiple samples in large metabolomics datasets.
- Untargeted metabolomics combined with targeted methods can effectively discern potential adulterations in botanical natural products.
- Contamination testing is recommended prior to conducting studies involving botanical natural products.

## 9. Identification, structure elucidation, and quantitative Analysis of bioactive compounds in the selected study material

Once a natural product has been selected for *in vitro*, non-clinical *in vivo*, or clinical studies, it should be thoroughly



characterized, and concentrations and identities of key bioactive constituents (*i.e.*, those responsible for the reported biological activity) should be determined. Such knowledge can be used to adjust dosage and conduct quality control. The challenge is that identities of bioactive constituents relevant to the activity being studied are often not known and are likely to differ depending on the biological activity evaluated.<sup>62,128,129</sup> Assigning bioactive constituents is not trivial, as a given botanical extract may contain hundreds or thousands of constituents.<sup>130</sup> The collective activity of the extract may be due to the combined action of these constituents acting in an additive, synergistic, or antagonistic manner.<sup>131–135</sup> Despite these challenges, focusing solely on one or two constituents in an extract is not prudent (discussed in Sections 6.3–6.5). For some of the well characterized botanicals, nearly complete characterization of major constituents can sometimes be possible. For example, in our work with green tea, 15 commercially available pure standards were obtained that represented major and minor peaks in a LC-MS chromatogram (Fig. 2). The sections below highlight key points to consider when conducting qualitative and quantitative analysis of botanical extracts and assigning biologically active components. Based on practical experience, a summary of common pitfalls during quantitative and qualitative analysis of botanical extracts is provided (Table 2).

### 9.1 Determining which constituents of a botanical natural product are biologically active

The gold standard approach for identifying bioactive mixture components is bioassay-guided fractionation, in which the extract is subjected to successive rounds of fractionation and purification (see Section 9.2), with each fraction prioritized for the next stage based on biological assay data. Such an approach has been used in countless studies, with perhaps the most well-known being discovery of the cancer chemotherapeutic agent taxol (aka paclitaxel) from the Pacific yew tree.<sup>136,137</sup> An alternate approach to bioassay-guided fractionation when targets are known and available in sufficient quantity is pulsed ultrafiltration mass spectrometry,<sup>138,139</sup> the advantages and disadvantages of which are detailed elsewhere.<sup>140</sup> In general, we find that bioassay-guided fractionation is the most effective strategy for identifying bioactive mixture components when the specific targets and mechanism of action of a given botanical natural product are not known, and when those activities can be modeled effectively *in vitro*. Bioassay-guided fractionation can be enhanced using statistical approaches to integrate biological assay data with chemical metabolite (metabolomics) profiles, a process often referred to as ‘biochemometrics’.<sup>77,141</sup>

**9.1.1 Advantages and limitations of *in vitro* assays.** A critical element of bioassay-guided fractionation is biological evaluation. Biological activity of a given botanical extract constituent is typically assessed using an *in vitro* assay.<sup>21,142,143</sup> Isolated compounds with promising activity may eventually be tested in non-clinical (animal) models and potentially in clinical studies. However, it is often infeasible to conduct such *in vivo* studies with the speed and scale necessary to facilitate

bioassay-guided fractionation; thus, fractionation and isolation efforts are typically guided by *in vitro* evaluation.

*In vitro* assays are used to assess the potential for natural product constituents as lead compounds for a diverse array of illnesses, including cancer, bacterial infections, and diabetes.<sup>144–146</sup> Cell-based assays can be used to guide future *in vivo* studies to test for safety and efficacy of natural products,<sup>147</sup> to give insight regarding potential interactions between natural products and conventional drugs,<sup>148–150</sup> and to predict how food matrices could affect the pharmacokinetics of natural product constituents.<sup>151</sup> For studies evaluating interactions between natural products and conventional drugs, mechanistic insight gained from *in vitro* studies is imperative to guide further evaluation *via* mathematical (*e.g.*, physiologically-based pharmacokinetic) modeling or clinical studies.<sup>6</sup> However, *in vitro* assay data may not always translate to *in vivo* activity, partially due to the inherent complexity of biological systems, and the effects of other biological factors (*e.g.*, microbiota) that are not accounted for in most *in vitro* models.<sup>152</sup> Additionally, many botanical natural products are used for purported health benefits that are non-specific and difficult to model *in vitro*, such as “improving immune health” or “enhancing adaptability to stress”. In light of these challenges, it is important to keep in mind that the extent to which any bioassay-guided fractionation experiment is useful is fundamentally limited by the availability of a relevant, robust and translatable biological assay.

**9.1.2 Approaches for integrating *in vitro* data with chemical composition.** Visual inspection of chromatographic data is often used to guide isolation efforts as part of bioassay-guided fractionation experiments. Limitations of this approach include a tendency to focus on constituents that are most abundant, most responsive to the detector, or most easily isolated. In addition, re-isolation of known active compounds has plagued traditional bioassay-guided fractionation experiments. This problem of re-isolation can be addressed using various “dereplication” approaches, which often rely on comparing experimentally measured spectroscopic data with databases for known compounds.<sup>158–161</sup> When the active constituents of a botanical natural product are not known, several methods are available to correlate metabolite profiles with bioactivity data and guide isolation of constituents most likely to be active. Use of statistical approaches to integrate biological assay data with measurements of chemical composition is termed “biochemometrics”.<sup>141</sup> Partial least-squares (PLS) modeling and associated variants (partial least-squares discriminate analysis (PLS-DA) and orthogonal partial least-squares discriminant analysis (OPLS-DA)) have become major statistical methods for biochemometric analysis.<sup>162–164</sup> PLS modeling has been adapted to identify individual (or several) metabolites that are predicted to be responsible for the bioactivity of a complex natural product mixture.<sup>77,165,166</sup> For the NaPDI Center study of green tea, a biochemometric approach was used to predict which catechins were responsible for the *in vitro* inhibition of intestinal UDP-glucuronosyltransferases (UGTs). The selectivity ratio<sup>167–169</sup> was used as a metric to demonstrate the extent to which a given mixture constituent was associated with biological activity.<sup>170</sup> Five catechins were identified as major



**Table 2** Common pitfalls in quantitative and qualitative interpretation of data from botanical extracts**(1) Assuming that relative peak intensities in a chromatogram represent relative concentrations in the sample**

Do not assume that a large peak means a high concentration of analyte in an LC-MS or LC-UV chromatogram. Instead, recognize that they reflect the responsiveness of the analyte to the detector, *i.e.* its ionization efficiency (for MS) or its molar absorptivity (UV). One approach to circumvent this problem is to use an evaporative light scattering detector (ELSD) or charged aerosol detector (CAD).<sup>153,154</sup> The response of these detectors closely reflects relative abundance of analytes in the sample, but they are less sensitive than MS or UV

**(2) Attempting to quantify analytes without reference standards**

Because response of the detector (mass spectrometer or spectrophotometer) depends on structure, an authentic standard should be used to quantify the unknown of interest

**(3) Assuming that the absence of a constituent in the data means an absence of the compound in the sample**

Failure to observe a compound in the data means only that it was not detectable by the technique used above a specific stated limit of detection. It does not mean that the analyte was absent from the sample. Including a positive control can help address this problem. If the analyte of interest is detectable in the positive control but not in the sample, the analyte likely is absent from the sample (above the limit of detection)

**(4) Overstating the accuracy to which a concentration is known**

Absolute quantitation (knowledge of the true amount of a given compound in a botanical sample) is difficult to achieve. Factors such as inefficient extraction, lack of purity in the standard, and drift in instrument response make knowledge of true absolute concentration extremely difficult to assess. Avoid chasing perfect and absolute quantitation when relative or approximate quantitation will be sufficient. On the other hand, knowing that most quantitative measures are not as absolute as the reports may imply, avoid the pitfall of comparing samples from different runs or different laboratories when the results may be different due to factors inherent in the analysis. Include appropriate replicates, references and controls in each experiment such that comparison among experiments is possible. Report the concentration of a given analyte with its associated uncertainty and to the correct significant figures

**(5) Assuming confirmation of identity with MS data**

Even if appropriate standards are used to match retention time and fragmentation patterns of unknown compounds in a sample, mass spectrometry data does not confirm configuration of stereoisomers. NMR is needed for such confirmations. However, depending on the needs of the study, knowledge of configuration of stereoisomers may not be necessary, and MS data may be sufficient

**(6) Extrapolating results beyond the linear range of the calibration curve**

Always dilute samples so that the analyte of interest falls within the linear range of the calibration curve. Failure to do so may result in serious underestimation of analyte concentration. Errors also result from attempting to quantify analytes present at too low of a concentration, *i.e.* below the limit of quantification for the analytical method

**(7) Failing to account for matrix interference**

Matrix interference occurs when the matrix (everything but the analyte in the sample) alters the response of the analyte. This issue can be particularly pronounced with mass spectrometry as an analytical method. The best strategy for avoiding matrix interference is to dilute the sample as much as possible and subject it to chromatographic

**Table 2 (Contd.)**

separation prior to analysis. A validation check that involves spiking the sample with a standard and comparing its response in matrix and solvent is necessary to check for matrix interference<sup>95</sup>

**(8) Assuming a sample is “pure” based on LC-UV data**

When interpreting LC-UV data, it is important to remember that contaminants will not be detectable unless they absorb light in the region used for the analysis. Quantitative NMR represents an alternate method to determine purity<sup>155</sup>

**(9) Assuming that each peak in an LC-UV chromatogram represents a single compound**

Multiple compounds in a complex sample may coelute in what appear as a single peak. Thus, one peak does not necessarily mean one compound. MS detection can be used to identify coeluting compounds if they differ in mass. Examination of UV spectra at multiple retention times across a single chromatographic peak can also aid in detecting coeluting compounds

**(10) Assuming that each peak in the mass spectrum represents a different ion**

Clustering and in-source fragmentation in mass spectrometric analysis often lead to multiple masses that represent a single ion. Identifying the true “molecular ion” can be difficult. Software packages such as RamClust<sup>156</sup> and IntelliXtract<sup>157</sup> can help assign identities of clusters and fragments and group associated ions

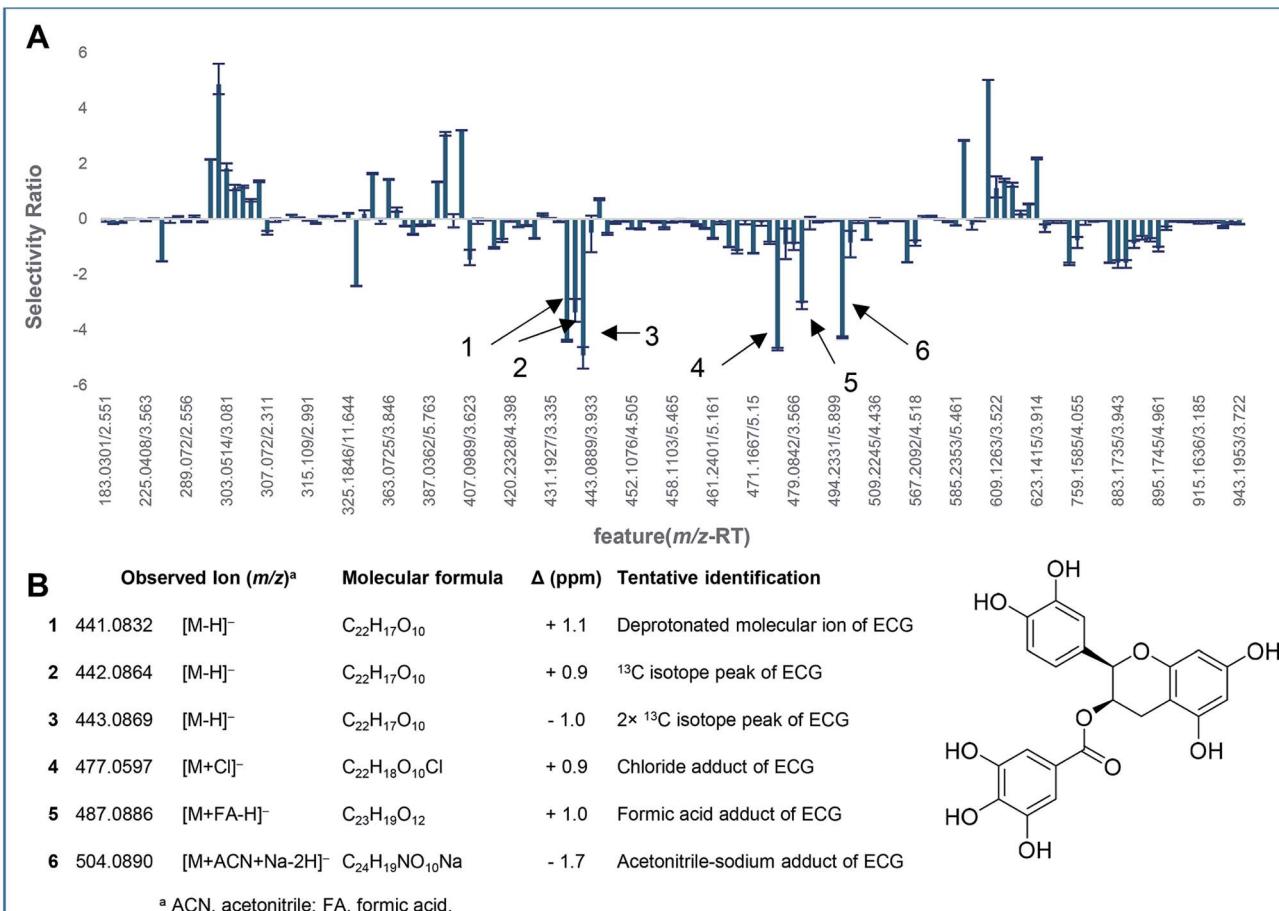
constituents in a selected bioactive subfraction. Selectivity ratio analysis predicted (–)-epicatechin gallate (ECG) to be a major contributor to inhibition of UGTs (Fig. 8). This prediction was confirmed with follow up *in vitro* studies using commercially available catechins and the clinically relevant intestinal UGT substrate raloxifene.<sup>170</sup>

## 9.2 Obtaining “pure” standards

Efforts to identify and quantify constituents of botanical extracts often rely on the availability of purified standards. Several commercial suppliers provide pure samples of individual botanical compounds and include Sigma-Aldrich (St. Louis, MO, USA), ChromaDex (Irvine, CA, USA), NIST (Gaithersburg, MD, USA), and the United States Pharmacopeia (Rockville, MD, USA). The modest cost of these standards typically far outweighs the time and effort involved in obtaining standards *via* independent isolation methods. A standard comes with a “certificate of analysis” that reports identity and purity of the sample. Additional characterization (by LC-MS and/or NMR) is recommended to confirm accuracy of information included in the certificate of analysis.<sup>155</sup>

When standards of the compounds of interest are not available commercially, isolating these compounds from the complex starting material is necessitated.<sup>171</sup> A number of separation techniques can be used for this purpose, including solid phase extraction,<sup>172</sup> counter current chromatography,<sup>173,174</sup> HPLC,<sup>175</sup> or thin layer chromatography (TLC).<sup>176</sup> The most common strategy for isolation of constituents from natural products is a combination of liquid–liquid partitioning and column chromatography approaches. A typical fractionation





**Fig. 8** Biochemometric analysis of the inhibition of UDP-glucuronosyltransferase (UGT) activity by green tea constituents. Mass spectrometry-based metabolomics was correlated with bioactivity to generate a partial least squares (PLS) model for eight green tea subfractions. This model was used to determine the selectivity ratio for each feature detected in the green tea extract (A), the x-axis represents sample components detected as unique  $m/z$ -retention time (TR) pairs (features) in the extract. The more negative values on the y-axis represent more significant contribution to the observed UGT inhibitory activity. All six detected ions (spectral variables) represent isotope peaks and cluster ions of ( $-$ )-epicatechin gallate (ECG) (B) molecular ion, suggesting that ECG is the dominant bioactive ion in the green tea matrix. Adapted with permission from Tian et al. (2018).<sup>170</sup>

scheme involves several steps, as illustrated in Fig. 9. The sample is extracted in methanol, partitioned between chloroform and water or ethyl acetate/water, and subsequently subjected to an additional partitioning step between methanol : acetonitrile and hexanes.<sup>177</sup> These steps could be modified using acid/base conditions if the target metabolites are alkaloids. The resulting residue is dried under nitrogen and subjected to several stages of normal phase flash chromatography. Finally, pure compounds are isolated using reversed phase preparative HPLC. The solvents and stationary phases used for the separation can be varied to optimize separation of the compound of interest. Isolation of milligrams of pure material often takes weeks or months (or may be unsuccessful) depending on the difficulty of the separation and the skill of the analyst.<sup>178</sup> As such, isolation of all constituents in a given botanical extract is typically unfeasible, and it is helpful to guide isolation efforts with biological assay data, as described in Section 9.1.

### 9.3 Structure elucidation of constituents of botanical natural products

Identification of metabolites in botanical extracts has been reviewed previously,<sup>179,180</sup> and structure elucidation of natural product constituents is detailed in several textbooks.<sup>181,182</sup> Two approaches are commonly used: isolation and subsequent structure elucidation *via* NMR or characterization of mixture constituents without isolation *via* LC-MS. Conclusive assignments of structure are made *via* NMR and it is the preferred technique when sufficient material (typically 0.5 to 1 mg) has been isolated.<sup>183</sup> Structure elucidation *via* NMR is conducted in combination with other spectroscopic approaches, notably high resolution MS, which enables determination of molecular formula.<sup>180</sup> Accepted guidelines from peer-reviewed journals (e.g., *Journal of Natural Products*) recommend that assigned  $^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra be presented for review, as well as molecular formulae calculated by relying on accurate mass measurements with MS, UV absorptivity, and any additional

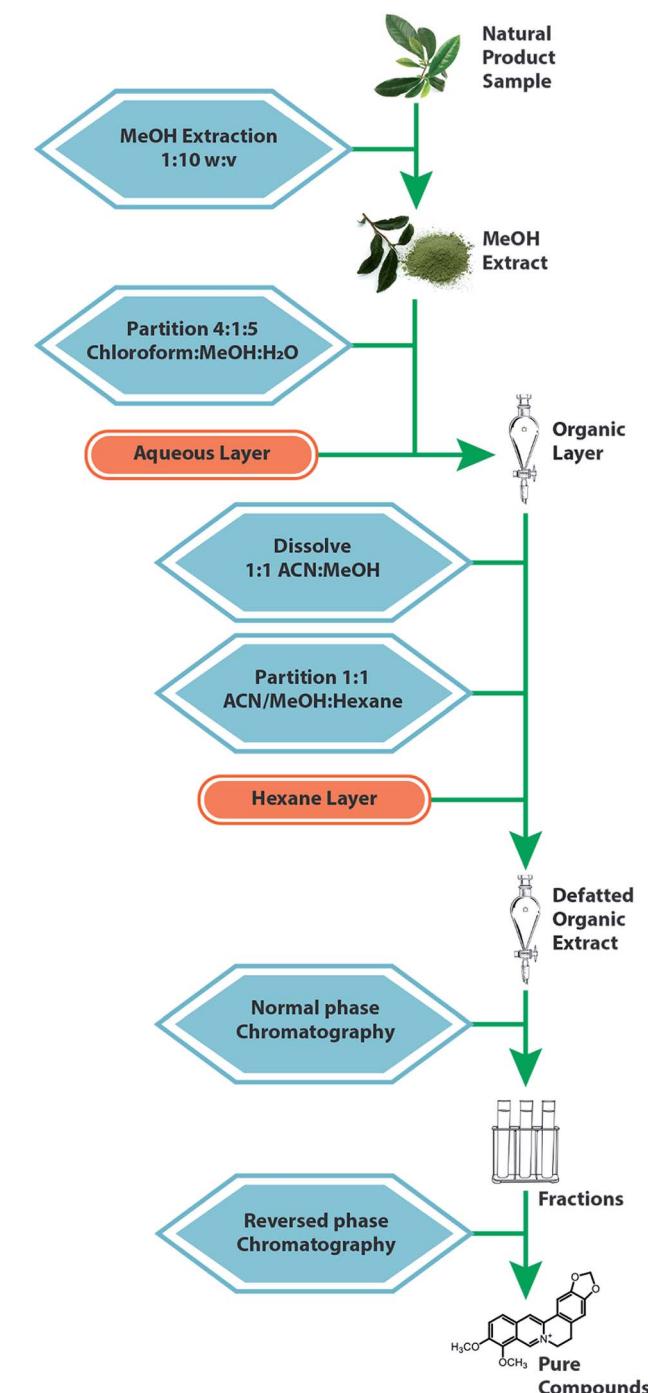


Fig. 9 Standard extraction and fractionation scheme for a natural product. The sample material is macerated with an organic solvent, usually methanol, then subjected to repeated liquid–liquid partitions, yielding a final organic extract. This residue is fractionated with a normal-phase flash chromatography system, after which the fractions are further separated via a reverse-phase HPLC separation to yield purified compounds.<sup>77,185</sup>

stereoscopic information as needed.<sup>10</sup> In the very near future, we anticipate that many journals will require the deposition of raw NMR data associated with the structure elucidation of organic molecules, akin to the way DNA data are deposited in

GenBank; a recent review in *Natural Products Reports*, co-authored by over 70 scientists, advocates for this practice.<sup>184</sup>

For comprehensive characterization of mixtures without isolation, LC-MS (or GC-MS for volatile samples), is the most effective approach. Challenges, pitfalls, and strategies to identify botanical natural product metabolites by mass spectrometry have been detailed previously.<sup>62</sup> If high resolution mass spectrometry is used, accurate measurements of mass are possible to tentatively identify extract components based on calculations of molecular formula. Mass spectrometers with the ability to measure mass with sufficient accuracy include those with Orbitrap, Fourier transform inductively coupled resonance (FT-ICR) or quadrupole time-of-flight (Q-ToF) mass analyzers. The confidence with which structure can be assigned based on mass spectrometric data is increased when retention time and fragmentation patterns for the component of the unknown sample can be compared to those of pure, isolated standards.<sup>186–188</sup> However, mass spectrometry is not a conclusive technique for assigning configuration of stereoisomers.

#### 9.4 Quantitative analysis

The goal of quantitative analysis is to determine the absolute concentration of a known constituent (analyte) of a complex botanical extract. Effective quantitative analysis requires an isolated standard identical to the constituent to be quantified (see Section 9.2, Obtaining “pure” standards). Dilutions of this standard are prepared and analyzed with the extract. A calibration curve of detector response (most commonly chromatographic peak area under the curve) *vs.* analyte concentration is plotted. The calibration curve is analyzed *via* regression over the linear range (range in which response increases linearly with increasing concentration). The response (peak area) for the analyte in the extract is substituted into the equation for the best fit line, and the concentration in the extract and the original plant sample is back calculated by adjusting for all necessary dilutions.<sup>189</sup> To improve the accuracy of quantitative analyses, methods using isotopically labeled standards (when available) can also be employed.<sup>190–192</sup>

The most popular techniques for quantitative analysis of botanical natural products are GC or GC-MS (for volatile sample components) and LC-UV/VIS or LC-MS (for nonvolatile sample components). The relative merits of these techniques are detailed elsewhere.<sup>62</sup> Briefly, LC-MS often (but not always) provides lower limits of detection compared to LC-UV/VIS. An additional advantage of LC-MS is that it can be used to obtain orthogonal data about analyte mass and fragmentation pattern that is helpful for deconvoluting chromatograms and confirming identification.<sup>158</sup> LC-UV/VIS, on the other hand, is typically less susceptible to matrix interference and for some applications demonstrates improved linearity, particularly at the high end of the calibration curve.<sup>95</sup> Instruments with UV/VIS detectors are also less expensive and available to a broader base of analysts. Having access to instruments equipped with both UV and MS detectors, such that identity of the quantified constituents can be confirmed using more than one data type, is ideal. When both UV and MS data are available, quantitative analysis



can be accomplished with whichever technique provides the most linear data in the region of interest.

When conducting quantitative analysis, analyzing the extract (botanical natural product) at multiple dilutions is recommended. A typical analysis includes a series of 2-fold or 10-fold dilutions of the extract, with the highest concentration in the range of  $1.0 \text{ mg mL}^{-1}$  (expressed as dry weight of extract per volume of dilution solvent). It is often desirable to quantify more than one constituent, and the ideal dilution to achieve a concentration within the linear range of the calibration curve will vary from constituent to constituent. Thus, analyzing the extract at multiple dilutions is helpful. A series of 2-fold or 10-fold dilutions of the standard is also included in the same analysis with the extract to generate a calibration curve for quantitative analysis. The resulting calibration curve should span the maximum linear range, and several dilutions above and below this range should be included to enable the full range to be determined. AOAC INTERNATIONAL (The Association of Analytical Communities) recommends including 6–8 data points within the linear range.<sup>193</sup>

One question that often arises when conducting quantitative analysis is the issue of whether (or how) a method has been validated. Validation is a process by which a given method is optimized for parameters related to accuracy, precision, limits of detection and quantitation, linear range and robustness.<sup>90</sup> Other elements of method evaluation in the context of botanical natural product research include measurements of extraction efficiency and checking for potential matrix interference. Methods can be validated both within a single laboratory (intralaboratory validation)<sup>95,194</sup> or across multiple laboratories (interlaboratory validation).<sup>67,90</sup> The International Council for Harmonisation of Technical Requirements for Pharmaceuticals for Human Use (ICH) provides general guidelines for validation of analytical methods,<sup>195</sup> and more specific guidelines and definitions related to validation of constituents of botanicals have been drafted by a subcommittee of AOAC INTERNATIONAL.<sup>13,193</sup> When selecting a method for analysis of botanicals, we recommend reviewing multiple published methods (if available), with particular attention to those that have been validated. However, for practical reasons, including differences in available expertise, samples and instrumentation, it may be difficult to precisely replicate even a validated method for a new study. This is particularly true if the method has not already been subjected to interlaboratory validation. For this reason, it is typically useful to validate the selected method as part of the given project. One excellent strategy for checking accuracy is to obtain a botanical reference sample for which concentrations of the bioactives of interest are known, then compare the results obtained with the selected method to those reported on the certificate of analysis for the reference material. Such validation can be performed using reference standards from NIST (if available) for which marker compound concentrations have been reported. To avoid misleading results due to matrix interference, it is important to check the accuracy of the method in the botanical matrix of interest. For example, our methods for quantifying catechins as part of the NaPDI Center's green tea study were tested by analyzing a NIST green tea standard reference material, not

isolated catechins.<sup>28</sup> If no commercial reference material of the botanical is available, alternate techniques for quantifying metabolites are available. One technique is to conduct a “spike recovery” study, in which a standard is spiked into the botanical matrix at known concentrations, and the concentration is back-calculated to evaluate accuracy.<sup>95,196–198</sup> Quantitative NMR and pre- or post-column derivatization<sup>199</sup> also can be used to quantify metabolites without known standards.<sup>200</sup>

We end this section with a word of caution, which is that it is easy to become so focused on method validation that one is distracted from the overall research question (which typically relates to safety and/or efficacy of a botanical). It is often true that variability in biological response is so significant that it is unnecessary (and even a waste of time) to pin down concentrations in a botanical extract to the highest level achievable by a trained analytical chemist. The extent to which validation is necessary should be carefully considered in the context of the research question being evaluated.

#### Box 6 Summary of recommendations: identification, structure elucidation, and quantitative analysis of bioactive constituents

- Knowledge of which constituents in the botanical extract are biologically active is needed to establish quality control procedures and determine dosage.
- Biological activity may be due to more than one constituent, and the active constituent profile may vary depending on which biological activity is tested.
- Because of the inherent complexity of botanical extracts, isolating all constituents and testing them individually for biological activity is infeasible.
- Bioassay-guided fractionation, which can be enhanced with biostatistical analyses, can be used to direct isolation efforts towards identifying compounds likely to be active, provided that a relevant and robust *in vitro* assay is available.
- Results of bioassay-guided fractionation experiments should be interpreted with consideration of the potential limitations of translatability to clinical studies.
- Quantitative analysis of active compounds can be accomplished using isotopically labeled standards or calibration curves prepared from standards of the compound of interest in the relevant matrix.

## 10. Conclusions

The process of selecting botanical natural products for *in vitro*, non-clinical *in vivo*, and clinical studies is complex, and the ideal approach must be tailored on a case by case basis. Nonetheless, as a result of extensive experience, and much trial and error, we have identified a few common themes, pitfalls, and questions that almost always arise. In the preceding sections, we sought to address these by presenting examples from work conducted as a part of the NaPDI Center in the context of current literature. While constructing the recommendations provided in this report, we have considered the need to balance the ideal, *i.e.* what would be done in a situation where time, person power, and funding were unlimited, with the practical, *i.e.* what is possible subject to realistic constraints. We provide Table 3 as a summary of many of the recommendations from the text, and to highlight a few



**Table 3** Practical tips for selecting botanical natural products prior to conducting *in vitro*, non-clinical *in vivo*, and clinical studies**Screen many products**

Acquire and chemically evaluate as many different products as feasible. Most studies reviewed selected >20 products,<sup>28,29,86,94,201</sup> although lower thresholds may be acceptable depending on limitations due to funding, logistics, and availability. Products selected for a given study or studies should be representative of consumer usage

**Confirm identity claims**

Verify all claims on labels *via* an independent analysis (e.g., the researcher's laboratory or a contract laboratory). This is important both for commercial botanical products and reference standards

**Authenticate thoroughly**

Authenticate using multiple orthogonal techniques and by comparing with trustworthy, vouchered samples. While this may seem obvious, avoid the wasted effort of conducting a research project starting with the wrong botanical

**Obtain vouchers**

Collect a voucher specimen and submit it to a publicly accessible collection at an herbarium. When a commercial product is used and a voucher specimen is not accessible, compare the chemical profile of the botanical to be used in the study to that of a vouchered reference material

**Collect plant specimens using paper bags**

Plant specimens should be stored in paper, rather than plastic, bags after collection. Plastic bags can contaminate the specimens with polymers that may mistakenly be attributed to the specimen. Plastic bags also prevent gas exchange, allowing moisture to accumulate rapidly, and accelerating fungal growth and potential mycotoxin contamination

**Reserve samples**

When preparing an extract, always save a labelled and catalogued sample of the starting product (extract, supplement, or dried plant material). When purifying an extract for analysis (*i.e.* conducting preparative chromatography or liquid–liquid partitioning), save a small sample of the extract and of each successively purified fraction. This reserved aliquot can be re-analyzed as necessary for chemical composition and biological activity. It is better to err on the side of saving too much of these materials than to save too little (or none). Relative to the cost of the future studies, the price of starting with well more than is needed is modest

**Extract with methanol**

Use methanol to prepare samples for chemical analysis and *in vitro* evaluation. Methanol is an effective extraction solvent because it facilitates reproducible production and stable storage of extracts. If necessary, conduct a parallel extraction with other solvents to ensure relevance to consumer usage. In the authors' experience, selection of extraction solvent typically does not change the profile of small molecule metabolites in a sample to a large extent, albeit relative amounts of these compounds may be altered. Methanol has a lower boiling point than ethanol or water, which facilitates its evaporation at lower temperatures, thus minimizing the chance of decomposition of the sample during evaporation. It is also our experience that the difficulty in working with and storing extracts prepared in other solvents introduces artefacts and unexpected complications into a research study, and some other solvents (such as DMSO) can contribute to the oxidation of extract constituents<sup>88</sup>

**Include controls in analyses**

Botanical reference standards aid in the detection of adulterated products and enable botanical authentication, whereas chemical

**Table 3 (Contd.)**

standards aid in quantification and provide the opportunity to monitor the performance of the analytical method

**Purchase or collect more of a single lot than needed for *in vitro* and/or clinical studies**

When using a commercial product, purchase double or triple the amount of product needed for the given study or studies. Where possible, purchase all of this material from a single lot to ensure consistency within and between studies

**Characterize fully (not just for marker compounds)**

Obtain the most comprehensive profile of metabolites possible when characterizing an extract. Techniques that focus on a few metabolites (or purported "marker compounds") may produce misleading results and may unintentionally imply similarity among products that may possess different biological activities

**Consider the limitations of the analytical technique**

A few common errors in interpreting data analysis of botanical extracts are listed in Table 2. Careful data analysis and inclusion of appropriate controls and replicates is important to avoid these errors

**Be cognizant of potential endotoxin contamination**

Lipopolysaccharides (LPS, also referred to as endotoxins) are bacterial components that can profoundly influence *in vitro* assays, particularly those evaluating immunomodulatory activity. Because botanical extracts contain endophytic and epiphytic bacteria (and other microbes), it is virtually impossible to avoid contamination with LPS in extracts prepared from botanicals. Even ethanol extracts can contain sufficient LPS content to alter the results of biological assays<sup>202</sup>

**Store samples cool, dark, and dry**

Never store extracts in liquid form, even in the freezer. Freeze/thaw cycles, especially in solvents such as DMSO, can result in accelerated degradation. Storing samples dry and in dark and cool conditions is the best way to prevent degradation. Cooler temperatures, such as refrigeration or freezing, are indicated to prolong the activity and chemical stability of prepared extracts.<sup>203</sup> However, the type and extent of degradation that occurs will differ depending on the natural product being evaluated. Thus, it is advisable to monitor for potential degradation products by collecting chromatographic profiles immediately after the sample is prepared, and then at regular intervals or prior to each experiment in which it will be evaluated for biological activity

additional points for consideration. It is our expectation that consideration of the topics addressed herein will enable the design of rigorous and reproducible studies with complex botanical natural products, and prevent the wasted time, funds, and effort that goes into studies where product selection has not been carefully considered.

## 12. Conflicts of interest

There are no conflicts to declare.

## 13. Acknowledgements

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## 14. References

- 1 C. H. Wu, C. C. Wang, M. T. Tsai, W. T. Huang and J. Kennedy, *J. Evidence-Based Complementary Altern. Med.*, 2014, **2014**, 872320.
- 2 K. M. Job, T. K. L. Kiang, J. E. Constance, C. M. T. Sherwin and E. Y. Enioutina, *Expert Rev. Clin. Pharmacol.*, 2016, **9**, 1597–1609.
- 3 B. T. Gufford, J. T. Barr, V. González-Pérez, M. E. Layton, J. R. White, N. H. Oberlies and M. F. Paine, *CPT: Pharmacometrics Syst. Pharmacol.*, 2015, **4**, 701–710.
- 4 T. Smith, K. Kawa, V. Eckl, C. Morton and R. Stredney, *HerbalGram*, 2017, **115**, 56–65.
- 5 M. A. Klein, R. L. Nahin, M. J. Messina, J. I. Rader, L. U. Thompson, T. M. Badger, J. T. Dwyer, Y. S. Kim, C. H. Pontzer, P. E. Starke-Reed and C. M. Weaver, *J. Nutr.*, 2010, **140**, 1192S–1204S.
- 6 M. F. Paine, D. D. Shen and J. S. McCune, *Drug Metab. Dispos.*, 2018, **46**, 1041–1045.
- 7 J. Dwyer, P. Coates and M. Smith, *Nutrients*, 2018, **10**, 41.
- 8 Center for Drug Evaluation and Research, *Botanical Drug Development: Guidance for Industry*, U.S. Department of Health and Human Services, 2015.
- 9 National Center for Complementary and Integrative Health (NCCIH), *NCCIH Policy: Natural Product Integrity*, National Institutes of Health, 2016, <https://nccih.nih.gov/research/policies/naturalproduct.htm> (accessed February 18, 2018).
- 10 Journal of Natural Products, Preparation and Submission of Manuscripts, [http://pubs.acs.org/paragonplus/submission/jnprdf/jnprdf\\_authguide.pdf](http://pubs.acs.org/paragonplus/submission/jnprdf/jnprdf_authguide.pdf), (accessed June 8, 2017).
- 11 Phytochemistry Letters, Guide for Authors, <https://www.elsevier.com/journals/phytochemistry-letters/1874-3900/guide-for-authors>, (accessed June 8, 2017).
- 12 Planta Medica, Guidelines for Authors, [https://www.thieme.de/statics/dokumente/thieme/final/de/dokumente/zw\\_thieme-en/PlantaMedica-Authors-December-2016.pdf](https://www.thieme.de/statics/dokumente/thieme/final/de/dokumente/zw_thieme-en/PlantaMedica-Authors-December-2016.pdf), (accessed June 8, 2017).
- 13 AOAC International, *J. AOAC Int.*, 2012, **95**, 268–272.
- 14 D. J. Kroll and N. H. Oberlies, *Focus. Altern. Complement. Ther.*, 2003, **8**, 302–306.
- 15 Office of Dietary Supplements, *Dietary Supplement Label Database*, <https://www.dsld.nlm.nih.gov/dsld/index.jsp>, (accessed September 16, 2018).
- 16 Centers for Disease Control and Prevention (CDC), *National Health and Nutrition Examination Survey Data*, U.S. Department of Health and Human Services, 2017, <https://www.cdc.gov/nchs/nhanes/index.htm> (accessed August 23, 2017).
- 17 Anon., *Consumer Reports*, 2003, vol. 68, pp. 40–43.
- 18 J. Health, Green Tea Supplement Reviews, <http://www.benefitsofgreentea.info/2007/green-tea-supplement-reviews.php>, (accessed November 11, 2015).
- 19 T. Smith, K. Kawa, V. Eckl and J. Johnson, *HerbalGram*, 2016, **111**, 67–73.
- 20 Amazon.com, Amazon Best Sellers, <http://www.amazon.com/Best-Sellers-Grocery-Gourmet-Food-Green-Tea-Beverages/zgbs/grocery/16318471>, (accessed November 11, 2015).
- 21 R. B. van Breemen, *J. Med. Chem.*, 2015, **58**.
- 22 National Park Service (NPS), *Preparing and storing herbarium specimens*, U.S. Department of the Interior, 2009, <https://www.nps.gov/museum/publications/conserveogram/11-12.pdf> (accessed September 19, 2017).
- 23 W. T. Stearn, *Biol. J. Linn. Soc.*, 1971, **3**, 225–233.
- 24 M. Greve, A. M. Lykke, C. W. Fagg, R. E. Gereau, G. P. Lewis, R. Marchant, A. R. Marshall, J. Ndayishimiye, J. Bogaert and J.-C. Svenning, *S. Afr. J. Bot.*, 2016, **105**, 317–323.
- 25 Phytochemistry, Author Information Pack, <https://www.elsevier.com/journals/phytochemistry/0031-9422?generatepdf=true>, (accessed August 31, 2017).
- 26 C. E. Piersen, N. L. Booth, Y. Sun, W. Liang, J. E. Burdette, R. B. v. Breemen, S. E. Geller, C. Gu, S. Banuvar, L. P. Shulman, J. L. Bolton and N. R. Farnsworth, *Curr. Med. Chem.*, 2004, **11**, 1361–1374.
- 27 R. B. van Breemen, H. H. S. Fong and N. R. Farnsworth, *Chem. Res. Toxicol.*, 2007, **20**, 577–582.
- 28 J. J. Kellogg, T. N. Graf, M. F. Paine, J. S. McCune, O. M. Kvalheim, N. H. Oberlies and N. B. Cech, *J. Nat. Prod.*, 2017, **80**, 1457–1466.
- 29 E. D. Wallace, N. H. Oberlies, N. B. Cech and J. J. Kellogg, *Food Chem. Toxicol.*, 2018, **120**, 439–447.
- 30 J. M. Harnly, D. Luthria and P. Chen, *J. AOAC Int.*, 2012, **95**, 1579–1587.
- 31 P. Posadzki, L. Watson and E. Ernst, *Eur. J. Clin. Pharmacol.*, 2013, **69**, 295–307.
- 32 D. J. Kroll, H. S. Shaw, J. M. Mathews and N. H. Oberlies, *Acta Hortic.*, 2005, **720**, 137–147.
- 33 S. Ding, E. Dudley, S. Plummer, J. Tang, R. P. Newton and A. G. Brenton, *Phytochemistry*, 2008, **69**, 1555–1564.
- 34 P. Chen, M. Ozcan and J. Harnly, *Anal. Bioanal. Chem.*, 2007, **389**, 251–261.
- 35 W. Li, Y. Sun, J. F. Fitzloff and R. B. Van Breemen, *Chem. Res. Toxicol.*, 2002, **15**, 1174–1178.
- 36 E. A. Abourashed and I. A. Khan, *J. Pharm. Sci.*, 2001, **90**, 817–822.
- 37 S. M. Henning, C. Fajardo-Lira, H. W. Lee, A. A. Youssefian, V. L. W. Go and D. Heber, *Nutr. Cancer*, 2003, **45**, 226–235.



38 M. Friedman, C. E. Levin, S. H. Choi, E. Kozukue and N. Kozukue, *J. Food Sci.*, 2006, **71**, C328–C337.

39 D. D. Soejarto, *J. Ethnopharmacol.*, 1996, **51**, 1–15.

40 D. D. Soejarto, H. H. S. Fong, G. T. Tan, H. J. Zhang, C. Y. Ma, S. G. Franzblau, C. Gyllenhaal, M. C. Riley, M. R. Kadushin, J. M. Pezzuto, L. T. Xuan, N. T. Hiep, N. V. Hung, B. M. Vu, P. K. Loc, L. X. Dac, L. T. Binh, N. Q. Chien, N. V. Hai, T. Q. Bich, N. M. Cuong, B. Southavong, K. Sydara, S. Bouamanivong, H. M. Ly, T. V. Thuy, W. C. Rose and G. R. Dietzman, *J. Ethnopharmacol.*, 2005, **100**, 15–22.

41 H. Zedan, *Handbook of the Convention on Biological Diversity*, Earthscan Publications Ltd., 2001.

42 T. A. Kursar, C. C. Caballero-George, T. L. Capson, L. Cubilla-Rios, W. H. Gerwick, M. P. Gupta, A. Ibañez, R. G. Linington, K. L. McPhail, E. Ortega-Barría, L. I. Romero, P. N. Solis and P. D. Coley, *BioScience*, 2006, **56**, 1005–1012.

43 Nagoya Protocol, *Nagoya protocol on access to genetic resources and the fair and equitable sharing of benefits arising from their utilization to the convention on biological diversity*, Convention on Biological Diversity, 2011, [www.cbd.int/abs/text/default.shtml](http://www.cbd.int/abs/text/default.shtml) (accessed November 12, 2017).

44 International Institute for Environment and Development, Nagoya Protocol on Access to Genetic Resources and Benefit-Sharing, <https://biocultural.iied.org/nagoya-protocol-access-genetic-resources-and-benefit-sharing>, (accessed June 1, 2018).

45 M. C. Leal, A. Hilario, M. H. G. Munro, J. W. Blunt and R. Calado, *Nat. Prod. Rep.*, 2016, **33**, 747–750.

46 H. A. Raja, T. R. Baker, J. G. Little and N. H. Oberlies, *Food Chem.*, 2017, **214**, 383–392.

47 R. Upton, A. Graff, G. Jolliffe, R. Langer and R. Williamson, *American Herbal Pharmacopoeia: Botanical Pharmacognosy - Microscopic Characterization of Botanical Medicines*, CRC Press, Boca Raton, FL, 2016.

48 N. V. Ivanova, M. L. Kuzmina, T. W. A. Braukmann, A. V. Borisenko and E. V. Zakharov, *PLoS One*, 2016, **11**, e0156426.

49 D. P. Little, *Genome*, 2014, **57**, 513–516.

50 D. F. Coutinho Moraes, D. W. Still, M. R. Lum and A. M. Hirsch, *Planta Med.*, 2015, **81**, 687–695.

51 New York State Office of the Attorney General, A. G. Schneiderman asks major retailers to halt sales of certain herbal supplements as DNA tests fail to detect plant materials listed on majority of products tested, <https://ag.ny.gov/press-release/ag-schneiderman-asks-major-retailers-halt-sales-certain-herbal-supplements-dna-tests>.

52 I. Parveen, S. Gafner, N. Techén, S. J. Murch and I. A. Khan, *Planta Med.*, 2016, **82**, 1225–1235.

53 P. M. Hollingsworth, D.-Z. Li, M. van der Bank and A. D. Twyford, *Philos. Trans. R. Soc. B*, 2016, **371**, 20150338.

54 P. M. Hollingsworth, S. W. Graham and D. P. Little, *PLoS One*, 2011, **6**, e19254.

55 W. L. Clement and M. J. Donoghue, *BMC Evol. Biol.*, 2012, **12**, 73.

56 A. J. Fazekas, P. R. Kesanakurti, K. S. Burgess, D. M. Percy, S. W. Graham, S. C. Barrett, S. G. Newmaster, M. Hajibabaei and B. C. Husband, *Mol. Ecol. Resour.*, 2009, **9**, 130–139.

57 N. Pei, D. L. Erickson, B. Chen, X. Ge, X. Mi, N. G. Swenson, J.-L. Zhang, F. A. Jones, C.-L. Huang and W. Ye, *Sci. Rep.*, 2015, **5**, 15127.

58 A. J. Fazekas, M. L. Kuzmina, S. G. Newmaster and P. M. Hollingsworth, *Methods Mol. Biol.*, 2012, **858**, 223–252.

59 M. Blumenthal and W. R. Busse, *The complete German Commission E monographs*, American Botanical Council, Austin, Texas, 1998.

60 U. S. Pharmacopeia, *Herbal Medicines Compendium*, <http://hmc.usp.org/>, (accessed September 19, 2017, 2017).

61 V. E. Tyler, *Herbs of choice: the therapeutic use of phytomedicinals*, Pharmaceutical Products Press (imprint of Haworth Press, Inc.), 1994.

62 N. B. Cech, *LCCG North Am.*, 2013, **31**, 938–947.

63 C. Simmler, J. G. Napolitano, J. B. McAlpine, S.-N. Chen and G. F. Pauli, *Curr. Opin. Biotechnol.*, 2014, **25**, 51–59.

64 L. Vaclavik, A. J. Kryniotsky and J. I. Rader, *Anal. Bioanal. Chem.*, 2014, **406**, 6767–6790.

65 J. Zhu, X. Fan, Y. Cheng, R. Agarwal, C. M. V. Moore, S. T. Chen and W. Tong, *PLoS One*, 2014, **9**, e87462.

66 A. Navarrete, B. Avula, Y.-W. Choi and I. A. Khan, *J. AOAC Int.*, 2006, **89**, 8–15.

67 M. A. Markus, J. Ferrier, S. M. Luchsinger, J. Yuk, A. Cuerrier, M. J. Balick, J. M. Hicks, K. B. Killday, C. W. Kirby, F. Berrue, R. G. Kerr, K. Knagge, T. Gödecke, B. E. Ramirez, D. C. Lankin, G. F. Pauli, I. Burton, T. K. Karakach, J. T. Arnason and K. L. Colson, *Planta Med.*, 2014, **80**, 732–739.

68 J. Harnly, P. Chen, J. Sun, H. Huang, K. L. Colson, J. Yuk, J.-A. H. McCoy, D. T. H. Reynaud, P. B. Harrington and E. J. Fletcher, *Planta Med.*, 2016, **82**, 250–262.

69 B. Avula, Y.-H. Wang, G. Isaac, J. Yuk, M. Wrona, K. Yu and I. A. Khan, *Planta Med.*, 2017, **83**, 1097–1308.

70 D. Kumar, *Crit. Rev. Anal. Chem.*, 2016, **46**, 400–412.

71 J. Yuk, K. L. McIntyre, C. Fischer, J. Hicks, K. L. Colson, E. Lui, D. Brown and J. T. Arnason, *Anal. Bioanal. Chem.*, 2013, **405**, 4499–4509.

72 H. G. Gika, G. A. Theodoridis, R. S. Plumb and I. D. Wilson, *J. Pharm. Biomed. Anal.*, 2014, **87**, 12–25.

73 T. F. Jorge, J. A. Rodrigues, C. Caldana, R. Schmidt, J. T. van Dongen, J. Thomas-Oates and C. António, *Mass Spectrom. Rev.*, 2016, **35**, 620–649.

74 J. L. Markley, R. Brüschweiler, A. S. Edison, H. R. Eghbalmia, R. Powers, D. Raftery and D. S. Wishart, *Curr. Opin. Biotechnol.*, 2017, **43**, 34–40.

75 T. Sangster, H. Major, R. Plumb, A. J. Wilson and I. D. Wilson, *Analyst*, 2006, **131**, 1075–1078.

76 W. B. Dunn, I. D. Wilson, A. W. Nicholls and D. Broadhurst, *Bioanalysis*, 2012, **4**, 2249–2264.

77 J. J. Kellogg, D. A. Todd, J. M. Egan, H. A. Raja, N. H. Oberlies, O. M. Kvalheim and N. B. Cech, *J. Nat. Prod.*, 2016, **79**, 376–386.



78 J.-E. Lee, B.-J. Lee, J.-O. Chung, J.-A. Hwang, S.-J. Lee, C.-H. Lee and Y.-S. Hong, *J. Agric. Food Chem.*, 2010, **58**, 10582–10589.

79 Y. Fujimura, K. Kurihara, M. Ida, R. Kosaka, D. Miura, H. Wariishi, M. Maeda-Yamamoto, A. Nesumi, T. Saito, T. Kanda, K. Yamada and H. Tachibana, *PLoS One*, 2011, **6**, e23426.

80 J.-E. Lee, B.-J. Lee, J.-O. Chung, H.-J. Shin, S.-J. Lee, C.-H. Lee and Y.-S. Hong, *Food Res. Int.*, 2011, **44**, 597–604.

81 J.-E. Lee, B.-J. Lee, J.-A. Hwang, K.-S. Ko, J.-O. Chung, E.-H. Kim, S.-J. Lee and Y.-S. Hong, *J. Agric. Food Chem.*, 2011, **59**, 10579–10585.

82 L.-S. Lee, J. H. Choi, N. Son, S.-H. Kim, J.-D. Park, D.-J. Jang, Y. Jeong and H.-J. Kim, *J. Agric. Food Chem.*, 2013, **61**, 332–338.

83 J.-E. Lee, B.-J. Lee, J.-O. Chung, H. N. Kim, E.-H. Kim, S. Jung, H. Lee, S. J. Lee and Y.-S. Hong, *Food Chem.*, 2015, **174**, 452–459.

84 P. K. Gupta, G. W. Barone, B. Gurley, E. K. Fifer and H. P. Hendrickson, *Drug Metab. Dispos.*, 2015, **43**, 534–552.

85 P. Bogdanski, J. Suliburska, M. Szulinska, M. Stepien, D. Pupek-Musialik and A. Jablecka, *Nutr. Res.*, 2012, **32**, 421–427.

86 C. Bergeron, J. F. Livesey, D. V. C. Awang, J. T. Arnason, J. Rana, B. R. Baum and W. Letchamo, *Phytochem. Anal.*, 2000, **11**, 207–215.

87 H. Wohlmuth, K. Savage, A. Dowell and P. Mouatt, *Phytomedicine*, 2014, **21**, 912–918.

88 T. G. McCloud, *Molecules*, 2010, **15**, 4526–4563.

89 B. J. Gurley, *Clin. Pharmacol. Ther.*, 2011, **89**, 915–919.

90 P. N. Brown and M. C. Roman, *J. AOAC Int.*, 2008, **91**, 694–701.

91 E. M. Mudge, J. M. Betz and P. N. Brown, *Adv. Nutr.*, 2016, **7**, 390–398.

92 J. G. Napolitano, T. Gödecke, M. F. Rodríguez-Brasco, B. U. Jaki, S.-N. Chen, D. C. Lankin and G. F. Pauli, *J. Nat. Prod.*, 2012, **75**, 238–248.

93 D. L. Luthria, S. Mukhopadhyay, R. J. Robbins, J. W. Finley, G. S. Banuelos and J. M. Harnly, *J. Agric. Food Chem.*, 2008, **56**, 5457–5462.

94 M. Wang, J. Zhao, B. Avula, Y.-H. Wang, C. Avonto, A. G. Chittiboyina, P. L. Wylie, J. F. Parcher and I. A. Khan, *J. Agric. Food Chem.*, 2014, **62**, 12103–12111.

95 T. N. Graf, N. B. Cech, S. J. Polyak and N. H. Oberlies, *J. Pharm. Biomed. Anal.*, 2016, **126**, 26–33.

96 M. Coimbra, B. Isacchi, L. van Bloois, J. S. Torano, A. Ket, X. Wu, F. Broere, J. M. Metselaar, C. J. Rijcken, G. Storm, R. Bilia and R. M. Schiffelers, *Int. J. Pharm.*, 2011, **416**, 433–442.

97 B. Lapornik, M. Prošek and A. Golec Wondra, *J. Food Eng.*, 2005, **71**, 214–222.

98 J. Azmir, I. S. M. Zaidul, M. M. Rahman, K. M. Sharif, A. Mohamed, F. Sahena, M. H. A. Jahurul, K. Ghafoor, N. A. N. Norulaini and A. K. M. Omar, *J. Food Eng.*, 2013, **117**, 426–436.

99 J. J. Kellogg, E. D. Wallace, T. N. Graf, N. H. Oberlies and N. B. Cech, *J. Pharm. Biomed. Anal.*, 2017, **145**, 604–610.

100 N. D. Yuliana, A. Khatib, R. Verpoorte and Y. H. Choi, *Anal. Chem.*, 2011, **83**, 6902–6906.

101 E. Saccenti, H. C. J. Hoefsloot, A. K. Smilde, J. A. Westerhuis and M. M. W. B. Hendriks, *Metabolomics*, 2014, **10**, 361–374.

102 H. Abdi and L. J. Williams, *Wiley Interdiscip. Rev. Comput. Stat.*, 2010, **2**, 433–459.

103 R. A. LaBudde and J. Harnly, *J. AOAC Int.*, 2012, **95**, 273–285.

104 M. I. López, M. P. Callao and I. Ruisánchez, *Anal. Chim. Acta*, 2015, **891**, 62–72.

105 D. A. Sheen, W. F. C. Wocha, K. A. Lippa and D. W. Bearden, *Chemom. Intell. Lab. Syst.*, 2017, **162**, 10–20.

106 D. Grapov, K. Wanichthanarak and O. Fiehn, *Bioinformatics*, 2015, **31**, 2757–2760.

107 C. V. Rider, G. E. Dinse, D. M. Umbach, J. E. Simmons and R. C. Hertzberg, in *Chemical Mixtures and Combined Chemical and Nonchemical Stressors*, Springer, 2018, pp. 235–270.

108 C. V. Rider, N. J. Walker and S. Waidyanatha, *Clinical Pharmacology & Therapeutics*, 2018.

109 E. Ernst, *J. Intern. Med.*, 2002, **252**, 107–113.

110 M. Tims, *Botanical Adulterants Bulletin*, 2016, June, pp. 1–6.

111 P. Geng, J. M. Harnly, J. Sun, M. Zhang and P. Chen, *Anal. Bioanal. Chem.*, 2017, **409**, 2591–2600.

112 N. Sarma, G. Giancaspro and J. Venema, *Drug Test. Anal.*, 2016, **8**, 418–423.

113 J. M. Betz, P. N. Brown and M. C. Roman, *Fitoterapia*, 2011, **82**, 44–52.

114 C. Simmler, J. G. Graham, S.-N. Chen and G. F. Pauli, *Fitoterapia*, 2018, **129**, 401–414.

115 P. Geng, J. M. Harnly and P. Chen, *J. Agric. Food Chem.*, 2015, **63**, 6189–6211.

116 N. P. Mncwangi, A. M. Viljoen, J. Zhao, I. Vermaak, W. Chen and I. A. Khan, *Phytochemistry*, 2014, **106**, 104–115.

117 A. Pengelly, K. Bennett, K. Spelman and M. Tims, *Goldenseal: An Appalachian Plant Monograph*, Appalachian Center for Ethnobotanical Studies, 2012, <https://www.frostburg.edu/fsu/assets/File/ACES/Hydrastis%20canadensis%20for%20ACES%20website.pdf> (accessed October 25, 2017).

118 H. A. Weber, M. K. Zart, A. E. Hodges, H. M. Molloy, B. M. O'Brien, L. A. Moody, A. P. Clark, R. K. Harris, J. D. Overstreet and C. S. Smith, *J. Agric. Food Chem.*, 2003, **51**, 7352–7358.

119 A. A. Grippo, B. Hamilton, R. Hannigan and B. J. Gurley, *Am. J. Health-Syst. Pharm.*, 2006, **63**.

120 J. W. Wong, K. Zhang, K. Tech, D. G. Hayward, A. J. Kryniotsky, I. Cassias, F. J. Schenck, K. Banerjee, S. Dasgupta and D. Brown, *J. Agric. Food Chem.*, 2010, **58**, 5884–5896.

121 C. Witschi and E. Doelker, *Eur. J. Pharm. Biopharm.*, 1997, **43**, 215–242.

122 Z. Veprikova, M. Zachariasova, Z. Dzuman, A. Zachariasova, M. Fenclova, P. Slavikova, M. Vaclavikova, K. Mastovska, D. Hengst and J. Hajslova, *J. Agric. Food Chem.*, 2015, **63**, 6633–6643.



123 P. Raman, L. C. Patino and M. G. Nair, *J. Agric. Food Chem.*, 2004, **52**, 7822–7827.

124 A. C. Brown, *Food Chem. Toxicol.*, 2017, **107**, 449–471.

125 Food and Drug Administration (FDA), *Guidance for Industry: Current Good Manufacturing Practice in Manufacturing, Packaging, Labeling, or Holding Operations for Dietary Supplements; Small Entity Compliance Guide*, U.S. Department of Health and Human Services, 2010, <https://www.fda.gov/Food/GuidanceRegulation/GuidanceDocuments/RegulatoryInformation/DietarySupplements/ucm238182.htm> (accessed October 4, 2017).

126 R. Danko, *Testing Ingredient and Components for Identity and Quality*, Natural Product Insider, 2016, vol. 21, p. 48.

127 R. Dowlatabadi, F. Farshidfar, Z. Zare, M. Pirali, M. Rabiei, M. R. Khoshayand and H. J. Vogel, *Metabolomics*, 2017, **13**, 1–11.

128 R. Alvarez-Zapata, A. Sánchez-Medina, M. Chan-Bacab, K. García-Sosa, F. Escalante-Erosa, R. V. García-Rodríguez and L. M. Peña-Rodríguez, *J. Chromatogr. Aa*, 2015, **1422**, 213–221.

129 C. Simmler, D. Kulakowski, D. C. Lankin, J. B. McAlpine, S.-N. Chen and G. F. Pauli, *Adv. Nutr.*, 2016, **7**, 179–189.

130 C. G. Enke and L. J. Nagels, *Anal. Chem.*, 2011, **83**, 2539–2546.

131 H. A. Junio, A. A. Sy-Cordero, K. A. Ettefagh, J. T. Burns, K. T. Micko, T. N. Graf, S. J. Richter, R. E. Cannon, N. H. Oberlies and N. B. Cech, *J. Nat. Prod.*, 2011, **74**, 1621–1629.

132 B. G. Chiari, J. A. Severi, D. Pauli-Credendio, P. Abackerli, C. M. d. Sylos, W. Vilegas, M. A. Correa and V. L. B. Isaac, *Int. J. Pharm. Pharm. Sci.*, 2012, 331–336.

133 H. Wagner, *Fitoterapia*, 2011, **82**, 34–37.

134 K. Niemeyer, I. R. Bell and M. Koithan, *J. Herb. Med.*, 2013, **3**, 112–119.

135 J. Burns, L. Zhao, E. W. Taylor and K. Spelman, *Toxicology*, 2010, **278**, 140–159.

136 N. H. Oberlies and D. J. Kroll, *J. Nat. Prod.*, 2004, **67**, 129–135.

137 M. C. Wani, H. L. Taylor, M. E. Wall, P. Coggon and A. T. McPhail, *J. Am. Chem. Soc.*, 1971, **93**, 2325–2327.

138 R. B. van Breemen, C. R. Huang, D. Nikolic, C. P. Woodbury, Y. Z. Zhao and D. L. Venton, *Anal. Chem.*, 1997, **69**, 2159–2164.

139 Z. Yang, Y. Zhang, L. Sun, Y. Wang, X. Gao and Y. Cheng, *Anal. Chim. Acta*, 2012, **719**, 87–95.

140 L. Ciesla and R. Moaddel, *Nat. Prod. Rep.*, 2016, **33**, 1131–1145.

141 H. Martens, S. W. Bruun, I. Adt, G. D. Sockalingum and A. Kohler, *J. Chemom.*, 2006, **20**, 402–417.

142 M. Balouiri, M. Sadiki and S. K. Ibnsouda, *J. Pharm. Anal.*, 2016, **6**, 71–79.

143 G. Brusotti, I. Cesari, A. Dentamaro, G. Caccialanza and G. Massolini, *J. Pharm. Biomed. Anal.*, 2014, **87**, 218–228.

144 B. M. Dietz, A. Hajirahimkhan, T. L. Dunlap and J. L. Bolton, *Pharmacol. Rev.*, 2016, **68**, 1026–1073.

145 A. L. Harvey, R. Edrada-Ebel and R. J. Quinn, *Nat. Rev. Drug Discovery*, 2015, **14**, 111.

146 D. J. Newman and G. M. Cragg, *J. Nat. Prod.*, 2016, **79**, 629–661.

147 W. Chun-Guang, Y. Jun-Qing, L. Bei-Zhong, J. Dan-Ting, W. Chong, Z. Liang, Z. Dan and W. Yan, *Eur. J. Pharmacol.*, 2010, **627**, 33–41.

148 B. T. Gufford, G. Chen, A. G. Vergara, P. Lazarus, N. H. Oberlies and M. F. Paine, *Drug Metab. Dispos.*, 2015, **43**, 1353–1359.

149 G. Li, K. Huang, D. Nikolic and R. B. Van Breemen, *Drug Metab. Dispos.*, 2015, **43**, 1670–1678.

150 P. Posadzki, L. Watson and E. Ernst, *Br. J. Clin. Pharmacol.*, 2013, **75**, 603–618.

151 C. A. Van Loo-Bouwman, T. H. J. Naber, M. Minekus, R. B. van Breemen, P. J. M. Hulshof and G. Schaafsma, *J. Agric. Food Chem.*, 2014, **62**, 950–955.

152 S. Possemiers, S. Bolca, W. Verstraete and A. Heyerick, *Fitoterapia*, 2011, **82**, 53–66.

153 J. M. Price, J. G. Little and T. R. Baker, *Planta Med.*, 2014, **80**, CL12.

154 K. L. Krivos, B. T. Regg, J. M. Price, D. A. McMillan and T. R. Baker, *Planta Med.*, 2014, **80**, PPL19.

155 J. G. Napolitano, T. Gödecke, D. C. Lankin, B. U. Jaki, J. B. McAlpine, S.-N. Chen and G. F. Pauli, *J. Pharm. Biomed. Anal.*, 2014, **93**, 59–67.

156 C. D. Broeckling, F. A. Afsar, S. Neumann, A. Ben-Hur and J. E. Prenni, *Anal. Chem.*, 2014, **86**, 6812–6817.

157 N. D. Paguigan, T. El-Elimat, D. Kao, H. A. Raja, C. J. Pearce and N. H. Oberlies, *J. Antibiot.*, 2017, **70**, 553–561.

158 T. El-Elimat, M. Figueroa, B. M. Ehrmann, N. B. Cech, C. J. Pearce and N. H. Oberlies, *J. Nat. Prod.*, 2013, **76**, 1709–1716.

159 M. Wang, J. J. Carver, V. V. Phelan, L. M. Sanchez, N. Garg, Y. Peng, D. D. Nguyen, J. Watrous, C. A. Kapono and T. Luzzatto-Knaan, *Nat. Biotechnol.*, 2016, **34**, 828.

160 T. Hoffmann, D. Krug, S. Hüttel and R. Müller, *Anal. Chem.*, 2014, **86**, 10780–10788.

161 S. P. Gaudêncio and F. Pereira, *Nat. Prod. Rep.*, 2015, **32**, 779–810.

162 D. G. Cox, J. Oh, A. Keasling, K. L. Colson and M. T. Hamann, *Biochim. Biophys. Acta*, 2014, **1840**, 3460–3474.

163 N. D. Yuliana, A. Khatib, Y. H. Choi and R. Verpoorte, *Phytother. Res.*, 2011, **25**, 157–169.

164 F. Qiu, G. Cai, B. U. Jaki, D. C. Lankin, S. G. Franzblau and G. F. Pauli, *J. Nat. Prod.*, 2013, **76**, 413–419.

165 E. R. Britton, J. J. Kellogg, O. M. Kvalheim and N. B. Cech, *J. Nat. Prod.*, 2018, **81**, 484–493.

166 L. K. Caesar, J. J. Kellogg, O. M. Kvalheim, R. A. Cech and N. B. Cech, *Planta Med.*, 2018.

167 M. Farrés, S. Platikanov, S. Tsakovski and R. Tauler, *J. Chemom.*, 2015, **29**, 528–536.

168 T. Rajalahti, R. Arneberg, F. S. Berven, K.-M. Myhr, R. J. Ulvik and O. M. Kvalheim, *Chemom. Intell. Lab. Syst.*, 2009, **95**, 35–48.



169 T. Rajalahti, R. Arneberg, A. C. Kroksveen, M. Berle, K.-M. Myhr and O. M. Kvalheim, *Anal. Chem.*, 2009, **81**, 2581–2590.

170 D.-D. Tian, J. J. Kellogg, N. Okut, N. H. Oberlies, N. B. Cech, D. D. Shen, J. S. McCune and M. F. Paine, *Drug Metab. Dispos.*, 2018, **46**, 552–560.

171 F. Bucar, A. Wube and M. Schmid, *Nat. Prod. Rep.*, 2013, **30**, 525–545.

172 A. Michalkiewicz, M. Biesaga and K. Pyrzynska, *J. Chromatogr. A*, 2008, **1187**, 18–24.

173 J. B. McAlpine, J. B. Friesen and G. F. Pauli, in *Natural Products Isolation*, ed. S. D. Sarker and L. Nahar, Humana Press, Totowa, NJ, 2012, pp. 221–254, DOI: 10.1007/978-1-61779-624-1\_9.

174 N.-C. Kim, N. H. Oberlies, D. R. Brine, R. W. Handy, M. C. Wani and M. E. Wall, *J. Nat. Prod.*, 2001, **64**, 251–253.

175 N. Pfoze, B. Myrboh, Y. Kumar and M. R. Rohman, *Journal of Medicinal Plants Studies*, 2014, **2**, 48–57.

176 S. Agatonovic-Kustrin, D. W. Morton and A. P. Yusof, *Mod. Chem. Appl.*, 2015.

177 T. El-Elimat, H. A. Raja, T. N. Graf, S. H. Faeth, N. B. Cech and N. H. Oberlies, *J. Nat. Prod.*, 2014, **77**, 193–199.

178 T. N. Graf, M. C. Wani, R. Agarwal, D. J. Kroll and N. H. Oberlies, *Planta Med.*, 2007, **73**, 1495–1501.

179 N. Bross-Walch, T. Kuhn, D. Moskau and O. Zerbe, *Chem. Biodivers.*, 2005, **2**, 147–177.

180 A. Bouslimani, L. M. Sanchez, N. Garg and P. C. Dorrestein, *Nat. Prod. Rep.*, 2014, **31**, 718–729.

181 P. Crews, J. Rodriguez and M. Jaspars, *Organic Structure Analysis 2nd edn*, Oxford University Press, New York, NY, USA, 2009.

182 R. G. Linington, P. G. Williams and J. B. MacMillan, *Problems in Organic Structure Determination: A Practical Approach to NMR Spectroscopy*, CRC Press, Boca Raton, FL, USA, 2015.

183 M. Halabalaki, K. Voungogiannopoulou, E. Mikros and A. L. Skaltsounis, *Curr. Opin. Biotechnol.*, 2014, **25**, 1–7.

184 J. B. McAlpine, S.-N. Chen, A. Kutateladze, J. B. MacMillan, G. Appendino, A. Barison, M. A. Beniddir, M. W. Biavatti, S. Bluml, A. Boufridi, M. S. Butler, R. J. Capon, Y. H. Choi, D. Coppage, P. Crews, M. T. Crimmins, M. Csete, P. Dewapriya, J. M. Egan, M. J. Garson, G. Genta-Jouve, W. H. Gerwick, H. Gross, M. K. Harper, P. Hermanto, J. M. Hook, L. Hunter, D. Jeannerat, N.-Y. Ji, T. A. Johnson, D. G. I. Kingston, H. Koshino, H.-W. Lee, G. Lewin, J. Li, R. G. Linington, M. Liu, K. L. McPhail, T. F. Molinski, B. S. Moore, J.-W. Nam, R. P. Neupane, M. Niemitz, J.-M. Nuzillard, N. H. Oberlies, F. M. M. Ocampos, G. Pan, R. J. Quinn, D. S. Reddy, J.-H. Renault, J. Rivera-Chávez, W. Robien, C. M. Saunders, T. J. Schmidt, C. Seger, B. Shen, C. Steinbeck, H. Stuppner, S. Sturm, O. Taglialatela-Scafati, D. J. Tantillo, R. Verpoorte, B.-G. Wang, C. M. Williams, P. G. Williams, J. Wist, J.-M. Yue, C. Zhang, Z. Xu, C. Simmler, D. C. Lankin, J. Bisson and G. F. Pauli, *Nat. Prod. Rep.*, 2018, in press.

185 M. Figueroa, A. K. Jarmusch, H. A. Raja, T. El-Elimat, J. S. Kavanaugh, A. R. Horswill, R. G. Cooks, N. B. Cech and N. H. Oberlies, *J. Nat. Prod.*, 2014, **77**, 1351–1358.

186 H. Xu, H. Niu, B. He, C. Cui, Q. Li and K. Bi, *Molecules*, 2016, **21**, 664.

187 E. P. Ekanayaka, M. D. Celiz and A. D. Jones, *Plant Physiol.*, 2015, **167**, 1221–1232.

188 T. Xie, Y. Liang, H. Hao, A. Jiye, L. Xie, P. Gong, C. Dai, L. Liu, A. Kang and X. Zheng, *J. Chromatogr. Aa*, 2012, **1227**, 234–244.

189 K. A. Reynertson and K. Mahmood, *Botanicals: Methods and techniques for quality & authenticity*, CRC Press, Boca Raton, FL, USA, 2015.

190 S. Arrivault, M. Guenther, S. C. Fry, M. M. Fuenfgeld, D. Veyel, T. Mettler-Altmann, M. Stitt and J. E. Lunn, *Anal. Chem.*, 2015, **87**, 6896–6904.

191 O. Chahrour, D. Cobice and J. Malone, *J. Pharm. Biomed. Anal.*, 2015, **113**, 2–20.

192 K. Habler and M. Rychlik, *Anal. Bioanal. Chem.*, 2016, **408**, 307–317.

193 AOAC International, *AOAC Official Methods of Analysis*, 2012, pp. 1–32.

194 K. M. VanderMolen, N. B. Cech, M. F. Paine and N. H. Oberlies, *Phytochem. Anal.*, 2013, **24**, 654–660.

195 ICH Harmonized Tripartite Guideline, Q2 (R1), 2005, 1.

196 M. C. Roman, J. M. Betz and J. Hildreth, *J. AOAC Int.*, 2007, **90**, 68–81.

197 A. K. Jarmusch, A. M. Musso, T. Shymanovich, S. A. Jarmusch, M. J. Weavil, M. E. Lovin, B. M. Ehrmann, S. Saari, D. E. Nichols and S. H. Faeth, *J. Pharm. Biomed. Anal.*, 2016, **117**, 11–17.

198 K. M. VanderMolen, G. R. Ainslie, M. F. Paine and N. H. Oberlies, *J. Pharm. Biomed. Anal.*, 2014, **98**, 260–265.

199 Y. Lv, X. Yang, Y. Zhao, Y. Ruan, Y. Yang and Z. Wang, *Food Chem.*, 2009, **112**, 742–746.

200 Y. Liu, S.-N. Chen, J. B. McAlpine, L. L. Klein, J. B. Friesen, D. C. Lankin and G. F. Pauli, *J. Nat. Prod.*, 2014, **77**, 611–617.

201 A. Booker, A. Suter, A. Krnjic, B. Strassel, M. Zloh, M. Said and M. Heinrich, *J. Pharm. Pharmacol.*, 2014, **66**, 811–822.

202 D. A. Todd, T. V. Gulleedge, E. R. Britton, M. Oberhofer, M. Leyte-Lugo, A. N. Moody, T. Shymanovich, L. F. Grubbs, M. Juzumaitė, T. N. Graf, N. H. Oberlies, S. H. Faeth, S. M. Laster and N. B. Cech, *PLoS One*, 2015, **10**, e0124276.

203 J. Dai, A. Gupte, L. Gates and R. Mumper, *Food Chem. Toxicol.*, 2009, **47**, 837–847.

