

Synergy-Directed Fractionation of Botanical Medicines: A Case Study with Goldenseal (*Hydrastis canadensis*)

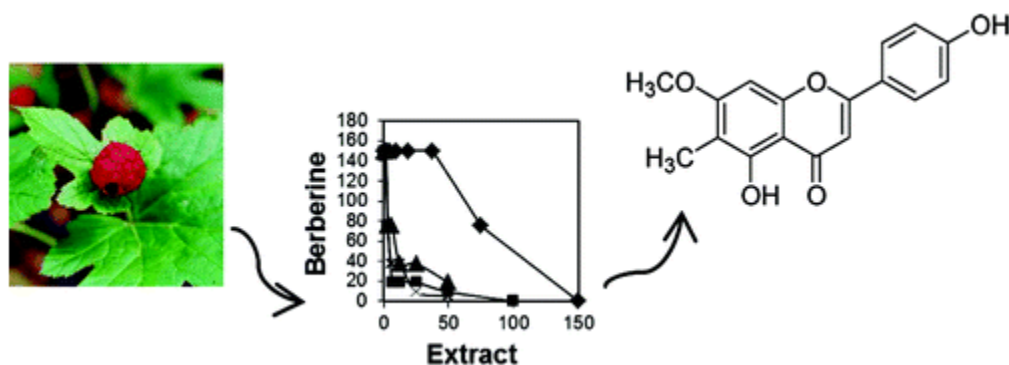
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“Synergy-Directed Fractionation of Botanical Medicines: A Case Study with Goldenseal (*Hydrastis canadensis*).” Hiyas A. Junio, Arlene A. Sy-Cordero, Keivan A. Ettefagh, Johnna T. Burns, Kathryn T. Micko, Tyler N. Graf, Scott J. Richter, Robert E. Cannon, Nicholas H. Oberlies, and and Nadja B. Cech *Journal of Natural Products*, 2011, 74, 1621-1629.

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Abstract:



It is often argued that the efficacy of herbal medicines is a result of the combined action of multiple constituents that work synergistically or additively. Determining the bioactive constituents in these mixtures poses a significant challenge. We have developed an approach to address this challenge, synergy-directed fractionation, which combines comprehensive mass spectrometry profiling with synergy assays and natural products isolation. The applicability of synergy-directed fractionation was demonstrated using the botanical medicine goldenseal (*Hydrastis canadensis*) as a case study. Three synergists from goldenseal were identified, sideroxylin (1), 8-desmethyl-sideroxylin (2), and 6-desmethyl-sideroxylin (3). These flavonoids synergistically enhance the antimicrobial activity of the alkaloid berberine (also a constituent of *H. canadensis*) against *Staphylococcus aureus* by inhibition of the NorA multidrug resistance pump. The flavonoids possess no inherent antimicrobial activity against *S. aureus*; therefore, they could have been missed using traditional bioactivity-directed fractionation. The flavonoid synergists are present at higher concentration in extracts from *H. canadensis* leaves, while the

antimicrobial alkaloid berberine is present at higher levels in *H. canadensis* roots. Thus, it may be possible to produce an extract with optimal activity against *S. aureus* using a combination of goldenseal roots and leaves.

Keywords: herbal medicines | *Hydrastis Canadensis* | *Staphylococcus aureus* | synergy-directed fractionation

Article:

Investigators who study botanical medicines face a formidable challenge. The effects of such preparations can result from the interaction of an array of chemically diverse components, whose identity and concentration may vary from one preparation to the next. Proponents of the medicinal use of botanicals often argue that their components interact synergistically, such that the combined effect is greater than the sum of the parts. Indeed, there are many examples in the literature whereby constituents are more effective in the phytochemical matrix than when isolated.(1, 2) However, the specific components responsible for these effects, and the fundamental mechanisms by which they interact, are rarely known. Methods are needed to unravel the complexity of botanical medicines and identify their key bioactive constituents. Such information would in turn enable better quality control and more meaningful and reproducible clinical trials.

A common approach for studying complex plant preparations is bioactivity-guided fractionation. With this technique, extracts are screened in pursuit of those that contain biologically active compounds. The active extracts are then partitioned and purified, with the ultimate goal of identifying single active compounds. A limitation of bioactivity-guided fractionation for studying botanical medicines is that it may not facilitate the identification of synergists. One type of synergy, potentiation, occurs when a compound possesses no activity on its own, but enhances the activity of another active compound.(3) It is possible with bioactivity-guided fractionation for potentiators to end up in inactive fractions throughout the isolation process, and thereby to be overlooked. With the studies presented here, we sought to develop a new method, synergy-directed fractionation, which would facilitate identification of an array of biologically active compounds, including synergists, in a complex botanical medicine.

The botanical medicine goldenseal, *Hydrastis canadensis* L. (Ranunculaceae), is presented as a case study for the development of the methods presented herein. A recent survey ranked goldenseal among the 20 most popular herbal supplements used worldwide,(4) and it is reported as the sixth most commonly used herbal preparation by children under 18 years of age.(5) Goldenseal is recommended as a treatment for a number of ailments, including viral and bacterial infections and as an antidiarrheal.(6) This use dates back to traditional application by Native Americans; the Cherokee employed the plant topically for eye and skin disease, and it was used by the Iroquois for gastrointestinal issues.(7) There is some justification in the scientific literature for the effectiveness of goldenseal and its constituents. Crude goldenseal root(8-12) and leaf(10) extracts, and their most abundant alkaloid berberine,(13-15) have been shown to possess antimicrobial activity in vitro. Berberine has also demonstrated efficacy in vivo against severe diarrhea(16) and eye infections caused by *Chlamydia trachomatis*.(17)

While the antibacterial activity of goldenseal is generally attributed to alkaloids, several investigators have suggested a role for additional compounds that synergistically or additively enhance the effectiveness of the alkaloids.(8, 9, 18) Most recently, we reported that extracts from the leaves of goldenseal contain synergists that increase the antimicrobial activity of the alkaloid berberine against *Staphylococcus aureus*.(19) Berberine alone has only weak antimicrobial activity (MIC ~150 µg/mL).(20) This low activity can be attributed, at least in part, to berberine being a substrate for bacterial multidrug resistant (MDR) efflux pumps, which function to extrude the compound from bacterial cells, rendering it inactive at low concentrations.(20) We have determined that goldenseal leaf extracts contain unidentified constituents that appear to act as inhibitors of NorA, the major chromosomal multidrug resistance efflux pump expressed by *S. aureus*.(19) These findings are significant given that a major mechanism by which bacteria become resistant to treatment is by the overexpression of efflux pumps.(21) Thus, it may be possible, using extracts that contain both efflux pump inhibitors and antimicrobial agents, to achieve efficacy against drug-resistant bacteria and to limit the development of resistance. A major goal of these studies was to employ synergy-directed fractionation to identify the compounds in *H. canadensis* leaf extracts that synergize the antimicrobial activity of berberine. In so doing, we sought to demonstrate the applicability of synergy-directed fractionation as a method to study botanical medicines. Additional goals were to investigate efflux pump inhibition as the likely mode of action for synergists from *H. canadensis* and to compare the abundance of active compounds in extracts from the roots and leaves of this plant.

RESULTS AND DISCUSSION

The approach for synergy-directed fractionation developed as a result of these studies is outlined in Figure 1. Crude extracts are subjected to synergy testing to identify those likely to contain synergists. Active extracts are fractionated, and each fraction is profiled using liquid chromatography–mass spectrometry (LC-MS) and again subjected to a synergy assay. LC-MS profiles are compared to bioactivity data, and statistical correlations are used to identify potential bioactive compounds (which could include both synergists and those responsible for direct activity). The process of fractionation, synergy testing, and analysis with LC-MS is repeated iteratively until sufficient pure material is obtained for structure elucidation with NMR and MS.

The synergy-directed fractionation approach is perhaps best illustrated by example. In the following section, we demonstrate its application to identify synergists from the botanical medicine goldenseal (*H. canadensis*). Leaf material was chosen for this study because previous studies showed greater synergistic antimicrobial activity of goldenseal leaf extracts than goldenseal root extracts.(19) The first step in the process of identifying the compounds responsible for this effect was to select an appropriate method to assay synergy. The checkerboard assay,(22) which has been widely applied to study antimicrobial synergists, was chosen. To conduct the checkerboard assay, a crude *H. canadensis* leaf extract was tested at a range of concentrations in combination with the known antimicrobial agent berberine (also a component of *H. canadensis*). Minimum inhibitory concentrations (MIC) were measured for each combination of berberine/goldenseal extract concentrations, and an isobologram was plotted (Figure 2). Wagner has presented an excellent review of the use and interpretation of isobolograms for evaluation of synergy among phytochemical compounds.(2) Briefly, an isobologram is a plot where each x,y data pair represents a combination of concentrations at

which a desired activity is obtained (i.e., growth of bacteria is completely inhibited). The shape of the isobologram is indicative of either synergistic, additive, or antagonistic interactions among the compounds or extracts tested. As a quantitative measure of synergy, fractional inhibitory concentrations (FIC) can also be calculated using the same data on which the isobologram is based, as described elsewhere.(19) An FIC cutoff of <0.5 is usually applied to indicate synergy,(23) while an antagonistic effect is characterized by $FIC >4$, and FIC values between 0.5 and 4 indicate no interaction.(22) In this study, the crude *H. canadensis* extract demonstrated synergism when combined with berberine, as indicated by the convex shape of the isobologram (Figure 2) and the FIC value of 0.19 (Table 1).

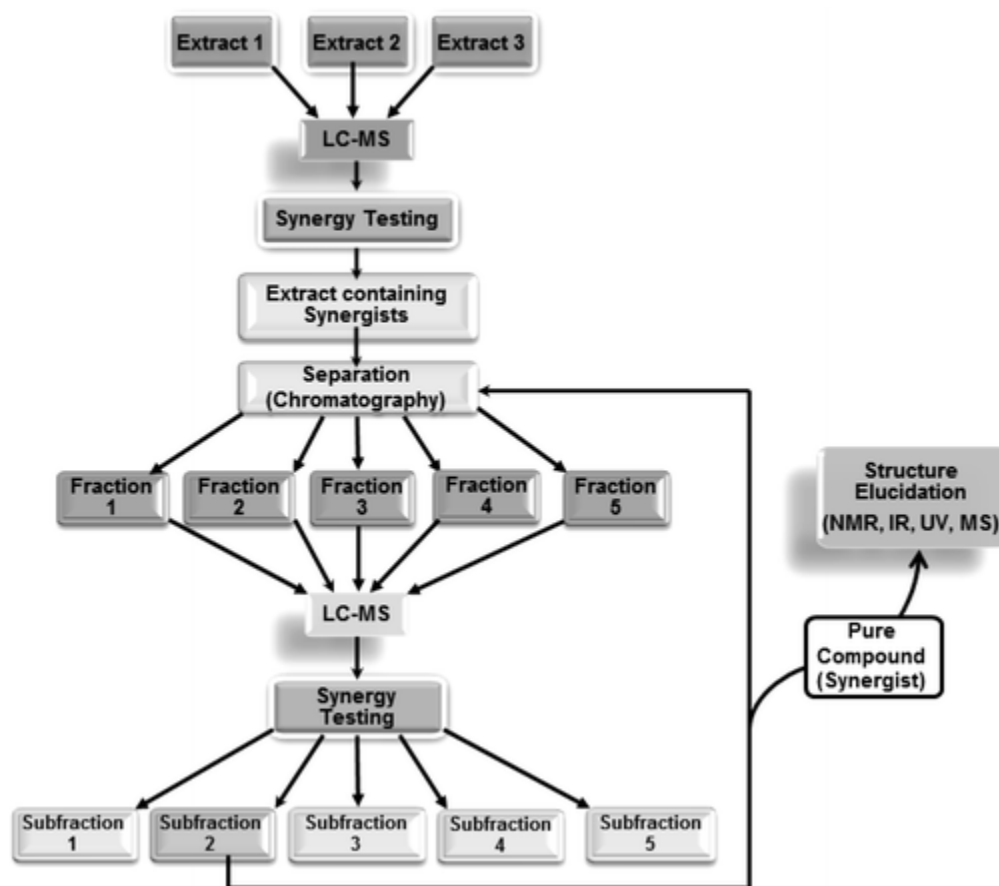


Figure 1. Synergy-directed fractionation. Synergy testing is used to identify extracts and fractions likely to contain combinations of compounds that work together. These extracts are then fractionated, and LC-MS profiles are compared to bioassay data to identify potential bioactive compounds. The process is repeated iteratively as necessary to obtain pure compounds (synergists), the structures of which are elucidated using NMR and other spectroscopic techniques.

Once synergy had been demonstrated for the crude *H. canadensis* extract, the next step was to identify the compounds responsible for this activity. Toward this goal, the extract was fractionated (see Figure S1 provided as Supporting Information), and the activity of each fraction

in combination with berberine was evaluated. Because of limited time and materials, rather than construct a full isobologram, the activity of each fraction was measured at a single concentration (75 $\mu\text{g}/\text{mL}$) in combination with a range of concentrations of berberine. The MIC of berberine in the presence of the fraction was then compared to the MIC of berberine alone, and fractions were deemed “active” if they enhanced the activity of the berberine (decreased its MIC). Note that this activity does not necessarily imply synergy; a fraction could enhance the activity of berberine due to *either* potentiation or an additive antimicrobial effect. Several of the fractions collected from the first-stage fractionation did, indeed, decrease the MIC of berberine (Figure 3A). The most active of these was fraction 4, which decreased the MIC of berberine 16-fold, from 75 $\mu\text{g}/\text{mL}$ (berberine alone) to 4.7 $\mu\text{g}/\text{mL}$ (berberine in combination with fraction 4).

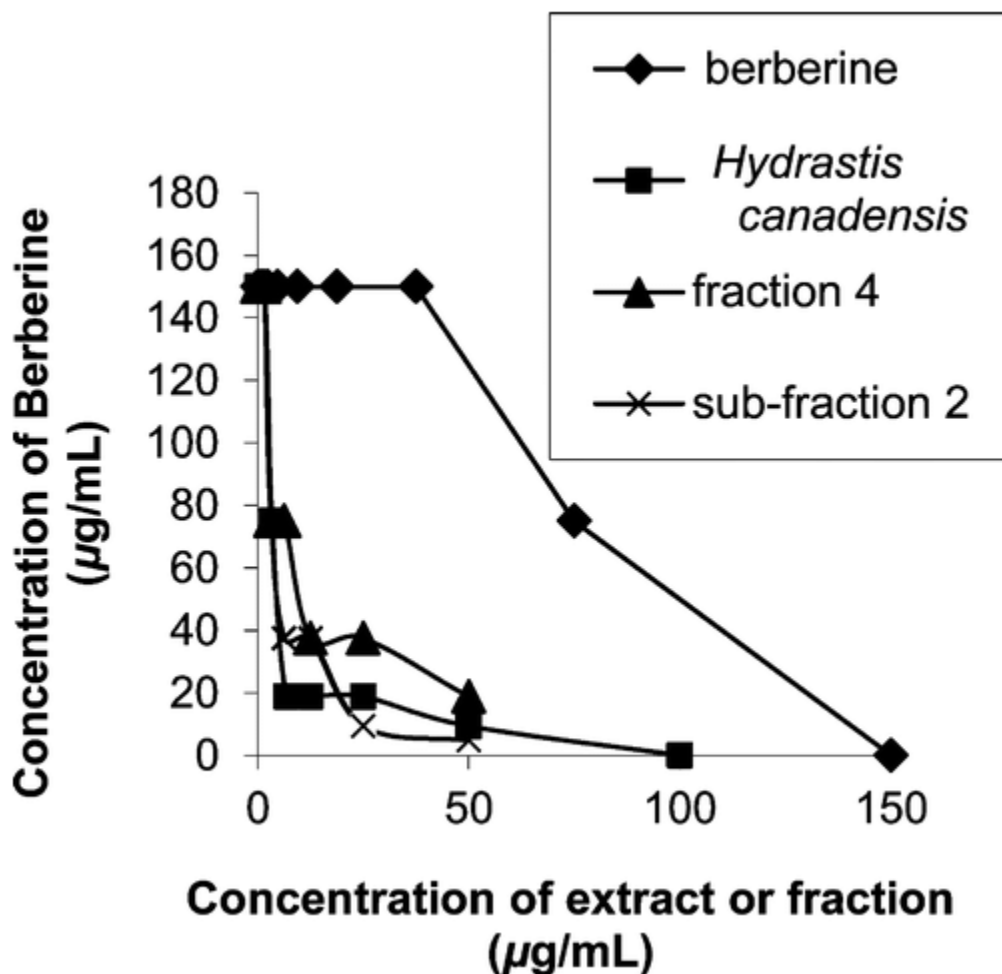


Figure 2. Isobolograms for berberine, the tannin-free *Hydrastis canadensis* extract (after liquid–liquid extraction, see Supporting Information, Figure S1), the most active fraction from the first stage of the separation (fraction 4), and the most active fraction from the second stage of the separation (subfraction 2). All were tested in combination with berberine. The crude extract and its two fractions synergistically enhanced the antimicrobial activity of berberine, as demonstrated by the convex shape of the isobolograms, and the FIC values are reported in Table 1.

Table 1. MIC and FIC Values (Indicative of Synergy) for a *Hydrastis canadensis* Extract and Fractions against Wild-Type *Staphylococcus aureus* (NCTC8325-4)

treatment	MIC ($\mu\text{g/mL}$)	FIC index
berberine	150	1.00
<i>Hydrastis canadensis</i> extract ^a	100	0.19
fraction 4 ^b	>50	0.13
subfraction 2 ^c	>50	0.03

^a The tannin-free *H. canadensis* extract after liquid–liquid partitioning (see Supporting Information, Figure S1) was tested. ^b Fraction 4 was the most active of the fractions collected from the first-stage separation of the tannin-free *H. canadensis* extract. Due to limited solubility, the maximum concentration tested was 50 $\mu\text{g/mL}$. No antimicrobial activity was observed at this concentration. ^c Subfraction 2 was the most active of the fractions collected from the second-stage separation (using flash chromatography over silica gel with fraction 4 as starting material).

A limitation of the simplified approach employed for screening fractions (constant concentration of fraction in combination with a range of concentrations of berberine) is that it does not enable additive effects to be distinguished from synergistic effects. Thus, the observed activity of fraction 4 was tested further with an in-depth synergy assay. A checkerboard assay identical to that employed for the crude extract was conducted using a range of combinations of fraction 4 with berberine. The resulting isobologram (Figure 2) and the FIC value of 0.13 (Table 1) were indicative of synergy.

Once it had been determined that fraction 4 did, indeed, contain synergist(s), LC-MS profiles were compared to the bioactivity data (Figure 3) to determine whether activity could be attributed to specific known or unknown compounds. LC-MS analysis revealed the presence of several ions that were not known constituents of *H. canadensis* in fraction 4. These ions were identified initially based on their m/z values ($M - H^-$ at 311 and 297). NMR analysis of pure isolated compounds eventually enabled their structures to be elucidated as a group of three flavonoids (**1**, **2**, and **3**), as described below.

It was possible, using LC-MS, to verify that the activity of fraction 4 was not due to the presence of the three major known alkaloids in *H. canadensis*, berberine (**4**), hydrastine, and canadine. From Figure 3C, it is apparent that fractions 9, 10, and 11 contained the highest levels of berberine. The weak activity of these fractions (2-fold decrease in MIC) was attributed to this compound, and they were not investigated further. Hydrastine and canadine were present at higher concentrations in fraction 3 than in fraction 4, but fraction 4 was significantly more active. This suggested that other compounds were responsible for the activity of fraction 4, a finding that was not surprising given that hydrastine and canadine have been shown to be inactive against *S. aureus*, both alone and in combination with berberine.(19)

Visual inspection of the data in Figure 3 revealed separate correlations with the presence of compounds at m/z 297 and 311. The strength of this correlation was evaluated jointly using a statistical method. A multiple correlation analysis was conducted to examine the extent of linear

association between peak areas of ions 297 and 311 and the MIC of the fractions obtained from the first-stage separation. A moderate correlation was observed ($R = 0.77$, $p = 0.02$), which suggests that the joint presence of ions 297 and 311 explains approximately 60% (0.77^2), but does not completely explain the activities observed for the fractions. Presumably, other compounds that play a role in the overall activity of the crude extract are also present. Indeed, visual inspection of the data could also have led to this conclusion, given that fractions 5 and 6 demonstrated pronounced activity (8-fold decrease in MIC of berberine) even without significant levels of berberine or of the ions with m/z 297 and 311.

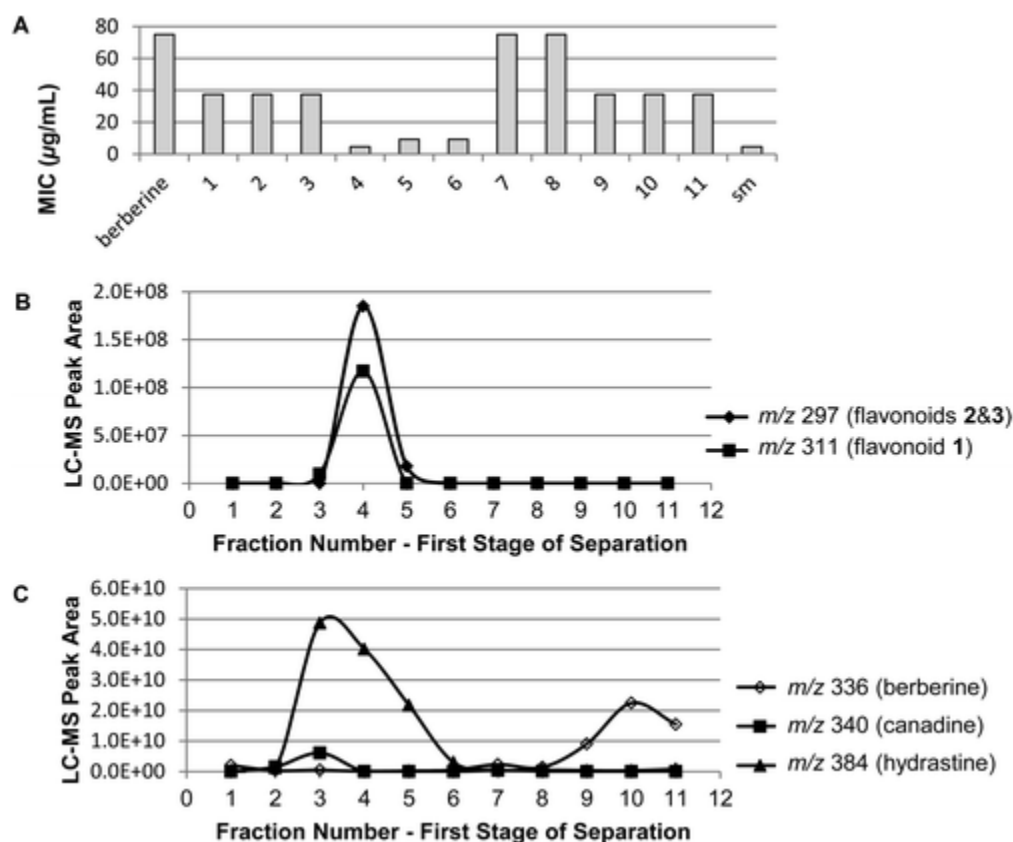


Figure 3. Panel A shows minimum inhibitory concentration (MIC) of starting material (sm), berberine alone, and berberine in combination with 11 fractions (each at a fixed concentration of 75 $\mu\text{g/mL}$) from the first stage of separation (flash chromatography over silica gel with $\text{CHCl}_3/\text{MeOH}$ gradient) of the *Hydrastis canadensis* leaf extract. The MIC of berberine (75 $\mu\text{g/mL}$) was reduced significantly by combining with the starting material and with several of the fractions, with the most active fraction being fraction 4. Panels B and C show the distribution of flavonoids (measured with negative-ion LC-MS) and alkaloids (measured with positive-ion LC-MS), respectively, in the fractions. Fraction 4 contains the highest levels of the three flavonoids. There is a moderate multiple correlation between MIC and flavonoid peak area ($R = 0.77$, $p = 0.016$).

Fraction 4 was subjected to a second stage of separation using flash chromatography with a hexane/ethyl acetate gradient (see fractionation scheme, Supporting Information Figure S1). Mass spectrometry profiles were compared to MIC data (Figure 4), and again the compounds

with m/z 297 and 311 were present at highest concentration in the most active fraction (subfraction 2). Subfraction 2 was also subjected to the synergy assay (Figure 2), and an even more pronounced synergistic enhancement of the activity of berberine was observed (FIC = 0.03). The second stage of purification yielded multiple active fractions (Figure 4), which explains the weak multiple correlation observed between MIC and peak area of compounds with m/z 297 and 311 for the second stage of separation ($R = 0.55$, $p = 0.37$). This weak correlation implies that the compounds with m/z 297 and 311 do not fully account for the activity of fraction 4 and that, again, additional active compounds were present in this fraction.

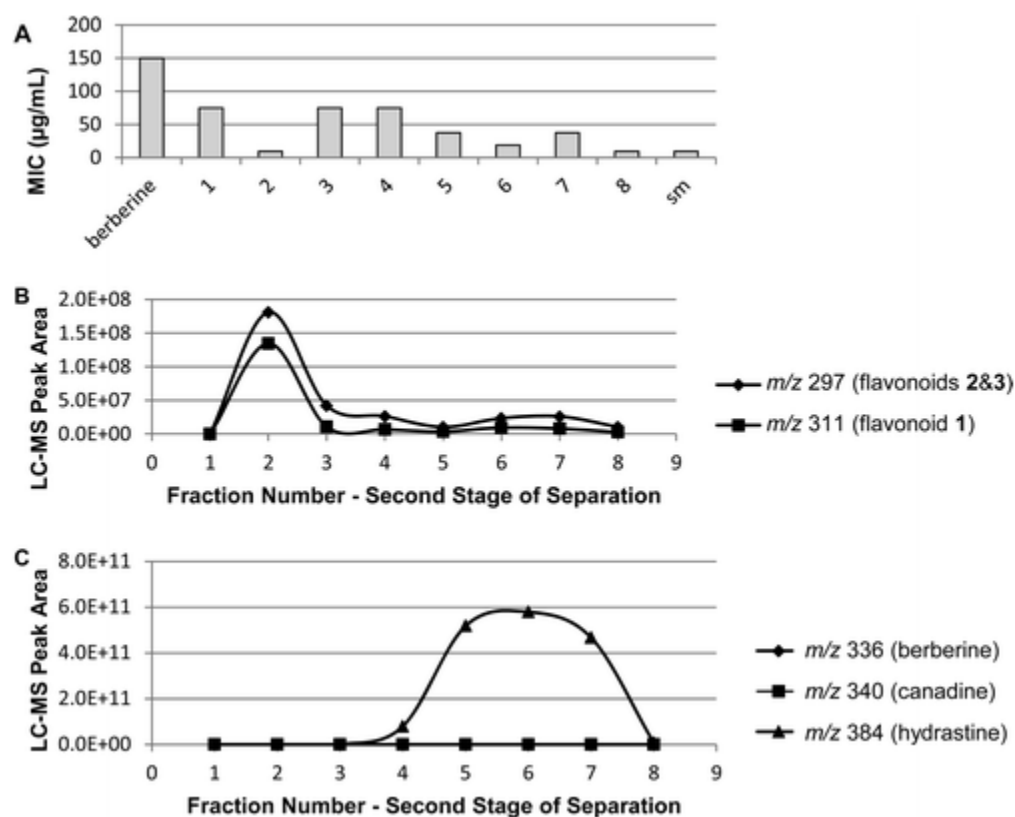
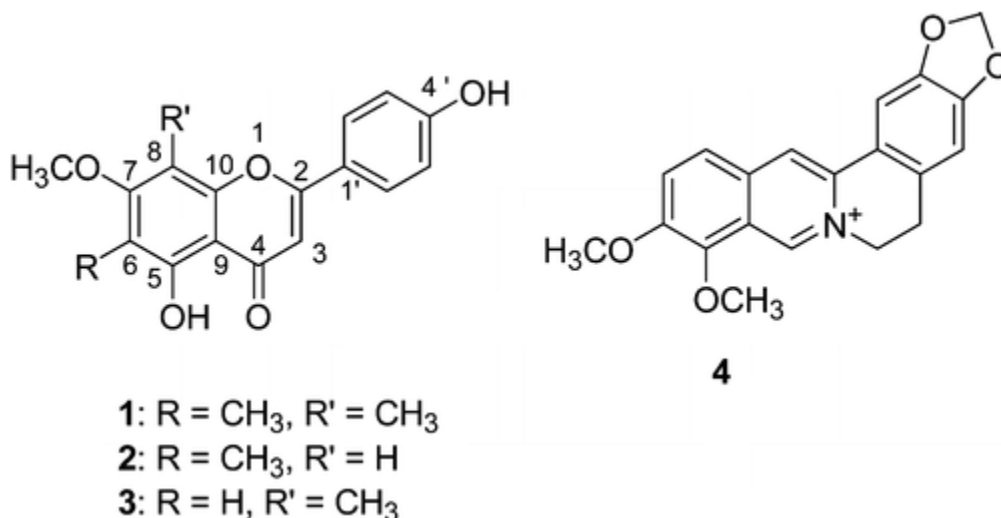


Figure 4. Comparison between MIC (A) and distribution of flavonoids (B) and alkaloids (C) after the second stage of the separation (flash chromatography over silica gel with hexane/EtOAc gradient) of the *Hydrastis canadensis* leaf extract. Subfraction 2 contained the highest levels of the three flavonoids and was one of the most active fractions. Note, however, that other compounds appear to play a role in the activity of the starting material (sm); several fractions with little to no flavonoid content have activity on the same order as subfraction 2. The presence of additional active compounds would also explain the weak multiple correlation between activity and flavonoid content after this second stage of separation ($R = 0.55$, $p = 0.37$).

Two rounds of preparative HPLC starting with subfraction 2 ultimately led to the isolation of the flavonoids sideroxylin (**1**), 6-desmethyl-sideroxylin (**2**), and 6-desmethyl-sideroxylin (**3**). These flavonoids were the same compounds observed in the LC-MS spectra at m/z 311 (sideroxylin) and 297 (isomeric 8-desmethyl-sideroxylin and 6-desmethyl-sideroxylin). The flavonoids have been reported previously as constituents of *Eucalyptus* spp. (**1** and **2**)(24-26) and *Dracaena*

cochinchinensis (**3**, referred to in the reference by the name 4',5-dihydroxy-7-methoxy-8-methylflavone).⁽²⁷⁾ Ours is the first publication to report these flavonoids as constituents of *H. canadensis*.⁽²⁸⁾ Other flavonoids have, however, been identified in *H. canadensis*,⁽⁹⁾ and, as previously mentioned, it is well known that the plant contains the alkaloid berberine (**4**). Both ¹H and ¹³C NMR chemical shift data (Supporting Information, S2) were in excellent agreement with those reported for the flavonoids,⁽²⁴⁻²⁷⁾ and high-resolution mass spectrometry measurements confirmed the molecular formulas of C₁₈H₁₆O₅ for sideroxylin and C₁₇H₁₄O₅ for the isomers 8-desmethyl-sideroxylin and 6-desmethyl-sideroxylin.



Although the crude extract and early stage fractions were quite soluble in the antimicrobial assay medium (Müller-Hinton broth with 2% DMSO), the isolated flavonoids demonstrated very poor solubility (<13 µg/mL) in this same medium. As such, it was not possible to observe an influence of the pure flavonoids on the MIC of berberine against wild-type *S. aureus*. However, activity was observed for 8-desmethyl-sideroxylin against a NorA efflux pump overexpressing strain of *S. aureus* (*S. aureus* K2378, *norA*⁺⁺, Table 2), which is more sensitive to inhibitors. Note that although the reported assay concentration of 8-desmethyl-sideroxylin was 75 µg/mL (Table 2), the actual concentration was likely much lower, due to the aforementioned poor flavonoid solubility. The flavonoid demonstrated no antimicrobial activity alone (MIC >300 µg/mL) against the *norA*⁺⁺ *S. aureus*, but decreased the MIC of berberine 2-fold. Thus, it can be said that 8-desmethyl-sideroxylin (**2**) synergistically enhances, or potentiates, the antimicrobial activity of berberine against *norA*⁺⁺ *S. aureus*.

Table 2. MIC of Berberine Alone and in Combination with 8-Desmethyl-sideroxylin (2) against *S. aureus* K2378 (a *norA*⁺⁺ Strain)^a

treatment	MIC ($\mu\text{g/mL}$)
berberine (4)	300
8-desmethyl-sideroxylin (2)	>300
berberine (4) + 8-desmethyl-sideroxylin (2) (75 $\mu\text{g/mL}$)	150
berberine (4) + reserpine (75 $\mu\text{g/mL}$)	75

^a Reserpine served as a positive control.

Previous literature has shown flavonoids to act as efflux pump inhibitors,(20, 29, 30) and our group has demonstrated efflux pump inhibitory activity for goldenseal leaf extracts.(19) Thus, we hypothesized efflux pump inhibition as a likely mode of action for the *H. canadensis* flavonoids. To test this hypothesis, the activity of each of the three individual flavonoids was evaluated with an ethidium bromide efflux assay (Figure 5). Ethidium bromide is a substrate of the *S. aureus* efflux pump NorA and fluoresces strongly (due to intercalation with DNA) inside bacterial cells.(20) With the ethidium bromide efflux assay, cells are loaded with ethidium bromide and fluorescence is monitored over time. A decrease in fluorescence indicates efflux of ethidium bromide from the cells, and in the presence of an efflux pump inhibitor, fluorescence should decrease more slowly. The ethidium bromide efflux assay has the advantages of being more sensitive than the checkerboard synergy assay and also having a greater tolerance for DMSO (10% DMSO was used in this assay, as compared to 2% DMSO in the checkerboard assays). These features enabled the solubility problems with the pure flavonoids to be circumvented when using the efflux assay.

Figure 5A shows the results of the ethidium bromide efflux assay with wild-type *S. aureus*(NCTC 8325-4). The positive control, carbonyl cyanide *m*-chlorophenylhydrazone (CCCP), and compounds **1**, **2**, and **3** inhibited efflux of ethidium bromide. Significant differences between vehicle control (10% DMSO) and treatment with sideroxylin (**1**) ($p = 0.02$), 8-desmethyl-sideroxylin (**2**) ($p = 0.0002$), 6-desmethyl-sideroxylin (**3**) ($p = 0.002$), and CCCP ($p = 0.0002$) were observed at $t = 300$ s. When the same experiment was repeated using the *norA* deletion mutant *S. aureus* (K1758), neither CCCP nor any of the flavonoids significantly inhibited efflux of ethidium bromide (Figure 5B). The results demonstrate that all three flavonoids from *S. aureus* inhibit the NorA efflux pump of *S. aureus*.

Although all three flavonoids demonstrated some efflux pump inhibitory activity, they differed in potency (Figure 5A). Sideroxylin (**1**), the dimethyl analogue, was a less effective inhibitor than either of the monomethyl analogues, 8-desmethyl-sideroxylin (**2**) and 6-desmethyl-sideroxylin (**3**). It appears that substitution on both the 6 and 8 positions decreases efflux pump inhibitory activity.

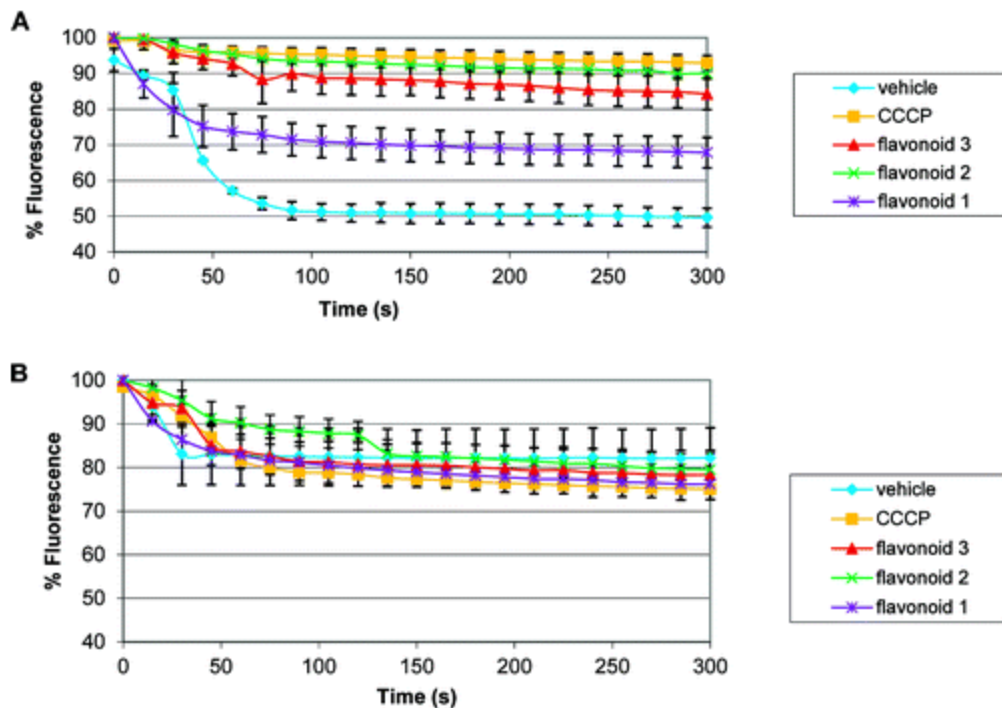


Figure 5. The three flavonoids from *Hydrastis canadensis* inhibit the NorA efflux pump of *Staphylococcus aureus*. Decrease in fluorescence over time is due to efflux of ethidium bromide, which is blocked by the positive control (CCCP) and all three flavonoids, sideroxylin (1), 8-desmethoxy-sideroxylin (2), and 6-desmethoxy-sideroxylin (3). Sideroxylin is the least active of the three flavonoids. The effect is observed with wild-type *Staphylococcus aureus* (NCTC8325-4) (A) but not a NorA deletion mutant (K1758) (B). All of the flavonoids and controls were tested at a final concentration of 100 μ M in Müller-Hinton broth containing 10% DMSO (vehicle). Each point is an average percent fluorescence from three independent measurements, with error bars equal to \pm standard error (SE).

A relevant question to the quality control of dietary supplements from *H. canadensis* aerial portions is whether the flavonoids and alkaloids are present at detectable levels in such preparations. To address this question, ethanolic extracts were analyzed that had been prepared using a method based on that employed in the manufacture of dietary supplements.⁽³¹⁾ Both flavonoids and alkaloids were detected in the ethanolic extracts. Retention times and CID fragmentation patterns for these compounds (see Supporting Information, Table S2) matched those of standards. In agreement with previous literature, the alkaloids were present at higher levels in extracts prepared from goldenseal roots than leaves.⁽¹⁸⁾ Conversely, the flavonoids were detected only at very low levels in the root/rhizome extracts, but at much higher levels in the leaf extracts (Table 3). Goldenseal leaf extracts contained an average of 1.9 ± 0.4 mM 8-desmethoxy-sideroxylin/6-desmethoxy-sideroxylin (quantified as a mixture) and 73.3 ± 0.9 μ M sideroxylin. Levels in the root extracts were more than 50 times lower. Notably, the same leaf extracts for which data are presented in Table 3 have been shown previously to have efflux pump inhibitory activity,⁽¹⁹⁾ which is likely attributable, at least in part, to the presence of flavonoids 1, 2, and 3.

Table 3. Concentrations (\pm SE, $N = 6$) of Alkaloids (Berberine (4), Hydrastine, Canadine) and Flavonoids [Sideroxylin (1), 8-Desmethyl-sideroxylin (2), and 6-Desmethyl-sideroxylin (3)] in Extracts from the Roots/Rhizomes and Leaves of *Hydrastis canadensis*^a

	berberine (mM) \pm SE	hydrastine (mM) \pm SE	canadine (mM) \pm SE	flavonoids 2/3 ^b (mM) \pm SE	flavonoid 1 (mM) \pm SE
goldenseal (<i>H. canadensis</i>) root/rhizome	36.4 \pm 6.4	3.14 \pm 0.50	0.0687 \pm 0.0035	0.0378 \pm 0.011	0.00084 \pm 0.00033
goldenseal (<i>H. canadensis</i>) leaf	7.78 \pm 0.60	1.12 \pm 0.12	0.4500 \pm 0.0040	1.85 \pm 0.41	0.0733 \pm 0.0089

^a Data are averages of measurements made from five separate leaf extracts and five separate root/rhizome extracts (each from a single plant). ^b Flavonoids 2 and 3 were quantified as a mixture because the two isomers were not resolved using reversed-phase HPLC-MS with a C-18 stationary phase.

The finding that goldenseal leaf extracts have higher levels of synergists while root extracts contain higher levels of alkaloids suggests the potential benefit of using a mixture of root and leaf material in the production of dietary supplements from goldenseal. Further studies would, however, be needed to evaluate the safety and efficacy of goldenseal leaf extracts in vivo. If goldenseal leaf material was shown to be efficacious, this could add value to cultivated goldenseal crops. In addition, the use of goldenseal leaf material has the added advantage of reducing impact on wild goldenseal populations. Goldenseal leaves can be harvested in the fall, after the berries have dropped, without killing the plants. Sustainable production of goldenseal would be particularly desirable given that it is listed as threatened by CITES in much of its native habitat.(32)

In terms of the broader relevance of this study, it has been shown that synergy-directed fractionation can be employed to identify both compounds with direct antimicrobial activity (berberine) and efflux pump inhibitors that synergistically enhance berberine's activity (the three flavonoids) from a complex botanical extract. The results highlight three important elements that distinguish synergy-directed fractionation from bioactivity-guided fractionation. First, the application of synergy testing to the crude extract made it possible to determine that it contained synergists even prior to fractionation. Second, integration of synergy testing with the fractionation process allowed synergists to be tracked as they were purified. This process ultimately facilitated the isolation of synergists that possess no inherent antimicrobial activity, compounds that may have been missed in inactive fractions with traditional bioactivity-guided fractionation. Third, comparison of LC-MS data and biological activity data after each stage of separation allowed potential active compounds to be identified and tracked throughout the isolation process. This establishes an important linkage between biological activity of the crude extract and the presence of specific active compounds. In the example presented here, it was possible based on LC-MS profiles to identify ions corresponding to the active flavonoids even after only one stage of separation. Importantly, it was not necessary to have prior knowledge of the identities of the compounds detected in the active fractions to correlate their presence with biological activity. These compounds were initially identified solely on the basis of characteristic m/z values, and it was only after their likely role in the biological activity of the extract had been established that they were isolated and their identities determined via NMR.

Visual inspection and statistical analyses of the data after the first and second stages of separation suggest the presence of multiple active compounds, including some that have not yet been identified. By no means does this finding negate the importance of the flavonoids and alkaloids in the activity of *H. canadensis*. Rather, the results suggest that the antimicrobial activity of the crude *H. canadensis* extract is due to multiple compounds, including antimicrobial alkaloids, flavonoids that synergistically enhance the antimicrobial activity of the alkaloids, and

several additional compounds whose identities and modes of action have yet to be determined. This finding is consistent with the claims often made for botanical medicines: that their activity results from the combined (and perhaps synergistic) action of an array of chemically diverse constituents.

EXPERIMENTAL SECTION

General Experimental Procedures

NMR spectra were acquired with a JEOL ECA-500 spectrometer (500 MHz for ^1H and 125 MHz for ^{13}C) using $\text{DMSO-}d_6$. LC-MS measurements were made using an LCQ Advantage (Thermo) coupled to an Agilent HP1100 HPLC with a Prevail C18 column (3 μm ; 110 \AA , 50 \times 2.1 mm). HRESIMS data were acquired using an LTQ-Orbitrap (Thermo). Flash chromatography separations were accomplished using an automated CombiFlash RF system (Teledyne-Isco). HPLC separations were performed on a Varian HPLC system (ProStar 210 pumps, ProStar 710 fraction collector, ProStar 335 photodiode array detector) with Galaxie Chromatography Workstation software (version 1.9.3.2). Preparative-scale HPLC separations employed a C-18 YMC-Pack ODS-A column (5 μm , 120 \AA ; 250 \times 20 mm, Waters) or a Luna PFP(2) column (pentafluorophenyl propyl, 5 μm , 100 \AA ; 250 \times 21.2 mm, Phenomenex). Analytical-scale HPLC separations were accomplished with YMC ODS-A (5 μm , 120 \AA ; 150 \times 4.6 mm, Waters) and Luna PFP(2) (5 μm , 100 \AA ; 150 \times 4.6 mm, Phenomenex) columns. For antimicrobial assays, OD_{600} was read using a POLARstar Optima microplate reader (BMG Labtech, Inc.). Fluorescence was monitored in efflux assays using a Spex FluoroMax-2 spectrofluorometer (Instruments S. A., Inc.). Müller-Hinton broth, carbonyl cyanide *m*-chlorophenylhydrazone, berberine (**4**) (purity >98% by HPLC), (1*R*,9*S*)-(-)- β -hydrastine (purity >98% by HPLC), and reserpine (purity >98% by TLC) were purchased from Sigma Aldrich, and canadine (tetrahydroberberine, purity >98% by HPLC, stereochemistry unconfirmed) was obtained from Chromadex. Other reagents were purchased from Fisher Chemical, except EtOH (95%) and HPLC-grade MeOH, which were obtained from Pharmco-AAPER. Nanopure water was prepared with a Nanodiamond water purification system (Barnstead).

Plant Material

Individual goldenseal (*Hydrastis canadensis*) plants and bulk leaf material were collected in July 2010 from William Burch in Hendersonville, North Carolina (NC, N 35°24.277', W 082°20.993', 702.4 m elevation). The plants were cultivated in their native environment, a hardwood forest understory, and were at least 5 years old at time of harvest. A voucher specimen was deposited at the Herbarium of the University of North Carolina at Chapel Hill (NCU583414) and authenticated by Dr. Alan S. Weakly. Samples were air-dried at 37 °C before extraction. This temperature was chosen because it was sufficient to dry plant material without causing discoloration of the leaves.

Extraction

Ethanollic *H. canadensis* extracts (50:50 EtOH/H₂O) were prepared via a procedure used for the preparation of botanical dietary supplements, as described previously.⁽¹⁹⁾ Extracts were

prepared separately for the root and leaf material from five individual goldenseal plants (for a total of 10 extracts, five leaf and five root). An additional larger scale extraction (2.3 kg of leaf material) was performed to facilitate isolation. Leaves were homogenized and percolated in MeOH overnight, and the MeOH extract was concentrated in vacuo and subjected to liquid–liquid extraction, as described previously.(33) Briefly, the extract was defatted by partitioning between hexane and 10% aqueous MeOH (1:1). The dried aqueous MeOH fraction was partitioned further between 4:1:5 CHCl₃/MeOH/H₂O, and the organic extract was washed with 1% saline solution to remove hydrosoluble tannins.(34)

Isolation

The isolation scheme is provided as Supporting Information (Figure S1). The first stage of normal-phase flash chromatography (330 g silica gel column) was conducted at a 60 mL/min flow rate with a 110 min hexane/CHCl₃/MeOH gradient. The most active fraction from the first stage separation (fraction 4) was subjected to a second stage of normal-phase flash chromatography (120 g silica gel column), which employed an 85 mL/min flow rate and a 50 min hexane/EtOAc/MeOH gradient. The most active fraction from the second-stage separation (subfraction 2) was purified using reversed-phase preparative HPLC with a YMC ODS-A column at a 8.0 mL/min flow rate. A MeOH/H₂O gradient was employed, which progressed linearly from 70:30 to 85:15 over 60 min. Compound **1** eluted at 44.85 min (22.2 mg, 98.6% purity, 0.00096% yield), and compounds **2** and **3** eluted as a mixture (53.4 mg) between 36.8 and 37.3 min. The mixture of **2** and **3** was subjected to an additional stage of preparative HPLC using a Luna PFP(2) (pentafluorophenyl propyl) stationary phase with a 21.2 mL/min flow rate. An CH₃CN/H₂O gradient was used, initiating at 45:55 (isocratic for 12 min), increasing linearly to 58:42 from 12.1 to 15.0 min, and from 58:42 to 60:40 from 15.1 to 20.0 min. Compound **2** eluted at 17.4 min (14.8 mg, 99% purity, 0.00064% yield), and compound **3** eluted at 15.21 min (2.40 mg, 98.9% purity, 0.0001% yield).

Sideroxylin (1):

white solid; HRESIMS m/z 311.0922 [M – H[–]] (calcd for C₁₈H₁₅O₅[–], 311.0925); ¹H NMR (500 MHz DMSO-*d*₆) and ¹³C NMR (125 MHz DMSO-*d*₆) chemical shifts were in agreement with literature values(24-26) and are provided as Supporting Information (Table S1 and Figure S2).

8-Desmethyl-sideroxylin (2):

white solid; HRESIMS m/z 297.0765 [M – H[–]] (calcd for C₁₇H₁₃O₅[–], 297.0768); ¹H NMR (500 MHz DMSO-*d*₆) and ¹³C NMR (125 MHz DMSO-*d*₆) were in agreement with literature values,(24-26) as indicated by chemical shift data (Table S1) and NMR spectra (Figure S2) provided as Supporting Information.

6-Desmethyl-sideroxylin (3):

white solid; HRESIMS m/z 297.0761 [M – H[–]] (calcd for C₁₇H₁₃O₅[–], 297.0768). ¹H NMR (500 MHz DMSO-*d*₆) and ¹³C NMR (125 MHz DMSO-*d*₆)(7) data were in agreement with

literature(27) and are provided for reference as Supporting Information (Table S1 and Figure S2).

LC-MS

Alkaloids and flavonoids were quantified and identified using LC-MS with an ion trap mass spectrometer. Reversed-phase HPLC was used for the separation with a Prevail C18 column and a gradient employing 1% aqueous acetic acid in acetonitrile. The gradient was as follows (ratios represent CH₃CN/H₂O): 0 to 5 min, 0:100 to 32:68; 5.1 to 20 min, 32:68 to 100:0; 20 to 25 min, isocratic 100:0; 25 to 25.10 min, 100:0 to 0:100. Extracts were analyzed both in the positive-ion mode (for detection and quantification of alkaloids) and the negative-ion mode (for detection and quantification of flavonoids). Standards were analyzed over a concentration range of 7.8×10^{-7} to 1.0×10^{-4} M (flavonoids) and 5.0×10^{-7} to 1.0×10^{-4} M (alkaloids). Calibration curves were plotted as the area of the relevant selected ion trace (as detected by the LC-MS) versus concentration. Extracts were diluted so that constituents fell within the linear range of the calibration curves. The alkaloids berberine, hydrastine, and canadine were identified by comparison of retention time, *m/z*, and CID fragmentation patterns with those of commercial standards (see Table S2 provided as Supporting Information), as described previously.(19) The flavonoids were identified in the same fashion using compounds **1**, **2**, and **3**, isolated as part of this study, as standards.

Antimicrobial Assays

Two strains of *S. aureus* were employed, wild-type (NCTC 8325-4)(35) and NorA overexpressing (K2378, *norA*⁺⁺).(36) Minimum inhibitory concentration was evaluated according to Clinical Laboratory Standards Institute guidelines.(37) A single colony inoculum of *S. aureus* was grown to log phase in Mueller-Hinton broth and was adjusted to a final assay dilution of 1.0×10^5 cfu/mL based on absorbance at 600 nm (OD₆₀₀). The negative control consisted of 2% DMSO in broth (vehicle), and the known efflux pump inhibitor reserpine(38) served as the positive control. Triplicate wells were prepared with all treatments and controls. Additional wells were included containing the samples without bacteria, for the purpose of background subtraction. OD₆₀₀ was read after incubation for 20 h at 37 °C. MIC was defined as the concentration at which no statistically significant difference was observed (via ANOVA) between the negative control and treated samples.

Synergy Testing

Broth microdilution antimicrobial checkerboard assays(19, 22) were performed to evaluate synergy for the crude extracts. Extracts were tested in combination with berberine over a concentration range of 5 to 300 µg/mL. The antibiotic norfloxacin (positive control) demonstrated an MIC value of 2 µg/mL, consistent with the literature.(39) The vehicle control consisted of 2% DMSO in broth. To evaluate synergistic interactions, isobolograms were plotted and FIC indices calculated, as described previously.(19) After each round of fractionation, a simplified version of the synergy assay was used to enable rapid screening of the large number of fractions generated. This involved testing berberine at a range of concentrations (5 to 300 µg/mL) in combination with a constant concentration of each of the extract fractions (75 µg/mL).

The known efflux pump inhibitor reserpine,(38) also at 75 µg/mL, was employed as a positive control for these experiments. Fractions were deemed active if they enhanced the activity of berberine (reducing its MIC against *S. aureus* by at least 2-fold) and were advanced to the next stage of separation. Using the simplified synergy assay, it is not possible to distinguish synergistic effects from additive effects, so the most active fractions were also subjected to the synergy checkerboard assay as described for the crude extracts.

Efflux Pump Inhibition Assay

The efflux pump inhibition assay was conducted using a previously published method.(19) Wild-type *S. aureus* (NCTC 8325-4)(35) and an isogenic NorA deletion mutant (K1758 *norA*⁻)(40) were grown to OD₆₀₀ = 0.74 in Mueller-Hinton broth. Ethidium bromide (25 µM) and carbonyl cyanide *m*-chlorophenylhydrazone (100 µM) were added, and the bacteria were incubated at 20 °C with agitation (300 rpm) for 20 min. This solution was diluted to OD₆₀₀ = 0.40 with broth containing ethidium bromide and CCCP at 25 and 100 µM, respectively. Aliquots (1 mL) of this solution were centrifuged at 20 °C for 5 min at 13000g in a Spectrafuge 24D centrifuge (Labnet). The pellets were stored on ice for <1 h. Prior to measurement, the pellets were thawed for 5 min and 1 mL of fresh broth containing DMSO (10% final assay concentration) and treatments (CCCP or flavonoid) was added. Fluorescence of these solutions was measured every second for 300 s with $\lambda_{\text{ex}} = 530 \text{ nm}$, $\lambda_{\text{emiss}} = 600 \text{ nm}$, and slit widths of 5 mm.

Statistics

Statistical significance of means versus control was evaluated using two-factor ANOVA with $p < 0.05$ considered significant. Multiple (canonical) correlations were computed to assess the degree of association of MIC jointly with the peak area of several ions (flavonoid **1**, m/z 311, and the flavonoid **2/3** mixture, m/z 297). The peak areas appeared to be non-normally distributed, and sample sizes were small, so p -values for assessing the statistical significance of the multiple correlations were calculated using permutation tests.

SUPPORTING INFORMATION

¹³C and ¹H NMR data, a fractionation scheme, LC-MS fragmentation data, and retention times for alkaloids and flavonoids.

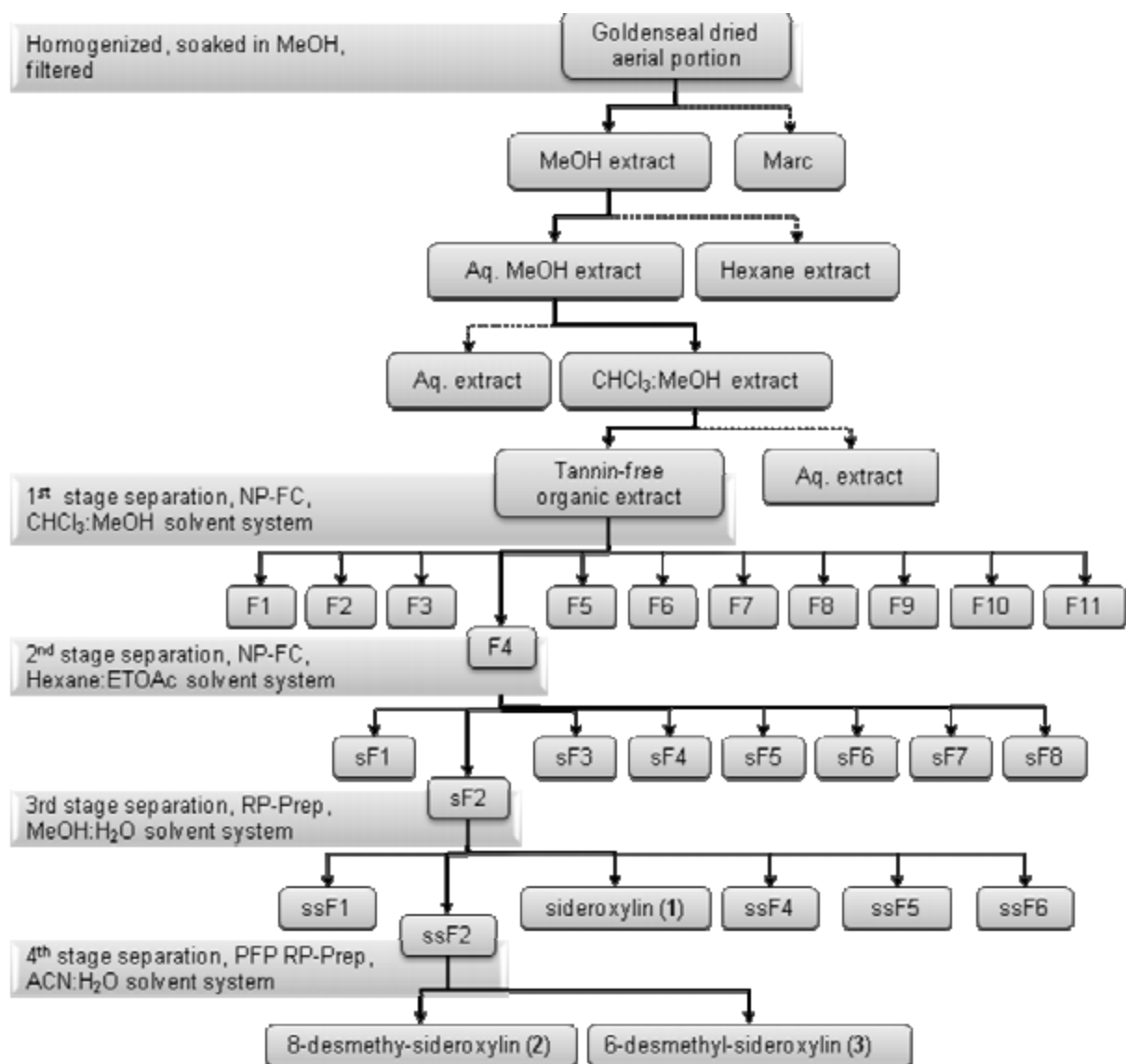


Figure S1. Fractionation scheme. *H. canadensis* leaves were percolated in MeOH, subjected to liquid/liquid partitioning, and separated using four subsequent stages of chromatography, two rounds of flash chromatography over silica gel, followed by two rounds of preparatory scale HPLC. After each round of separation, each fraction was profiled using LC-MS and tested for antimicrobial activity, as shown in Figures 3 and 4 of the main text.

Table S1. NMR spectroscopic data (500 MHz for ^1H and 125 MHz for ^{13}C , DMSO-d₆) for *H. canadensis* flavonoids sideroxylin (1), 8-demethyl-sideroxylin (2), and 6-demethyl-sideroxylin (3).

Position #	1		2		3	
	δ_{H} , m (J in Hz)	δ_{C}	δ_{H} , m (J in Hz)	δ_{C}	δ_{H} , m (J in Hz)	δ_{C}
2	-	164.2	-	163.8	-	163.6
3	6.89, s	102.7	6.82, s	103	6.80, s	101.4
4	-	182.6	-	181.9	-	181.8
5	-	156.3	-	157.5	-	159.0
6	-	113.1	-	107.5	6.50, s	94.5
7	-	162.1	-	163	-	162.1
8	-	108.7	6.83, s	90.3	-	102.5
9	-	152.3	-	155.4	-	153.2
10	-	103.4	-	104.4	-	103.5
1'	-	120.4	-	121	-	123.9
2'	7.96, d (10)	128.6	7.93, d (9)	128.5	7.93, d (9)	128.1
3'	6.94, d (10)	116.2	6.89, d (9)	116	6.90, d (9)	115.9
4'	-	161.7	-	161.5	-	162.1
5'	6.94, d (10)	116.2	6.89, d (9)	116	6.90, d (9)	115.9
6'	7.96, d (10)	128.6	7.93, d (9)	128.5	7.93, d (9)	128.1
6-CH ₃	2.10, s	8.1	1.97, s	7.3	-	-
8-CH ₃	2.34, s	8.3	-	-	2.20, s	7.2
7-OCH ₃	3.75, s	60.4	3.89, s	56.3	3.87, s	55.8

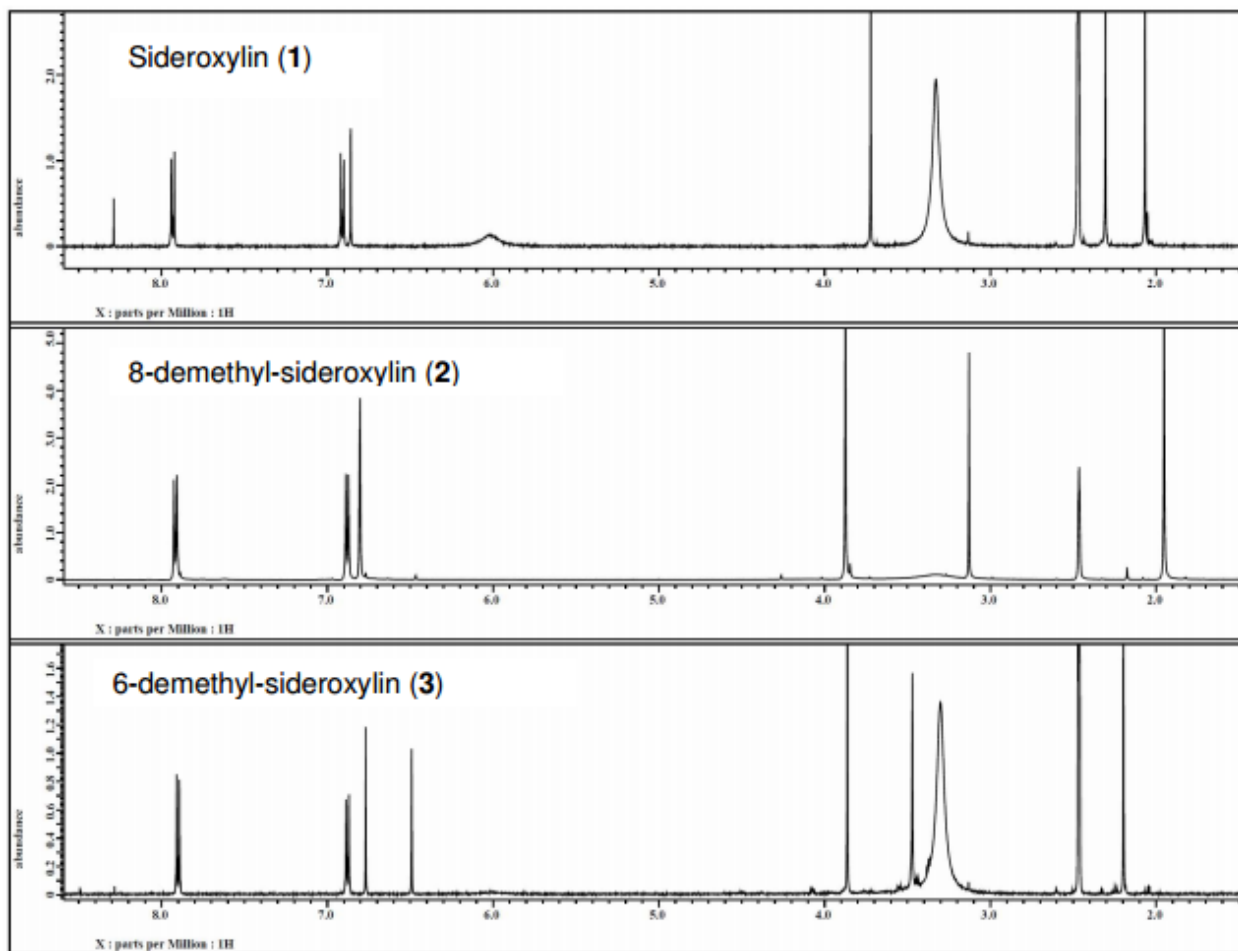


Figure S2. ¹H NMR spectra (500 MHz, DMSO-d₆) for *H. canadensis* flavonoids **1**, **2**, and **3**.

Compound	t _R ^a (min)	molecular ion (m/z)	CID Fragment Ions ^c
Sideroxylin (1)	16.92	311.14 (M-H ⁻)	296.14, 311.11, 150.48, 183.05, 228.23
8-demethyl-sideroxylin (2)	16.26	297.15 (M-H ⁻), 299.21 (MH ⁺)	297.10, 282.16, 300.17, 347.44
4',5-dihydroxy-7-methoxy-8-methylflavone (3)		297.15 (M-H ⁻), 299.21 (MH ⁺)	Co-elutes with 8-demethyl-sideroxylin, indistinguishable fragmentation
Berberine (4) ^d	10.65	336.11 (M ⁺)	321.08, 336.07, 292.21, 306.14
Hydrastine (5)	7.28	384.02 (MH ⁺)	322.99, 190.11, 335.00, 366.01
Canadine (6)	9.45	340.18 (MH ⁺)	176.10, 340.09, 149.02

- a. The retention time is for a reversed phase HPLC separation with an CH₃CN:H₂O gradient.
- b. The flavonoids (**1**, **2**, and **3**) could be detected both as MH⁺ in the positive ion mode and M-H⁻ in the negative ion mode.
- c. Berberine is inherently charged, so is detected as M⁺ rather than MH⁺
- d. CID fragment ions for the alkaloids are for the positive ion mode using the MH⁺ ion (or M⁺ in the case of berberine) as the precursor and 35% relative collision energy. CID fragment ions for flavonoids are for the negative ion mode using the M-H⁻ ion as the precursor. Ions are shown in order of decreasing relative abundance.

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