

Medicinal plant extracts with efflux inhibitory activity against Gram-negative bacteria

Mark I. Garvey, Mukhlesur Rahman, Simon Gibbons, Laura J.V. Piddock

► To cite this version:

Mark I. Garvey, Mukhlesur Rahman, Simon Gibbons, Laura J.V. Piddock. Medicinal plant extracts with efflux inhibitory activity against Gram-negative bacteria. International Journal of Antimicrobial Agents, 2011, 37 (2), pp.145. 10.1016/j.ijantimicag.2010.10.027 . hal-00659900

HAL Id: hal-00659900 https://hal.science/hal-00659900

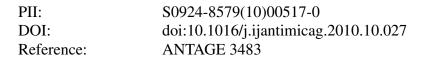
Submitted on 14 Jan2012

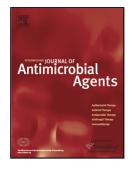
HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

Accepted Manuscript

Title: Medicinal plant extracts with efflux inhibitory activity against Gram-negative bacteria

Authors: Mark I. Garvey, Mukhlesur Rahman, Simon Gibbons, Laura J.V. Piddock





To appear in: International Journal of Antimicrobial Agents

 Received date:
 31-8-2010

 Revised date:
 21-10-2010

 Accepted date:
 22-10-2010

Please cite this article as: Garvey MI, Rahman M, Gibbons S, Piddock LJV, Medicinal plant extracts with efflux inhibitory activity against Gramnegative bacteria, *International Journal of Antimicrobial Agents* (2010), doi:10.1016/j.ijantimicag.2010.10.027

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

Medicinal plant extracts with efflux inhibitory activity against Gramnegative bacteria

Mark I. Garvey ^a, Mukhlesur Rahman ^b, Simon Gibbons ^{b,*}, Laura J.V. Piddock ^{a,*}

 ^a Antimicrobial Agents Research Group, School of Immunity and Infection, College of Medical and Dental Sciences, University of Birmingham, Birmingham B15 2TT, UK
 ^b Department of Pharmaceutical and Biological Chemistry, The School of Pharmacy, University of London, 29–39 Brunswick Square, London WC1N 1AX, UK

ARTICLE INFO

Article history:

Received 31 August 2010

Accepted 22 October 2010

Keywords:

Efflux pump inhibitor

Antibiotic resistance

Synergistic interactions

* Corresponding authors. Tel.: +44 121 414 6966; fax: +44 121 414 6819 (L.J.V.

Piddock); tel.: +44 207 753 5913; fax: +44 207 753 5964 (S. Gibbons).

E-mail addresses: I.j.v.piddock@bham.ac.uk (L.J.V. Piddock);

simon.gibbons@pharmacy.ac.uk (S. Gibbons).

ABSTRACT

It was hypothesised that extracts from plants that are used as herbal medicinal products contain inhibitors of efflux in Gram-negative bacteria. Extracts from 21 plants were screened by bioassay for synergy with ciprofloxacin against Salmonella enterica serotype Typhimurium, including mutants in which acrB and tolC had been inactivated. The most active extracts, fractions and purified compounds were further examined by minimum inhibitory concentration testing with five antibiotics for activity against Enterobacteriaceae and Pseudomonas aeruginosa. Efflux activity was determined using the fluorescent dye Hoechst 33342. Eighty four extracts from 21 plants, 12 fractions thereof and 2 purified molecules were analysed. Of these, 12 plant extracts showed synergy with ciprofloxacin, 2 of which had activity suggesting efflux inhibition. The most active extract, from Levisticum officinale, was fractionated and the two fractions displaying the greatest synergy with the five antibiotics were further analysed. From these two fractions, falcarindiol and the fatty acids oleic acid and linoleic acid were isolated. The fractions and compounds possessed antibacterial activity especially for mutants lacking a component of AcrAB-TolC. However, no synergism was seen with the fractions or purified molecules, suggesting that a combination of compounds is required for efflux inhibition. These data indicate that medicinal plant extracts may provide suitable lead compounds for future development and possible clinical utility as inhibitors of efflux for various Gram-negative bacteria.

1. Introduction

Over the last decade there has been a dramatic reduction in the number of pharmaceutical companies developing new antimicrobial agents [1]. In parallel, the number of antibiotic-resistant Gram-negative bacteria has increased [2]. With the reduction in the number of new agents and in antibiotic development, there has been a resurgence of interest in the search for compounds that will restore the activity of licensed antimicrobial agents that until recently had excellent activity against Gramnegative bacteria. In 1998 it was shown that plant-derived compounds have activity against Gram-positive bacteria, in particular Staphylococcus aureus [3]. Several compounds, such as reserpine, behave as if they inhibit efflux pumps and hence have become known as efflux pump inhibitors (EPIs) [4]. Since that time, numerous phytochemicals have been shown to have activity against S. aureus or other Grampositive bacteria, or to act as potential EPIs with antimicrobials for Gram-positive bacteria [5–11]. Unfortunately, most of the plant-derived compounds have little or no activity with antibiotics against Gram-negative bacteria and it was suggested that, as many plant pathogens are Gram-negative bacteria, plants may not produce molecules effective against these organisms [5]. None the less, there are many medicinal plants that have traditional usage in various parts of the world in the treatment of infections caused by Gram-negative bacteria [12,13].

Gram-negative bacteria have innate multidrug resistance to many antimicrobial compounds owing to the presence of efflux pumps. In the Enterobacteriaceae, the efflux pump most commonly associated with this innate multidrug resistance is the AcrAB–ToIC efflux system.

Homologues of this pump are found in other Gram-negative bacteria, including *Pseudomonas aeruginosa, Acinetobacter* spp. and *Campylobacter jejuni* [14]. The presence of these pumps and their broad substrate profile is the cause of the innate resistance to many of the agents that have good antimicrobial activity against Grampositive bacteria. Overproduction of these efflux pumps confers clinically relevant resistance to many antimicrobial agents, including ciprofloxacin and tigecycline, in Enterobacteriaceae [14].

We hypothesised that extracts from plants that are used as natural herbal medicines contain molecules that act as EPIs against the efflux pumps of Gram-negative bacteria. Therefore, 84 extracts from 21 medicinal plants were investigated for synergistic activity with ciprofloxacin against *Salmonella enterica* serotype Typhimurium. These plants were chosen as they all have a use in various systems of medicine and some are used as topical antimicrobials, for example *Melissa officinalis*. Extracts were screeened for differential activity against strains that either expressed or overproduced the AcrAB–TolC pump but had no activity for mutants in which a component of this pump had been inactivated. In this way, it was hoped to identify extracts that contained an EPI specific for this pump. For those extracts where data suggested active components, the effect upon growth of *Salmonella* spp. as well as inhibition of efflux activity was determined. Two extracts showing the greatest synergy with ciprofloxacin were fractionated and the active molecules were identified.

2. Materials and methods

2.1. Bacterial strains, storage and growth

All bacteria used in this study are listed in Table 1. Construction of mutants derived from *S*. Typhimurium has been described previously [16,19,21,22]. *Salmonella* Typhimurium L3 is a pre-therapy human clinical isolate and L10 is a post-therapy human clinical isolate that overexpresses *acrAB*, both of which have been previously described [20,23]. All bacteria were stored on Protect[™] beads (Technical Service Consultants Ltd., Heywood, UK) at –80 °C. Identification of each species was confirmed by Gram-stain and analytical profile index (API 20E; bioMérieux, Marcy-l'Étoile, France).

2.2. Media and chemicals

Bacteria were grown on Luria–Bertani (LB) agar plates (Oxoid Ltd., Basingstoke, UK) and in LB broth (Oxoid Ltd.). All antibiotics and other compounds were obtained from Sigma (Poole, UK).

2.3. Extraction of plants

In total, 21 plants were used in this study (Table 2). All plants except *Catha edulis* (purchased locally in London, UK) were purchased from Proline Botanicals (Essendine, UK). Each plant (200–400 g) was extracted at 21 °C sequentially with 300–600 mL of chloroform, methanol and water.

2.4. Fractionation of extracts

Following assessment of bioactivity, the chloroform extract of *Levisticum officinale* (10 g) was fractionated by vacuum liquid chromatography (VLC) on silica gel 60G (Merck Ltd., Lutterworth, UK) using solvent systems of increasing polarity. Fractions were eluted with hexane containing increasing 10% or 20% increments of ethyl acetate (EA) to give fractions A–H (A, 10% EA; B, 20% EA; C, 30% EA; D, 40% EA; E, 50% EA; F, 60% EA; G, 80% EA; and H, 100% EA). Fraction I was eluted with 5% methanol in EA, fraction J with 10% methanol in EA, fraction K with 20% methanol in EA and finally fraction L with 50% methanol in EA.

2.5. Purification of active compounds

Bioassay-directed isolation indicated that the VLC fraction eluted with 30% EA was active. This was purified by preparative thin-layer chromatography (Si Gel; Merck) (mobile phase 20% EA in toluene; double development) yielding falcarindiol (**1**; 450 mg) and levistolide A (**2**; 69 mg) (Fig. 1), which were identified by comparison with literature data [24,25].

2.6. Bioassay screening

Screening with ciprofloxacin in the absence or presence of plant extracts (Table 2), fractions of *L. officinale* and *M. officinalis*, and purified compounds from fractions of the chloroform extract of *L. officinale* for synergistic activity was carried out by bioassay essentially as described previously [26,27]. Six rows of 6 wells (total 36 wells) each of 8-mm diameter were cut out of each agar plate using an agar cutter (Fisher Scientific, Loughborough, UK). Then, 200 μ L of sterile Iso-SensitestTM broth

(Oxoid Ltd.) containing ciprofloxacin (0.5, 1 and 2 mg/L) was added to three sets of wells. This volume and concentration gave a zone of inhibition \geq 15 mm in diameter. To determine test compound synergism with ciprofloxacin, 200 µL of a solution containing ciprofloxacin (0.5, 1 and 2 mg/L) and the plant extract (100 mg/L) in broth was added to three separate wells. The plates were incubated overnight at 37 °C and the diameters of the inhibition zones were measured. Each experiment contained three technical repeats and was repeated on three separate days. A zone of inhibition of the antibiotic plus test plant extract larger than that of the antibiotic alone was taken to suggest synergy. Positive controls were ciprofloxacin plus L-phenylalanyl-L-arginyl- β -naphthylamide (PA β N) (100 mg/L) or carbonyl cyanide *m*-chlorophenylhydrazone (CCCP) (100 µM), and negative controls were Iso-Sensitest broth alone and plant extract alone to determine whether the plant extract had inherent antimicrobial activity.

2.7. Determination of the minimum inhibitory concentration (MIC) of antibiotics \pm synergising extracts, fractions and compounds

The MICs of ciprofloxacin, tetracycline, chloramphenicol, erythromycin and ethidium bromide (EtBr) in the absence or presence of plant extracts and fractions (at 100 mg/L) were determined by the agar doubling dilution method and by the microbroth dilution method [28,29]. MIC determinations were repeated at least three times in independent experiments. Antibiotics, compounds and EPIs were made up and used according to the manufacturer's instructions.

Page 7 of 35

2.8. Determination of the fractional inhibitory concentration index (FICI)

Activity of antibiotics plus the plant extract/fraction/purified compound was determined using the checkerboard technique [30]. The first antibiotic (ciprofloxacin) of a combination was serially diluted along the ordinate of a 96-well microtitre tray; the second agent (e.g. plant extract) was diluted along the abscissa. An inoculum equal to a 0.5 McFarland turbidity standard was prepared from each strain in LB broth. Each well in the microtitre tray was inoculated with 100 mL of the bacterial suspension (at 5×10^5 colony-forming units/mL) and the plates were incubated at 37 °C for 24 h under aerobic conditions. The resulting checkerboard contained each combination of ciprofloxacin and test agent with wells that contained the highest concentration of each agent at opposite corners. The MIC was defined as the lowest concentration of antibiotic that completely inhibited growth of the organism as detected with the naked eye [28]. Synergy was defined as an FICI (FICI) of ≤ 0.5 , no interaction was defined as an FICI of >0.5-4 and antagonism was defined as an FICI >4 [31].

2.9. Accumulation of Hoechst 33342 \pm plant extracts by Salmonella Typhimurium Efflux activity of five S. Typhimurium strains [L3, L10, SL1344, ATCC 15277 (LT2) and ATCC 14028s] was determined by measuring accumulation of the fluorescent dye Hoechst 33342 (bisbenzimide; 2.5 μ M) in the absence or presence of known efflux inhibitors (CCCP 100 μ M and PA β N 100 mg/L) and in the absence or presence of those plant extracts (at 100 mg/L) revealed to have activity in the bioassay and MIC tests. Measurements were taken at excitation and emission wavelengths of 350 nm and 460 nm, respectively, over 30 min using a FLUOstar

OPTIMA (BMG Labtech, Aylesbury, UK) as previously described [32]. Differences in accumulation in the absence of the plant extracts were analysed for statistical significance using the two-tailed Student's *t*-test. *P*-values \leq 0.05 were considered significant.

2.10. Growth kinetics ± plant extracts

The growth kinetics of L3, L10, SL1344, ATCC 15277 (LT2) and ATCC 14028s in the absence or presence of plant extracts was determined by monitoring the optical density at 600 nm every 10 min at 37 °C for 24 h using a FLUOstar OPTIMA (BMG Labtech). *Levisticum officinale* chloroform extract and *M. officinalis* methanol extract (at 100 mg/L) were added to the bacterial cultures at mid-logarithmic growth phase (2 h). Samples were also removed and visualised microscopically to detect any gross changes to cell morphology, i.e. filamentation. A two-tailed Student's *t*-test was used to compare the generation times. *P*-values ≤ 0.05 were considered significant.

3. Results

3.1. Medicinal plant extracts synergise with ciprofloxacin

Plants were extracted with chloroform, methanol and water and were screened to identify those that showed synergy with ciprofloxacin and could be putative EPIs.

Ciprofloxacin was chosen as increased MICs of this agent are often associated with overproduction of the AcrAB–ToIC pump. The positive control was PA β N at 100 mg/L, which was the lowest concentration to elicit an effect in this assay. Initial

experiments were performed with *Salmonella* Typhimurium as a model Gramnegative bacterium and because a defined isogenic set of mutants that lack or overproduce the AcrAB–TolC pump was available. Eighty-four extracts from 21 plants were screened for synergistic activity with ciprofloxacin against *S*. Typhimurium SL1344, ATCC 15277 (LT2), ATCC 14028s, L3 and L10. Of these, 12 extracts showed synergy with ciprofloxacin for these strains, whereas 72 showed little or no activity.

The most active antibiotic-potentiating extracts were those from *L. officinale* (extracted with chloroform, methanol and water), *M. officinalis* (chloroform and methanol), *Cinnamomum aromaticum* (chloroform), *Pimpinella anisum* (chloroform) and *Pulmonaria officinalis* (chloroform and methanol) (Table 3) as well as the chloroform extracts of *Angelica sinensis*, *Tussilago farfara* (methanol) and *Commiphora molmol* (chloroform) (data not shown). Of the extracts investigated, those from *M. officinalis* and *L. officinale* were the most active against strains that either had wild-type levels of AcrAB–TolC or that overproduced AcrAB (Table 4). Chloroform and methanol extracts of *L. officinale* and *M. officinalis* (both 100 mg/L) had no effect on the growth kinetics of the five *Salmonella* strains tested (data not shown), and by MIC determination the extract of *L. officinale* had synergistic activity with ciprofloxacin against most of the strains (Table 4).

3.2. Plant extracts synergise with several antibiotics

Owing to their promising activity, the MICs of the chloroform extract of *L. officinale* and the methanol extract of *M. officinalis* (i.e. those extracts that gave the largest zones of inhibition in the bioassay) were determined in combination with tetracycline,

chloramphenicol, erythromycin and EtBr. All extracts showed better activity in liquid media than in agar, with MIC values one to two dilutions lower from the microbroth dilution procedure than with agar (data not shown). No antimicrobial activity was detected at 100 mg/L of either extract; likewise, chloroform and methanol had no activity at the concentration used. Synergy between the extracts from *M. officinalis* and L. officinale with tetracycline, chloramphenicol, erythromycin or EtBr was seen for fewer strains than when the plant extracts were combined with ciprofloxacin. None the less, EPI-like activity was detected for both extracts combined with tetracycline, erythromycin and EtBr against wild-type Salmonella and the strain in which AcrAB was overproduced. The MIC experiments with the chloroform extract of L. officinale indicated that it had synergistic activity with ciprofloxacin and tetracycline for the salmonellae and two Klebsiella pneumoniae strains (Table 4). Of interest, the chloroform extract of L. officinale also had synergistic activity with ciprofloxacin and EtBr for Salmonella Typhimurium in which ToIC was not produced and with EtBr for S. Typhimurium in which AcrAB was not produced (Table 4). The M. officinalis extract had synergistic activity with most antibiotics for the salmonellae and K. pneumoniae strains (Table 4).

The extracts of the other plants showing synergistic activity with ciprofloxacin in the bioassay did not reduce the MIC of ciprofloxacin as much as the extracts from *L. officinale* or *M. officinalis*, and when synergy was observed it was seen for fewer bacterial strains. For these reasons, no further experiments were performed with these extracts.

3.3. Activity of fractions derived from the chloroform extract of Levisticum officinale As the chloroform extract of *L. officinale* showed synergistic activity with more agents and bacteria than that from *M. officinalis*, it was fractionated as the first step to the identification of the active component(s). The chloroform extract of *L. officinale* was subjected to chromatography and 12 fractions (A–L) were obtained. MICs of ciprofloxacin and tetracycline in the absence or presence of each fraction (100 mg/L) were determined. Compared with the chloroform extract of *L. officinale*, few of the fractions had the same synergistic activity with ciprofloxacin. Of interest, fractions A, C, F, G and K synergised with ciprofloxacin for the *Salmonella* mutants in which AcrAB and TolC were not produced, and A, C, G and K also synergised with ciprofloxacin for the two *K. pneumoniae* strains. Fractions A, G and K also had synergistic activity with ciprofloxacin for wild-type *Salmonella* and the clinical isolate that overproduced AcrAB (Table 5). Two fractions (C and F) also had antimicrobial activity alone for the mutant *Salmonella* in which a component of AcrAB–TolC was not produced (Table 5); this was most marked for fraction C.

3.4. Activity of compounds purified from fraction C of Levisticum officinale

As fraction C showed synergistic activity in combination with ciprofloxacin and also antimicrobial activity alone especially against the strains in which *acrB* and *tolC* had been inactivated, this fraction was analysed to identify the active compounds. Nuclear magnetic resonance (NMR) revealed that it contained predominantly the antibacterial fatty acids oleic acid and linoleic acid plus falcarindiol and levistolide A (Fig. 1). These compounds were further purified and the spectral data were in close agreement to those published [24,25]. The absolute stereochemistry of falcarindiol

Page 12 of 35

was determined using a modified Mosher's ester method as the 3(S), 8(S) diastereomer (Fig. 1).

MICs of linoleic acid and falcarindiol were >500 mg/L for the wild-type *Salmonella* SL1344 and ATCC 14028s. For the ToIC mutants of these strains the MIC of linoleic acid was 500 mg/L, and the MIC of falcarindiol was 75 mg/L and 125 mg/L, respectively. When linoleic acid was combined with ciprofloxacin MICs of 0.008 mg/L and 0.004 mg/L were obtained for the ToIC mutants, and for falcarindiol in combination with ciprofloxacin MIC values <0.001 mg/L were obtained.

Synergistic activity between ciprofloxacin and the compounds for the mutants was confirmed in checkerboard assays, with FICI values of 0.14–0.27 for the combination and the TolC mutants (Table 6). The FICI for ciprofloxacin in combination with the methanol-soluble extract from *M. officinalis* against L3 (pre-therapy isolate) was lower than that for L10 (that overexpresses *acrB*). This suggests that a higher concentration of the putative EPI is required to inhibit growth of L10 owing to the greater production of AcrB by L10. No synergy was seen with this combination for the parental strains, despite it being clearly demonstrated with the extract itself. These data suggest that an EPI-like molecule(s) was lost during the separation process or was inactivated during purification.

3.5. Efflux activity of plant extracts

Accumulation of Hoechst 33342 has been used as a marker for efflux as it is fluorescent and is a substrate of several efflux pumps including AcrAB–TolC [17,32]. An increase in the amount of Hoechst 33342 accumulated in the presence of an EPI

such as PAβN indicates inhibition of efflux [17]. To determine whether synergistic activity displayed by the extracts of *L. officinale* with various antibiotics was due to inhibition of efflux, accumulation of Hoechst 33342 in the presence and absence of 100 mg/L of the chloroform and methanol extracts of *L. officinale* was carried out for five strains of *S.* Typhimurium [SL1344, ATCC 15277 (LT2), ATCC 14028s, L3 and L10]. For the four antibiotic-susceptible wild-type strains of *S.* Typhimurium, both extracts of *L. officinale* increased the amount of Hoechst 33342 accumulated (i.e. reduced efflux) by 1.5–2-fold (Fig. 2). The efflux activity of the strain that overexpressed AcrB (L10) was also reduced by 1.5-fold.

4. Discussion

Many conventional drugs have arisen from natural products, including plants. However, few antimicrobial agents have come from this source, with the vast majority in clinical use derived from products naturally produced by microorganisms [33]. None the less, the use of medicinal and herbal plant remedies to treat infectious diseases is common in many countries [34]. Owing to the continued clinical pressure for novel approaches to combat antibiotic-resistant bacterial infections, there is a need to identify new agents to treat such infections. One approach is to screen for natural products from plants. Plants have already successfully yielded compounds with activities suggesting that they inhibit efflux pumps of Gram-positive bacteria [3,5]. Therefore, in this study we sought to identify medicinal plants that could provide compounds for further antimicrobial drug development. In addition, as there are many clinically licensed antibacterial agents for the treatment of infections by Gram-negative bacteria, but which are effluxed by the various pumps possessed by these bacteria, we sought to screen for activity that suggested efflux inhibition. One

desirable property of a putative EPI is that it should synergise with antibiotics for bacteria with wild-type or overproduction of AcrAB-ToIC but have no affect on a strain in which AcrAB or ToIC are produced. As ciprofloxacin is a substrate of many bacterial efflux pumps [14], the initial experiments used a simple bioassay to identify plant extracts that synergised with this agent [27]. This allowed many plant extracts made under different conditions to be screened with speed. In a prior study, we sought to identify compounds with EPI-like activity and to use this as an aid to predict the common structural features of EPIs and so guide the type of plant from which to make extracts for this study [27]. Of the 84 extracts from the 21 plants, 2 extracts, an ethanol extract from *M. officinalis* (lemon balm) and a chloroform extract of L. officinale (lovage root), had the greatest activity in terms of antibiotic potentiation with either the antibiotics ciprofloxacin or tetracycline or the dye EtBr and these data suggest that these extracts contain an inhibitor of efflux. The extract from L. officinale had the greatest EPI-like activity and so this was taken forward and fractionated as an initial step to identify the compounds with EPI-like activity. It was surprising to find that none of the 12 fractions of the chloroform extract of L. officinale had EPI-like activity, as those fractions that displayed synergy with ciprofloxacin did so in all strains irrespective of the presence or absence of a functional AcrAB-ToIC system. These data suggest an interaction with other pumps or another mechanism of synergy. Most interesting was that two fractions (C and F) had some antimicrobial activity alone against all strains including those that lacked TolC. This is highly unusual as the majority of plant extracts display activity only towards Gram-positive bacteria. Fraction C, which had the greatest antimicrobial activity, was further fractionated to identify the compounds conferring the observed antimicrobial activity. Of these, falcarindiol (1) had the greatest antimicrobial activity and a strong additive

effect to the activity of ciprofloxacin. Falcarindiol has previously been shown to possess antibacterial activity, but for mycobacteria and Gram-positive bacteria and not Gram-negative bacteria [24,35–37]. It is possible that the activity in the fraction was potentiated by the co-isolated fatty acids, and these could have membrane permeabilising activity. As these are ubiquitous in all organisms, this does not entirely explain the data or why so few plant extracts are active against Gramnegative bacteria. Rather, the data suggest that falcarindiol is a substrate of ToIC and this may be why it has not been previously identified as having antimicrobial activity for Gram-negative bacteria.

Despite not successfully obtaining the compound(s) that conferred the EPI-like activity in the original plant extract, we postulate that the putative inhibitor of efflux was either lost or was inactivated during fractionation or that it acts in combination with another compound in the plant extract from which it was separated during the fractionation process. Further work to examine reconstituting fractions is required to understand whether a synergistic phenomenon is associated with selected fractions. This will also allow us to conclude whether a putative EPI is being lost in the separation process or is unstable.

Gram-negative plant pathogens such as *Pseudomonas syringae* can cause enormous damage to crop plants [38], and *P. syringae* has been shown to possess a homologue of the tripartite resistance–nodulation–cell division (RND) efflux system MexAB–OprM [39]. This would suggest that plants may not produce chemicals that are active against efflux pumps of Gram-negative plant pathogens. However, the current data suggest that plants do produce molecules with activity against Gram-

negative bacteria. Furthermore, we postulate that these molecules are substrates of ToIC (and homologues thereof) and associated efflux pumps. Therefore, for plants to withstand infection by these pathogens they need to produce EPIs for the natural antimicrobial agents to be active. It is tempting to speculate that falcarindiol is produced alongside a compound(s) that inhibits efflux.

Lorenzi et al. workers [40] have recently shown that extracts from plants can also provide compounds with EPI-like activity or antibacterial activity per se. They also found that an extract of an essential oil from a Corsican plant, *Helichrysum italicum*, was able to reduce the MIC of chloramphenicol for Enterobacter aerogenes, Acinetobacter baumannii and P. aeruginosa. Two fractions of this essential oil and one compound within this oil, geraniol, synergised with chloramphenicol both for the wild-type strain of *E. aerogenes* and its AcrAB mutant. Another recent study by Adonizio et al. [41] showed that medicinal plant extracts can attenuate the virulence of *P. aeruginosa* when explored in the *Caenorhabditis elegans* model system. Although the extracts were from medicinal plants previously used in herbal medicine, the active compounds were not identified. The current data taken together with that of Adonizio et al. [41] and Lorenzi et al. [40] indicate that plants as a source of new and/or novel antimicrobial compounds with activity for Gram-negative bacteria are an underexplored resource. Plant extracts such as those described by ourselves and others provide lead compounds for further exploration and development as antimicrobial agents, as agents to inhibit efflux or for combination with licensed antimicrobials.

Acknowledgments

The authors thank Dr Peter Puplampu for technical support, and Prof. Kim Lewis for helpful discussions while writing this manuscript.

Funding

This work was supported by The Leverhulme Trust (project F/00/094/AR).

کر ک

Competing interests

None declared.

Ethical approval

Not required.

References

- [1] Boucher HW, Talbot GH, Bradley JS, Edwards JE, Gilbert D, Rice LB, et al. Bad bugs, no drugs: no ESKAPE! An update from the Infectious Diseases Society of America. Clin Infect Dis 2009;48:1–12.
- [2] Livermore D. Has the era of untreatable infections arrived? J Antimicrob Chemother 2009;64(Suppl 1):i29–36.
- [3] Hseih PC, Siegel SA, Rogers B, Davis D, Lewis K. Bacteria lacking a multidrug efflux pump: a sensitive tool for drug discovery. Proc Natl Acad Sci U S A 1998;95:6602–6.
- [4] Neyfakh AA, Bidnenko VE, Chen LB. Efflux-mediated multidrug resistance in Bacillus subtilis: similarities and dissimilarities with the mammalian system. Proc Natl Acad Sci U S A 1991;88:4781–5.
- [5] Tegos G, Stermitz FR, Lemovskaya O, Lewis K. Multidrug pump inhibitors uncover remarkable activity of plant antimicrobials. Antimicrob Agents Chemother 2002;10:3133–41.
- [6] Stermitz FR, Cashman KK, Halligan KM, Morel C, Tegos GP, Lewis K. Polyacylated neohesperidosides from *Geranium caespitosum*: bacterial multidrug resistance pump inhibitors. Bioorg Med Chem Lett 2003;13:1915–8.
- [7] Belofsky G, Percivill D, Lewis K, Tegos GP, Ekart J. Phenolic metabolites of Dalea versicolor that enhance antibiotic activity against model pathogenic bacteria. J Nat Prod 2004;67:481–4.
- [8] Belofsky G, Carreno R, Lewis K, Ball A, Casadei G, Tegos GP. Metabolites of the 'smoke tree', *Dalea spinosa*, potentiate antibiotic activity against multidrugresistant *Staphylococcus aureus*. J Nat Prod 2006;69:261–4.

- [9] Cherigo L, Pereda-Miranda R, Fragoso-Serrano M, Jacobo-Herrera N, Kaatz GW, Gibbons S. Inhibitors of bacterial multidrug efflux pumps from the resin glycosides of *Ipomoea murucoides*. J Nat Prod 2008;71:1037–45.
- [10] Kumar A, Khan IA, Koul S, Koul JL, Taneja SC, Ali I, et al. Novel structural analogues of piperine as inhibitors of the NorA efflux pump of *Staphylococcus aureus*. J Antimicrob Chemother 2008;61:1270–6.
- [11] Timmer A, Günther J, Rücker G, Motschall E, Antes G, Kern WV. *Pelargonium sidoides* extract for acute respiratory tract infections. Cochrane Database Syst Rev 2008;(3):CD006323.
- [12] Rüegg T, Calderón AI, Queiroz EF, Solís PN, Marston A, Rivas F, et al. 3-Farnesyl-2-hydroxybenzoic acid is a new anti-*Helicobacter pylori* compound from *Piper multiplinervium*. J Ethnopharmacol 2006;103:461–7.
- [13] Sawer IK, Berry MI, Brown MW, Ford JL. The effect of cryptolepine on the morphology and survival of *Escherichia coli*, *Candida albicans* and *Saccharomyces cerevisiae*. J Appl Bacteriol 1995;79:314–21.
- [14] Piddock LJV. Clinically relevant chromosomally encoded multidrug resistance efflux pumps in bacteria. Clin Microbiol Rev 2006;19:382–402.
- [15] Wray C, Sojka WJ. Experimental Salmonella typhimurium infection in calves. Res Vet Sci 1978;25:139–43.
- [16] Buckley A, Webber MA, Cooles S, Randall LP, La Ragione RM, Woodward MJ, et al. The AcrAB–TolC efflux system of *Salmonella enterica* serovar
 Typhimurium plays a role in pathogenesis. Cell Microbiol 2006;8:847–56.
- [17] Blair JM, La Ragione RM, Woodward MJ, Piddock LJ. The membrane fusion protein AcrA has a distinct role in antibiotic resistance and virulence of

Salmonella enterica serovar Typhimurium. J Antimicrob Chemother 2009;64:965–72.

- [18] Tindall BJ, Grimont PA, Garrity GM, Euzéby JP. Nomenclature and taxonomy of the genus Salmonella. Int J Syst Evol Microbiol 2005;55:521–4.
- [19] Nishino K, Latifi T, Groisman EA. Virulence and drug resistance roles of multidrug efflux systems of *Salmonella enterica* serovar Typhimurium. Mol Microbiol 2006;59:126–41.
- [20] Piddock LJV, White DG, Gensberg K, Pumbwe L, Griggs DJ. Evidence for an efflux pump mediating multiple antibiotic resistance in *Salmonella typhimurium*. Antimicrob Agents Chemother 2000;44:3118–21.
- [21] Eaves DJV, Ricci V, Piddock LJV. Expression of *acrB*, *acrF*, *acrB*, *marA* and *soxS* in *Salmonella enterica* serovar Typhimurium: role in multiple antibiotic resistance. Antimicrob Agents Chemother 2004;48:1145–50.
- [22] Webber MA, Bailey AM, Blair JMA, Morgan E, Stevens MP, Hinton JC, et al. The global consequence of disruption of the AcrAB–TolC efflux pump in *Salmonella enterica* includes reduced expression of SPI-1 and other attributes required to infect the host. J Bacteriol 2009;191:4276–85.
- [23] Piddock LJV, Griggs DJ, Hall MC, Jin YF. Ciprofloxacin resistance in clinical isolates of *Salmonella typhimurium* obtained from two patients. Antimicrob Agents Chemother 1993;37:662–6.
- [24] Lechner D, Stavri M, Oluwatuyi M, Pereda-Miranda R, Gibbons S. The antistaphylococcal activity of *Angelica dahurica* (Bai Zhi). Phytochemistry 2004;65:331–5.

- [25] Kaouadji M, De Pachtere F, Pouget C, Chulia, AJ, Lavaitte S. Three additional phthalide derivatives, an epoxymonomer and two dimers from *Ligusticum wallichii* rhizomes. J Nat Prod 1986;49:872–7.
- [26] Lund ME, Blazevic DJ, Matsen JM. Rapid gentamicin bioassay using a multiple-antibiotic-resistant strain of *Klebsiella pneumoniae*. Antimicrob Agents Chemother 1973;4:569–73.
- [27] Piddock LJV, Garvey MI, Rahman MM, Gibbons S. Natural and synthetic compounds such as trimethoprim behave as inhibitors of efflux in Gram-negative bacteria. J Antimicrob Chemother 2010;65:1215–23.
- [28] Andrews JM. Determination of minimum inhibitory concentrations. J Antimicrob Chemother 2001;48(Suppl 1):5–16. J Antimicrob Chemother 2002;49:1049.
- [29] Andrews JM. Determination of minimum inhibitory concentrations. http://www.bsac.org.uk/susceptibility_testing/guide_to_antimicrobial_susceptibility _testing.cfm. BSAC; 2006 [accessed 23 November 2010].
- [30] Eliopoulos GM, Moellering R. Antimicrobial combinations. In: Lorian V, editor. Antibiotics in laboratory medicine. 4th ed. Baltimore, MD: Lippincott Williams and Wilkins; 1996. p. 330.
- [31] Odds FC. Synergy, antagonism, and what the chequerboard puts between them. J Antimicrob Chemother 2003;52:1–3.
- [32] Coldham NG, Webber MA, Woodward MJ, Piddock LJ. A 96-well plate fluorescence assay for assessment of cellular permeability and active efflux in *Salmonella enterica* serovar Typhimurium and *Escherichia coli*. J Antimicrob Chemother 2010;65:1655–63.

- [33] Newman D, Cragg G. Natural products as a source of new drugs over the last25 years. J Nat Prod 2007;70:461–77.
- [34] Ankli A, Heinrich M, Bork P, Wolfram L, Bauerfeind P, Brun R, et al. Yucatec
 Mayan medicinal plants: evaluation based on indigenous uses. J Ethnopharmacol 2002;79:43–52.
- [35] Stavri M, Gibbons S. The antimycobacterial constituents of dill (*Anethum graveolens*). Phytother Res 2005;19:938–41.
- [36] Schinkovitz A, Stavri M, Gibbons S, Bucar F. Antimycobacterial polyacteylenes from *Levisticum officinale*. Phytother Res 2008;22:681–4.
- [37] Chou SC, Everngam MC, Sturtz G, Beck JJ. Antibacterial activity of components from *Lomatium californicum*. Phytother Res 2006;20:153–6.
- [38] Arnold DL, Jackson RW, Waterfield NR, Mansfield JW. Evolution of microbial virulence: the benefits of stress. Trends Genet 2007;23:293–300.
- [39] Stoitsova SO, Braun Y, Ullrich MS, Weingart H. Characterization of the RNDtype multidrug efflux pump MexAB–OprM of the plant pathogen *Pseudomonas syringae*. Appl Environ Microbiol 2008;74:3387–93.
- [40] Lorenzi V, Muselli A, Bernardini FA, Berti L, Pagès JM, Amaral L, et al. Geraniol restores antibiotic activities against multidrug-resistant isolates from Gram-negative species. Antimicrob Agents Chemother 2009;53:2209–11.
- [41] Adonizio A, Leal SM, Ausubel FM, Mathee K. Attenuation of *Pseudomonas* aeruginosa virulence by medicinal plants in a *Caenorhabditis elegans* model system. Antimicrob Agents Chemother 2008;57:809–13.

Fig. 1. Structures of falcarindiol (1) and levistolide A (2).

Fig. 2. Accumulation of Hoechst 33342 in the absence or presence of extracts of *Levisticum officinale*. White bars, Hoechst 33342 alone; dotted bars, Hoechst 33342 with chloroform (CHCl₃) extract of *L. officinale*; dark grey bars, Hoechst 33342 with methanol (MeOH) extract of *L. officinale*. A value of 1 represents accumulation of Hoechst 33342 in the absence of plant extract; a value >1 indicates increased accumulation of Hoechst 33342 (i.e. reduced efflux). Error bars indicate standard deviations.

Table 1

Bacterial strains used in this study

Strain	Species	Description	Reference/source
L354	S. Typhimurium	SL1344	[15]
L109	S. Typhimurium	SL1344 tolC::aph	[16]
L110	S. Typhimurium	SL1344 acrB::aph	[16]
L884	S. Typhimurium	SL1344 acrA::aph	[17]
L785	S. Typhimurium	ATCC 15277 (LT2)	[18]
L828	S. Typhimurium	ATCC 14028s	[18]
L829	S. Typhimurium	ATCC 14028s ∆tolC	[19]
L830	S. Typhimurium	ATCC 14028s ∆ <i>acrB</i>	[19]
L831	S. Typhimurium	ATCC 14028s ∆acrAB	[19]
L3	S. Typhimurium	Human pre-therapy clinical isolate, antibiotic-susceptible	[20]
L10	S. Typhimurium	Human post-therapy clinical isolate; MDR and overproducing acrAB	[20]
A1	Enterobacter cloacae	NCTC 10005	NCTC
B14	Serratia marcescens	NCTC 2847	NCTC
G1	Pseudomonas aeruginosa	NCTC 10662	NCTC
H42	Klebsiella pneumoniae	NCTC 10896	NCTC
H43	K. pneumoniae	NCTC 9633	NCTC
1114	Escherichia coli	NCTC 10538	NCTC
J29	Morganella morganii	NCTC 235	NCTC

S. Typhimurium, *Salmonella enterica* serotype Typhimurium; ATCC, American Type Culture Collection; MDR, multidrug-resistant; NCTC, National Culture Type Collection (Health Protection Agency, Colindale, London, UK).

Table 2

List of medicinal plants used in this study

Plant name	Common name	Family
Anemone nemorosa	Wood anemone	Ranunculaceae
Angelica sinensis	Female ginseng	Apiaceae
Apium graveolens	Dill	Apiaceae
Artemisia abrotanum	Southernwood	Asteraceae
Asclepias tuberosa	Butterfly weed	Apocynaceae
Boswellia serrata	Salai	Burseraceae
Catha edulis	Khat	Celastraceae
Centella asiatica	Gotu kola	Apiaceae
Cinnamomum aromaticum	Cassia cinnamon	Lauraceae
Citrus aurantium	Bitter orange	Rutaceae
Commiphora molmol	Myrrh	Burseraceae
Daucus carota	Wild carrot	Apiaceae
Ephedra funereal	Death Valley ephedra	Ephedraceae
Glycyrrhiza glabra	Liquorice	Leguminosae
Levisticum officinale	Lovage	Apiaceae
Melissa officinalis	Lemon balm	Lamiaceae
Papaver somniferum	Opium poppy	Papaveraceae
Pimpinella anisum	Anise	Apiaceae
Pulmonaria officinalis	Lungwort	Boraginaceae
Thymus vulgaris	Thyme	Lamiaceae
Tussilago farfara	Coltsfoot	Asteraceae

Table 3

Plant extracts with greatest synergy with ciprofloxacin

Strain ^a	ΡΑβΝ	Lo	Lo	Lo	М	М	С	Pi	Pu	Pu
		MeOH	$CHCI_3$	Water	MeOH	CHCl ₃	CHCI ₃	$CHCI_3$	CHCI ₃	MeOH
S. Typhimurium SL1344	S	NE	NE	NE	NE	NE	NE	NE	NE	NE
S. Typhimurium ATCC 15277 (LT2)	S	S	S	S	S	NE	S	S	S	S
S. Typhimurium ATCC 14028s	S	S	S	S	S	S	S	S	S	S
Human pre-therapy isolate L3	S	S	S	S	S	S	S	S	S	S
Human post-therapy isolate L10	S	S	S	S	S	S	NE	S	S	S

^a Details of strains are given in Table 1.

PAβN, L-phenylalanyl-L-arginyl-β-naphthylamide; Lo, *Levisticum officinale*; M, *Melissa officinalis*; C, *Cinnamomum aromaticum*; Pi, *Pimpinella anisum*; Pu, *Pulmonaria officinalis*; MeOH, methanol extraction; CHCl₃, chloroform extraction; water, water extraction; S, synergy (zone of inhibition reproducibly larger than that of antibiotic alone); NE, no effect.

Table 4

Minimum inhibitory concentrations (MICs) of ciprofloxacin (CIP), tetracycline (TET), chloramphenicol (CHL), erythromycin (ERY) and ethidium bromide (EtBr) in the absence and presence of *Levisticum officinale* and *Melissa officinalis* extracts against Gramnegative bacteria

Strain ^a	MIC (r	ng/L) ^b													
	CIP	CIP +	CIP +	TET	TET	TET	CHL	CHL	CHL	ERY	ERY	ERY	EtBr	EtBr	EtBr
		Lo	Μ		+ Lo	+ M		+ Lo	+ M		+ Lo	+ M		+ Lo	+ M
S. Typhimurium	0.03	0.008	0.015	2	1	1	8	8	8	512	512	512	2048	2048	2048
L354															
S. Typhimurium	0.03	0.008	0.008	16	0.5	0.5	8	8	8	512	256	512	2048	1024	2048
L828															
S. Typhimurium	0.008	0.004	0.008	4	0.5	0.5	1	1	1	8	8	8	16	8	16
L829															
S. Typhimurium	0.008	0.008	0.008	4	0.5	0.5	2	8	8	64	64	64	64	32	64
L831															
S. Typhimurium	0.008	0.002	0.008	4	0.5	0.25	8	4	8	512	512	512	512	512	512
L3															
S. Typhimurium	0.06	0.03	0.06	8	4	4	32	16	32	512	256	512	2048	1024	2048
L10															

Enterobacter	0.12	0.06	0.12	8	4	8	4	2	4	512	256	512	2048	2048	2048
cloacae A1															
Serratia	0.06	0.06	0.06	64	64	64	4	4	4	256	256	256	2048	2048	2048
marcescens B14															
Pseudomonas	1	1	1	64	64	64	64	64	64	512	512	512	2048	2048	2048
aeruginosa G1															
Klebsiella	0.06	0.03	0.03	16	2	4	128	64	128	512	256	512	2048	1024	1024
pneumoniae															
H42															
K. pneumoniae	0.12	0.06	0.06	8	1	2	4	4	4	512	512	512	2048	1024	2048
H43															
Escherichia coli	0.06	0.03	0.06	4	2	4	4	4	4	512	512	512	256	256	256
1114															
Morganella	0.015	0.008	0.015	8	8	8	16	16	16	512	512	512	2048	2048	2048
morganii J29															

S. Typhimurium, Salmonella enterica serotype Typhimurium; Lo, chloroform extract of L. officinale extract; M, methanol extract of

M. officinalis.

^a Details of strains are given in Table 1.

^b Bold text indicates synergistic combinations (i.e. MICs for the combination lower than for the antibiotic alone).

Table 5

Minimum inhibitory concentrations (MICs) of ciprofloxacin (CIP) in the absence and presence of the six most active fractions

Strain ^b	MIC (r	ng/L) ^b								
	CIP	CIP + Lo	CIP + A	CIP + C	C alone	CIP + F	F alone	CIP + G	CIP + J	CIP + K
S. Typhimurium L354	0.03	0.008	0.03	0.03	250	0.03	250	0.03	0.03	0.03
S. Typhimurium L884	0.008	0.008	0.002	0.002	64	0.008	250	0.004	0.008	0.004
S. Typhimurium L110	0.008	0.008	0.002	0.004	128	0.008	250	0.004	0.008	0.004
S. Typhimurium L109	0.015	0.008	0.002	0.002	8	0.008	128	0.004	0.0015	0.004
S. Typhimurium L828	0.03	0.008	0.015	0.03	250	0.03	250	0.03	0.03	0.03
S. Typhimurium L829	0.008	0.004	<0.001	<0.001	8	0.004	128	0.004	0.015	0.004
S. Typhimurium L830	0.008	0.008	<0.001	<0.001	128	0.004	250	0.004	0.008	0.004
S. Typhimurium L831	0.008	0.008	<0.001	<0.001	64	0.004	250	0.004	0.008	0.004
S. Typhimurium L3	0.008	0.002	<0.001	<0.001	250	0.008	250	0.008	0.008	0.008
S. Typhimurium L10	0.06	0.03	0.03	0.12	250	0.12	250	0.03	0.12	0.03
Enterobacter cloacae A1	0.12	0.06	0.12	0.12	250	0.12	250	0.12	0.12	0.12
Serratia marcescens B14	0.06	0.06	0.06	0.06	250	0.06	250	0.06	0.06	0.06
Pseudomonas aeruginosa G1	1	1	1	1	250	1	250	1	1	1
Klebsiella pneumoniae H42	0.06	0.03	0.06	0.015	250	0.015	250	0.015	0.015	0.015
K. pneumoniae H43	0.12	0.06	0.12	0.03	250	0.03	250	0.06	0.03	0.03

Escherichia coli 1114	0.06	0.03	0.06	0.06	250	0.06	250	0.06	0.06	0.06
Morganella morganii J29	0.015	0.008	0.015	0.015	250	0.015	250	0.015	0.015	0.015

S. Typhimurium, Salmonella enterica serotype Typhimurium; Lo, chloroform extract of L. officinale; CIP + letter, indicates the

fraction of the chloroform extract of L. officinale.

^a Details of strains are given in Table 1.

^b Bold text indicates synergistic combinations (i.e. MICs for the combination lower than for the antibiotic alone).

Table 6

Fractional inhibitory concentration index (FICI) values of ciprofloxacin (CIP) with plant extracts, fractions and purified molecules for

Strain ^a	FICI ^b									
	CIP + Lo	CIP + M	CIP + FC	CIP + FF	CIP + Lin	CIP + Fal				
SL1344	0.94	0.97	0.99	0.97	1.34	0.97				
SL1344 tolC::aph			0.27	0.38	0.17	0.26				
ATCC 15277 (LT2)	0.46	0.59								
ATCC 14028s	0.48	0.41	1.01	0.98	1.09	1.05				
ATCC 14028s ∆tolC			0.32	0.33	0.14	0.14				
Human pre-therapy isolate L3	0.32	0.13								
Human post-therapy isolate L10	0.31	0.22								

Salmonella enterica serotype Typhimurium

Lo, Levisticum officinale chloroform extract; M, Melissa officinalis methanol extract; FC, fraction C from L. officinale; FF, fraction F

from L. officinale; Lin, linoleic acid; Fal, falcarindiol.

^a Details of strains are given in Table 1.

^b Bold text indicates synergy [31].

