

Inhibitors of efflux pumps of Gram-negative bacteria inhibit Quorum Sensing

Leonard Amaral^{1,2*}, Joseph Molnar^{2,3}

¹ Grupo de Micobacterias, Unidade de Microbacteriologia, Centro de Malaria e Doenças Tropicais (CMDT), Instituto de Higiene e Medicina Tropical, Universidade Nova de Lisboa, Lisbon, Portugal

² Cost Action BM0701 (ATENS) of the European Commission/European Science Foundation

³ Department of Medical Microbiology and Immunobiology, University of Szeged, Szeged, Hungary

Corresponding Author & Address:

Leonard Amaral*

Unidade de Microbacteriologia, Instituto de Higiene e Medicina Tropical, Universidade Nova de Lisboa Rua Junqueira 100, 1349-008 Lisbon, Portugal; Tel +3513652600; Fax +351 213632105; Email: lamaral@ihmt.unl.pt

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ABSTRACT

Quorum Sensing (QS) systems of bacteria consist of a producer of the QS signal and the responder. The generation of a QS signal provides the means by which a population can behave in a concerted manner such as swarming, swimming and secretion of biofilm, etc. Because concerted behaviour bestows protection to the bacterial species, and hence factors involved in the severity of an infection such as virulence are products of QS systems, compounds that inhibit the QS system have significant clinical relevance. Recent evidence suggests that the secretion of QS signals takes place via the efflux pump system of the producer of the signal. Interestingly, compounds such as phenothiazines and trifluoromethyl ketones (TFs) that inhibit proton motive force (PMF) activities such as swarming and swimming also inhibit the PMF dependent efflux pump systems of bacteria and their QS systems. This review discusses the relationship between the efflux pump, the QS system and the compounds that affect both. Lastly, suggestions are made regarding classes of compounds that have been shown to inhibit PMF dependent efflux pumps and the need to evaluate them for QS inhibitory properties.

INTRODUCTION

The role of efflux pumps in multi-drug resistance

Efflux pumps of bacteria provide protection from noxious agents that are present in the

environment in which they exist. Noxious agents may be naturally occurring compounds present in environments outside and within the human. With respect to infectious disease, the universe of external environments may be aqueous reservoirs where compounds are added to reduce or even eliminate infectious agents, the home where

attempts to reduce the bacterial load present in the kitchen, bathroom, bedroom, play areas, swimming pool, etc., facilities used by children (child care, schools, etc) and on occasion, the property (area of septic systems, etc.) are often needed, and transportation facilities (cruise ships) that may require fumigation or disinfectants. However, although the aforementioned sites are important for reducing infectious disease, it is the environmental sites of health care areas where the problem of potential infectious disease is far more difficult to control. At these sites, patients presenting with an infectious disease may infect others even when no longer present, leaving behind bacterial loads in nasal discharges, saliva and other body fluids. Transmission of bacteria to patients in a health care facility may be from contaminated surfaces and microdroplets of expired air and certainly from health care workers who are attending the patient as well as those who are ancillary to health care such as orderlies who transport the patient, maids who change soiled bed linen, etc. A large proportion of patients that enter health care facilities for non-infectious pathology acquire a bacterial infection [1-3]. These nosocomial infections pose serious threats to the patient since it is highly probable that the infectious agent has its origins in a patient that receives antibiotic therapy. Due to a variety of reasons which will be discussed, therapy of the bacterial infection places the bacterium under stress and the bacterium responds by invoking a series of mechanisms that render it resistant to one or more antibiotics-hence it is multi-drug resistant (MDR). MDR infections are problematic for therapy and as of today, nosocomial infections such as those produced by methicillin resistant *Staphylococcus aureus* are more prevalent in most hospitals world-wide than methicillin susceptible strains of *Staphylococcus aureus* (MSSA) [4, 5]. Consequently, in order to prevent nosocomial infections, hospitals devote much energy towards maintaining as sterile environment as possible, and to accomplish this disinfectants are used in great quantities. Agents used to prevent infectious disease in the house and its property, in the health care facilities as well as agents used to treat bacterial infections, to the bacterium, they are "one of the same", namely, noxious agents that threaten their survival, and their response to these agents in part involves the over-expression of genes that regulate and code for efflux pumps that

recognise such agents and extrude them to the environment before they reach their intended targets [6-10]. Because over-expressed efflux pumps render antibiotic therapy problematic, an intense search for agents that inhibit specific efflux pumps of specific bacteria has been conducted during the past decade [9].

The role of Quorum Sensing (QS) in infection

Communication between bacteria of the same strain or species and between species contributes to their survival [11-13]. Communication involves the secretion of signals that invoke a specific response from the responder [11-13]. This communication process is termed Quorum sensing (QS). When it takes place between strains of the same species, communication is directed towards the reduction of population growth and hence, reducing the possibility of exceeding the nutritional support of the environment [13]. Other signals may involve a population response that involves the secretion of bioactive molecules that inhibit the replication of a competing population species [14-16] or even kill [biocidins] [17-21] or promote a swarming effect that recruits members of the same species to migrate to a specific location [22-24] similar to swarming by insects subsequent to signals indicating site of food [example bees]. Other responses involving "community action" are also invoked and these result in secretion of materials that will protect the bacterium from external danger. These materials, termed biofilm, encase the bacteria at distances from each other [25-29] and within the matrix of this biofilm are channels used for further communication [30]. Biofilms are produced in the wild, at sites such as surfaces of rocks which maintain the bacterial population *in situ* [31] and are also produced at sites of the human colonised by infecting bacteria [32, 33]. Biofilms are also problematic with prosthetic devices in that subsequent to their placement, an infection may take place and the surfaces of the device are suitable for the production and maintenance of biofilm which has bacteria embedded [34]. Agents that inhibit the QS response of the infecting bacterium are obviously important and hence, the search for such agents that inhibit the QS system and biofilm formation has been in effect for the past two decades [11-13].

Relationship between Efflux Pumps, QS and Biofilm secretion

There is a relationship between efflux pumps (EP), QS and biofilm (BF) secretion which has come to the forefront only recently [13]. Control of this relationship is critical for successful therapy of MDR bacterial infections which have become rather commonplace. It is the intent of this review to identify agents which may serve to interfere with the complex system of EP-QS-BF interaction.

Bacterial Efflux pumps (EPS)

Bacterial efflux pumps (EPs) are proteins that are localised and imbedded in the plasma membrane of the bacterium and whose function is to recognise noxious agents that have penetrated the protective cell wall of the organism and have reached the periplasm or cytoplasm, and extrude the agents prior to their reaching their intended targets [6-10]. However, they also recognise toxic compounds produced from their own metabolic processes and hence perform excretory functions as well [35, 36]. In other words, they are transporters of noxious compounds from within the bacterial cell to the external environment. With the possible exception of excretory functions, efflux pumps utilize sources of energy for their function inasmuch as they transport compounds against a concentration gradient. In general there are two types of efflux pumps that are distinguished with respect to the immediate source of energy that is utilized for export functions: ABC transporters [37-40] and proton motive force dependent transporters [6-10, 35, 36, 41]. ABC transporters consist of two domains-one that is embedded in the plasma membrane and the other is on the medial side of the plasma membrane. Their structure may consist of 6 or 12 helical sequences parts of which face the medial side of the cytoplasmic membrane. For ABC transporters that consist of 6 helices, there are two types of binding sites; one for the agent (substrate) that is to be transported and another for the binding of ATP. Subsequent to the recognition of the noxious agent and its binding, ATP is bound and hydrolysed and the energy liberated affords a conformational change of the transporter that promotes the movement of the noxious agent to the environment [37-42]. The precise structural changes that take place in the transporter that lead to the extrusion of the noxious agent as well as the

means by which the transporter recognises structurally unrelated compounds is not yet completely understood. Proton motive force (PMF) dependent transporters obtain their energy for function from the proton motive force. The proton motive force is the result of cellular metabolism which yields protons that are not used for coupling with molecular oxygen and which are exported to the surface of the cell [43-45] where they are distributed and bound to components of the protective lipopolysaccharide layer that covers the cell and constitutes a part of the outer cell wall of Gram-negative [46] and the cell wall of Gram-positive bacteria [47]. The larger the concentration of protons (hydronium ions) on the surface of the cell with respect to their lower concentration on the medial side of the cytoplasmic membrane creates an electrochemical gradient that is termed the proton motive force (PMF) [48]. In addition, the binding of these hydronium ions promotes a pH on the surface of the cell that is two to three units lower than the pH of the bulk milieu [43-45]. Because hydronium ions cannot penetrate the cell wall or the membrane, they may re-enter the cell only through channels such as porins in general [49, 50]. The movement of these hydromium ions from the surface of the cell to the periplasm or cytoplasm is predicated upon systems that use the PMF as source of energy-namely the resistance nodulation division (RND) family of transporters. However, before this movement of hydronium ions is discussed, the structure and mechanism of the main efflux pump of *Escherichia coli* needs addressing.

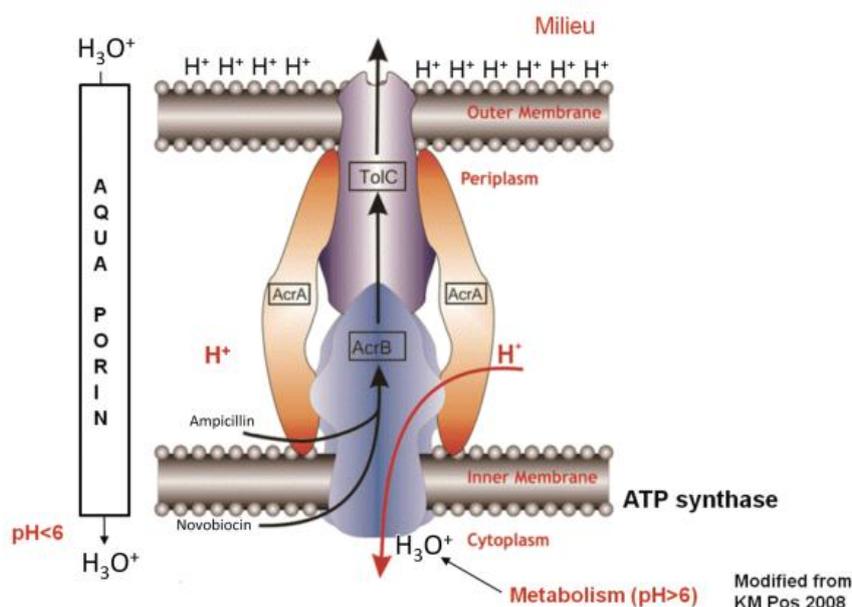
E. coli has a multiplicity of efflux pumps that may exceed 30 in number [51]. However, the main efflux pump of this organism is the AcrAB-TolC efflux pump [52, 53] which when deleted, its function is replaced by the AcrEF-TolC efflux pump [51]. Both efflux pumps are members of the resistance nodulation division family of transporters [51] and consist of three proteins: the transporter AcrB coded by the gene *acrB* and is intimately attached to the plasma membrane; two fusion proteins AcrA coded by the gene *acrA* that flank the AcrB transporter and are believed to assist the movement of a substrate through the AcrB transporter [35]; and, TolC which is also part of other tri-unit efflux pumps of the organism [35], is contiguous with the AcrB transporter and provides a conduit for the extrusion of the

substrate [38]. The structure of the tripartite AcrAB-TolC efflux pump is in the plasma membrane is not yet defined. The AcrAB-TolC efflux pump has been studied for three decades and has been shown to extrude a large variety of unrelated compounds with widely different structures [35]. Although the means for the recognition of the substrate to be extruded appears to involve a pocket within the transporter, it appears to be defined by a phenylalanine residue [54]. Nevertheless, studies employing fluorochromes recognised by the AcrB transporter indicate that the binding and release of the substrate are pH dependent [55]. At low pH the dissociation of the substrate is high and at high pH it is very slow. In a physiological environment of ca. pH 7, if the dissociation of the substrate is slow or not at all, then the effectiveness of the pump to extrude a noxious agent would be nullified. However, since the pump functions at this pH, conditions that

result in the dissociation of the substrate needed for continuous pump action must involve a decrease of the pH of the internal cavity of the pump to which the substrate is bound. It has been postulated that the lowering of the pH takes place by the generation of hydronium ions from metabolism [6] which pass from the cytoplasmic side of the plasma membrane through the transporter. At lower pH, there is no need for the generation of metabolically derived hydronium ions since these ions can be diverted by the PMF from the surface of the cell to the periplasm via porins. Whether hydronium ions are to be generated from the hydrolysis of ATP at high pH or used for the synthesis of ATP at low pH is a special function of ATP synthase [56-58].

Nevertheless, a model of the RND AcrAB-TolC efflux pump of Gram-negatives and suggested hypothesis for its mechanism of action is presented by Figure 1.

Figure 1. Model of the AcrAB-TolC efflux pump of a Gram-negative bacterium.



Hypothesis. At near neutral pH, Hydronium ions from hydrolysis of ATP by ATP synthase pass through the AcrB transporter, reduce the pH to a point that causes the release of the substrate. When the hydronium ions reach the surface of the cell they are distributed over that surface and bind to lipopolysaccharides and basic amino acids. When there is a need for hydronium ions for activity of the efflux pump and the pH is lower than neutral, and the hydrolysis of ATP is not favoured, hydronium ions from the surface of cell via the PMF mobilize through the Aqua porins and reach the transporter where they are pushed through the transporter by the peristaltic action caused by the fusion proteins. Substrates bound to the transporter dissociate when the pH is reduced by the flow of hydronium ions and are carried out by the flow of water.

Inhibitors of bacterial efflux pumps

The compound phenylalanine-arginine-β-naphthylamide (MC-207,110), otherwise known as

PAβN, was created by Lomovskaya et al. [59] and reported to significantly decrease intrinsic and acquired resistance of *Pseudomonas aeruginosa* to fluoroquinolones [59]. Because acquired resistance

was related to the over-expression of the organism's main efflux pumps MexAB-OprM, MexCD-OprJ, MexEF-OprN, PA β N rapidly became known as an inhibitor of efflux pumps whenever resistance to given antibiotics is increased as a consequence of over-expression of efflux pumps. PA β N therefore was claimed to inhibit the efflux pumps of *Escherichia coli* [60, 61], *Enterobacter aerogenes* [61], *Acinetobacter baumannii* [62], *Brucella melitensis* [63], *Salmonella enterica* serovar Typhimurium [64, 65], *Yersinia pestis* [66] and all encountered Gram-negatives [6, 7, 8, 9, 35, 39, 65-71]. However, PA β N is not an inhibitor of an efflux pump but rather a competitor of substrates used by the given efflux pump [72-74] and recent evidence has suggested to indicate that it is also a permeabilising agent [75]. If we use the term "efflux pump inhibitor" (EPI) to mean a compound that affects the efflux activity of an efflux pump, then PA β N and other compounds that affect the activity of an efflux pump could be called an EPI. In this respect, phenothiazines [6, 7, 9, 41, 51, 73, 74, 76-83] and hydantoin compounds [84] are EPI's since they reduce the activity of intrinsic- and over-expressed efflux pumps. However, it must be noted that with respect to the phenothiazines chlorpromazine (CPZ) and thioridazine (TZ), they express their anti-efflux pump activity by inhibiting the binding of Ca²⁺ to enzymes involved in metabolism [85]. It should also be noted that a chelating agent such as EDTA also can inhibit the activity of an efflux pump by denying Ca²⁺ to energy systems dependent upon this cation for activity [85]. How hydantoins exert their inhibitory activity against the efflux pump system of Gram-negatives is not yet known. However, because they are also powerful inhibitors of the human ABC transporter Pgp-1 [86-88], they probably have an effect on an ABC transporter of the Gram-negative bacterium that recognises the ethidium bromide substrate used in the efflux pump assay.

Phenothiazines are known to inhibit the motility of bacteria [89-90] and act synergistically with an inhibitor of the proton motive force (PMF) to inhibit motility [91]. Trifluoromethyl ketones (TFs) are known to be inhibitors of motility by inhibiting access to the energy provided by the PMF for flagellae movement [90-95] and because phenothiazines also affect the PMF [6], the synergistic combination of a phenothiazine with TFs suggested that TFs might also have activity

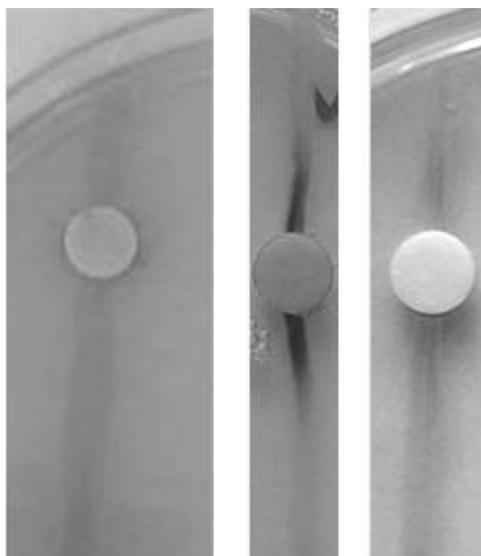
against PMF dependent efflux pumps of a Gram-negative bacterium such as *Escherichia coli*. Moreover, because TFs have similar structures to those of omeprazol, lansopresol, etc. all of which are strong inhibitors of the PMF, evaluation of 12 TFS for activity against the efflux pump of *Escherichia coli* provided evidence that indeed, some TFS were similarly powerful inhibitors of the PMF dependent efflux pump-supposedly, the main efflux pump AcrAB-Tol [13].

Inhibitors of the QS of bacteria

Inhibitors of a bacterial QS system must be distinguished from their activity on the producer of the signal, the responder to a QS signal or both. Quorum sensing inhibitors (QSIs) are compounds that specifically block QS systems without affecting bacterial growth. The QS system of bacteria that has received the greatest attention is the acylated homoserine lactone (AHL) system which produces and secretes AHL [96-113]. AHL acts as a communication molecule which regulates the behavior of the members of the bacterial population. Concerted behaviour such as swarming [99], production of surfactant which facilitates movement on surfaces [99] and virulence factors by *Pseudomonas aeruginosa* [97], production and secretion of biofilms [104, 105] are examples among a growing list of concerted behaviours [100-103]. Obviously, the regulation of bacterial behavior and population density by AHLs suggests activity at the genetic level. The AHLs density-dependent regulatory systems rely on two proteins, an AHL synthase, most commonly a member of the LuxI family of proteins, and an AHL receptor protein belonging to the LuxR family of transcriptional regulators. Low population density cells produce a basal level of AHL that is dependent on an AHL synthase. With increase of population density, AHL accumulates in the medium and when it reaches a critical threshold concentration, the AHL molecule binds to its cognate receptor. The binding of AHL to its receptor promotes induction or repression of AHL-regulated genes. The genes which are regulated are responsible for a large number of functions such as bioluminescence [106], plasmid conjugal transfer [107], biofilm formation [104, 105], motility [101, 105], antibiotic biosynthesis [108-113], production of virulence factors [96], etc. It should be noted that *in vitro* experiments utilizing the cell free culture medium for purposes of studying the presence of QS signal

molecules, are difficult to carry out because of technical problems such as the very low concentration of these QS signal molecules present in the media, the instability of the QS signal molecule, and with attempts to concentrate the medium such as with lyophilisation, selective filtration, etc., the components of the concentrate cause formation of complexes that tie up or interfere with the QS signal molecule. Nevertheless, these difficulties can be overcome with selective design of the experimental protocol. An example of such a system is provided by [Figure 2](#). In this example, a compound (a trifluoromethyl ketone) that is being studied for an effect on the responder of the QS signal, is impregnated onto a paper disk that contains the QS signal (acylated hydroxyl lactone-AHL) and is then placed directly on the agar streaked responder environmental bacterium CV026. As noted by [Figure 2](#), the TF employed inhibits the production of colour that is generated by exposure to the QS signal AHL.

Figure 2. The effect of a trifluoromethyl ketone (TF) on the response of CV026 to the QS signal acylated hydroxyl lactone (AHL).



Legends:

- A:** Control disk with MM4 (20µg) alone (no colour).
- B:** Control disk with AHL (10ng) alone (deep purple coloration)
- C:** Disk with MM4 (20µg/disc) and AHL (10 ng) (very light purple coloration)

The inhibition of the QS system of pathogenic bacteria is a universal goal of drug discovery [114, 115]. Because the AHL system of Gram-negatives has been extensively studied and because in general, Gram-negatives due to their

more complex cell envelope have greater resistance to current antibiotics as opposed to Gram-positives, the review of QS inhibitors will be mostly on pathogenic Gram-negatives.

Perhaps the best known inhibitors of the AHL QS system are homologues of AHL. N-butanoyl-L-homoserine lactone (C4-HSL) and N-hexanoyl-L-homoserine lactone (HHL) are produced in cultures of *Serratia liquefaciens* at a ratio of 10:1, respectively [116]. These analogues of AHL serve as autoinducers of swarming [116]. However, analogues of C4-HSL such as N-(3-oxododecanoyl)-L-homoserine lactone (3-oxo-C(12)-HSL) produced by *Pseudomonas aeruginosa* that do not affect cell growth of its own population as well as that of *Proteus mirabilis*, *Escherichia coli*, *Alcaligenes faecalis* and *Stenotrophomonas maltophilia*, inhibit completely the growth of *Legionella pneumophila* as well as the formation of its biofilm [117]. Among compounds that inhibit production and secretion of AHL are sulphur-containing AHL-analogues such as N-(heptylsulfanylacetyl)-L-homoserine lactone (HepS-AHL) which reduces production of protease by *Aeromonas salmonicida* [96]. Protease production is a virulence factor which when its production and release are blocked, render the bacterium less virulent [96].

Although it is obvious that analogues of QS signals such as AHLs may readily inhibit secretion and synthesis of QS signals, there may be a plethora of compounds that can similarly serve as inhibitors of QS signals, either generally or specifically. Among the general inhibitors of QS are phenothiazines [12, 118]. Because phenothiazines inhibit many energy dependent systems of bacteria such as motility [89, 90, 95], and these phenothiazines also inhibit efflux pumps of bacteria [6, 7, 9, 41, 51, 73, 74, 76-83], there seems to be a correlation between an active efflux pump system and a functional QS system. That this assumption is correct, recent evidence has been provided showing that the efflux pumps of the AHL responding environmental *Chromobacterium violaceum* (CV026) bacterium and that of *E. coli* are inhibited by the phenothiazine thioridazine (TZ) [12]. Because TZ is known to inhibit genes that regulate and code for efflux pumps of bacteria [41, 119, 120], it is possible that the inhibition of the responding CV0126 bacterium to AHLs [12] involves the inhibition of genes that code and regulate the efflux pump of the responder which is assumed to

recognise the AHL signal as an noxious agent and hence would extrude it to the environment [12]. Moreover, given the results that show that the phenothiazine structural scaffold is the component of the phenothiazine derivative that is responsible for the inhibition of QS [118], attention should be given to phenothiazines as parental compounds for derivatives with anti-QS properties. It should be noted that the secretion of biofilm takes place via the efflux pumps system of the bacterium [121, 122]. Consequently, the inhibition of an efflux pump should manifest itself as an inhibitor of the QS component responsible for biofilm formation.

During the past two decades drug development and discovery has focused on plants as sources of bioactive compounds [123-141]. However, as noted by the reviews cited, most of the focus has been on discovery of compounds from plant sources that have bioactivity in human pathology. Nevertheless, because of the successes encountered with compounds from plant sources and due to the ever increasing problem of multi-drug resistance that emanates from over-expressed efflux pumps [6-9], plants as sources of antimicrobial compound have during recent years assumed an important part in drug development and discovery [67, 142-158]. In particular, since the discovery of berberine a powerful inhibitor of bacterial efflux pumps [159], plants have become sources of inhibitors of efflux pumps [160-164]. Given that efflux pumps and the QS of bacteria have an intimate relationship as described in this review, attention has been focused on plants for potential sources of inhibitors of efflux pumps and QS systems. Essential oils from Columbian plants have yielded a large number of compounds that inhibit the QS system of responding bacteria such as limonene-carvone, the citral (geranial-neral) (isolated from *Lippia alba*), α -pinene (from *Ocotea* sp.), β -pinene (from *Swinglea glutinosa*), cineol (from *Elettaria cardamomum*), α -zingiberene (from *Zingiber officinale*) and pulegone (from *Mintostachys mollis*) [165]. Several other essential oils, in particular were shown to present promising inhibitory properties for the short chain AHL quorum sensing (QS) system in *Escherichia coli* containing the biosensor plasmid pJBA132, in particular *Lippia alba*. Citral was the only essential oil that presented some activity for the long chain AHL QS system in *Pseudomonas putida* containing the plasmid pRK-C12 [165]. Other essential oils

from rose, geranium, lavender and rosemary oils were shown to have potent inhibitory activity against the QS system of bacteria whereas essential oils from *Eucalyptus* and citrus oils presented moderate activity and those from chamomile, orange and juniper oils were ineffective [11]. Essential oil from *Syzygium aromaticum* (Clove) has also been shown to inhibit the QS systems of *hromobacterium violaceum* (CV12472 and CVO26) and *Pseudomonas aeruginosa* (PAO1) whereas essential oils from cinnamon, lavender and peppermint had moderate activity [166].

To our knowledge, no other work has been conducted for studies of essential oils as potential sources of inhibitors of the QS system. Moreover, given that an essential oil has now been shown to inhibit the efflux pump of antibiotic resistant Gram-negative bacteria [167], essential oils shown to inhibit the QS systems of bacteria are good candidates for evaluation for inhibitory activity against efflux pumps of multi-drug resistant Gram-negative bacteria.

The relationship between efflux pumps and the QS of bacteria seems well established. Given that efflux pumps of Gram-negative bacteria that bestow multi-drug resistant are dependent upon the PMF for activity [6, 8, 168-170], and since TFs inhibit the efflux pump of bacteria [13] supposedly by the inhibition of the PMF energy source [6, 8, 13], TFs were evaluated for activity against the QS system of bacteria [13]. As expected, the TFs that had the greatest specific activity against the efflux pump were the most active as inhibitors of the QS response [13]. These latter studies serve to support the intimate connection between efflux pumps and the QS system of bacteria. Consequently, compounds that affect both are good candidates for inhibitory properties against the secretion of biofilm matrix which is dependent upon the QS system [31-34] and the efflux pump for secretion of the biofilm [121, 122].

CONCLUDING REMARKS

The essence of this review is to correlate the relationship of the efflux pump system to the QS system of bacteria via the use of compounds that inhibit both systems. Simply put, inhibitors of the efflux pump system also, when studied, inhibit the QS system as well. Because the PMF dependent

efflux pump system of Gram-negatives that is over-expressed is responsible for the multi-drug phenotype of the bacterium, compounds that affect the PMF of the bacterium are candidates that will inhibit the activity of the pump. Consequently, this inhibition will inhibit the secretion of biofilm, and because biofilm is a deterrent to the action of antibiotics, compounds that affect the efflux pump system are promising candidates for clinical evaluation. The effect of compounds that inhibit the efflux pump on the QS system is at this time not completely understood. However, if we consider that QS signals directed at another bacterial species constitutes a noxious agent, then the extrusion of the QS signal by an efflux pump may be significantly affected by the inhibitor of the pump. With respect to concerted action by a bacterial population such as swarming that is brought about by a QS signal, the inhibitory qualities of an anti-QS compound may lie in the inhibition of the secretion of the QS signal via the efflux pump itself. Consequently, we should distinguish the effect of a compound on the producer of the QS signal versus the response of the same or different species of bacteria. At this time, little work has actually been reported concerning the QS of bacteria and the effect of compounds from different sources. Regardless, future studies are suggested to include a wider gamut of activities when studying the QS system of

bacteria. Therefore, within one study, it may be very significant that any compound deemed worthy of consideration for QS study, be studied for effects on the producer of the QS signal, the responder to the QS signal, the efflux pump of the producer of the signal and the efflux pump of the responder. In this manner, a more complete understanding of how a compound can mediate its effects at the antibacterial level can be more fully understood.

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CONFLICT OF INTERESTS

There is no conflict of interest.

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