

Deciphering the role of coumarin as a novel quorum sensing inhibitor suppressing virulence phenotypes in bacterial pathogens

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Abstract The rapid unchecked rise in antibiotic resistance over the last few decades has led to an increased focus on the need for alternative therapeutic strategies for the treatment and clinical management of microbial infections. In particular, small molecules that can suppress microbial virulence systems independent of any impact on growth are receiving increased attention. Quorum sensing (QS) is a cell-to-cell signalling communication system that controls the virulence behaviour of a broad spectrum of bacterial pathogens. QS systems have been proposed as an effective target, particularly as they control biofilm formation in pathogens, a key driver of antibiotic ineffectiveness. In this study, we identified coumarin, a natural plant phenolic compound, as a novel QS inhibitor, with potent anti-virulence activity in a broad spectrum of pathogens. Using a range of biosensor systems, coumarin was active against short, medium and long chain *N*-acyl-homoserine lactones, independent of any effect on growth. To determine if this suppression was linked to anti-virulence activity, key virulence systems were studied in the nosocomial pathogen *Pseudomonas aeruginosa*. Consistent with suppression of QS, coumarin inhibited biofilm, the production of phenazines and swarming motility in this organism potentially linked to reduced expression of the *rhII* and *pqsA* quorum sensing genes. Furthermore, coumarin significantly inhibited biofilm

formation and protease activity in other bacterial pathogens and inhibited bioluminescence in *Aliivibrio fischeri*. In light of these findings, coumarin would appear to have potential as a novel quorum sensing inhibitor with a broad spectrum of action.

Keywords Coumarin · Natural compound · Quorum sensing inhibition · Anti-biofilm · Bacterial pathogens

Introduction

The emergence of multidrug-resistant bacteria over the last few decades has become a global health issue for medical professionals and public health systems throughout the world (Tanwar et al. 2014). Once the cornerstone of medical intervention to treat microbial infections, the indiscriminate use of antibiotics has contributed to the evolution of broad spectrum resistance mechanisms (Fernández and Hancock 2012). In most cases, the selective pressure underpinning the emergence of resistant strains has arisen due to the mode of action of antibiotics, i.e. inhibiting bacterial growth. In the last decade, there has been a sharp decline in the development of new classes of antibiotics, with the result that resistance is continuing to increase unchecked. Furthermore, the realisation that microbial communities exist predominantly as multicellular antibiotic-refractive aggregates called biofilms in situ has further emphasised the challenge facing antibiotic therapies in the clinic. Therefore, there is an urgent need for the development of alternative therapies that (1) can target these multidrug-resistant isolates in the biofilm state and (2) are less likely to select for resistant strains (Cegelski et al. 2008; Cooper and Shlaes 2011; LaSarre and Federle 2013).

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One of the new generation of promising targets to emerge with the capacity to fulfil both criteria is the bacterial quorum sensing system (QS) (Atkinson and Williams 2009; Njoroge and Sperandio 2009). QS is a population-density-dependent cell-cell communication system affording bacteria the capacity for multicellular control of gene expression linked to virulence and pathogenesis. A broad spectrum of QS systems have been reported, with Gram-negative and Gram-positive bacteria favouring distinct signalling components. Gram-negative bacteria use a spectrum of signalling molecules as autoinducers for cell-cell communication, the principal among these being the *N*-acyl-homoserine lactones (AHLs). Gram-positive bacteria typically employ autoinducing peptides (AIPs) for this role. QS systems are central to the regulation of the biofilm mode of growth and the expression of virulence phenotypes (Ng and Bassler 2009; Schertzer et al. 2009). Biofilm formation is a major virulence factor of clinical human pathogens, and this mode of growth can lead to bacteria being 10–1000-fold more resistant to conventional antibiotics (Høiby et al. 2011). As it has been previously reported, QS systems are not directly involved in essential cellular processes (Rasmussen and Givskov 2006). Thus, molecules that inhibit the QS system, and by extension biofilm formation, instead of bacterial growth, commonly called quorum quenching (QQ) molecules or quorum sensing inhibitors (QSi) have been suggested as a powerful strategy for disease control (Kalia 2013; Rasmussen and Givskov 2006).

Since the first published studies on QQ (Dong et al. 2000; Manefield et al. 1999), several studies have been undertaken to discover novel QSi isolated from different organisms and different environments (Kalia 2013; Kalia and Purohit 2011; Koh et al. 2013), while new QSi have also been synthesised based on the structure of natural known QS inhibitors (Kim et al. 2008). In this sense, the plant kingdom has been described as a rich source of potent molecules to develop new drugs for medicine (Hanson 2003; Koh et al. 2013; Savoia 2012), and some plant extracts have been shown to have QS inhibition activity (Choo et al. 2006; Chu et al. 2013; Vattem et al. 2007). Some of the most promising bioactive molecules found in plants are from the coumarin family, a large structurally diverse family of plant phenolic compounds characterised by their pharmacological properties (Venugopala et al. 2013). The most important medical applications of these compounds have been related to their antitumour (Devji et al. 2011; Jamier et al. 2014; Vanamala et al. 2006), anti-inflammatory (Bucolo et al. 2009; Fylaktakidou et al. 2004) and anticoagulant activities (Golfakhrabadi et al. 2014; Payá et al. 1992). Several studies have been conducted analysing the role of some of these compounds against bacterial pathogens (Fig. 1). Some coumarin derivatives have been described for their antimicrobial activities (Ojala et al. 2000; de Souza et al. 2005), for their inhibition of virulence phenotypes such as biofilm and motility (Lee et al. 2014; Zeng et al. 2008), and

in the case of 2 furanocoumarins (dihydroxybergamottin and bergamottin) for their inhibition of the AI-1 and AI-2 QS systems (Girenavar et al. 2008). Therefore, we selected coumarin, the prototype molecule of the family, to analyse its ability to inhibit QS signalling in a range of pathogenic organisms, focusing particularly on the classical AHL-based system of Gram-negative bacteria. We also extended this analysis to assess the capacity for coumarin to inhibit different virulence phenotypes in bacterial pathogens and to link these inhibition effects with its possible QS inhibition mode of action. The growth-independent QQ activity of coumarin was found to be particularly potent towards the respiratory pathogen *Pseudomonas aeruginosa* and extended to a range of Gram-negative organisms. The anti-biofilm activity of coumarin towards this pathogen is highly significant in light of the global morbidity and mortality associated with respiratory infection, particularly the chronic biofilm infections that are characteristic of a range of respiratory diseases.

Materials and methods

Bacterial strains and culture conditions The bacterial strains used in the present study are summarised in Table 1. Luria-Bertani (LB, agar or broth) was routinely used to grow *Edwardsiella tarda*, *Escherichia coli* MUH, *P. aeruginosa* PA14, *Salmonella typhimurium* C5369, *Staphylococcus aureus* NCDO949 and *Stenotrophomonas maltophilia*. *Listeria monocytogenes* and *Aliivibrio fischeri* MJ11 were grown on brain-heart infusion (BHI, agar or broth) and LB salt medium (agar or broth), respectively. All of the strains used in this study were grown at 37 °C, with the exception of *A. fischeri* MJ11, which was routinely grown at 23 °C. Culture media were supplemented with the appropriate antibiotics when necessary (carbenicillin 200 µg ml⁻¹, gentamicin 30 µg ml⁻¹ and tetracycline 25 µg ml⁻¹). Coumarin was purchased from Sigma-Aldrich (UK) and was dissolved in methanol. Different AHLs used in this work were purchased from Sigma-Aldrich (UK) and were dissolved in DMSO.

Quorum quenching biosensor assay The following quorum quenching reporter strains were used to identify quorum quenching ability of coumarin: *Serratia marcescens* SP15 for the detection of short AHLs (Slater et al. 2003), *Chromobacterium violaceum* DSM30191 for the detection of medium AHLs (McClellan et al. 1997) and *Agrobacterium tumefaciens* NTL4 for the detection of long AHLs (contains a plasmid pZLR4 carrying a *traG::lacZ* reporter fusion; Farrand et al. 1996; Yin et al. 2012). Different amounts of coumarin (150, 125, 100, 75, 50 and 25 µg) were inoculated on a TLC Silica Gel 60 (Millipore, UK) and overlaid with soft LB agar (0.5 % agar) inoculated with the different biosensors strains at an OD 600 nm of 0.5. Methanol was used as a control. The

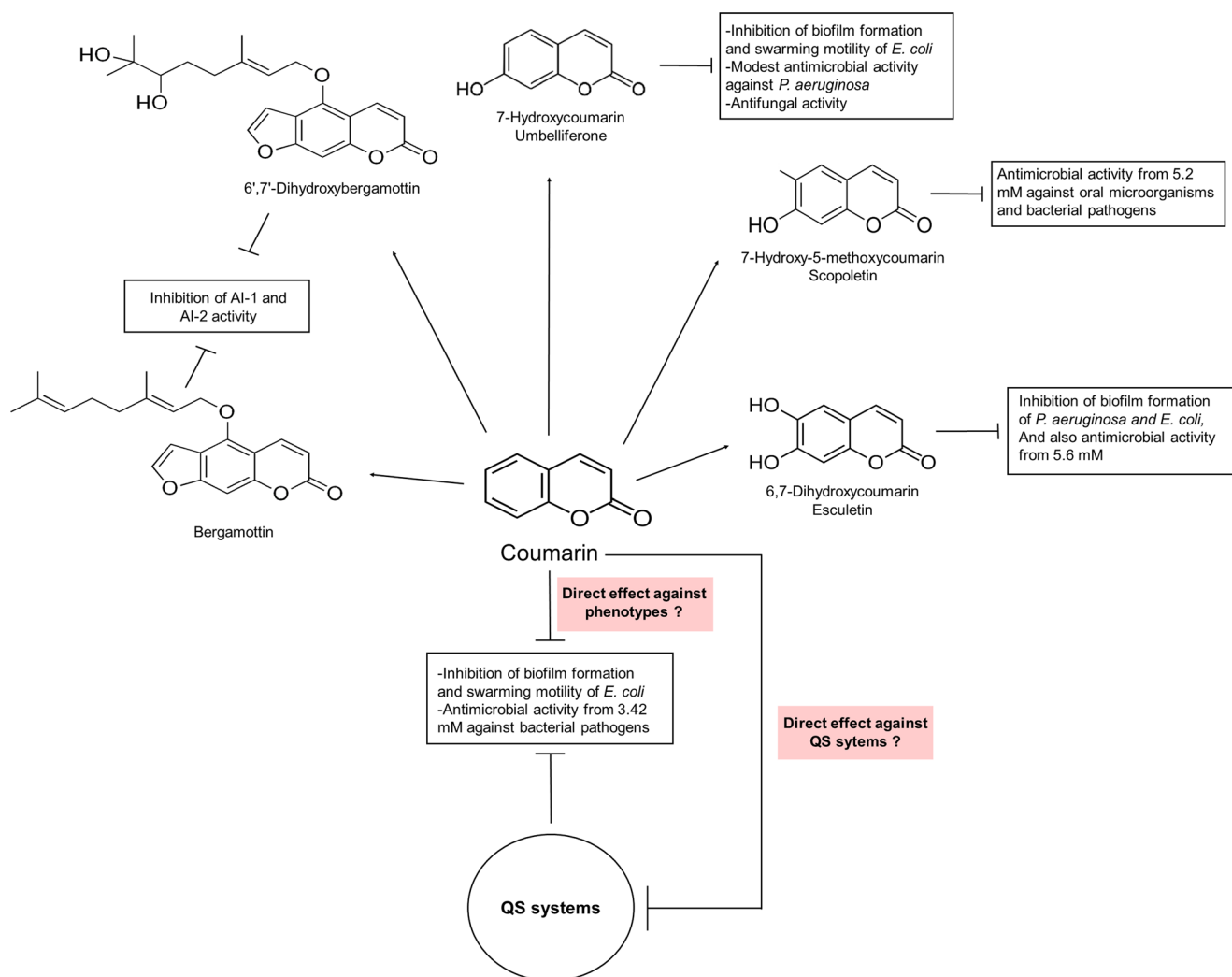


Fig. 1 Chemical structures of the prototype molecule of the coumarin family, coumarin and some derivative molecules

previous plating protocol was used for the different biosensor strains with the exception that soft LB agar for *A. tumefaciens* NTL4 was supplemented with 60 $\mu\text{g ml}^{-1}$ of 5-bromo-4-chloro-3-indolyl- β -D-galactopyranoside and 10 and 40 μM of 3OC10, 3OC12 and 3OC14 homoserine lactones (Sigma-Aldrich, UK), as this strain does not produce its own AHL.

Range of quorum quenching activity against different *N*-acyl homoserine lactones To elucidate the range of quorum quenching AHL specificity of coumarin, the reporter strains commonly used to detect quorum sensing activity were used. *S. marcescens* SP19 (Poulter et al. 2010) was used to detect the QQ activity of coumarin against short chain AHLs (C4-HSL and C6-HSL). *C. violaceum* CV026 was used to detect the QQ activity of coumarin against medium chain AHLs (C6-HSL and 3OC8-HSL) (McClean et al. 1997). *A. tumefaciens* NTL4, the same strain used in the quorum quenching biosensor assay, was used to detect QQ activity against long chain AHLs. LB agar plates inoculated with

S. marcescens SP19 and *C. violaceum* CV026 at an OD 600 nm of 0.5 were prepared. Wells were made in the agar plates, and 20 μl of a 200 μM concentration of C4-HSL, C6-HSL and 3OC8-HSL were added. One hundred and 125 μg of coumarin were also added. To determine the effect of coumarin against long AHLs, a different protocol was used. In 96-well microtitre plates (Starstedt, UK), the wells were filled with LB agar supplemented with 60 $\mu\text{g ml}^{-1}$ of 5-bromo-4-chloro-3-indolyl- β -D-galactopyranoside, with the biosensor strain *A. tumefaciens* NTL4 at an OD 600 nm of 0.5, with 20 μl of a 50 and 100 μM concentration of the different long AHLs (3OC10-HSL, 3OC12-HSL and 3OC14-HSL) and with 100 and 125 μg of coumarin. Methanol was used as control in both protocols.

β -Galactosidase assay *P. aeruginosa* PAO1 transformants containing relevant promoter fusions of the quorum sensing genes *lasI*, *rhlI* and *pqsA* (Table 2) were analysed. Overnight cultures were diluted to OD 600 nm 0.02 in 20 ml LB in the

Table 1 Bacterial strains used in this study

Strains	Relevant characteristics	Reference/source
Gram negative		
<i>Burkholderia cepacia</i> NCTC10743	Opportunistic human pathogen	NCTC
<i>Edwardsiella tarda</i>	Fish pathogen and opportunistic human pathogen	UCC/Biomerit collection
<i>Escherichia coli</i> MUH76317	Multidrug-resistant pathogen	MUH
<i>Pseudomonas aeruginosa</i> PA14, DSM19882	Opportunistic human pathogen	DSMZ
<i>Salmonella typhimurium</i> C5369	food-borne pathogen, causes listeriosis	UCC/Biomerit collection
<i>Stenotrophomonas maltophilia</i>	Uncommon pathogen in humans	A. van der Sar, Vumc, Amsterdam
<i>Vibrio anguillarum</i>	Salmonid fish pathogen	UCC/Biomerit collection
<i>Allivibrio fischeri</i> MJ11	Isolated from <i>Monocentris japonica</i> , bioluminescent strain, symbiont	Ruby 1996
Gram positive		
<i>Listeria monocytogenes</i> LO28	Serotype 1/2c	C. Hill lab, UCC
<i>Staphylococcus aureus</i> NCDO949	Serotype 3	NCDO

NCTC National Collection of Type Cultures, Culture Collection of Public Health England; MUH Mercy University Hospital, Cork, Ireland; DSMZ DSMZ-German Collection of Microorganisms and Cell Cultures, Leibniz Institute; NCDO National Collection of Dairy Organisms, Reading, UK

presence and absence of coumarin (1.36 mM) grown at 37 °C with shaking at 150 rpm. Beta-galactosidase activity was measured at different times: 2, 4, 6, 8.5 and 24 h, as described previously by Miller (1992).

Biofilm formation and attachment Biofilm formation assays were performed as described previously (O'Toole and Kolter 1998) with minor modifications. *P. aeruginosa* PA14 was inoculated from a fresh overnight culture into 500 µl of LB broth in 24-well microtitre plates (Starstedt, UK) at an OD 600 nm of 0.05. Different concentrations of coumarin were tested: 10.0, 5.0, 2.0, 1.71, 1.36 and 1.0 mM; LB broth with methanol was used as control. These concentrations of coumarin were equivalent in coumarin content to those spotted on the TLC plates (e.g. 150, 125, 100 and 75 µg), in a final volume of 500 µl. All biofilms were incubated at 37 °C for 24 h statically. Attachment assays were conducted as described previously (McCarthy et al. 2014) with minor modifications. PA14 was inoculated into 500 µl of LB broth in 24-well microtitre plates (Starstedt, UK) at an OD 600 nm of 0.25. Different concentrations of coumarin were tested: 1.36 and 1.71 mM. Cultures were incubated at 37 °C for 3 h statically. Biofilm and attachment were

measured by solubilisation of crystal violet with 96 % ethanol and quantified at 570 nm.

In vitro biofilm formation of PA14 Sterile slides were placed into 50 ml Falcon tubes containing 15 ml of LB broth, with different concentrations of coumarin (1.36 and 1.71 mM). Overnight cultures of *P. aeruginosa* PA14 were used to inoculate 15 ml of LB at an OD 600 nm of 0.05. Tubes were incubated for 24 h at 37 °C without shaking. Slides were gently recovered from the tubes and were washed with sterile distilled water. A simple staining with crystal violet was used to visualise the biofilm formation at the liquid-air interface.

Growth analysis of PA14 Growth curves of *P. aeruginosa* PA14 were carried out in LB broth at 37 °C using different concentrations of coumarin (2.0, 1.71, 1.36 and 1.0 mM; LB broth with methanol was used as control). PA14 was inoculated from a fresh overnight culture at an OD 600 nm of 0.05. Ten-fold serial dilutions in sterile phosphate buffer were carried out every 2 h. Aliquots (100 µl) of each 10-fold serial dilution were plated onto LB agar. The plates were incubated

Table 2 *Pseudomonas aeruginosa* transformants used for promoter expression analysis

<i>P. aeruginosa</i> strains	Relevant characteristics	Source
PAO1 pME3853::PlasI	PAO1 wild type with 174 bp <i>lasI</i> upstream fragment and translational <i>lasI</i> :: <i>lacZ</i> fusion in pME6010	Pessi et al. (2001)
PAO1 pMP220::PrhII	PAO1 wild type with <i>rhII</i> - <i>lacZ</i> promoter fusion in pMP220	Bayse et al. (2005)
PAO1 pLP0996::PpqsA	PAO1 wild type with <i>pqsA</i> - <i>lacZ</i> promoter fusion in pLP170	McGrath et al. (2004)

at 37 °C for 24 h. The number of colony forming units (CFU) per ml were estimated from LB counts.

Antimicrobial activity Aliquots (1 ml) from an overnight fresh culture of *P. aeruginosa* PA14 were washed three times and resuspended in 1 ml of sterile distilled water. The OD 600 nm was measured, and the inoculum was prepared at an OD 600 nm of 0.1. Subsequently, a cotton swab was used to spread the inoculum on the complete surface of the agar plate. Sterile filter paper discs of 6 mm of diameter were used to determine the antimicrobial activity of coumarin. Different amounts of coumarin were inoculated onto the filter paper discs (150, 125, 100 and 75 µg) and were then placed on LB agar plates that were previously inoculated with PA14 as described above.

Phenazine production Overnight cultures of *P. aeruginosa* PA14 were inoculated into 5 ml of LB broth with different concentrations of coumarin (2.0, 1.71, 1.36 and 1.0 mM; LB broth with methanol was used as control) at an OD 600 nm of 0.05. Cultures were incubated at 37 °C for 24 h at 180 rpm. The total volume of the culture was then centrifuged at 4000g for 10 min. Three millilitre of chloroform was added to the supernatant and vortexed. The bottom phase was then recovered following centrifugation at 4000g for 5 min, and 2 ml of 0.2 M HCl was added. The samples were then vortexed for 20 s and allowed to settle, and the absorbance of the pink layer upper phase was measured at 570 nm.

Motility assays Swimming and swarming motility of *P. aeruginosa* PA14 in the presence of coumarin were assayed. For swim plates, LB 0.3 % (w/v) agar was used. For swarm plates, 0.8 % (w/v) nutrient broth Eiken, 0.5 % (w/v) glucose and 0.6 % (w/v) Eiken agar was used. Different concentrations of coumarin were tested in both types of motility (1.36 and 1.71 mM; methanol was used as control). The bacterial cells were gently inoculated using a toothpick at the centre of the agar surface, and the plates were incubated at 37 °C for 24 h.

Bioluminescence of *Aliivibrio fischeri* *A. fischeri* MJ11 strain was grown in LB salt agar at 23 °C for 24 h in the presence of different concentrations of coumarin (2.0, 1.71, 1.36 and 1.0 mM; LB salt agar with methanol was used as control). The results were based on the loss of luminescence in darkness.

Biofilm formation in different bacterial strains The same protocol described above for PA14 was used with minor modifications. Different culture media and times were tested for the different bacterial strains (Table 1). The 1.36 and 1.71 mM concentrations of coumarin were used. LB broth was used for *P. aeruginosa* PA14 and *S. maltophilia* for 24 h; LB salt for *A. fischeri* MJ11 for 24 h; BHI broth for *L. monocytogenes* for

24 h; TSB for *E. coli* MUH, *S. typhimurium* C5369 and *V. anguillarum* for 24 h, and TSB for 48 h was used for *E. tarda* and *S. aureus* NCDO949.

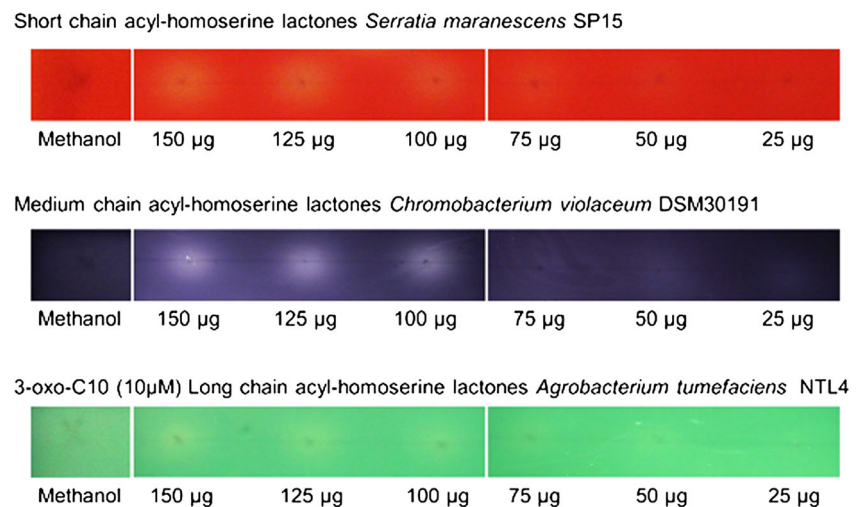
Determination of protease activity Overnight broth cultures of the different strains used in this study (Table 1) were prepared at an OD 600 nm of 1.0. Aliquots (5 µl) were placed onto different agar media supplemented with 1 % (w/v) skimmed milk with different concentrations of coumarin (1.71 and 1.36 mM; methanol was used as control). TSA+ 2 % (w/v) of NaCl was used for *A. fischeri* MJ11, TSA was used for *S. aureus* NCDO949 and *S. maltophilia*, BHI agar was used for *L. monocytogenes*, and LB agar was used for the rest of the strains. Protease activity was evaluated by measuring the zone of clearing around the inoculum spot after 72 h.

Statistical analysis Three independent biological replicates were performed for all experiments. Statistical analysis was performed using two-tailed paired Student's *t* test. Differences were considered significant when the *p* value was ≤0.05.

Results

Coumarin blocks the AHL-QS system Members of the coumarin family of plant phenolic compounds have previously been shown to possess promising pharmacological properties, with evidence of interference with the AI-1 and AI-2 QS systems. However, the structurally distinct coumarin molecule itself has not received attention in this regard. Therefore, to establish whether coumarin can target the classical AHL-based QS signal molecules, a range of QS inhibition experiments were performed. In the first approach, the effect of coumarin against the QQ biosensor reporter strains (*S. marcescens* SP15, *C. violaceum* DSM30191 and *A. tumefaciens* NTL4) was analysed (Fig. 2). Coumarin exhibited a strong QS inhibition activity against all three biosensors; the first evidence that this compound has QQ activity. Both SP15 and DSM30191 displayed pigmentation inhibition at 150, 125 and 100 µg, suggesting that coumarin could be blocking short and medium chain AHL-QS signalling. Inhibition was also observed with the long chain AHL biosensor, *A. tumefaciens* NTL4, which (unlike the other two biosensors) requires the addition of exogenous AHLs for activation. Pigmentation inhibition was detected in the presence of three different long chain AHLs (10 and 40 µM) upon addition of 150, 125 and 100 µg of coumarin to the medium. The greatest inhibition was at the lowest concentration of AHL (10 µM) for 3OC10-HSL. Importantly, to confirm that the reduction of pigment production was related to the inhibition of the QS system, and not any growth defect, re-isolation of the biosensor strains from the inhibition spots resulted in characteristic

Fig. 2 Quorum quenching activity of coumarin. Different amounts of coumarin were spotted onto a TLC plate and then overlaid with the three different quorum quenching biosensor strains: *S. marcescens* SP15 for the detection of short AHLs, *C. violaceum* DSM30191 for the detection of medium AHLs and *A. tumefaciens* NTL4 for the detection of long AHLs. Pigmentation inhibition indicates that coumarin had a quorum sensing inhibition activity



pigment production after 24 h of growth in the absence of coumarin (data not shown).

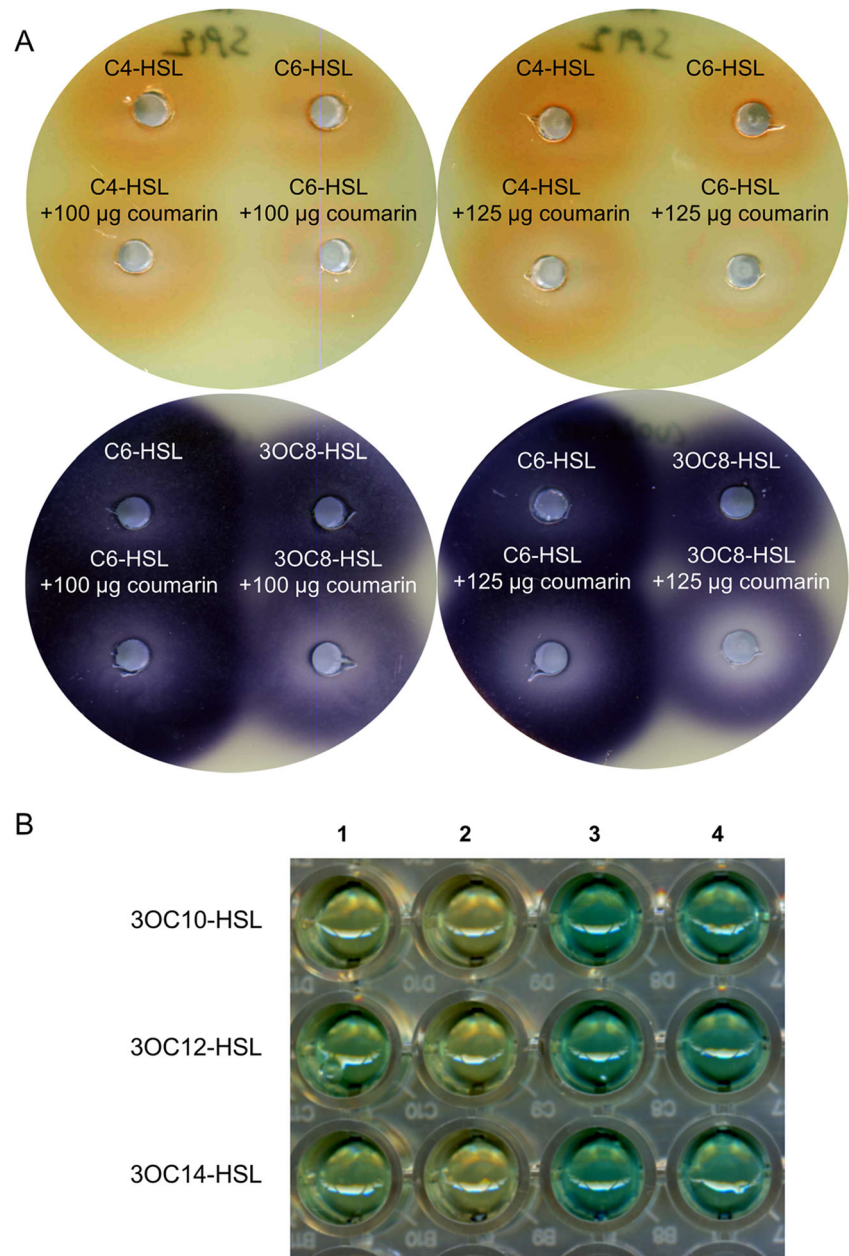
In order to determine whether the activity of coumarin was AHL specific, or broad spectrum, the QS biosensor reporter strains (*C. violaceum* CV026, *S. marcescens* SP19 and *A. tumefaciens* NTL4) were used. Coumarin at 100 and 125 µg was able to inhibit QS system-associated pigment production on all three biosensor strains (Fig. 3). Coumarin at both concentrations displayed a similar inhibition effect against the different AHLs tested (C4-HSL, C6-HSL and 3OC8-HSL). As with the previous result using *Agrobacterium* to test the long AHLs (Fig. 2), coumarin showed the highest inhibition activity against 3OC10-HSL. These results suggested that coumarin could have a broad spectrum of action against AHL-QS signal molecules.

Quorum sensing systems are repressed in the presence of coumarin Having established that coumarin was capable of interfering with QS in all three classical biosensor strains, the next step was to test the activity in a QS active pathogen. For this, the serious nosocomial pathogen *P. aeruginosa* was selected encoding the LasIR and RhlIR AHL-based signalling systems. In addition, *P. aeruginosa* also encodes an alkyl quinolone signalling system (*Pseudomonas* quinolone signal) that interfaces with both AHL systems in a QS hierarchical structure that is characteristic of many pathogenic bacteria. Time course expression analysis of the promoters of the *pqsA* (Fig. 4a) and *rhII* (Fig. 4b) genes revealed that coumarin (at 1.36 mM) decreases the expression of both. The expression of the *pqsA* promoter was decreased in early and late log phase (Fig. 4a), while promoter expression of the *rhII* gene was decreased by coumarin at the stationary phase (Fig. 4b). Interestingly, expression of the *lasI* promoter was unaffected in the presence of coumarin, suggesting that some specificity exists in its interaction with the QS sensory proteins (Fig. 4c).

Coumarin inhibits biofilm formation of *P. aeruginosa* PA14 without inhibiting growth and attachment As QS is central to the control of biofilm and virulence in pathogenic bacteria, the ability of coumarin to interfere with these phenotypes in *P. aeruginosa* was examined. Addition of coumarin led to a significant decrease in biofilm formation of *P. aeruginosa* PA14 using the standard static biofilm formation assay. From the different concentrations tested (from 1.0 to 10 mM), the minimal concentration of coumarin that had an effect in biofilm formation inhibition was 1.36 mM (Fig. 5a). Subsequently, an in vitro biofilm formation assay (Fig. 5b) on slides confirmed this inhibition, enabling visualisation of the attached cells and supporting the previous results obtained by the static biofilm assay. Nevertheless, the presence of coumarin did not alter the cell attachment at any of the concentrations tested (Fig. S1a), suggesting that mature biofilm formation and not initial attachment was the target of the compound. In addition, bacterial growth curves were performed using different concentrations of coumarin (1.0, 1.36, 1.71 and 2.0 mM), to test if coumarin had an effect on bacterial growth (Fig. S1b). The cultures showed typical bacterial growth curves including the different phases (lag, exponential and stationary) during 10 h. No significant differences were observed under the different concentrations of coumarin. In addition, an antimicrobial activity assay on LB agar plates using coumarin from 1.0 to 2.0 mM revealed that coumarin did not show antimicrobial activity against *P. aeruginosa* PA14 at these concentrations (Fig. S2a). Therefore, coumarin suppresses biofilm formation in *P. aeruginosa* in a growth independent manner.

Coumarin is capable of inhibiting additional QS-regulated virulence phenotypes in *P. aeruginosa* Phenazines have been shown to act as an important virulence factor against bacteria, fungi and mammals (Lau et al. 2004; Price-Whelan et al. 2006). Importantly, phenazine production is under the control of QS in *P. aeruginosa*. Consistent with the QSi activity of

Fig. 3 Coumarin presented a broad spectrum of action against different AHLs. **a** The quorum sensing biosensor strains, *S. marcescens* SP19 and *C. violaceum* CV026, were used to detect coumarin activity against short and medium AHLs. **b** *A. tumefaciens* NTL4 was used to detect the specific activity of coumarin against long AHLs. 1, 20 μ l of a 100 μ M of AHL+100 μ g of coumarin; 2, 20 μ l of a 100 μ M of AHL+125 μ g of coumarin; 3, 20 μ l of a 100 μ M of AHL; 4, 20 μ l of a 100 μ M of AHL+methanol



coumarin, phenazine production was inhibited by coumarin in a concentration-dependent manner in *P. aeruginosa* PA14 at 24 h (Fig. 6). Significant differences were observed between the control with methanol and each coumarin concentration tested.

Another multicellular virulence marker in *P. aeruginosa*, swarming motility has also been described as affecting biofilm architecture in this species (Shrout et al. 2006). Swarming motility of *P. aeruginosa* PA14 with and without different concentrations of coumarin (Fig. 7a) was evaluated by measuring the length of dendrites from the centre (Fig. 7b). The dendrite length of cells grown with 1.36 and 1.71 mM of coumarin were 56.6 ± 2.80 and 32.6 ± 2.51 mm respectively.

Significant differences were observed when compared with the control in swarm plates with methanol (69.3 ± 6.8 mm). Thus, the swarming motility of *P. aeruginosa* PA14 was decreased by the addition of coumarin. However, swimming motility was unaffected by the presence of coumarin (Fig. S2b). Taken together, coumarin exhibits bioactivity towards distinct QS-associated virulence systems in *P. aeruginosa*.

Coumarin exhibits broad spectrum species-dependent inhibition of QS-associated phenotypes The AHL-based QS system is widespread among Gram-negative bacteria. Therefore, the analysis of coumarin QSi activity was extended to a range of phylogenetically distinct bacteria. The QS paradigm was first

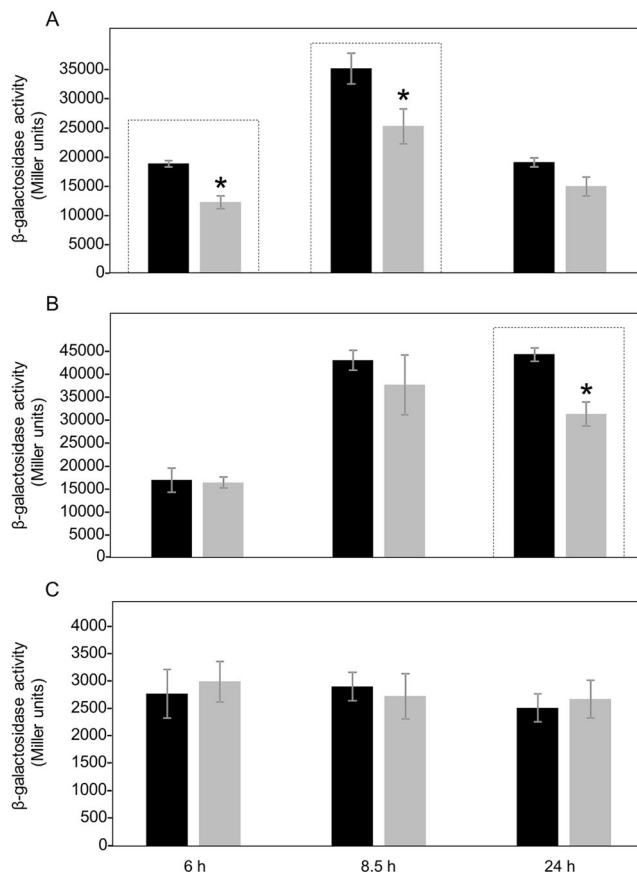


Fig. 4 QS is repressed in response to coumarin. **a** Promoter activity of the *pqsA* gene was significantly reduced in the presence of 1.36 mM of coumarin. The expression of the *pqsA* promoter was decreased in early and late log phase. **b** Promoter activity of the *rhII* quorum sensing gene was significantly reduced by the addition of coumarin at 1.36 mM. The expression of the *rhII* promoter was decreased at the stationary phase. **c** Promoter activity of the *lasI* quorum sensing gene was unaffected in the presence of coumarin. All experiments were performed in triplicate. Black bars: control treated with methanol and grey bars: 1.36 mM of coumarin. Significant differences with respect to the control were represented by asterisk

reported in bioluminescent *A. fischeri*, a phenotype that is regulated under the LuxI/R QS system. Consistent with the QSi activity of coumarin, bioluminescence was reduced in all concentrations assayed (from 2.0 to 1.0 mM) (Fig. 8a), without any impact on the bacterial growth observed directly on agar plates.

As with *P. aeruginosa*, biofilm formation is under the control of QS in many pathogens. Therefore, the effect of coumarin on the biofilm formation on different opportunistic human pathogens, fish pathogens and symbiotic bacteria was analysed (Table 1). Of the 10 bacterial strains analysed, biofilm was significantly reduced in five of them (*P. aeruginosa* PA14, *E. coli* MUH, *V. anguillarum*, *E. tarda* and *S. aureus* NCDO949) (Fig. 8b). Interestingly, of these, only two possess homologues of AHL genes, with others encoding AI-2 or even AIP systems. *E. coli* utilises the AI-2 QS signal molecule to

control the biofilm formation (González Barrios et al. 2006), while *S. aureus*, a Gram-positive pathogen, uses AIP signaling. Therefore, the QSi activity of coumarin may extend beyond the classical AHL system, although we cannot rule out an AHL-responsive sensor protein in these organisms.

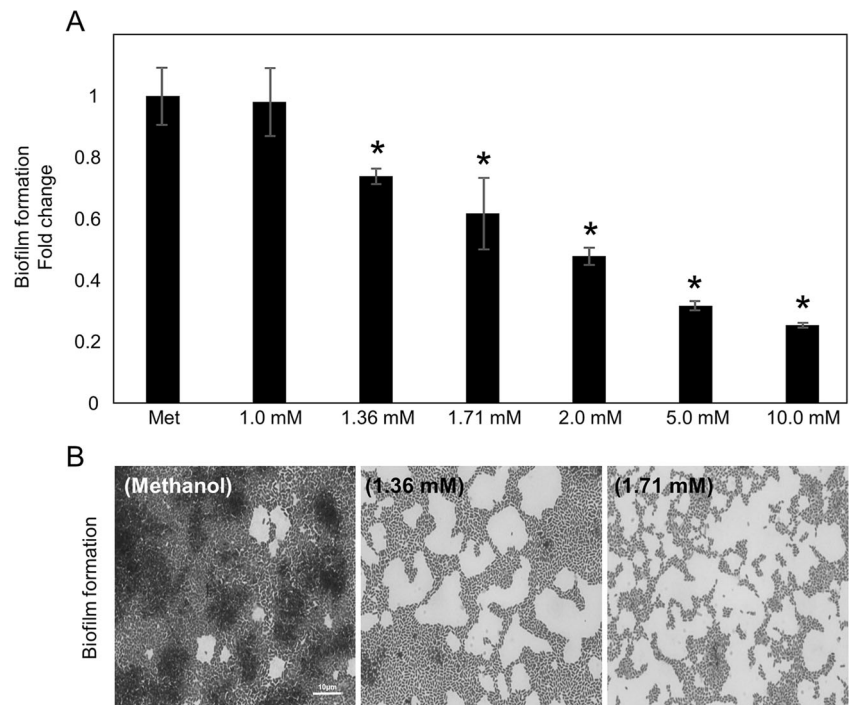
The impact of coumarin on QS-associated protease activity was investigated in the same 10 bacterial strains (Table 1). Five of the bacterial strains tested were positive for protease activity: *P. aeruginosa* PA14, *V. anguillarum*, *S. maltophilia*, *B. cepacia* and *S. aureus*. *P. aeruginosa* PA14 and *V. anguillarum* displayed the highest protease activity, which was unaffected by the addition of coumarin, as was the case for *S. aureus* (data not shown). In contrast, coumarin was able to inhibit protease activity in *S. maltophilia* and *B. cepacia* (Fig. 8c), suggesting that the control of protease activity by QS may be species dependent.

Discussion

The emergence of multidrug-resistant bacteria has become a major global concern, with predictions of the perfect storm receiving increased attention in the face of sharp declines in alternative drug development (Cooper and Shlaes 2011; Gould 2009). Research efforts have shifted from ‘killing’ compounds to the search for new anti-virulence anti-biofilm compounds, providing an effective alternative to controlling bacterial infections (Kalia 2013; LaSarre and Federle 2013; Rasmussen and Givskov 2006). In this study, we describe a new member of a growing class of next generation compounds which exploit a key virulence switch in bacterial pathogens termed quorum sensing.

Coumarin is a natural molecule derived from plants, whose pharmacological safety has been established. Exposure to coumarin (from food or cosmetics products) does not pose any risk for human health, although high doses have been associated with liver toxicity in rodents (Felter et al. 2006; Lake 1999). Furthermore, coumarin has been recently shown to possess anti-biofilm properties against *E. coli* (Lee et al. 2014). Taken together with the fact that the biofilm mode of growth is under the control of QS systems in Gram-negative bacteria (Solano et al. 2014), these findings suggested that coumarin could have potential as an effective QSi anti-virulence compound. Therefore, to establish whether or not coumarin had QSi activity, the ability of the pure compound to inhibit AHL-QS systems was investigated using classical AHL QQ biosensor reporter strains. Coumarin was able to inhibit pigment production on all three QQ biosensor reporter strains and subsequently displayed a broad range of inhibition activity of different AHLs (Figs. 2 and 3). These data suggest that coumarin may inhibit native AHL-QS systems in bacterial pathogens and by extension, therefore, could be an effective

Fig. 5 Reduction of biofilm formation in *P. aeruginosa* at 24 h by the addition of different concentrations of coumarin. Significant differences with respect to the control are represented by an asterisk



inhibitor of virulence traits in these organisms. Importantly, all QSi effects were growth independent, and concentrations were below the MIC for this compound (de Souza et al. 2005).

In this study, coumarin was able to inhibit biofilm formation (Fig. 5), phenazine production (Fig. 6) and swarming motility (Fig. 7) without impairing *P. aeruginosa* growth capacities (Fig. S1b). These constitute three of the major QS-regulated virulence phenotypes in *P. aeruginosa* and other pathogens (Daniels et al. 2004; De Kievit 2009; Köhler et al. 2000). Biofilm formation presents a major challenge to the clinical management of infection, with chronic and medical device infections proving particularly refractive to conventional therapeutics. Initiated by the formation of microcolonies on a surface, maturation and subsequent dispersal underpin the persistence of a broad spectrum of serious pathogens. Both

phenazine production and swarming motility are also multicellular phenotypes which are central to the pathophysiology of infection. It is worth noting that both phenotypes are intrinsically linked to biofilm formation in *P. aeruginosa*. Swarming motility has been reported to contribute to the early stages of *P. aeruginosa* biofilm formation (Shrout et al. 2006), playing an essential role in the surface colonisation contributing to the formation of the biofilms (Verstraeten et al. 2008). The role of phenazines in the formation of mature biofilms has also been described (Ramos et al. 2010). Together, modulation of both virulence systems may contribute to the suppression of biofilm formation by coumarin.

In order to gain an insight into the possible mode of QS inhibition by coumarin, *P. aeruginosa* was used as a model organism. *P. aeruginosa* encodes three different QS systems:

Fig. 6 Coumarin displays an inhibition of phenazine production. Phenazine production was calculated with respect to the methanol control, and significant differences are represented by an asterisk

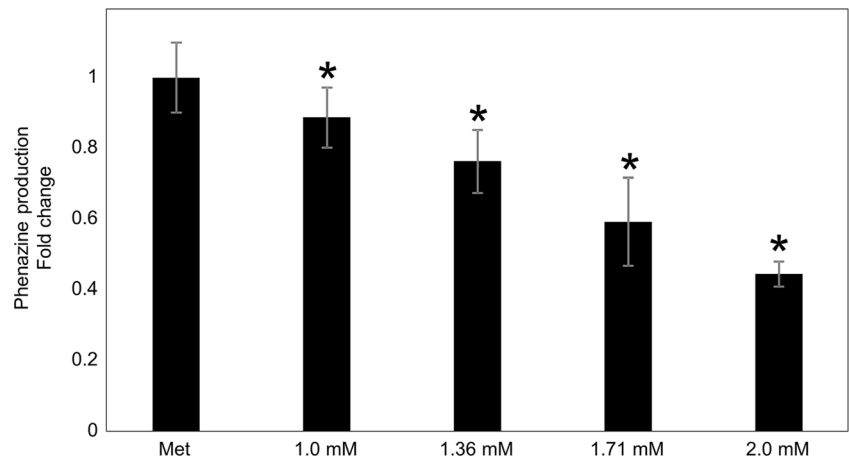
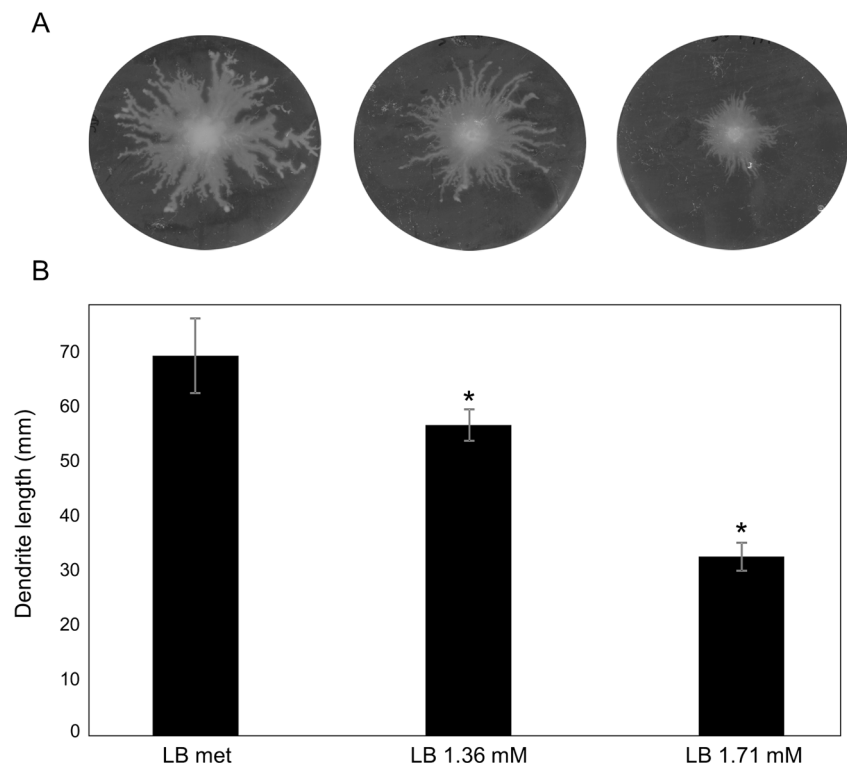


Fig. 7 Impact of coumarin on swarming motility. **a** Coumarin inhibited swarming motility of *Pseudomonas aeruginosa* PA14; from the left to the right: swarm plate+methanol, swarm plate+1.36 mM of coumarin and swarm plate+1.71 mM of coumarin. **b** Swarming motility was measured by the length of dendrites from the centre of the plate. Significant differences with respect to the control are represented by an asterisk



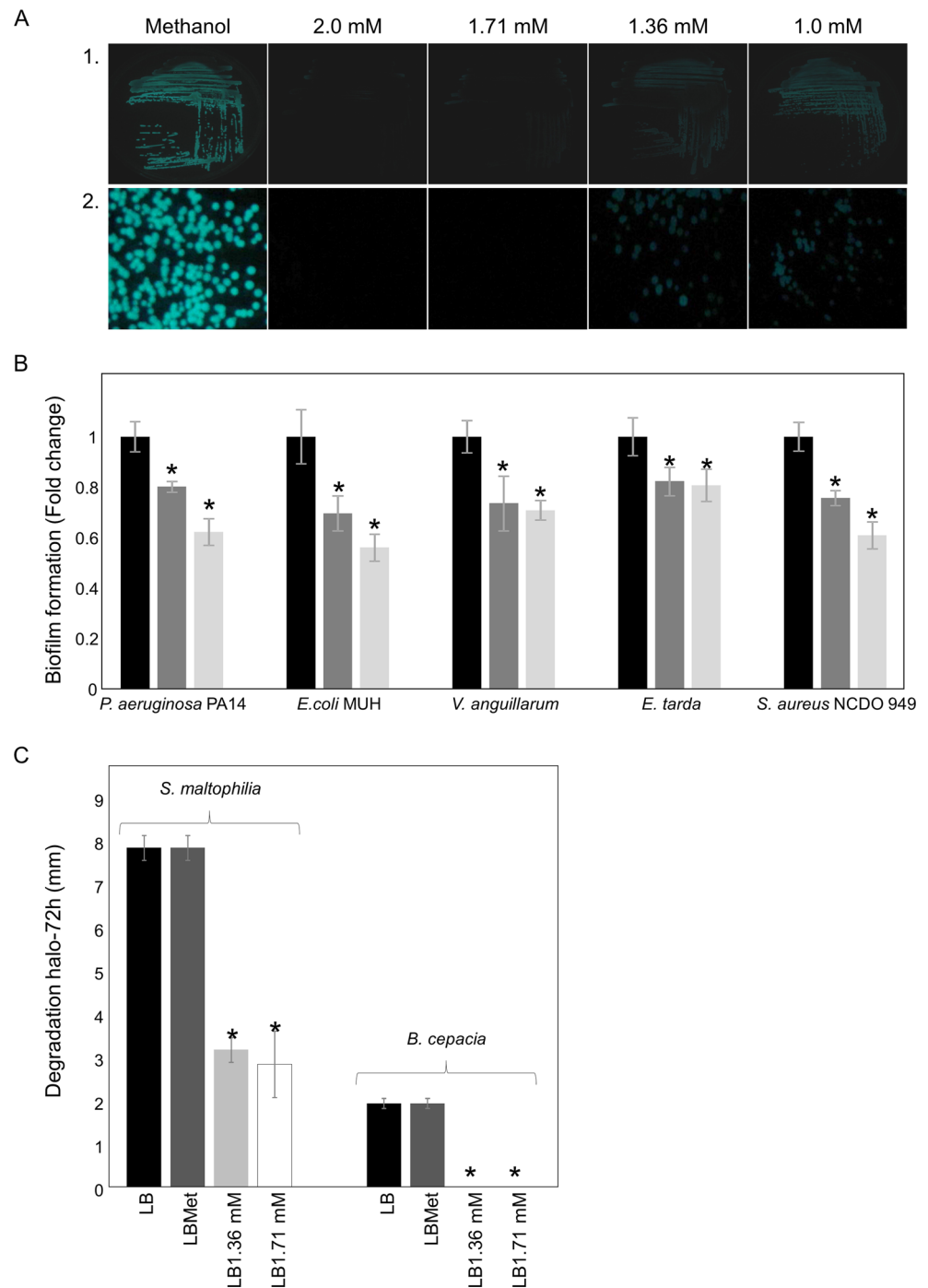
LasI/R, RhII/R which are the classical AHL-based systems and a species-specific *Pseudomonas* quinolone signal (PQS) system, all of which regulate biofilm and virulence phenotypes in this species (Bacalso et al. 2011; Bjarnsholt et al. 2010; Häussler and Becker 2008). These systems in the QS network of *P. aeruginosa* are arranged hierarchically and control the transcription of hundreds of genes (Dietrich et al. 2006; Wilder et al. 2011). All three QS systems have been shown to play a role in the formation of mature biofilms, with RhII/R and PQS in particular associated with the latter stages of development (Davies 1998; Favre-Bonté et al. 2003; Yang et al. 2009). It was perhaps unsurprising, therefore, that reduced promoter activity of *rhII* and *pqsA* was observed in the presence of coumarin. Phenazine production in *P. aeruginosa* is under the control of the PQS signalling system (Déziel et al. 2004). Therefore, the inhibition of phenazine production by coumarin (Fig. 6) is perhaps a result of the reduced expression of the *pqsA* promoter (Fig. 4a). Swarming motility in *P. aeruginosa* is modulated by RhII/R-associated biosurfactant rhamnolipids (Caiazza et al. 2005; Ochsner et al. 1994), with the reduced expression of the *rhII* promoter (Fig. 4b) offering a possible mechanism for its suppression. Transcriptional promoter activity of *lasI* was not influenced (Fig. 4c), suggesting that the activity of coumarin is linked to the interaction of the signal molecule with the LasR receptor and not the production of the signal per se.

In establishing coumarin as a QSi compound, the question remained as to its spectrum of activity. Importantly, while the AHL-based QS system exhibits a degree of conservation in

Gram-negative bacteria, its circuitry and downstream targets can be quite diverse. Therefore, further experiments analysing the effect of coumarin on different quorum sensing regulated phenotypes in different bacterial species were undertaken. The first AHL molecule and its regulatory circuit were discovered in the bioluminescent marine bacterium *A. fischeri* (formerly *Vibrio fischeri*) (Ruby 1996). This bacterium is found in symbiosis with marine animals and has excellent bioluminescent properties, controlling itself through the LuxI/R QS system, homologous to the LasI/R and RhII/R systems of *P. aeruginosa* (Ruby 1996; Rutherford and Bassler 2012). Coumarin was capable of inhibiting bioluminescence in *A. fischeri* MJ11 (Fig. 8a), supporting a role for coumarin against the LuxI/R QS system. Protease activity is another important virulence factor (Lebrun et al. 2009) that has been reported to be under the control of QS systems in different bacteria (Hentzer et al. 2003; Suppiger et al. 2013). Coumarin inhibited protease activity in *S. maltophilia* and *B. cepacia* (Fig. 8c).

Biofilm formation was investigated in a range of bacterial pathogens. Some but not all of the bacterial species tested showed inhibition of biofilm formation upon addition of coumarin (Fig. 8b). Coumarin was effective in reducing biofilm formation of *E. tarda* and *V. anguillarum*, both of which encode AHL-QS homologue genes (Milton et al. 1997; Morohoshi et al. 2004). These data suggest that the inhibition effect could be through blocking the AHL signalling system, at least in these species. Interestingly, coumarin also had an impact on biofilm formation in *E. coli* and *S. aureus*. It has

Fig. 8 Coumarin inhibits different QS-associated phenotypes in different bacterial strains. **a** Inhibition of bioluminescence in *Aliivibrio fischeri* by the addition of coumarin. *Aliivibrio fischeri* MJ11 was grown in LB salt at 23 °C at different concentrations of coumarin. **1** The bioluminescence of *Aliivibrio fischeri* was inhibited at all the concentrations of coumarin tested. **2** Image magnification where the decrease in bioluminescence was observed in isolated colonies. This experiment was carried out in triplicate. **b** Coumarin affected biofilm formation in different bacterial strains. Significant differences with respect to the control in each bacterial species were represented by an asterisk. Black bar: control treated with methanol; dark grey bar: 1.36 mM of coumarin; light grey bar: 1.71 mM of coumarin. **c** Coumarin reduces protease activity. Significant differences with respect to the control in each bacterial species are represented by an asterisk



previously been described that *E. coli* utilises a different QS signal molecule, the AI-2 to control the biofilm formation (González Barrios et al. 2006), and the anti-biofilm activity of coumarin towards *E. coli* was previously reported (Lee et al. 2014). The AI-2 signal molecule, produced by LuxS, is a communication signal used by both Gram-negative and Gram-positive bacteria (Surette et al. 1999). In this sense, our observation that coumarin also inhibited the biofilm formation in the Gram-positive pathogen *S. aureus* (Fig. 8b), a process regulated by the *agr* QS system (Periasamy et al.

2012; Yarwood et al. 2004), could support a role for coumarin as a universal QSi.

Inhibition of the QS cell-to-cell communication system is one of the most promising strategies to control bacterial diseases based on its ability to modulate virulence phenotypes in a broad spectrum of pathogenic bacteria. In this study, we have identified the QS inhibition mode of action of coumarin, a natural product isolated from a variety of plants. As we have described in this work, coumarin not only exhibits a potent inhibitory effect against AHL-QS systems but also displays

inhibition towards the species-specific PQS system in *P. aeruginosa*. Coumarin may have the potential for broad spectrum QSi activity, based on the inhibition of virulence phenotypes regulated by different QS systems in different bacterial species. Future work will focus on unravelling, at the molecular level, the mode of action of this promising molecule.

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