



## N-acylhomoserine lactone-degrading bacteria isolated from hatchery bivalve larval cultures

Marta Torres<sup>a</sup>, Manuel Romero<sup>c</sup>, Susana Prado<sup>c</sup>, Javier Dubert<sup>c</sup>, Ali Tahrioui<sup>a</sup>, Ana Otero<sup>c</sup>, Inmaculada Llamas<sup>a,b,\*</sup>

<sup>a</sup> Department of Microbiology, Faculty of Pharmacy, Cartuja Campus, University of Granada, 18071 Granada, Spain

<sup>b</sup> Institute of Biotechnology, University of Granada, 18071 Granada, Spain

<sup>c</sup> Department of Microbiology and Parasitology, CIBUS-Faculty of Biology, University of Santiago de Compostela, South Campus, 15782 Santiago de Compostela, A Coruña, Spain

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### ABSTRACT

Quorum sensing (QS) systems, which depend on *N*-acylhomoserine lactone (AHL) signal molecules, mediate the production of virulence factors in many pathogenic microorganisms. One hundred and forty-six bacterial strains, isolated from a bivalve hatchery, were screened for their capacity to degrade five synthetic AHLs [*N*-butyryl-DL-homoserine lactone (C<sub>4</sub>-HSL), *N*-hexanoyl-DL-homoserine lactone (C<sub>6</sub>-HSL), *N*-octanoyl-DL-homoserine lactone (C<sub>8</sub>-HSL), *N*-decanoyl-DL-homoserine lactone (C<sub>10</sub>-HSL) and *N*-dodecanoyl-DL-homoserine lactone (C<sub>12</sub>-HSL)] using well diffusion agar-plate assays with three biosensors, *Chromobacterium violaceum* CV026, *C. violaceum* VIR07 and *Agrobacterium tumefaciens* NTL4 (pZLR4). The results of these assays led to our choosing four strains (PP2-67, PP2-459, PP2-644 and PP2-663) that were able to degrade all five synthetic AHLs, thus showing a wide spectrum of quorum quenching (QQ) activity. We subsequently confirmed and measured the QQ activity of the four strains by high-performance liquid chromatography plus mass-spectrometry analysis (HPLC-MS). One of the strains which showed the highest AHL-degrading activity, PP2-459, identified as being a member of the genus *Thalassomonas* was chosen for further study. Finally, using thin-layer chromatography (TLC), we went on to confirm this strain's capacity to degrade the AHLs produced by other non-pathogenic and pathogenic bacteria not taxonomically related.

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### 1. Introduction

Bacterial diseases in bivalve hatcheries are a considerable limiting factor in commercial aquaculture and result in substantial economic losses. For many years treatment with antibiotics has been the only viable strategy to tackle the problem. Nevertheless, their use does nothing to guarantee the ultimate success of larval cultures and their overuse has resulted in the development of resistance. The seriousness of the situation has recently promoted the exploration of novel strategies to control marine pathogens, but only some of them, such as the use of probiotics, are applicable to bivalve aquaculture (Defoirdt et al. 2007a, 2011a; Prado et al. 2010; Verschuere et al. 2000). One of the most promising alternatives is based on the inhibition of the expression of virulence genes, regulated in many aquaculture pathogens by bacterial cell-to-cell signalling, known as quorum sensing (QS)

(Bjarnsholt et al. 2011; Dong et al. 2007; González and Keshavan 2006; Natrah et al. 2011). QS is a population-density-dependent gene-expression mechanism which involves the production of signal molecules known as autoinducers, a ubiquitous phenomenon in bacteria (González and Marketon 2003). The most thoroughly characterized Gram-negative, bacterial intraspecific autoinducers are *N*-acylhomoserine lactones (AHLs), which have been reported to accumulate in the culture medium as the density of the population increases, and on reaching a critical threshold concentration bind to an AHL-receptor protein belonging to the LuxR family of transcriptional regulators. The activated LuxR/AHL complex then binds specific DNA sequences, resulting in the activation or repression of target genes, including in many cases the activation of important virulence phenotypes (Eberhard et al. 1991; Fuqua et al. 1994; Natrah et al. 2011).

It has been reported that some aquatic organisms such as micro-algae, macro-algae, invertebrates and also other bacteria have the potential to disrupt QS by means of various different mechanisms (Natrah et al. 2011). One such mechanism involves the production of compounds that interfere with the detection of signal molecules; these compounds, known as quorum sensing inhibitors (QSIs), were first described in the red marine alga *Delisea*

\* Corresponding author at: Department of Microbiology, Faculty of Pharmacy, University of Granada, Cartuja Campus, 18071 Granada, Spain. Tel.: +34 958 241741; fax: +34 958 246235.

E-mail address: [illamas@ugr.es](mailto:illamas@ugr.es) (I. Llamas).



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