

THE ANTIMICROBIAL ACTION OF *PSEUDOMONAS AERUGINOSA* BY-PRODUCTS IN THE CONTROL OF SINGLE AND MIXED BIOFILMS

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KEYWORDS

Biofilm, Gram-negative and -positive bacteria, *Pseudomonas* aeruginosa, by-products, antimicrobial activity.

ABSTRACT

Since bacteria are continuously acquiring resistance to conventional chemical agents, it is urgently needed the development of new strategies for biofilm control. It is well recognised that certain microorganisms represent an important source of novel biologically active compounds, with pronounced antibacterial activity, as secondary metabolites. Such substances are accepted to be essential for their producers, inhibiting other bacteria that compete for common resources. The main goal of this work was to investigate the antimicrobial effect of secondary metabolites secreted by P. aeruginosa on planktonic and sessile growth of several pathogens, in order to later use those molecules as bio-regulation agents. P. aeruginosa supernatants had potential as anti-biofilm agents but only against staphylococcal biofilms since they failed in disturb other biofilm consortia that encompassed Grambacteria. This trait makes them quite ineffective chemical countermeasures against real biofilms.

INTRODUCTION

In nature, bacteria live by interacting and communicating with each other, regardless they belong to the same (intraspecies) or different species (interspecies). One of the major mechanisms of cell-cell communication in bacteria involves the synthesis and release of chemical molecules called diffusible signal molecules (Waters and Bassler, 2005). These signals can be cell-density related (quorum sensing - QS) or produced by bacteria at different stages of growth. Primary and secondary metabolites are recognized to contribute to a wealth of interactions between organisms (Duan *et al.*, 2009) and can include a variety of nutrients, toxic or neutral metabolic byproducts, antibiotics and other signaling molecules. Such products

are released and accumulated in the surrounding environment during bacterial growth (Fuqua and Greenberg, 2002) and can induce expression of certain genes and/or physiological changes in neighbouring cells (Fuqua et al., 1996; Parsek and Greenberg, 2005). The properties of these signals and the response elicited by them are important in ensuring bacterial survival and propagation in natural environments where hundreds of bacterial species coexist (Jayaraman and Wood, 2008) Responses of bacteria to chemical signals are quite varied and can include synergistic and/or antagonistic effects. Most research into interspecies bacterial interactions has focused on the beneficial aspects of these relationships that may include coaggregation (Rickard et al., 2003; Sharma et al., 2005) and conjugation (Ghigo, 2001). These positive interactions give advantages to microorganisms through the transfer of chemical signals, exchange of genetic information, growth promotion and increase of metabolic activity (Shank and Kolter, 2009), and protection from adverse environmental conditions (Leriche et al., 2003). Positive interactions among competitors can even contribute to biodiversity (Gross, 2008). However, not all interactions are beneficial, since antagonistic interactions play an important role in bacterial species predominance. Competition for substrate is considered to be the major evolutionary driving force in the microbial world (Simões et al., 2007). Negative interactions can give rise to sporulation, suppression of respiration (Hoffman et al., 2006), growth inhibition through the production, for instance, of antimicrobial compounds, as antibiotics (Tait and Sutherland, 2002; Rao et al., 2005).

In order to investigate the effect derived from the bacterial release of secondary metabolites, *P. aeruginosa* by-products were evaluated on planktonic and sessile growth of several pathogens, in order to later use those molecules as bio-regulation agents. The role of such molecules was evaluated in cell suspensions and in biofilms of single and dual-species cultures formed by important human-associated pathogens.

METHODS

Supernatants from two *Pseudomonas aeruginosa* planktonic cultures (isolated: PaI and from collection: Pa) were recovered, filtered and stored for further experiments.

These supernatants were then tested on their own and on *Staphylococcus aureus*, *Staphylococcus epidermidis*, and *Escherichia coli* lawns.

After observing the results from the disk diffusion tests, the antimicrobial action of the supernatants from both *P. aeruginosa* strains was also assessed in single *Staphylococcus* (aureus and epidermidis) biofilm formation and in polymicrobial biofilms formed by those *Staphylococcus* species together with Gram-bacteria (*P. aeruginosa* and *E. coli*).

Supernatants were differently applied: as biofilm growth media complement and as biofilm disruption agents.

RESULTS

Both supernatants inhibited only Gram+ species lawns, being the more remarkable inhibition halos obtained with the isolated *P. aeruginosa* supernatant.

Concerning biofilms, metabolites from both strains can be considered anti-staphylococcal biofilms agents, since their single and mixed biofilm growth was significantly disturbed by both supernatants, regardless their mode of application. However, when staphylococcal species are entrapped in polymicrobial biofilms with *E. coli* and *P. aeruginosa*, supernatants did not exhibit noticeable anto-biofilm activity, mainly hen applied against established biofilms. In general, all mixed biofilms accumulated more mass and had more metabolic activity when submitted to the supernatants aggression.

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AUTHOR BIOGRAPHIES



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