

Lack of AHL-based quorum sensing in *Pseudomonas fluorescens* isolated from milk

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Abstract

Numerous bacteria coordinate gene expression in response to small signalling molecules in many cases known as acylhomoserine lactones (AHLs), which accumulate as a function of cell density in a process known as quorum sensing. This work aimed to determine if phenotypes that are important to define microbial activity in foods such as biofilm formation, swarming motility and proteolytic activity of two *Pseudomonas fluorescens* strains, isolated from refrigerated raw milk, are influenced by AHL molecules. The tested *P. fluorescens* strains did not produce AHL molecules in none of the evaluated media. We found that biofilm formation was dependent on the culture media, but it was not influenced by AHLs. Our results indicate that biofilm formation, swarming motility and proteolytic activity of the tested *P. fluorescens* strains are not regulated by acyl-homoserine lactones. It is likely that AHL-dependent quorum sensing system is absent from these strains.

Key words: milk, *Pseudomonas fluorescens*, biofilm formation, quorum sensing, quorum quenching.

Introduction

Bacteria communicate using small diffusible signalling molecules, called autoinducers, to coordinate gene expression in response to population density in a mechanism known as quorum sensing (QS) (Fuqua and Winans, 1994; Whitehead *et al.*, 2001). Many different types of QS signals were identified and the fatty acid derivatives (N-acyl-homoserine lactones - AHLs), or autoinducer-1 (AI1), produced by Gram-negative bacteria are the best known and most studied (Fuqua *et al.*, 1996; Eberl, 1999; Whitehead *et al.*, 2001). Quorum sensing allows bacteria to control different functions such as surface colonization and motility, production of exopolymers and antibiotics, biofilm development, bioluminescence, cell differentiation, competence, pigment production, conjugation, sporulation, toxin production, virulence gene expression, and production of several hydrolytic enzymes (Smith *et al.*, 2004). Many of these

phenotypes including biofilm formation, motility, production of enzymes, and toxins are important to define microbial activity in foods.

Although signalling compounds are produced by bacteria in foods (Gram *et al.*, 1999; Cloak *et al.*, 2002; Gram *et al.*, 2002; Christensen *et al.*, 2003; Jay *et al.*, 2003; Bruhn *et al.*, 2004; Jay, 2005), the role of QS in food deterioration is unknown. Pinto and collaborators (Pinto *et al.*, 2007) demonstrated that AHL-production is common among many psychrotrophic bacteria isolated from raw milk. Since all these microorganisms were isolated from the same source, cross-communication is relevant and raises the question of the kinds of phenotypes that are regulated when they are all growing together, and the relation of those to spoilage. The understanding of the role of QS in regulating spoilage phenotypes in bacteria is relevant and may be used to create new ways to preserve food products (Pinto *et al.*, 2007). Pillai and Jesudhasan (2006) reinforce that un-

derstanding the mechanism of quorum sensing may hold the key to food preservation and prevention of pathogen and spoilage bacterial growth and persistence in foods.

Among Gram-negative psychrotrophic bacteria, *Pseudomonas* prevails in refrigerated raw milk (Wiedmann *et al.*, 2000; Dogan and Boor, 2003; Pinto *et al.*, 2006), since they present a well-established physiologic mechanism of growth at low temperatures (Jay *et al.*, 2003). In the *Pseudomonas* genus, *Pseudomonas fluorescens* constitutes the major milk deteriorative species (Wiedmann *et al.*, 2000; Dogan and Boor, 2003; Pinto *et al.*, 2006) due to its ability to produce thermostable proteases and lipases that hydrolyze casein and lipids decreasing yield and sensory quality of dairy products (Sørhaug and Stepaniak, 1997; Wiedmann *et al.*, 2000; Dogan and Boor, 2003).

Quorum sensing systems based on AHLs signalling molecules have been identified in several *P. fluorescens* strains, as it has been shown for strain NCIMB 10586 (El-Sayed *et al.*, 2001), 2-79 (Shaw *et al.*, 1997; Cha *et al.*, 1998; Khan *et al.*, 2005), F113 (Koerstgens *et al.*, 2001), 5064 (Cui *et al.*, 2005), 2P24 (Wei and Zhang, 2006), and 395 (Liu *et al.*, 2007). In strain NCIMB 10586, quorum sensing regulates the synthesis of the antibiotic mupirocin through a LuxR-LuxI homologous system from *Vibrio fischeri*; in strain F113, AHLs are synthesized by a novel synthase; in strain 5064 the system regulates biosurfactant production but the details have not been worked out; in strain 2P42 quorum sensing regulates root colonization, biofilm formation and plant disease-suppressive ability through a regulatory system homologous to LuxR-LuxI; and finally in strain 395, AHLs had a slight effect on *aprX* expression through a system not understood. However, several studies have shown that other strains of *P. fluorescens* do not produce AHLs, as it has been the case for strain 1855.344 (Cha *et al.*, 1998), B52 (Allison *et al.*, 1998), and SBW25 (Bruijn and Raaijmakers, 2009). Furthermore, AHL molecules did not influence growth and proteolytic activity of a strain isolated from milk (Pinto *et al.*, 2010) neither the production of antifungal metabolites by *P. fluorescens* 2P24 used as a biocontrol agent (Wei and Zhang, 2006). A better understanding of the role of QS on the spoilage potential of *P. fluorescens* strains from food sources is of great interest due to their importance in food deterioration.

Biofilm formation and bacterial food spoilage due to hydrolytic enzymes generate significant problems to the food industry (Gram *et al.*, 2002). Bacteria can also use swarming motility to colonize nutrient-rich environments, which facilitates colony spreading and accelerates biomass production (Fraser and Hughes, 1999). This work aimed to determine if phenotypes related to food spoilage such as biofilm formation, swarming motility and proteolytic activity of *P. fluorescens* strains isolated from refrigerated raw milk are regulated by AHL molecules.

Material and Methods

Bacterial strains and growth conditions

Two highly proteolytic *P. fluorescens* strains (07A and 041) previously isolated from raw milk (Martins *et al.*, 2005) were chosen for this study due to their spoilage potential of milk and dairy products (Pinto *et al.*, 2006). The transconjugant strains were obtained by cloning the gentamicin-3-acetyltransferase gene on broad-host-range expression vector which was transferred to the wild-type strains as described later. The strains and plasmids used in this study are listed in Table 1.

The strains were grown at 25 °C in Luria-Bertani (LB), King's B and TYEP medium. The transconjugant strains were grown in media supplemented with gentamicin and trimethoprim 20 µg mL⁻¹. *Escherichia coli* MT102 pSB403 was grown at 30 °C in Luria-Bertani (LB) medium supplemented with tetracycline 50 µg mL⁻¹ (Winson *et al.*, 1998). Growth of liquid cultures was monitored spectrophotometrically with an Ultrospec 3100 Pro spectrophotometer (Biochrom, Ltd., Cambridge, England) by measurement of the optical density at 600 nm.

DNA manipulations

Cloning, restriction enzyme analysis, and transformation of *E. coli* were performed using established procedures. PCR was performed with TaKaRa Ex Taq polymerase (TaKaRa Shuzo, Shiga, Japan). Plasmid DNA was isolated using the QIAprep Spin Miniprep kit, and chromosomal DNA was purified with the DNeasy tissue kit. DNA fragments were purified from agarose gels by using the QIAquick gel extraction kit (all kits from Qiagen, Hilden, Germany).

Construction of the *P. fluorescens* transconjugant strains

The gentamicin-3-acetyltransferase gene (GenBank accession number U25061) of pBBR1MCS-5 was amplified by using the primer pair Gem-F (5' ATT ATG CAT GAA CCT GAA TCG CCA GCG G 3') and Gem-R (5' ATT ATG CAT GTT GAA CGA ATT GTT AGG TGG C 3'). The introduced restriction site *Nsi*I is underlined. The amplicon was digested with *Nsi*I and ligated directionally into the broadhost-range expression vector pMLBAD-*aiiA*-Trm^r yielding pMLBAD-*aiiA*-Trm^r-Gm^r. This plasmid containing the *aiiA* gene was transferred to *E. coli* XL1-Blue by transformation. The *aiiA* gene encodes the lactonase enzyme that hydrolyzes the ester bond of the homoserine lactone ring of AHLs, thus inhibiting QS communication (Dong *et al.*, 2000).

Afterwards, plasmids were delivered to *P. fluorescens* strains by triparental mating as previously described (de Lorenzo and Timmis, 1994). Briefly, donor (*E. coli* XL1-Blue pMLBAD-*aiiA*-Trm^r-Gm^r) and recipient strains, as well as the helper strain *E. coli* HB101

Table 1 - Bacterial strains and plasmids used in this study.

Strain	Plasmid	Description	Reference or source
<i>P. fluorescens</i> 07A		Wild type	(Martins <i>et al.</i> , 2005)
<i>P. fluorescens</i> 07A-2	pMLBAD-aiiA-Trm ^f -Gm ^f	Transconjugant, express the lactonase enzyme	This study
<i>P. fluorescens</i> 041		Wild type	(Martins <i>et al.</i> , 2005)
<i>P. fluorescens</i> 041-3	pMLBAD-aiiA-Trm ^f -Gm ^f	Transconjugant, express the lactonase enzyme	This study
<i>E. coli</i> MT102	pSB403, Tc ^r	Monitor strain: exhibits the highest sensitivity for 3-oxo-C6-HSL. However, several other AHL molecules are detected by this sensor.	(Winson <i>et al.</i> , 1998; Viana, 2006)
<i>E. coli</i> XL1-Blue	pMLBAD-aiiA- Trm ^f	Donor of pMLBAD-aiiA, Trm ^f that codify the lactonase enzyme	(Wopperer <i>et al.</i> , 2006)
<i>E. coli</i> XL1-Blue	pMLBAD-aiiA- Trm ^f -Gm ^f	Donor of pMLBAD-aiiA, Gm ^f that codify the lactonase enzyme	This study
<i>E. coli</i> HB101	pRK600, Cm ^r	Helper	Laboratory of Microbiology, University of Zürich

(pRK600), were grown overnight in 5 mL of LB medium supplied with the appropriate antibiotics. After culturing to an optical density of 0.9 at 600 nm, 2 mL of cell culture were harvested, washed, and resuspended in 500 µL of LB medium. Donor and helper cells (100 µL each) were mixed and incubated for 10 min at room temperature. Then, 200 µL of the recipient cells were added and the mixture was spot inoculated onto the surfaces of pre-heated LB agar plates. After overnight incubation at 30 °C, the cells were plated on *Pseudomonas* Isolation Agar (PIA) (Becton Dickinson Biosciences, Sparks, MD) containing antibiotics for counter selection of the donor, helper, and untransformed recipient cells.

Quantification of AHLs

Since *E. coli* MT102 pSB403 is able to detect low amounts of AHL, it was used for AHL detection (Winson *et al.*, 1998). The plasmid pSB403 contains the *Vibrio fischeri luxR* gene with *luxI* promoter region as a transcriptional fusion to the bioluminescence genes *luxCDABE*. The *V. fischeri* quorum sensing system relies on 3-oxo-C6-HSL, so the sensor plasmid consequently exhibits the highest sensitivity towards this AHL molecule. However, several other AHL molecules are detected by the sensor strain, albeit with reduced sensitivity (Winson *et al.*, 1998). A volume of 1 mL of overnight culture of *E. coli* MT102 pSB403 was inoculated into 5 mL LB supplemented with tetracycline and incubated at 30 °C for 1 h. Then, 100 µL of filter-sterilized culture supernatants in LB of *P. fluorescens* 07A and 041 were added to 100 µL of an exponential culture of the sensor strain in the wells of a microtiter dish. After incubation at 30 °C for 3 h, the expression of the bioluminescence reporter genes was measured using the program KC4 (Bio-Tek Instruments, Highland Park, Box

998, Vermont, USA). AHL concentrations were determined by comparing bioluminescence signal intensities with a defined concentration (0.1 mg mL⁻¹) of pure 3-oxo-C6-HSL.

Extraction of putative signalling molecules from supernatants of *P. fluorescens*

P. fluorescens 07A and 041 wild-type and transconjugant strains (10⁴ CFU mL⁻¹) were inoculated in 600 mL of King's B, LB, and TYEP. The cultures were incubated with aeration at 25 °C for 20 h or until the population reached 10⁹ CFU mL⁻¹. Then, the cells were harvested by centrifugation at 10,000 g for 20 min at 4 °C and 250 mL of the cell free supernatants were mixed with 100 mL of dichloromethane stabilized with ethanol in a 1,000 mL separating funnel. The mixture was shaken for 3 min with aeration every 20 s. When the two phases were separated, the dichloromethane-phase was collected (lower phase). The upper phase (aqueous phase) was mixed with 100 mL of dichloromethane and shaken again as described above. Lower dichloromethane-phase was collected and mixed with the first one. These steps were repeated until finishing the 600 mL of supernatant. Then, the remaining water was removed with water free MgSO₄ and it was filtered using Whatman paper. The filtered extracts were concentrated in a rotary evaporator at 40 °C, resuspended in 250 µL ethyl acetate, and maintained at -20 °C.

Detection of putative signalling molecules in supernatant extracts of *P. fluorescens*

Thirty milliliter of overnight culture of *E. coli* MT102 pSB403 were inoculated in 150 mL of LB agar. The inoculated LB plates were allowed to solidify and then 6 µL of extracts obtained from the supernatant of King's B, LB, and

TYEP inoculated with *P. fluorescens* 07A were transferred as drops to the plate's surface. Aliquots of 0.6 μL of HHL 1 mg mL^{-1} were used as positive controls. The plates were incubated overnight and the activation of the AHL monitor strain *E. coli* MT102 pSB403 was observed into a dark box that contained a highly sensitive photon-counting camera (C2400-40; Hamamatsu Photonics Herrsching, Germany) as described by Steidle *et al.* (2001).

Phenotypic characterization of wild type and transconjugant strains

Biofilm formation in polystyrene microtiter dishes was assayed essentially as described previously (Pratt and Kolter, 1998) with a few modifications. Cells of *P. fluorescens* 07A and 041 wild-type and transconjugant were grown in the wells of microtiter dishes in 100 μL of LB, minimal medium salt (MMS) or ABC medium supplemented with 10 mM citrate for 48 h at 25 °C. Thereafter, the medium was removed, and 100 μL of a 1% (wt/vol) aqueous solution of crystal violet (CV) was added. After staining at room temperature for 20 min, the dye was removed, and the wells were washed thoroughly. For quantification of attached cells, the CV was solubilized in an 800:120 (v/v) mixture of ethanol and dimethyl sulfoxide, and the absorbance was determined at 570 nm. This assay was done in two biological replicates and five independent experiments.

The ability to form a swarming colony was tested by point inoculating the strains into ABC minimal medium supplemented with 0.1% casamino acids and solidified with 0.4% agar as previously described (Eberl *et al.*, 1996; Huber *et al.*, 2001).

Proteolytic activity was determined in ABC, MMS, and TYEP using azocasein assay as previously described (Christensen *et al.*, 2003). Briefly, this activity was investigated on azocasein by incubating 250 μL of 2% azocasein (w/v) with 150 μL sterile filtered culture supernatant. The mixture was incubated at 30 °C for 12 h. Subsequently, the

mixture was incubated at room temperature for 15 min with 1.2 mL of 10% (w/v) trichloroacetic acid (TCA), and centrifuged for 10 min at 15,000 g. Prior to spectroscopic measurement, 600 μL supernatant were rescued and mixed with 750 μL 1 M NaOH. The proteolytic activity was quantified by the determination of the OD₄₄₀ against a blank reaction mixture with 150 μL culture media or 75 μL Tris-HCl 20 mM, pH 8.0, CaCl₂ 5 mM instead of the enzyme solution. One unit of proteolytic activity was defined as the unit of enzyme activity per hour per μg of protein. The method of Bradford (1976), using bovine serum albumin as a standard, was used to quantify protein concentrations in supernatant of media. This assay also was done in two biological replicates and five independent experiments.

Results and Discussion

Detection of bioluminescence induced by *P. fluorescens*

Potential signalling molecules present in supernatants obtained from *P. fluorescens* 07A and 041 did not induce *E. coli* MT102 pSB403 (Table 2). Therefore, this result suggests that *P. fluorescens* 07A and 041 isolated from cooled raw milk did not produce AHLs able to induce the high sensitive biosensor *E. coli* MT102 pSB403. Pinto and collaborators (2010) used another highly sensitive bioassay strain (*Agrobacterium tumefaciens* KYC55) and did not detect AHL molecules from *P. fluorescens* 07A, confirming the present results. Other biosensor strains have also been used and were not able to detect AHLs in this strain (Pinto *et al.*, 2007). According to Winson *et al.* (1998), there is a significant advantage of using *lux* sensors since the sensitivity to AHL is in picomol to nanomol concentrations over a large linear range.

Table 2 - Values of bioluminescence produced by *E. coli* MT102 pSB403 at 175 nm, after growth in LB broth supplemented with supernatant of *P. fluorescens* strains and supplemented with 3-oxo-C6-HSL. Data represent average of triplicate experiments.

Dilution rate	<i>P. fluorescens</i> 07A	<i>P. fluorescens</i> 041	LB Negative control	3-oxo-C6-HSL Positive control
1/2	13532	15823	14820	Nd ¹
1/4	15261	13831	15003	Nd
1/8	13967	15360	13230	Nd
1/16	14401	16580	15340	Nd
1/32	16977	16036	16720	51802
1/64	15862	14159	14579	28723
1/128	17092	17619	16220	21817
1/256	15716	13420	14943	20942

¹Nd - not determined. The intensity of the signal was higher than the detection limit of the equipment.

Supplementation of LB inoculated with *E. coli* MT102 pSB403 with extracts obtained from different media inoculated with *P. fluorescens*

As no activity derived from signalling molecules was found in the supernatant of LB medium inoculated with *P. fluorescens* 07A and 041 on the microtiter dish assay (Table 2), a correlation between the growth media and the production of signalling compounds was tested, but no AHL was found in the supernatant extracts obtained from King's B, LB, and TYEP media inoculated with *P. fluorescens*. These data confirm that these strains of *P. fluorescens* do not produce AHLs, contrasting with studies that found other strains of *P. fluorescens* as AHL producers (Shaw *et al.*, 1997; Cha *et al.*, 1998; Laue *et al.*, 2000; El-Sayed *et al.*, 2001; Mcphee, 2001; Khan *et al.*, 2005).

Mcphee (2001) found that synthesis of AHLs by *Pseudomonas* was influenced by the composition of the growth medium and environmental factors. According to Mcphee (2001), *P. fluorescens* was found to up-regulate enzyme synthesis when it grew in a spent culture supernatant, presumably having already high levels of synthesized AHL.

Phenotypic characteristics of wild type and transconjugant *P. fluorescens* strains

Biofilm

After 48 h of incubation, it was observed that *P. fluorescens* 07A and 041 produced less biofilm in LB and MMS than in ABC medium. The strain 041 was able to bind better than 07A in polystyrene microtiter dishes (Figure 1). Viana (2006) also observed that different strains of *P. fluorescens* isolated from raw milk had different abilities to bind to polystyrene and that minimal medium enhanced attachment. The ABC minimal medium is rich in divalent ions such as Ca^{+2} , Mg^{+2} , and Fe^{+2} and that could explain this phenotype. According to Fletcher *et al.* (1988), divalent

ions as Ca^{+2} and Mg^{+2} can directly influence biofilm formation due to electrostatic interactions, and indirectly as enzyme cofactors that influence the adhesion dependent on the microorganism physiology. Additionally, the presence of ions such as Ca^{+2} improve cross-binding between cells and between cells and surfaces (Koerstgens *et al.*, 2001).

No significant difference ($p > 0.01$) was found when wild type and transconjugant strains were compared for their ability to produce biofilm (Figure 1). This result shows that the quorum quenching mechanism provided by the lactonase enzyme AiiA, which cleaves AHL molecules (Dong *et al.*, 2000), did not influence biofilm formation in these particular strains of *P. fluorescens*, presumably because they are unable to produce detectable levels of AHLs (Table 2). Allison *et al.* (1998) suggested that QS was involved in promoting cell attachment and biofilm formation in *P. fluorescens* B52, but short chain AHLs were not involved. A mutant of *P. fluorescens* incapable to produce QS signals was significantly defective in biofilm formation (Wei and Zhang, 2006).

Swarming motility

To test swarming motility, the strains 07A and 041 were point inoculated into medium containing 0.4% agar. Only *P. fluorescens* 07A was capable of swarming (Figure 2). When AiiA was expressed in *P. fluorescens* 07A, swarming motility was reduced. Since no other evaluated phenotype had been influenced by the presence of the plasmid expressing AiiA in these strains, we believe that this assay was somehow compromised due to the sensitivity of the *P. fluorescens* 07A transconjugant strain to the presence of gentamicin on the medium where the strain was previously grown. Additionally, an unknown factor required for swarming motility may have been compromised in this strain due to unexpected reasons.

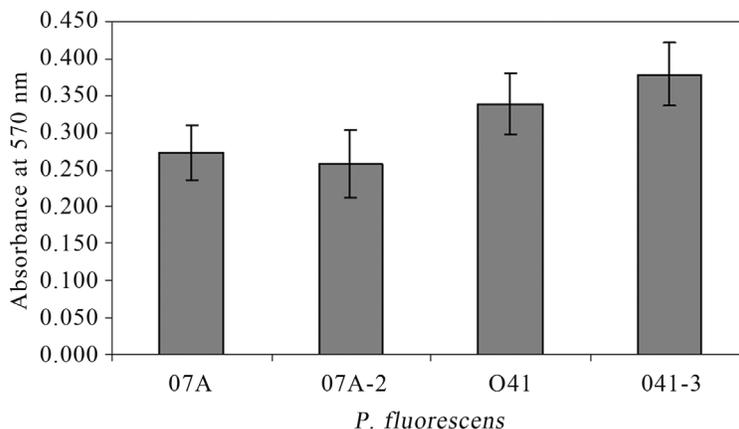


Figure 1 - Biofilm formation by *P. fluorescens* wild type (07A and 041) and transconjugant (07A-2 and 041-3) strains in ABC medium after incubation for 48 h at 25 °C, in polystyrene microtiter plates using crystal violet staining, and quantified based on the difference between the absorbance at 570 nm. Values are means ± standard errors (n = 5).

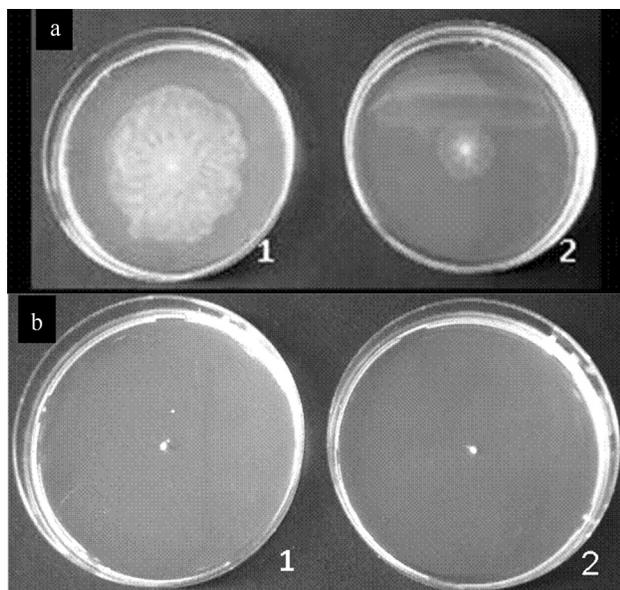


Figure 2 - Ability to form a swarming colony on ABC medium after incubation for 18 h at 25 °C. (a) 1, *P. fluorescens* 07A wild type; 2, *P. fluorescens* 07A-2 transconjugant. (b) 1, *P. fluorescens* 041 wild type; 2, *P. fluorescens* 041-3 transconjugant.

Extracellular protease

AHL-dependent QS systems control the production of extracellular proteolytic activity in many Gram-negative bacteria (Whitehead *et al.*, 2001; Kastbjerg *et al.*, 2007). However, quorum quenching provided by AiiA did not influence proteolytic activity in *P. fluorescens* 07A and 041 when they were grown in different broth media (Figure 3), indicating that the AHL-dependent regulation of this phenotype is not conserved in *P. fluorescens* strains. Pinto *et al.* (2010) also observed that synthetic AHLs or bacterial cell extracts obtained from *P. fluorescens* 07A added to the

medium did not influence growth or proteolytic activity suggesting that QS does not regulate protease production in that strain. The present study further confirms the results with strain 07A, using a totally different approach and expands the knowledge to another importantly spoilage strain (Martins *et al.*, 2005; Pinto *et al.*, 2006). Given that *P. fluorescens* are quite diverse, it is not surprising that some strains produce AHLs (Shaw *et al.*, 1997; Cha *et al.*, 1998; Laue *et al.*, 2000; El-Sayed *et al.*, 2001; Cui *et al.*, 2005; Khan *et al.*, 2005; Wei and Zhang, 2006; Liu *et al.*, 2007) and some others do not (Allison *et al.*, 1998; Cha *et al.*, 1998; Bruijn and Raaijmakers, 2009; Pinto *et al.*, 2010).

Our findings further highlight the diversity and complexity of *P. fluorescens* isolates and reiterate the importance of studies of this kind to improve our knowledge about this group of microorganisms. It would be interesting to analyze a wide collection of *P. fluorescens* strains isolated from raw milk in order to determine if AHL production is a common trait or an exception in strains predominating in this particular environment.

Conclusions

The production of AHLs by *P. fluorescens* 07A and 041 was not detected under any of the conditions used in this study. Both strains produce less biofilm in LB and MMS than in ABC minimal medium, and the strain 041 is better able to bind to polystyrene microtiter dishes than 07A. It was also verified that biofilm formation, swarming motility and proteolytic activity of *P. fluorescens* strains isolated from refrigerated raw milk are not regulated by AHLs. It is likely that the AHL-dependent quorum sensing system is absent from these strains.

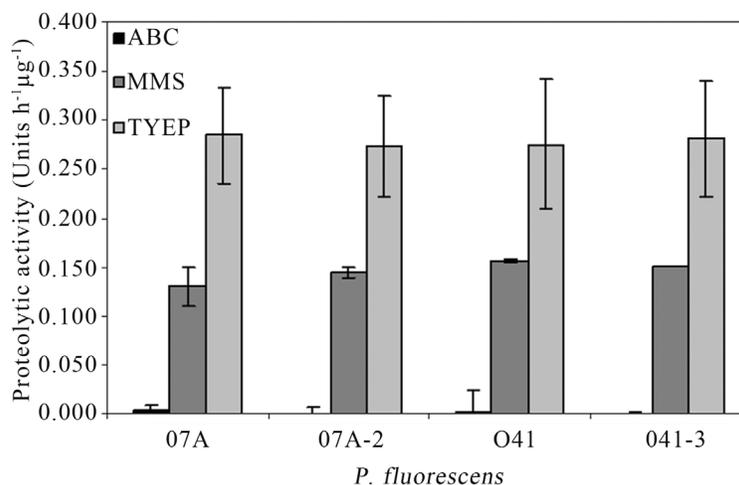


Figure 3 - Proteolytic activity on supernatant of ABC minimal medium, MMS minimal medium and TYEP medium inoculated with *P. fluorescens* wild type (07A and 041) and transconjugants (07A-2 and 041-3) after 24 h of incubation at 25 °C. Values are means ± standard errors (n = 5).

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