# Bacterial Quorum Sensing Molecule N-3-Oxo-Dodecanoyl-L-Homoserine Lactone Causes Direct Cytotoxicity and Reduced Cell Motility in Human Pancreatic Carcinoma Cells



## Ashwath S. Kumar<sup>1</sup>, Jeffrey N. Bryan<sup>1</sup>, Senthil R. Kumar<sup>1,2\*</sup>

1 Comparative Oncology and Epigenetics Laboratory, Veterinary Medicine and Surgery, University of Missouri, Columbia, Missouri, United States of America, 2 Harry S. Truman Veterans Hospital, Columbia, Missouri, United States of America

#### Abstract

In spite of chemotherapeutic and surgical advances, pancreatic cancer continues to have a dismal prognosis. Metastasis due to tumor cell migration remains the most critical challenge in treating pancreatic cancer, and conventional chemotherapy is rarely curative. In the quest for more novel molecules to fight this disease, we tested the hypothesis that the *Pseudomonas aeruginosa* quorum sensing signal molecule N-3-oxo-dodecanoyl-L-homoserine lactone (O-DDHSL) would be cytotoxic to and reduce mobility of pancreatic carcinoma cells (Panc-1 and Aspc-1). Results showed a decrease in cell viability from apoptosis, diminished colony formation, and inhibition of migration of the evaluated pancreatic carcinoma cell lines. Also, cell viability decreased in the presence of O-DDHSL when cells were grown in matrigel basement membrane matrix. While messenger RNA for *IQGAP-1* decreased in Panc-1 and HPDE cells upon exposure to O-DDHSL, no change was observed in Aspc-1 cells. Cofilin mRNA expression was found to be increased in both HPDE and Panc-1 cells with marginal decrease in Aspc-1 cells. *RhoC*, a Rho-family GTPase involved in cell motility, increased in the presence of O-DDHSL could be an effective biomolecule in eukaryotic systems with multimodal function for essential molecular targeting in pancreatic cancer.

Citation: Kumar AS, Bryan JN, Kumar SR (2014) Bacterial Quorum Sensing Molecule N-3-Oxo-Dodecanoyl-L-Homoserine Lactone Causes Direct Cytotoxicity and Reduced Cell Motility in Human Pancreatic Carcinoma Cells. PLoS ONE 9(9): e106480. doi:10.1371/journal.pone.0106480

Editor: Cheryl A. Nickerson, The Biodesign Institute, Arizona State University, United States of America

Received February 28, 2014; Accepted August 1, 2014; Published September 4, 2014

**Copyright:** © 2014 Kumar et al. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Funding: The authors have no funding or support to report.

Competing Interests: The authors have declared that no competing interests exist.

\* Email: kumars@missouri.edu

## Introduction

Pancreatic cancer is the fourth leading cause of cancer related mortality in the United States. According to 2013 statistics from the American Cancer Society, approximately 45,220 people (22,740 men and 22,480 women) were expected to be diagnosed with pancreatic cancer during that year. Patients are most commonly diagnosed with pancreatic cancer in an advanced stage and the disease is generally refractory to chemotherapy [1]. The most significant phenotypic change during the progression of cancer is the transition from a locally growing tumor to a metastatic phenotype. Pancreatic cancers harbor epigenetic alterations affecting phenotype, and their prevalence of these changes increases with higher grade carcinomas [2]. In spite of the discovery of gemcitabine, administered either alone or in combination with other drugs, the survival rate of 5% has remained unchanged for several decades now [3]. These facts indicate that novel compounds are required for treatment to combat this disease.

Quorum sensing (QS) is a process which enables bacterial population to communicate by means of secreted signaling molecules called autoinducers [4]. Human pathogenic bacteria such as *Pseudomonas aeruginosa* (*P. aeruginosa*), a Gram negative bacterium, make a range of acylhomoserine lactones (HSL) with N-acyl side chains from C4-C12 in length [5]. The *P. aeruginosa* bacteria produces a long chain HSL, N-3-oxo-dodecanoyl-L-homoserine lactone (O-DDHSL), and a short chain HSL, N-butyryl-L-homoserine (B-HSL) lactone, both of which influence the expression of virulence factors, swarming motility, and biofilm development [6]. The longer acyl side chain (eg: C12)-HSL molecules are more stable than their shorter chain counterparts (eg: C4)-HSL [7]. The shorter chain HSL can move in and out of cell membranes via free diffusion, while the longer acyl chain HSL is concentrated within the cell, possibly due to partitioning into bacterial membranes [8].

In a process called inter-kingdom signaling, bacterial QS molecules may modulate or influence the behavior of eukaryotic cells [9]. The lipophilic O-DDHSL molecule with an intact homoserine lactone ring interacts directly with phospholipids in model membrane systems and in Jurkat T-cell membranes [10]. The O-DDHSL molecule, upon entering mammalian cells [11,12], may activate nuclear peroxisome proliferator-activated receptors (PPAR) to influence transcriptional activity and NF- $\kappa$ B signaling [13]. It also appears that O-DDHSL can inhibit mammalian cell proliferation and cause cell death in certain cell

types, including cystic-fibrosis-airway epithelial cells [14], breast carcinoma cells [15], T-cells [16] and fibroblasts [17].

Based on existing reports that bacterial QS signals can modulate human cell behavior, we questioned whether O-DDHSL could affect pancreatic carcinoma cell phenotype and characteristics. The rationale for our studies is that pancreatic cancer patients have comparatively low survival rates and remain unresponsive to standard therapies; hence the quest for novel agents to treat pancreatic cancer is necessary. The mechanism of action of O-DDHSL in pancreatic carcinoma cells has yet to be tested. The elucidation of the mechanism of action of O-DDHSL could lead to the development of more effective analogs and novel therapeutic targets, leading to better therapeutic outcomes for pancreatic cancer patients.

The primary objective of our studies is to analyze the migration, viability and colony forming ability of pancreatic carcinoma cells *in vitro* and the effect of alteration of genes involved in these processes following O-DDHSL treatment. The central hypothesis is that O-DDHSL can modulate the genes primarily involved in pancreatic cell migration and proliferation, which include a small GTPase *Rho C* (ras homolog family member C), *cofilin* and *IQGAP-1* (IQ motif containing GTPase activating protein 1). It is expected that O-DDHSL will have multiple antitumor effects on pancreatic carcinoma cells.

## **Materials and Methods**

#### Materials

The pancreatic carcinoma cells Panc-1 and Aspc-1 were purchased from American Type Culture Collection (ATCC (CRL-1469 & CRL-1682)). Normal human pancreatic ductal epithelial cells HPDE6-C7 (HPDE) was kindly provided by Dr. Ming-Sound-Tsao, University of Toronto, Toronto, Canada (18). O-DDHSL and N-dodecanoyl-L-homoserine lactone-3-hydrazone-fluorescein (N-DD-HSL-3-HF) (Fig. 1A & B) were procured from Cayman chemicals, Ann Arbor, MI. N-(3-oxohexanoyl)-Lhomoserine lactone (O-HHSL) (Fig. 1C) was purchased from Sigma Chemical Company, St Louis, MO. Antibodies for *RhoC*, *cofilin*, *IQGAP-1* and  $\beta$ -*actin*, including an anti-rabbit secondary antibody, were purchased from Cell Signaling Technology (Danvers, MA). All other materials were purchased from Fisher Scientific unless mentioned otherwise.

#### Cell culture

Pancreatic carcinoma cells (Panc-1 and Aspc-1) were maintained in custom Roswell Park Memorial Institute (RPMI) medium containing 10% fetal bovine serum (FBS), gentamicin, L-glutamine, and pyruvate sodium. Both lines were grown at 37°C with 5% CO2. HPDE cells were grown in keratinocyte serum free media containing growth supplements at 37°C with 5% CO2.

#### Cell viability assay

Cell viability assays with and without HSL molecules were performed using a WST-1 assay (Roche Applied Science) according to the manufacturer's instructions. The HSL molecules were dissolved in 100% DMSO and further diluted in the cell growth media as required. The Panc-1, Aspc-1 and HPDE cells were seeded in 96-well plates at a density of  $6 \times 10^3$  cells per well and treated with O-DDHSL or O-HHSL (25–300  $\mu M)$  or DMSO vehicle. The cells were incubated for 24 or 48 hours at  $37^{\circ}C$  and 5% CO<sub>2</sub>. WST-1 reagent (8  $\mu$ l), a stable tetrazolium salt that is reduced by the production of NADPH on the surface of viable cells (Roche Applied Science, Indianapolis, IN), was added directly to the wells and incubated at 37°C for approximately one hour for color development. The formazan dve was quantitated by measuring the absorbance at 450 nm in a plate reader; measured absorbance directly correlated to the number of viable cells. The results are expressed as percent viable cells in both untreated and HSL treated samples. Each experiment was performed in triplicate.

#### Apoptosis assay

в

HPDE, Panc-1 and Aspc-1 cells were seeded in a 96-well plate at a density of  $6 \times 10^3$  cells per well and treated with O-DDHSL (300 µM and 200 µM), respectively; plates were incubated for 24 hours at 37°C and 5% CO2. Apoptosis was analyzed using a highthroughput fluorescence-based assay, using quenched (z-Asp-Glu-Val-Asp)<sub>2</sub>-rhodamine 110 for detection of caspase 3/7 activity (Cell Technology Inc, Mountain View, CA). Following media replacement, 50 µl of apoptosis reagent for color development was



Figure 1. Chemical structures of homoserinelactones. doi:10.1371/journal.pone.0106480.g001



N-dodecanoyl-L-homoserine lactone-3-hydrazone-fluorescein

added and cells were incubated for one hour according to the manufacturer's instructions. The plate was then read at an excitation/emission wavelength of 480/520 nm. Increase in fluorescence intensity compared to untreated cells (with DMSO only), corrected for cell number indicated cell apoptosis.

The phenotypic changes of HPDE, Panc-1 and Aspc-1 upon addition of O-DDHSL (300  $\mu$ M and 200  $\mu$ M) were analyzed 48 h after treatment. The plates were photographed in an Olympus CK40 microscope. The pictures were pseudocolored light brown to better visualize the cells.

Cell proliferation was performed by BRDU colorimetric assay according to the manufacturer's protocol (Roche Diagnostics, Indianapolis, IN). Briefly, (~10<sup>5</sup> cells/well) were added to 96-well plates and grown in the presence or absence of O-DDHSL (75 or 150  $\mu$ M) for 48 h at 37°C. Subsequently, BRDU was added to the wells and further incubated for 2 h. After removing the culture medium the cells were fixed and denatured. Finally, anti-BRDU antibody was added to detect the BRDU incorporated in the cells. The reaction product was quantified by measuring the absorbance at 450 nm.

## Effect of O-DDHSL on cells grown in 3D culture

The 3D culture of Panc-1, Aspc-1, and HPDE cells was performed according to a model described earlier [19]. Briefly, the cells were trypsinized from a monolayer to single cell suspension and pelleted by centrifugation. Prechilled chamber slides were coated with a thin layer of Growth Factor Reduced BD Matrigel matrix (BD Biosciences, Sane Jose, and CA) maintained at 4°C. The cells were suspended in serum free (DMEM/F12) media and added to matrigel (4:1 matrix to media) on ice. From this mixture a total of 60  $\mu$ l (10<sup>4</sup> cells/well) was added onto the matrigelprecoated chamber slides. After two weeks, O-DDHSL (200 µM) was added to cells in matrigel, and further incubated for 48 h at 37°C. Following incubation, Calcein AM dye (Ex/Em = 494/ 520 nm) (Biotium, Hayward, CA) was added to the cells and further incubated at 37°C for 2 h to check cell viability and membrane integrity. The cells were photographed using an Olympus CK40 fluorescent microscope.

#### Colony Forming Assay

Cell survival was analyzed using a colony forming assay [20]. Briefly, the cells were exposed to 150  $\mu$ M of O-DDHSL for 48 hours in standard cultures. After the drug treatment, the cells were washed with PBS, lifted from the culture plates and viability was tested with Trypan Blue. Live cell count was maintained the same in treated and untreated cells and transferred to 35 mm-culture dishes at a density of  $2 \times 10^4$  cells. After seven to ten days incubation at  $37^{\circ}$ C the cells were fixed with 70% methanol and stained with 0.5% crystal violet. The colonies were counted using Open CFU software [21]. Colonies, defined as groups of  $\geq 25$  cells, were identified and classified as a single colony. Clonogenic survival was expressed as a percentage relative to untreated controls and calculated as colonies per cell plated. Plating efficiency of untreated cells, based on the number of cells to number of colonies formed, was 72%.

#### In vitro migration assay

Cell migration ability was assessed using a wound healing assay [22]. Panc-1, or Aspc-1  $(2 \times 10^4)$  cells per well were seeded in 6-well plates and allowed to form a complete monolayer. The cells were treated with mitomycin-C for 2 h to block proliferation. Subsequently, a similar sized scratch was made with a sterile 200 µl pipette tip across the center of each well and immediately imaged at baseline, and 48 h, respectively, before and after

treatment with O-DDHSL 150  $\mu$ M (Panc-1 and HPDE) and 75  $\mu$ M (Aspc-1), respectively. The image was acquired using an Olympus CK40 phase contrast microscope. The measurement of the wound gap area was performed using Image J (NIH, USA) software. An arbitrary number of one was assigned to the wound area at 0 h. The values for 48 h are relative to baseline value. Three independent experiments were performed on separate days using different cell passages.

In order to detect O-DDHSL in cells, a fluorescent analog (N-Dd-HSL-3-HF, 10  $\mu M$ ) was added to the live cells grown in chamber slides (40–50% confluent) and treated for about 60 min at 37°C. Subsequently, the cells were fixed with paraformaldehyde. After washing with phosphate buffered saline (PBS), the slides were photographed using a confocal laser scanning microscopy.

## cDNA synthesis and qRT-PCR analysis

Total RNA was isolated from HDPE, Panc-1 and Aspc-1 cells treated with or without O-DDHSL (150 or 75  $\mu$ M for 48 h) using an RNeasy kit (Qiagen, Valencia, CA). First strand.

complimentary DNA (cDNA), obtained from the total RNA, was generated using iScript Reverse Transcription supermix (Bio-Rad, Hercules, CA). Resultant cDNA served as templates for PCR amplification with primers for RhoC (F-GGTCACACACCAG-CACTTTA, R-TTGGAGCCTGTAGCCTTTATTC), cofilin (F-GAGGTGAAGCGCAAGAA, R- GGTTGCATCATA-GAGGGCATAG) IQGAP-1 (F-CCACATCCAAGACAGG-CAATA, R-GGCATCCTCTGTGCTACTAAAG) and  $\beta$ -actin (F- GAAGTCCCTTGCCATCCTAAA, R- GTCTCAAGT-CAGTGTACAGGTAAG). Quantitative real-time PCR (qRT-PCR) was performed with CFX Connect (Bio-Rad, Hercules, CA) using iTaq Universal SYBR Green super mix. The quantification cycle values (Cq) method ( $\Delta\Delta$ Ct) was used for the relative quantification of gene expression. Relative gene expression was calculated in the cells before and after treatment with O-DDHSL. Statistical significance was calculated using one way Analysis of Variance (ANOVA). Differences in gene expression between O-DDHSL treated and untreated samples were considered significant when  $P \leq 0.05$ .

#### Western Blot analysis

The cell lysates from HPDE, Panc-1 and Aspc-1 was treated with M-PER mammalian protein extraction reagent (Thermo-Fisher, Rockford, IL) as per manufacturer's instruction. Protein concentration was determined using the Bicinchoninic acid (BCA) method (Thermo-Fisher, Rockford, IL). Equal amounts of protein  $(\sim 40 \ \mu g)$  were separated on a 10% SDS-polyacrylamide gel and transferred to a nitrocellulose membrane (Bio-Rad, Hercules, CA). The blots were blocked at room temperature for 1 h using Trissaline buffer (TBS) containing 0.1% Tween 20 and 10% nonfat milk. The membrane was further incubated with primary antibodies for cofilin, RhoC, IQGAP-1 or β-actin overnight in a cold chamber at 4°C in separate experiments. After being washed with TBS, the membrane was incubated with a horseradish peroxidase-labeled secondary antibody and visualized with Luminate Forte Western HRP substrate (Millipore, Billerica, MA,). The blot was imaged in a Kodak imaging station (Carestream Health, Rochester, NY).

#### Statistical analysis

The data was analyzed using an unpaired two-sided Student ttest. Differences between the control and drug treated samples were considered significant when  $P \le 0.05$ . Evaluation for normality was performed using the Shapiro-Wilk test.



**Figure 2. Effect of HSL on cell viability. A** - Panc-1 ( $6 \times 10^3$  cells per well) was treated with different concentrations of O-DDHSL for 24 and 48 h. Significant decrease in cell viability was observed with 200–300  $\mu$ M O-DDHSL (P $\leq$ 0.05) after 24 h. At 48 h, cell viability decrease was significant at O-DDHSL concentration at or above 100  $\mu$ M (P $\leq$ 0.02; n = 3). However, HPDE cell viability was not affected after 48 h except for a slight decrease at 300  $\mu$ M O-DDHSL. No effect was observed with O-HHSL. **B** – Aspc-1 ( $6 \times 10^3$  cells per well) was more sensitive to O-DDHSL exposure than Panc-1. A significant decrease (P $\leq$ 0.02) was observed in viability  $\geq$ 25  $\mu$ M O-DDHSL after 24 or 48 h (n = 3). Cell viability was not affected by O-HHSL. In both cases, DMSO (0.02%) was used as diluent control which did not affect the cell viability *per se.* **C**–**E** Panc-1, Aspc-1 and HPDE cells were plated on a layer of matrigel in chamber slides in serum free DMEM/F12 media and allowed to grow for two weeks. O-DDHSL (200  $\mu$ M) was added to the cells and incubated for 48 h at 37°C. Subsequently, a fluorescent dye Calcein AM was added and further incubated for 2 h for its uptake by the cells. C–E – Light microsopy pictures of cells growing on matrigel before and after addition of O-DDHSL showing morphological changes of apoptosis (top panel). Bottom panel shows the uptake of Calcein AM dye in the cells before and after treatment with O-DDHSL, showing dye uptake by viable cells and los: of cell viability resulting in the absence of dye uptake (40 ×). doi:10.1371/journal.pone.0106480.g002

## Results

#### Cell Viability

The cell viability of Panc-1 and Aspc-1 cells was tested in the absence and presence of HSL in a time-dependent manner. The decrease in cell viability of both cell lines was concentration and time-dependent (Fig. 2 A&B). O-DDHSL inhibited (P $\leq$ 0.02) Panc-1 viability at a concentration  $\geq$ 100 µM when incubated for 48 h (IC<sub>50</sub> = 120±5 µM) (n = 3). However, O-DDHSL had only minimal effect on HPDE cell viability. When parallel studies were conducted with O-HHSL a shorter chain HSL, no decrease in viability of Panc-1 cells was observed even at higher concentrations (300 µM). Aspc-1 cells, displayed more sensitivity (P $\leq$ 0.02) to O-DDHSL in low concentrations at both 24 and 48 h (IC<sub>50</sub> = 48±8 µM) (n = 3) while they also were not affected by O-HHSL. DMSO (0.02%), which was used as diluent control, did not affect the cell viability by itself.

The viability of cells grown in matrigel containing extracellular matrix components was tested in the presence of O-DDHSL. Panc-1, Aspc-1 and HPDE cells grew in matrigel as spheroids and retained the Calcein AM dye indicating viability (Figs. 2C–E).

Calcein AM dye is retained by the cells if only their membranes are intact. However, the O-DDHSL treated Panc-1 and Aspc-1 cells were not able to retain the dye due to lack of viable cells (Fig. 2C&D), indicating that O-DDHSL was capable of decreasing the viability of the cells grown in an extracellular matrix environment. HPDE cell viability at the tested O-DDHSL concentration appears to be less affected compared to carcinoma cells (Fig. 2E).

# O-DDHSL induces apoptosis of pancreatic carcinoma cells

Of the mechanisms for decreased survival of pancreatic carcinoma cells exposed to O-DDHSL, apoptosis was considered most likely. Caspase 3/7 activity was detected in HPDE (P = 0.059), Panc-1(P = 0.047) and Aspc-1 cells ( $P \le 0.02$ ) following exposure to O-DDHSL (300 and 200  $\mu$ M), after a 24 h time interval (Fig. 3A). The phenotypic changes are shown in Figure 3B. The cells become less viable and failed to proliferate. Altogether, the above studies indicate the O-DDHSL induces apoptosis in pancreatic carcinoma cells which results in decreased proliferation. Apoptosis was also observed in HPDE cells but to a



**Figure 3. Cell apoptosis in the presence of O-DDHSL. A** - Panc-1, Aspc-1 and HPDE-1 cells ( $6 \times 10^3$  cells per well) in triplicates was treated with 300 (Panc-1 and HPDE) or 200  $\mu$ M (Aspc-1) of O-DDHSL, respectively, for 24 h. Cell apoptosis was assayed using a fluorescent based caspase 3/7 activity detection kit. While apoptosis in both Panc-1 cells ( $P = 0.047^*$ ) and Aspc-1 cells ( $P = 0.02^{**}$ ) were significant, HPDE cell apoptosis was marginally significant ( $P = 0.059^{***}$ ) **B** – Light micrographs of Panc-1, Aspc-1 and HPDE cells before and after exposure to O-DDHSL at concentrations of 300  $\mu$ M (Panc-1 and HPDE) and 200  $\mu$ M (Aspc-1), respectively, for 48 h ( $20 \times$ ). **C** – Cell proliferation of HPDE, Panc-1 and Aspc-1cells upon exposure to O-DDHSL decreased by 35% ( $P = 0.042^*$ ), 47% ( $P \le 0.035^{***}$ ), and 52% ( $P \le 0.025^{***}$ ), respectively.

lesser extent compared to carcinoma cells. Cell proliferation was independently analyzed by BRDU assay. While the proliferation was reduced by 47% (Panc-1, (P $\leq$ 0.035) and 52% (Aspc-1, P $\leq$ 0.025) in pancreatic carcinoma cells only about 35% decrease (P=0.042) in proliferation was observed in HPDE cells (Fig. 3C).

## O-DDHSL treatment reduces colony forming ability and inhibits cell migration

The concentration of O-DDHSL was maintained slightly higher than respective IC<sub>50</sub> concentration to minimize apoptosis in the following studies. In Panc-1 and HPDE cells treated with O-DDHSL (150  $\mu$ M) the colony formation was reduced by 85% and 40%, respectively, compared to untreated cells (P≤0.05) (Fig. 4A &B). In Aspc-1 cells (O-DDHSL, 75  $\mu$ M) the colony formation decreased 70% compared to untreated cells.

For cell migration studies, one set of plates for each Panc-1 Aspc-1 and HPDE cells was treated with O-DDHSL (150 and 75  $\mu$ M) and monitored for wound closure after 48 h. The Panc-1

cells were photographed at 0 h, and 48 h (Fig. 5A). The wound area was measured at each time point to determine the migration rate which is depicted in Figure 5B. In untreated cells, the wound area closed completely (100%) by 48 h, while in O-DDHSL treated cells the capacity of cells to migrate and close the wound gap was only 25% (Panc-1) and 10% (Aspc-1) relative to untreated cells (Fig. 5A & B). Interestingly, in the HPDE cells though some cells migrate into the wound no complete wound closure was observed. A similar result was observed in O-DDHSL treated cells. Approximately, in both cases only 15–20% wound closure was observed after 48 h.

In order to visualize the presence of O-DDHSL in the cells, a fluorescent analog, N-Dd-HSL-3-HF, 10  $\mu$ M was added to the live cells grown in chamber slides. The results demonstrate uptake of fluorescent compound by the Panc-1, Aspc-1 and HPDE cells (Fig. 5C). When treated with O-HHSL, neither migration nor clonogenic ability of the cells was affected (data not shown).



**Figure 4. Colony formation upon treatment with O-DDHSL. A** - Colony forming assay was performed by plating  $2 \times 10^4$  cells per plate with or without treatment of Panc-1, Aspc-1 and HPDE cells with O-DDHSL (150  $\mu$ M) for 48 h. After seven to ten days incubation at 37°C the cells were fixed with 70% methanol and stained with 0.5% crystal violet. Cell colonies were counted using Open CFU software [21]. Colonies, defined as groups of  $\geq$  25 cells, were identified and classified as a single colony. **B** – Relative colony formation depicted as a percentage between O-DDHSL untreated and treated cells, Panc-1(P $\leq$ 0.029\*), Aspc-1(P $\leq$ 0.03\*\*) and HPDE (P $\leq$ 0.05\*\*\*). doi:10.1371/journal.pone.0106480.g004

#### Analysis of gene expression by qRT-PCR

As metastasis is a hallmark of pancreatic cancer, we tested whether addition of O-DDHSL would alter the gene expression associated with cell migration. RhoC, cofilin and IQGAP-1 genes are associated with cell migration and maintaining cell shape by interacting with the actin-based cytoskeleton. qRT-PCR was performed using specific gene primers for all the above mentioned genes (Fig. 6). Basal expression of these genes was found in HPDE cells. In HPDE and Panc-1 cells, increase in cofilin was two (P = 0.037) and one (P = 0.04) fold upon treatment with O-DDHSL. Similarly, the RhoC mRNA levels increased in Panc-1 by two fold (P = 0.032) and four fold in HPDE cells (P = 0.029), respectively. On the other hand, IQGAP-1 mRNA appeared to decrease one fold in Panc-1 cells and five fold (P = 0.024) in HPDE cells. In case of Aspc1 cells, while a decrease in cofilin (< one fold) and increased (two fold, P=0.036) RhoC mRNA expression was observed no change in IQGAP-1 mRNA level upon O-DDHSL treatment between untreated and treated cells was noted.

## Western blot analysis

The protein expression of *RhoC*, *cofilin* and *IQGAP-1* in Panc-1, Aspc-1 and HPDE cells was analyzed by Western blot following exposure to O-DDHSL (150  $\mu$ M or 75  $\mu$ M for 48 h) (Fig. 7). *β*-*actin* was used as a loading control. No qualitative changes in *RhoC* protein level was noted following O-DDHSL exposure in Panc-1 and Aspc-1 cells. In HPDE cells, RhoC protein expression increased compared to the wild type cells. No qualitative change in

*IQGAP-1* protein expression was noted in Panc-1 and Aspc-1 cells upon O-DDHSL treatment but a qualitative decrease was observed in HPDE cells. Cofilin was moderately decreased in Aspc1- cells while a slight increase was observed in Panc-1 cells. No changes was observed with HPDE cell *cofilin* expression after treatment with O-DDHSL.

## Discussion

Bacteria communicate with eukaryotic hosts in both pathogenic and symbiotic relationships by chemical means using QS molecules. Eukaryotes can detect and react to such bacterial signals. We explored the possibility that human pancreatic tumor cells could respond to QS molecules of *P. aeruginosa* bacteria, anticipating that such signaling molecules could modulate the growth and migration characteristics as well as gene expression of the cells. We observed that the long acyl chain QS molecule O-DDHSL was effective in modulating the pancreatic carcinoma cell properties, while O-HSSL with a short chain acyl group was not as effective.

Previous studies have reported that O-DDHSL was able to decrease the cell proliferation in breast carcinoma cells and murine T-cells [15,16]. Our studies indicate that O-DDHSL decreased the viability of pancreatic carcinoma cells in a concentration and time-dependent manner. Cells grown in 3D cultures provide a more physiologically relevant approach to the analysis of cell phenotypes and the influence of chemotherapy drugs (19). In the present study, we observed that O-DDHSL effectively decreased



**Figure 5. Cell migration by wound healing assay. A** - Panc-1, Aspc-1 and HPDE cells were plated and allowed to grow as a monolayer after which a similar sized scratch was made with a sterile tip in the culture plates. The plates were photographed at baseline (0 h). A set of plates were treated with O-DDHSL (150  $\mu$ M) and the gap closure was photographed after 48 h. In the presence of O-DDHSL, the migration of cells was considerably less, which resulted in non-closure of the wound gap compared to untreated cells (n = 3). Red arrows - Wound gap (20×). **B** - The wound area in the image was measured using Image J software and the baseline gap area was assigned an arbitrary number of 1. The untreated carcinoma cell wound closure values at 48 h (P = 0.027\*) was significant. The O-DDHSL treated cell wound closure value at 48 h for both Pan-1 and Aspc-1 was not significant (P≥0.05). In HPDE cells, the wound closure between treated and untreated cells was almost similar. **C** – Detection of a fluorescent analog of O-DDHSL, N-Dd-HSL-3-HF, 10  $\mu$ M, in different cells after 30 min at 37°C (40×).

the viability of carcinoma cells when grown in 3D cultures relative to normal pancreatic epithelial cells. Another, shorter acyl side chain, QS molecule O-HHSL, was not effective in decreasing cell viability (Fig. 1B). This could be due the structural instability of this molecule and opening of the lactone rings in the pH and temperature of cell culture conditions [7]. The HPDE cell viability was found to be less affected by O-DDHSL (Fig. 1A). A similar behavior in the presence of O-DDHSL was reported with normal non-tumorigenic breast epithelial cells [15] CCL-185 [23], Hep-2 [23] and Caco-2 epithelial cells [24] but not cystic fibrosis airway epithelial cells [25]. The O-DDHSL concentration used in our study is within reasonable limits because it has been reported that very high levels of QS molecules can accumulate in biofilms grown *in vitro*, resulting in O-DDHSL concentrations up to 300– 600  $\mu$ M [17].

Apoptosis induction by O-DDHSL has been reported in a variety of cells resulting in caspase activation [15,23]. O-DDHSL induced apoptosis in Panc-1, Aspc-1 and HPDE cells, possibly by activating caspases. The induction of apoptosis, however, was dependent on concentration and cell type. While Panc-1 and

HPDE cells required higher concentration of O-DDHSL to trigger apoptosis (300  $\mu$ M), Aspc-1 cells underwent apoptosis with lower concentrations (200  $\mu$ M). It is possible different cell types within the same type of cancer could vary in their sensitivity for O-DDHSL.

An important characteristic of pancreatic adenocarcinoma cells is their migration and invasive properties with nearly half of patients having metastasis at the time of diagnosis. We analyzed cell migration by a wound healing assay. We found that O-DDHSL exposure inhibited the migration of both Panc-1 and Aspc-1 cells, while untreated cells were able to completely close the wound gap. Based on these results, we hypothesized that the genes involved in cell motility could be a target for O-DDHSL. It is possible that migration could be affected by O-DDHSL induced apoptosis but the concentration of the compound added was reduced to near  $IC_{50}$  values to minimize apoptosis. It was difficult to assess the effect of O-DDHSL on HPDE cells as the wound healing ability was found to be similar in untreated and cells treated with O-DDHSL. It is likely that the ability of normal



**Figure 6. qRT-PCR analysis of genes** *Rho C, Cofilin* **and** *IQGAP-1* **in different cells.** In order to analyze the changes in mRNA expression of *RhoC, cofilin* and *IQGAP-1* in Panc-1, Aspc-1 and HPDE cells, upon treatment with O-DDHSL (150  $\mu$ M; 48 h), a qRT-PCR was performed with specific primers.  $\beta$ -actin was used as a housekeeping gene. Basal expression of all the above genes was observed in HPDE cells. *Cofilin* mRNA level increased in Panc-1 (P = 0.04\*) and HPDE cells (P = 0.037\*). Similarly, the RhoC mRNA levels increased in Panc-1 (P = 0.032\*\*) and HPDE cells (P = 0.029\*\*), respectively. The *IQGAP-1* mRNA level decreased in HPDE cells (P = 0.024\*\*\*). In Aspc-1 cells, no change was observed in *IQGAP-1* mRNA, but a doi:10.1371/journal.pone.0106480.q006



**Figure 7. Western blot for protein expression.** Equal amounts of protein (~40  $\mu$ g) was separated on a 10% SDS-polyacrylamide gel and transferred to a nitrocellulose membrane. The membrane was probed with primary antibodies for Rho-C (~42 kDa), cofilin (~18 kDa), *IQGAP-1* (~197 kDA) and,  $\beta$ -actin (~47 kDA) followed by incubation with a HRP-conjugated secondary antibody. The bands were detected using a chemiluminescent reagent. The blot image was acquired using a Kodak imaging software. doi:10.1371/journal.pone.0106480.g007

pancreatic epithelial cells to close the wound gap is rather a slow process compared to carcinoma cells.

To further gain insight into the inhibition of cell migration by O-DDHSL, we focused on three important genes which are essential for cell migration including *cofilin* [26] *IQGAP-1* [27] and the small GTPase *RhoC* [28]. *Cofilin* is an important regulator of actin cytoskeleton [24] and *IQGAP-1* localizes in the leading edge of migrating cells [27,29]. *RhoC* is a small GTPase which is an important effector of tumor cell motility [28] and is expressed in pancreatic tumors [30]. Similarly, *cofilin* is also present in pancreatic carcinoma tissues and possibly promotes its progression [31].

Upon treatment with O-DDHSL, the mRNA message of cofilin increased in HPDE and Panc-1 cells. The endogenous mRNA expression of *RhoC* increased in O-DDHSL treated Panc-1, Aspc-1 and HPDE cells. In HPDE cells, upon O-DDHSL treatment a decrease in IOGAP-1 was noted but the change was only marginal in Panc-1 and Aspc-1 cells. Protein expression studies indicated that in O-DDHSL treated cells no drastic changes were observed in case of IQGAP-1or RhoC in tumor cells except for HPDE cells. Altogether, O-DDHSL differentially modulates the gene expression of cofilin, RhoC and IQGAP-1, all involved in cell migration. IQGAP-1 was reported to be targeted by O-DDHSL in Caco-2 epithelial cells affecting their migration [24]. It has also been reported that more than one form of *cofilin* exist in pancreatic cancer tissues [31]. Cofilin-1 involved in motility and invasion is up-regulated in pancreatic carcinoma tissues while muscle cofilin-2 was reported to be downregulated [31]. At present, it is not clear whether O-DDHSL directly interact with these molecules, however, its effect seem to vary between the cells tested in this study.

Increase in RhoC gene expression is intriguing, because it was reported to promote tumor metastasis in melanoma [28]. While we observed increase in RhoC message, no significant changes in the protein expression was observed except in HPDE cells (Fig. 7). It is unclear whether RhoC is subjected to any posttranscriptional

#### References

- Almhanna K, Kim R (2008) Second-line therapy for gemcitabine-refractory pancreatic cancer: is there a standard? Oncology (Williston Park) 22: 1176–83.
- McCleary-Wheeler AL, Lomberk GA, Weiss FU, Schneider G, Fabbri M, et al. (2013) Insights into the epigenetic mechanisms controlling pancreatic carcinogenesis. Cancer Lett 328: 212–221.
- Fisher WE (2011) The promise of a personalized genomic approach to pancreatic cancer and why targeted therapies have missed the mark. World J Surg 35: 1766–1769.
- Henke JM, Bassler BL (2004) Three parallel quorum-sensing systems regulate gene expression in Vibrio harveyi. J Bacteriol 186: 6902–6914.
- Swift S, Downie JA, Whitehead NA, Barnard AM, Salmond GP, et al. (2001) Quorum sensing as a population-density-dependent determinant of bacterial physiology. Adv Microb Physiol 45: 199–270.
- Williams P, Cámara M (2009) Quorum sensing and environmental adaptation in Pseudomonas aeruginosa: a tale of regulatory networks and multifunctional signal molecules. Curr Opin Microbiol 12: 182–191.
- Yates EA, Philipp B, Buckley C, Atkinson S, Chhabra SR, et al. (2002) Nacylhomoserine lactones undergo lactonolysis in a pH-, temperature-, and acyl chain length-dependent manner during growth of Yersinia pseudotuberculosis and Pseudomonas aeruginosa. Infect Immun 70: 5635–5646.
- Pearson JP, Van Delden C, Iglewski BH (1999) Active efflux and diffusion are involved in transport of Pseudomonas aeruginosa cell-to-cell signals. J Bacteriol 181: 1203–1210.
- Pacheco AR, Sperandio V (2009) Inter-kingdom signaling: chemical language between bacteria and host. Curr Opin Microbiol 12: 192–198.
- Davis BM, Jensen R, Williams P, O'Shea P (2012) The interaction of Nacylhomoserine lactone quorum sensing signaling molecules with biological membranes: implications for inter-kingdom signaling. PLoS One 5: e13522.
- Ritchie AJ, Whittall C, Lazenby JJ, Chhabra SR, Pritchard DI, et al. (2007). The immunomodulatory Pseudomonas aeruginosa signalling molecule N-(3-oxododecanoyl)-L-homoserine lactone enters mammalian cells in an unregulated fashion. Immunol Cell Biol 85: 596–602.

regulation in the carcinoma cells [32,33]. Alternately, we speculate that an increase in *RhoC* gene upon modulation in one or other migration inducible genes such as cofilin or IQGAP-1 could be due to a compensatory pathway which reflects the plasticity of cell signaling networks [34,35]. However, increase in RhoC gene expression did not promote the cell migration in our studies as observed by failure of wound closure after 48 h in the presence of O-DDHSL. Previous studies reported that compensatory responses could be identified by gene expression and phosphoproteomic changes upon single drug treatment [34]. Studies with combinations of lapatinib and Ro31-8220 in KU-7 cells, a multi-kinase inhibitor, demonstrated compensatory cross-talk between p70S6 kinase and EGF receptor pathways, and the node of convergence was identified as phosphorylated BCL2-associated agonist of cell death (BAD) [35]. Further studies are required to understand the mechanism of RhoC increase, as this gene has been implicated as an independent mitogen from RhoA, and can induce metastasis in melanoma cells [28].

In conclusion, the combined findings from our study demonstrate that QS molecule O-DDHSL decreases cell viability, promotes apoptosis by activating caspases, and inhibits the wound healing process. O-DDHSL modulates genes responsible for migration such as *RhoC*, *cofilin* and *IQGAP-1*. However, we observed that O-DDHSL also affect some of the normal epithelial cells properties similar to that of tumor cells, which could be a limiting factor when it comes to the clinical application of this molecule. The potential for O-DDHSL as a possible chemotherapeutic agent for pancreatic cancer needs further *in vivo* evaluation.

#### **Author Contributions**

Conceived and designed the experiments: SRK. Performed the experiments: SRK ASK. Analyzed the data: SRK ASK JNB. Contributed reagents/materials/analysis tools: SRK JNB. Wrote the paper: SRK JNB.

- Shiner EK, Reddy S, Timmons C, Li G, Williams SC, et al. (2004) Construction of a bacterial autoinducer detection system in mammalian cells. Biol Proced Online 6: 268–276.
- Jahoor A, Patel R, Bryan A, Do C, Krier J, et al. (2008) Peroxisome proliferatoractivated receptors mediate host cell proinflammatory responses to Pseudomonas aeruginosa autoinducer. J Bacteriol 190: 4408–4415.
- Schwarzer C, Fu Z, Patanwala M, Hum L, Lopez-Guzman M, et al. (2012) Pseudomonas aeruginosa biofilm-associated homoserine lactone C12 rapidly activates apoptosis in airway epithelia. Cell Microbiol 14: 698–709.
- Telford G, Wheeler D, Williams P, Tomkins PT, Appleby P, et al. (1998) The Pseudomonas aeruginosa quorum-sensing signal molecule N-(3-oxododecanoyl)-L- homoserine lactone has immunomodulatory activity. Infect Immun 66: 36– 42.
- Li L, Hooi D, Chhabra SR, Pritchard D, Shaw PE (2004) Bacterial Nacylhomoserine lactone-induced apoptosis in breast carcinoma cells correlated with down-modulation of STAT3. Oncogene 23: 4894–902.
- Ortori CA, Dubern JF, Chhabra SR, Cámara M, Hardie K, et al. (2011) Simultaneous quantitative profiling of N-acyl-L-homoserine lactone and 2-alkyl-4(1H)- quinolone families of quorum-sensing signaling molecules using LC-MS/ MS. Anal Bioanal Chem 399: 839–850.
- Liu N, Furukawa T, Kobari M, Tsao M-S (1998) Comparative Phenotypic Studies of Duct Epithelial Cell Lines Derived from Normal Human Pancreas and Pancreatic Carcinoma. Am. J of Pathology 153: 263–269.
- Lee GY, Kenny PA, Lee EH, Bissell MJ (2007) Three-dimensional culture models of normal and malignant breast epithelial cells. Nat Methods 4: 359–365.
- Franken NA, Rodermond HM, Stap J, Haveman J, van Bree C (2006) Clonogenic assay of cells in vitro. Nat Protoc 1: 2315–2319.
- Geissmann Q (2013) OpenCFU, a New Free and Open-Source Software to Count Cell Colonies and Other Circular Objects. PLoS ONE 8(2): e54072: 1– 10.
- Schneider L, Cammer M, Lehman J, Nielsen SK, Guerra CF, et al. (2010) Directional cell migration and chemotaxis in wound healing response to PDGF-

AA are coordinated by the primary cilium in fibroblasts. Cell Physiol Biochem 25: 279–292.

- Tateda K, Ishii Y, Horikawa M, Matsumoto T, Miyairi S, et al. (2003) The Pseudomonas aeruginosa autoinducer N-3- oxododecanoyl homoserine lactone accelerates apoptosis in macrophages and neutrophils. Infect Immun 71: 5785– 5793.
- Karlsson T, Turkina MV, Yakymenko O, Magnusson KE, Vikström E (2012) The Pseudomonas aeruginosa N-acylhomoserine lactone quorum sensing molecules target *IQGAP-1* and modulate epithelial cell migration. PLoS Pathog 8: e1002953.
- Schwarzer C, Fu Z, Patanwala M, Hum L, Lopez-Guzman M, et al. (2012) Pseudomonas aeruginosa biofilm-associated homoserine lactone C12 rapidly activates apoptosis in airway epithelia. Cell Microbiol 14: 698–709.
- Delorme V, Machacek M, DerMardirossian C, Anderson KL, Wittmann T, et al.(2007) *Cofilin* activity downstream of Pak1 regulates cell protrusion efficiency by organizing lamellipodium and lamella actin networks. Dev Cell 13: 646–662.
- Noritake J, Fukata M, Sato K, Nakagawa M, Watanabe T, et al. (2004) Positive role of *IQGAP-1*, an effector of Rac1, in actin-meshwork formation at sites of cell-cell contact. Mol Biol Cell 15: 1065–76.
- Clark EA, Golub TR, Lander ES, Hynes RO (2000) Genomic analysis of metastasis reveals an essential role for *RhoC*. Nature 406: 532–535.

- Benseñor LB, Kan HM, Wang N, Wallrabe H, Davidson LA, et al. (2007) IQGAP-1 regulates cell motility by linking growth factor signaling to actin assembly. J Cell Sci 120: 658–669.
- Suwa H, Ohshio G, Imamura T, Watanabe G, Arii S, et al. (1998) Overexpression of the *rhoC* gene correlates with progression of ductal adenocarcinoma of the pancreas. Br J Cancer 77: 147–152.
- Wang Y, Kuramitsu Y, Ueno T, Suzuki N, Yoshino S, et al. (2011) Differential expression of up-regulated *cofilin-1* and down-regulated *cofilin-2* characteristic of pancreatic cancer tissues. Oncol Rep 26: 1595–1599.
- Rogers S1, Girolami M, Kolch W, Waters KM, Liu T, et al. (2008) Investigating the correspondence between transcriptomic and proteomic expression profiles using coupled cluster models. Bioinformatics 24: 2894–2900.
- 33. Vogel C1, Abreu Rde S, Ko D, Le SY, Shapiro BA, et al. (2010) Sequence signatures and mRNA concentration can explain two-thirds of protein abundance variation in a human cell line. Mol Syst Biol 6: 400.
- Gioeli D, Wunderlich W, Sebolt-Leopold J, Bekiranov S, Wulfkuhle JD, et al. (2011) Compensatory pathways induced by MEK inhibition are effective drug targets for combination therapy against castration-resistant prostate cancer. Mol Cancer Ther 10: 1581–1590.
- Axelrod M, Gordon VL, Conaway M, Tarcsafalvi A, Neitzke DJ, et al. (2013) Combinatorial drug screening identifies compensatory pathway interactions and adaptive resistance mechanisms. Oncotarget 4: 622–635.