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N-hexanoyl-L-homoserine lactone-degrading *Pseudomonas aeruginosa* PsDAHP1 protects zebrafish against *Vibrio parahaemolyticus* infection



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ABSTRACT

Four strains of N-hexanoyl-L-homoserine lactone (AHL)-degrading *Pseudomonas* spp., named PsDAHP1, PsDAHP2, PsDAHP3, and PsDAHP4 were isolated and identified from the intestine of *Fenneropenaeus indicus*. PsDAHP1 showed the highest AHL-degrading activity among the four isolates. PsDAHP1 inhibited biofilm-forming exopolysaccharide and altered cell surface hydrophobicity of virulent green fluorescent protein (GFP)-tagged *Vibrio parahaemolyticus* DAHV2 (GFP-VpDAHV2). Oral administration of PsDAHP1 significantly reduced zebrafish mortality caused by GFP-VpDAHV2 challenge, and inhibited colonisation of GFP-VpDAHV2 in the gills and intestine of zebrafish as evidence by confocal laser scanning microscope and selective plating. Furthermore, zebrafish receiving PsDAHP1-containing feed had increased phagocytic cells of its leucocytes, increased serum activities of superoxide dismutase and lysozyme. The results suggest that *Pseudomonas aeruginosa* PsDAHP1 could protect zebrafish from *V. parahaemolyticus* infection by inhibiting biofilm formation and enhancing defence mechanisms of the fish.

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1. Introduction

Diseases caused by biofilm formation of *Vibrio* spp. can be problematic in intensive shrimp farming and lead to economic loss, especially in nursery [1]. Current problem of acute hepatopancreatic necrosis disease caused by *Vibrio parahaemolyticus* results in a severe blow to the shrimp farming industry [2]. Skin haemorrhages, necroses and mortality have also been observed in *V. parahaemolyticus* infected fish [3]. The underlying pathogenesis caused by *Vibrio* infection in shrimp and fish are, however, not well understood. In experimentally-infected rabbits, it was reported that flagellae and pili of *V. parahaemolyticus* were adhered and formed biofilm in the gut epithelium of the animals [4]. The bacteria *V. parahaemolyticus* has been known to undergo reversible phase variation; and its architecture and integrity of strains form vigorous biofilms that are different from those of other *Vibrio*

species, which is due to the amounts of capsular polysaccharide and other cell surface molecules [5]. The polysaccharides excreted from the bacterial cells, or exopolysaccharide, plays an important role in the aggregation of the bacteria during biofilm formation [6]. Interfering with the biofilm formation of *Vibrio* spp. is one way to disrupt quorum sensing and serves as an alternative to the use of antibiotics [7]. Non-antibiotic molecules are naturally produced within bacterial communities; these include signalling molecules or surface active biosurfactants that interfere with biofilm formation [8].

Therefore, disruption of quorum sensing has been suggested as a new strategy to control bacterial infection in aquaculture [9]. For instance, *Pseudomonas* spp. produces bacteriocins, pyocin, and phenazines which are used as bioactive agents and two *Pseudomonas* strains, PAI-A and PAO1, were found to degrade 3-oxododecanoyl homoserine lactone and other long-acyl groups; the bacteria use the substance as an energy source [10].

Probiotics have been widely used in aquaculture to prevent bacterial infections [11], however, the mechanisms by which probiotics could control virulence of the pathogens are not well

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understood. Probiotics have been shown to enhance phagocytic activity, increase production of reactive oxygen species by macrophages, increase immunoglobulin-producing cells and acidophilic granulocytes in fish and shrimp [12]. It is also possible that the action of probiotics is on the disruption of the quorum sensing of bacterial pathogens, the mechanism of which has been under-investigated thus far. In an attempt to isolate viable and potential quorum quenching strains of bacteria, we report mechanisms by which an acylated homoserine lactone (AHL)-degrading *Pseudomonas aeruginosa*, PsDAHP1 strain, protected zebrafish against *V. parahaemolyticus* challenge.

2. Materials & methods

2.1. Culture of AHL-degrading *Pseudomonas* spp. and virulent *V. parahaemolyticus* tagged with green fluorescent protein (GFP)

The bacteria *Pseudomonas* spp. were isolated from intestine of healthy Indian shrimp *Fenneropenaeus indicus* collected from the Cuddalore coast, 250 km south of Chennai south-east coast of India. The bacterial strains were incubated on minimal media agar plates containing (10^6 CFU ml^{-1}) with 5 mg l^{-1} of N-hexanoyl-L-homoserine lactone (C6-HSL) as sole carbon and nitrogen sources for 48 h, at $30\text{ }^\circ\text{C}$ [13]. The bacteria were further cultured and maintained in Zobell marine agar (ZMA) and broth (ZMB) (Himedia Laboratories, India). Three virulent strains of *V. parahaemolyticus*: VpDAHV1 (HQ693275), VpDAHV2 (HQ625651), and VpDAHV3 (HQ693276), isolated from *F. indicus* with black-gill disease were used as challenging bacteria. A non-virulent *V. parahaemolyticus* strain, ATCC 17802, was used as reference. The isolated strains of *V. parahaemolyticus* were cultured in tryptic soy agar and broth at $30\text{ }^\circ\text{C}$ for 24 h. One strain of *V. parahaemolyticus*, VpDAHV2, was tagged with green fluorescent protein (GFP) by conjugal transfer of pVSV102 (Km^R), and GFP expression confirmed by confocal laser scanning microscopy (LSM 710, Carl Zeiss, Germany) [14]. Two strains of *Chromobacterium violaceum* (ATCC 12472 and CV026), being cultured in Luria-Bertani (LB) broth at $33\text{ }^\circ\text{C}$, were used to determine quorum sensing inhibitory activity.

2.2. Effect of cell-free *Pseudomonas* extracts on biofilm formation

Cell-free extracts of four strains of *Pseudomonas* spp. (PsDAHP1, PsDAHP2, PsDAHP3 and PsDAHP4) were tested for their ability to inhibit *V. parahaemolyticus* biofilm formation in microtitre plates [15]. The bacteria, $\sim 10^8$ cells ml^{-1} , were ultrasonicated using a UP100H ultrasonic processor (Hielscher, Germany) at 100 Watts for 1 min. They were centrifuged at $800\times g$ for 10 min, passed through a $0.2\text{ }\mu\text{m}$ pore-sized filter and stored at $-20\text{ }^\circ\text{C}$. The virulent strains of *V. parahaemolyticus* (VpDAHV1, VpDAHV2, VpDAHV3) and the reference one (ATCC 17802), at $\sim 10^6$ cells ml^{-1} , were inoculated into 96-well polystyrene microtitre plates containing $100\text{ }\mu\text{l}$ of Zobell marine broth and $50\text{ }\mu\text{l}$ of the *Pseudomonas* cell-free extracts for 24 h at $30\text{ }^\circ\text{C}$. The cultures were discarded and the wells gently rinsed twice with deionised water and air-dried. The wells were then stained with $210\text{ }\mu\text{l}$ of 0.1% (w/v) crystal violet for 10 min, rinsed with deionised water and air-dried. The dye was then eluted with $210\text{ }\mu\text{l}$ dimethyl sulfoxide and the amount of dye remained was determined at optical density of 595 nm, using a Bio-Rad enzyme-linked immunosorbent assay reader (California, USA).

To visualize biofilm, the four strains of *V. parahaemolyticus* (VpDAHV1, VpDAHV2, VpDAHV3 and ATCC 17802) were allowed to grow on two set of glass pieces ($1\times 1\text{ cm}$), which were placed in 24-well polystyrene plates supplemented with the cell-free extracts ($50\text{ }\mu\text{l}$) of each of the four isolates of *Pseudomonas* spp., and incubated for 24 h at $30\text{ }^\circ\text{C}$. The glass samples were stained with crystal

violet, and the visible biofilm examined under a light microscope (Nikon Inverted Research Microscope ECLIPSE Ti 100, Japan). Another set of glass pieces with biofilms grown was rinsed with PBS, stained with acridine orange (0.1%), and examined using a confocal laser scanning microscope (Carl Zeiss LSM 710) at 488 nm argon laser, with BP 500–640 band pass emission filter and Zen 2009 software (Carl Zeiss, Germany).

2.3. Degradation of N-hexanoyl-L-homoserine lactone (C6-HSL) assay

N-hexanoyl-L-homoserine lactone (C6-HSL) degradation activity of the cell-free extracts of the four isolates of *Pseudomonas* spp. (PsDAHP1, PsDAHP2, PsDAHP3 and PsDAHP4) was determined using *C. violaceum* CV026 as the reporter strain in a well-diffusion C6-HSL bioassay following the method described earlier [16]. The substance, C6-HSL, was used as a test compound because it is the major signalling molecule that regulates the expression of virulence factors, biofilm maturation and extracellular protease expression of pathogens such as *Aeromonas* sp., *Edwardsiella tarda* and *Vibrio* sp. [17,18].

Agar plates were prepared by mixing 3 ml overnight culture of CV026 in 20 ml LB agar which was then poured into petri dishes. After setting, a 5 mm diameter plug was removed from each plate to form a well for the assay. The AHL-lactonase reaction mixture containing $10\text{ }\mu\text{l}$ of the extracts and $190\text{ }\mu\text{l}$ of 24 nM C6-HSL in 50 mM phosphate buffer (pH 8.0) was incubated at $25\text{ }^\circ\text{C}$ for 45 min before termination by addition of $50\text{ }\mu\text{l}$ of 10% w/v sodium dodecyl sulphate (SDS). The reaction mixture was then transferred into the prepared plate wells, and the radius of CV026 growth after 24 h was used to determine residual C6-HSL levels. One unit of AHL-lactonase activity is defined as the amount of enzyme required to hydrolyse 1 nM C6-HSL per min under the conditions described. AHL-lactonase activity was reported as the degradation rate of AHL in $\text{mg l}^{-1}\text{ h}^{-1}$.

2.4. Microbial adhesion to hydrocarbon assay

Effects of the cell-free *Pseudomonas* extracts on cell surface hydrophobicity of GFP-VpDAHV2 were evaluated using a microbial adhesion to hydrocarbon assay. The bacteria GFP-VpDAHV2 (1 ml-OD530) were placed into glass tubes, and mixed with $100\text{ }\mu\text{l}$ of toluene, $50\text{ }\mu\text{l}$ of each of the extracts. The mixture was then allowed

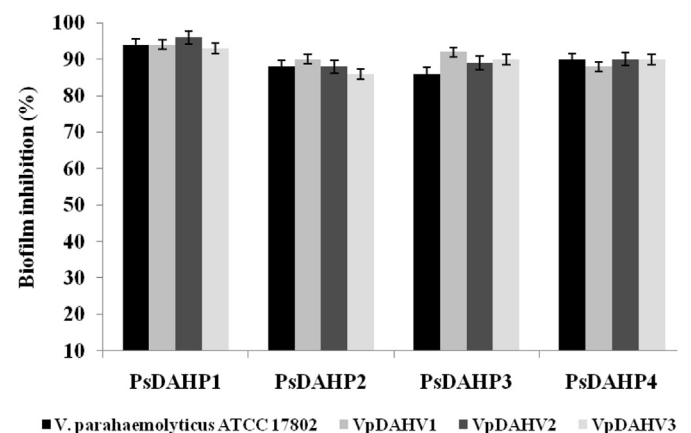


Fig. 1. Biofilm inhibition (%) of PsDAHP1, PsDAHP2, PsDAHP3 and PsDAHP4 against *V. parahaemolyticus* ATCC 17802, VpDAHV1, VpDAHV2 and VpDAHV3 at concentration $50\text{ }\mu\text{l}$ of extracts, quantified by crystal violet adsorption and measuring absorbance at 570 nm. Mean values of triplicate individual experiments and SDs are shown. Dunnett's test demonstrates significant difference between the treated and the control ($p < 0.05$).

to settle for 30 min, and optical density of the aqueous phase was determined by a spectrophotometer (UV1800 Shimadzu, Japan). Hydrophobicity index of GFP-VpDAHV2 was calculated based on the formula described [19].

2.5. Quantification of exopolysaccharide

The level of exopolysaccharide produced by *V. parahaemolyticus* was determined by carbohydrate assay. Sterile glass pieces were immersed in GFP-VpDAHV2 culture containing 50 μ l of each of the *Pseudomonas* extracts or Zobell broth (control) in 24-well polystyrene plates, and incubated for 24 h. The glass slides were rinsed with 0.9% NaCl solution and cell suspensions in the solution were transferred to test tubes with an equal volume of 5% phenol, to which 5 volumes of sulphuric acid containing 0.2% of hydrazine sulphate were added. The mixture was incubated in darkness for 1 h, centrifuged at 10,000 \times g for 10 min, and optical density of the supernatant determined at 490 nm.

2.6. Assessment of biofilm metabolic activity

To each of the microtitre plate wells containing *Pseudomonas* extracts and GFP-VpDAHV2 mature biofilm, 900 μ l of fresh ZMB broth, 90 μ l of XTT-2,3-bis (2-methoxy-4-nitro-5-sulphophenyl)-5-

[(phenyl amino) carbonyl]-2H-tetrazolium hydroxide salt solution (1 mg/ml) (Sigma Aldrich, St. Louis, MO), and 10 μ l menadione solution (1 mM) were added. The wells were incubated in darkness at 28 $^{\circ}$ C for 5 h. XTT tetrazolium salt was reduced to XTT formazan, and formed violet color. The wells were then placed on a spectrophotometer and the optical density measured at 490 nm [20].

2.7. Confocal laser scanning microscope observation of GFP-VpDAHV2 biofilm inhibition by cell-free extract of PsDAHPI

In 24-well polystyrene plates, GFP-VpDAHV2 was incubated with PsDAHPI cell-free extract (50 μ l) and glass pieces at 30 $^{\circ}$ C for 24 h. Glass pieces incubated with GFP-VpDAHV2 was rinsed twice with PBS and the biofilms quantified using a confocal laser scanning microscope to image GFP signal from cover-slip samples. Biofilms were assessed using COMSAT software.

2.8. Determination of the lethal dose of VpDAHV2 in zebrafish

Zebrafish (120 days of age, with an average weight and length of 400 mg and 2.5 cm, respectively) were allowed to acclimatise for 30 days and used to test the virulence of GFP-VpDAHV2. The bacteria were incubated in TSB at 30 $^{\circ}$ C for 24 h, centrifuged at 5000 \times g for 10 min, and the harvested cells serially diluted in sterile saline to

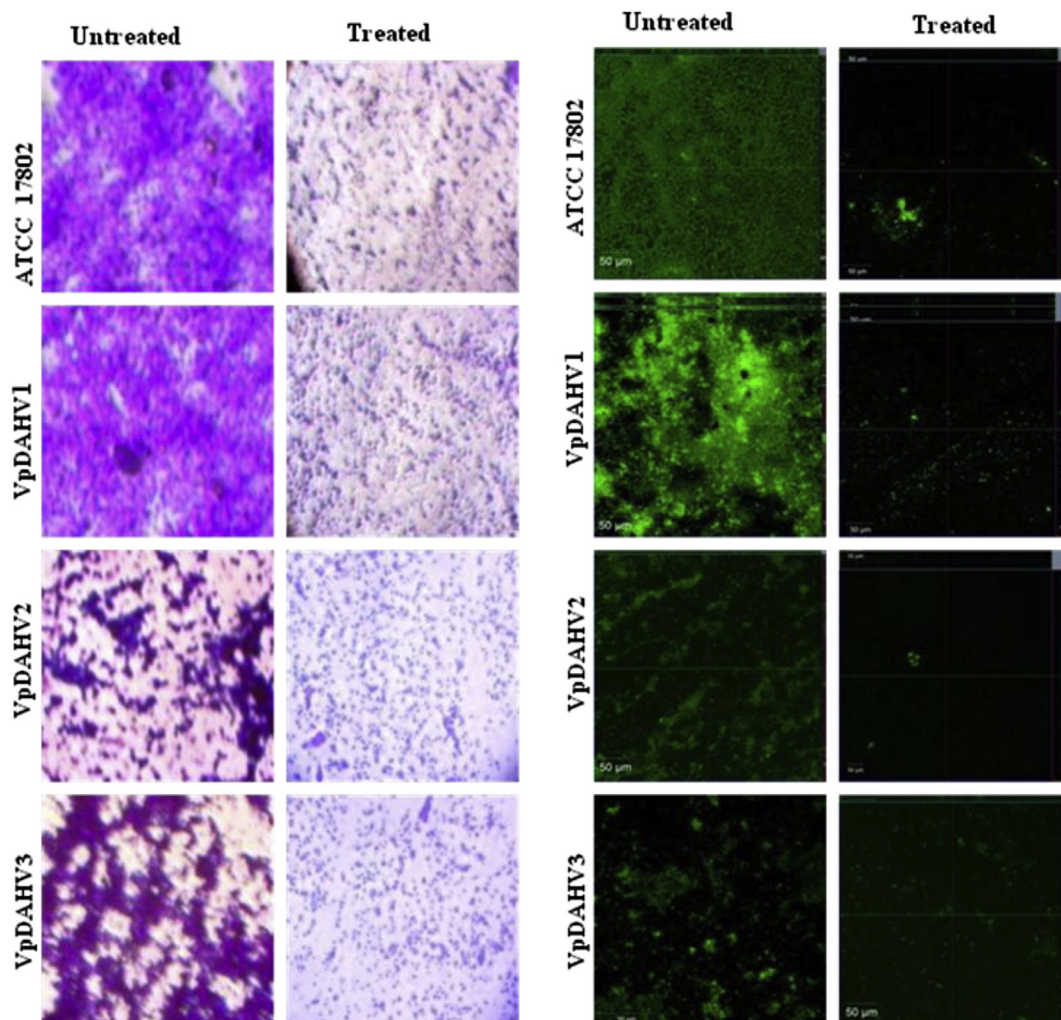


Fig. 2. Optical microscopic (40 \times) (left) and confocal laser scanning microscopic (right) visualization of anti-biofilm activity of PsDAHPI extract against *V. parahaemolyticus* ATCC 17802, VpDAHV1, VpDAHV2, and VpDAHV3.

obtain 10^3 to 10^7 colony-forming units (CFU) ml^{-1} . Aliquots of each dilution were plated on TSA to confirm the right concentrations in CFU ml^{-1} . Zebrafish were immersed for 24 h in duplicate vessels containing 10^3 , 10^4 , 10^5 , 10^6 or 10^7 CFU ml^{-1} of GFP-VpDAHV2. After the immersion, the fish were released into the fibreglass reinforced plastics tank and mortality was recorded for 10 days [21]. Virulence of GFP-VpDAHV2 was determined based on LD₅₀ which was calculated using the Probit method [22].

2.9. Preparation of GFP-VpDAHV2 and PsDAHP1-mixed feed

GFP-VpDAHV2 at 3×10^6 CFU ml^{-1} was found to be LD₅₀ for zebrafish; this bacterial density was employed for *V. parahaemolyticus* challenge. Zebrafish were immersed into the challenging tank for 24 h at 3×10^6 CFU ml^{-1} . After challenge the zebrafish were released into the fibreglass reinforced plastics tank and the water exchange was carried out every 4 days. Pure culture of PsDAHP1 from an overnight growth on ZMA plates containing 10 ml of ZMB were incubated at 30 °C for 24 h. After incubation, the cells were harvested by centrifugation at $2500 \times g$ for 20 min, washed twice with, and re-suspended in saline, in which the bacterial suspension was adjusted at 10^8 cells ml^{-1} . The PsDAHP1-mixed feed was prepared by gently spraying the required amount of bacterial suspension on the pre-mixed commercial feed, and mixed in a drum mixer to obtain a final concentration of 10^8 cells g^{-1} feed. The feed was dried in a drying cabinet using an air blower at 38 °C until moisture level was around 10%. After air-drying, it was sieved into pellets of appropriate size and stored at -20 °C until needed within seven days. The viability of the PsDAHP1 was determined by counting the number of colonies plated on ZMA.

2.10. Effect of PsDAHP1-mixed feed on VpDAHV2 infection of zebrafish

Zebrafish were randomly divided into four groups of 50 animals each, with 3 replicates. They were (1) Group A (*P. aeruginosa* control), being fed with PsDAHP1-containing feed and no challenge; (2) Group B (positive control), being fed with basal feed and challenged with GFP-VpDAHV2; (3) Group C (experimental group); being fed with PsDAHP1-containing feed and challenged with GFP-VpDAHV2; and (4) Group D (negative control), being fed with basal feed and no challenge. Fish mortality was recorded daily for 30-days [21], following the challenge; dead and moribund fish were removed and moribund fish examined for the presence of GFP-VpDAHV2 in the gills and intestine.

The dissected gills and intestine were washed twice with sterile phosphate buffer solution (PBS) and the tissues were homogenised individually in 1 ml sterile seawater and plated in ZMA medium

Table 1

N-hexanoyl-L-homoserine lactone (C6-HSL) degradation rate of selected AHL-degrading *Pseudomonas* isolates in LB medium supplemented with 5 mg l^{-1} C6-HSL. GenBank sequence accession numbers of partial 16S rRNA sequences of the isolates and related strains sharing 100% sequence identity with the sequences of the isolates (based on NCBI BLAST).

Strains	C6-HSL degradation ($\text{mg l}^{-1} \text{ h}^{-1}$)	Accession number	Related strains
PsDAHP1	1.5 ± 0.75	HQ400663	<i>Pseudomonas aeruginosa</i> LP8 (EU195558)
PsDAHP2	0.6 ± 0.3	HQ693274	<i>Pseudomonas aeruginosa</i> R9 (JQ660017)
PsDAHP3	0.5 ± 0.25	HQ693277	<i>Pseudomonas</i> sp QLD (JQ394932)
PsDAHP4	0.6 ± 0.3	HQ693272	<i>Pseudomonas aeruginosa</i> R9-771 (JQ660017)

with $100 \mu\text{g ml}^{-1}$ kanamycin and overlaid on LB agar supplemented with 1% NaCl. The plates were incubated at 30 °C and colonies counted after 48 h. Confocal laser scanning microscope was used to confirm the identity of colonies growing on selected plates as being GFP-VpDAHV2. The dissected gills and intestine were washed twice with PBS, placed in clean glass slide and validated for the colonisation of GFP-VpDAHV2, using 488 nm argon laser line scan for excitation with a multi-line argon laser and a band pass wavelength of 493–634 nm. For each experiment, an immersion dose of GFP-VpDAHV2 was plated as control.

2.11. Defence mechanisms in fish receiving PsDAHP1-mixed feed

In a parallel set-up, the effect of PsDAHP1 on parameters relating to defence mechanisms of zebrafish was examined. Five fish were taken randomly from each group after 10, 20 and 30 days of feeding. They were anaesthetised in 0.02% benzocaine solution (Sigma Inc., St. Louis, MO, USA). Individual blood samples (1.0–1.5 ml) were taken from the caudal vein using a heparinized syringe (25 G) fitted with a needle.

Blood (500 μl) from individual fish was mixed with 500 μl of AL medium (AIM-V medium) and Leibovitz's L 15 medium (GIBCO BRL, Gaithersburg, MD, USA), streptomycin and penicillin. Percoll (55%, Sigma Inc., St. Louis, MO, USA) was added and the mixture

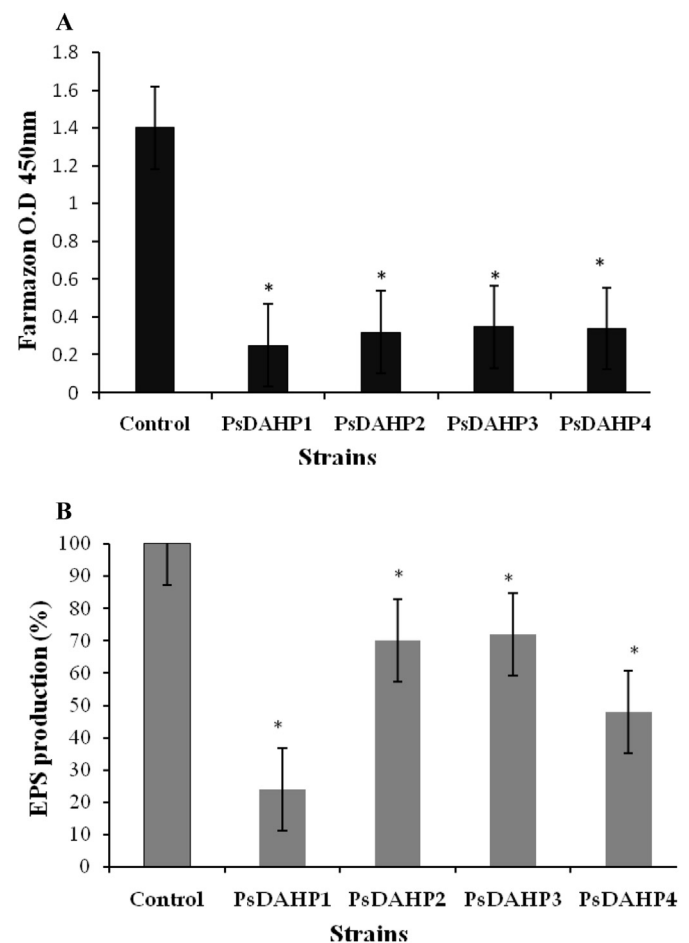


Fig. 3. PsDAHP1, PsDAHP2, PsDAHP3, and PsDAHP4 extracts inhibits the (A) metabolic activity and (B) Inhibition of exopolysaccharide (EPS) production of VpDAHV2 biofilm, as percentage of control. Experiments were performed in triplicates; mean \pm SD are shown. Dunnett's test demonstrates significant difference between the tests and the control, * indicates statistical value ($p < 0.05$).

centrifuged at $400\times g$ for 15 min at 10°C . Leucocytes were obtained from the interface and washed with AL medium by centrifugation at $600\times g$ for 10 min at 10°C . The leucocytes were then suspended in AL medium with 5.5 mM glucose. Cell viability was analysed by trypan blue (0.1%) with a haemocytometer.

For phagocytic cell test, 300 μl of leucocyte suspension (in L-15 medium) in 10 ml test tube, in triplicate, was added with 300 μl of formalin-killed *V. parahaemolyticus* (in PBS) and the mixture was incubated for 1 h at 17°C . Then, 900 μl of cold PBS was added, and the tubes were centrifuged at $300\times g$ for 5 min. The supernatants were discarded and the pellets were smeared on slides. The slides were air-dried, stained with Giemsa solution (Sigma Inc., St. Louis, MO, USA) and the leucocytes were visualized and the number of phagocytic cells per 100 adhered cells was counted microscopically.

For superoxide dismutase (SOD) and lysozyme activity assay, the blood samples were collected into an Eppendorf tube and centrifuged at $1500\times g$ for 10 min at 4°C . The serum samples were pooled and stored at -20°C .

Superoxide dismutase activity was determined by enzymatic assay method using a reagent kit (Sigma Inc., St. Louis, MO, USA). The reaction was based on its inhibitory effect on the rate of superoxide-dependant reduction of nitroblue tetrazolium (NBT) by xanthine–xanthine oxidase that was determined with a spectrophotometer at 550 nm. One unit of SOD activity was defined as the amount of enzyme necessary to produce 50% inhibition of the NBT reduction rate measured at 550 nm.

For lysozyme activity assay, 50 μl of the serum was mixed with 950 μl of *Micrococcus lysodeikticus* (Sigma Inc., St. Louis, MO, USA) in

0.05 M PBS (pH 6.2). The mixture was incubated at 25°C , and optical density determined spectrophotometrically after 1 min and 6 min at 530 nm. One unit of lysozyme activity was defined as the amount of enzyme producing a decrease in absorbance of $0.001\text{ min}^{-1}\text{ ml}^{-1}$ serum [23].

2.12. Statistical analysis

All assays were repeated at least three times, and all statistical analyses were performed using SPSS. Values were expressed as means \pm SD. Mean values were compared using one way ANOVA.

3. Results

3.1. In vitro screening of AHL-degrading *Pseudomonas* extract against *V. parahaemolyticus* biofilm

Four strains of AHL-degrading *Pseudomonas* spp. were identified based on 16S rRNA gene sequencing, namely *P. aeruginosa* HQ400663 (PsDAHP1), *P. aeruginosa* HQ693274 (PsDAHP2), *Pseudomonas* sp. HQ693275 (PsDAHP3) and *P. aeruginosa* HQ693272 (PsDAHP4) and the sequences were submitted to NCBI GenBank (accession numbers were HQ400663, HQ693274, HQ693277 and HQ693272) respectively. Cell-free extracts, 50 μl , of the four *Pseudomonas* spp. were screened for their ability to interrupt *V. parahaemolyticus* biofilm formation using crystal violet staining in a microtitre plate-based assay. All the extracts could suppress more than 80% of biofilm formation by ATCC 17802, VpDAHV1, VpDAHV2, and VpDAHV3 and highest suppression was obtained

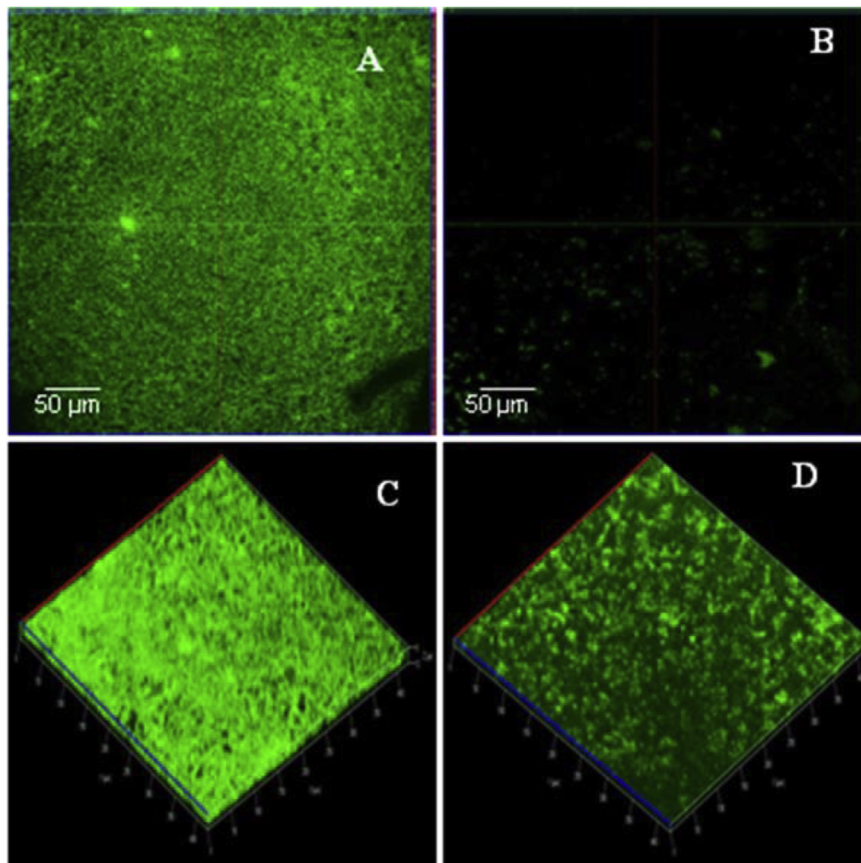


Fig. 4. Confocal laser scanning microscopy images of GFP-VpDAHV2 biofilms grown in the absence and presence of PsDAHP1 extract. (A) & (C), Control (absence of PsDAHP1 extract), 2D image; (B) & (D) GFP-VpDAHV2 + PsDAHP1 extract, 3D image.

Table 2
Mortalities of zebrafish with different dose of GFP-VpDAHV2.

Strain	Dose (CFU ml ⁻¹)	Mortality (%) (mean ± SD)	LD ₅₀ (CFU ml ⁻¹)
VpDAHV2	10 ³	13.2 ± 0.66	~ 3 × 10 ⁶
	10 ⁴	25.3 ± 1.26	
	10 ⁵	39.2 ± 1.96	
	10 ⁶	48.2 ± 2.41	
	10 ⁷	69.2 ± 3.46	
	10 ⁸	72 ± 3.6	

with PsDAHP1 extract (Fig. 1). Under light and confocal laser scanning microscopy, reduction in micro-colony formation in the four strains of *V. parahaemolyticus* was observed when treated with the cell-free extracts (Fig. 2). Among the four strains of *V. parahaemolyticus* tested, VpDAHV2 and ATCC 17802 were found to be potential biofilm formers, while the other two had lesser potential (data not shown). Therefore, in this study, VpDAHV2 was selected for further studies.

3.2. Degradation of C6-HSL assay

All the *Pseudomonas* cell-free extracts were able to degrade C6-HSL. Among the four, that of PsDAHP1 had maximum effect, with 1.5 mg C6-HSL degradation l⁻¹ h⁻¹ within 4 h, about 3 times higher than the extracts from other strains (Table 1).

3.3. Microbial adhesion to hydrocarbon and XTT assay (2, 3-Bis-(2-Methoxy-4-Nitro-5-Sulphophenyl)-2H-Tetrazolium-5-Carboxanilide)

Cell-free extracts of all the AHL-degrading *Pseudomonas* sp. (PsDAHP1, PsDAHP2, PsDAHP3, and PsDAHP4) were screened for their activities on hydrophobicity index. At 50 µl concentration, the extracts reduced the hydrophobicity index of the GFP-VpDAHV2 from 61 ± 0.94% to 21 ± 0.96%, 38 ± 0.84%, 41 ± 0.63%, 30 ± 0.6% in groups treated with PsDAHP1, PsDAHP2, PsDAHP3 and PsDAHP4 extracts, respectively. XTT assay confirmed that all the extracts inhibited the activity of GFP-VpDAHV2 that resulted in decreased

biomass in the GFP-VpDAHV2 biofilm (Fig. 3A), and maximum inhibition was observed in the PsDAHP1 activity.

3.4. Estimation of exopolysaccharide production

AHL-degrading *Pseudomonas* cell-free extracts were screened for their inhibitory activities on GFP-VpDAHV2 exopolysaccharide production. All the extracts inhibited exopolysaccharide production, with maximum inhibition by PsDAHP1 (Fig. 3B).

3.5. Inhibition of GFP-VpDAHV2 biofilm

Under confocal laser scanning microscope, with 2D and 3D images, PsDAHP1 extract showed significant inhibition of GFP-VpDAHV2 biofilm formation, whereas, no inhibition was detected in the untreated GFP-VpDAHV2 group (Fig. 4). The thickness and biovolume of biofilm formed in the control slides were 48.5 ± 2.45 µm and 44.1 ± 2.20 µm³, respectively, whereas the values in the treated slides were 15.2 ± 0.76 µm and 0.8 ± 0.04 µm³, respectively.

3.6. Effect of PsDAHP1-mixed feed on GFP-VpDAHV2 infection in the zebrafish

Virulence of GFP-VpDAHV2 in zebrafish was confirmed by immersion challenge and the LD₅₀ value was 3 × 10⁶ CFU ml⁻¹ (Table 2). Using the LD₅₀ dose to challenge the fish, mortality of the negative control and PsDAHP1-treated fish was found to be zero during the 30 days challenging period (Fig. 5). The positive control fish, treated with GFP-VpDAHV2, had mortality more than 70% at day 20 after the challenge, while the experimental group, receiving PsDAHP1-mixed feed, had mortality of 20% up to day 29 after the challenge.

After 30 days in culture, colonisation of GFP-VpDAHV2 was observed in the moribund zebrafish being challenged with GFP-VpDAHV2 in the gills and intestine of the fish, as observed under confocal laser scanning microscope (Fig. 6). In the gills, GFP-VpDAHV2 was mostly colonised in the lamellar region; whereas in the intestine, the colonisation was mostly in entire region. The

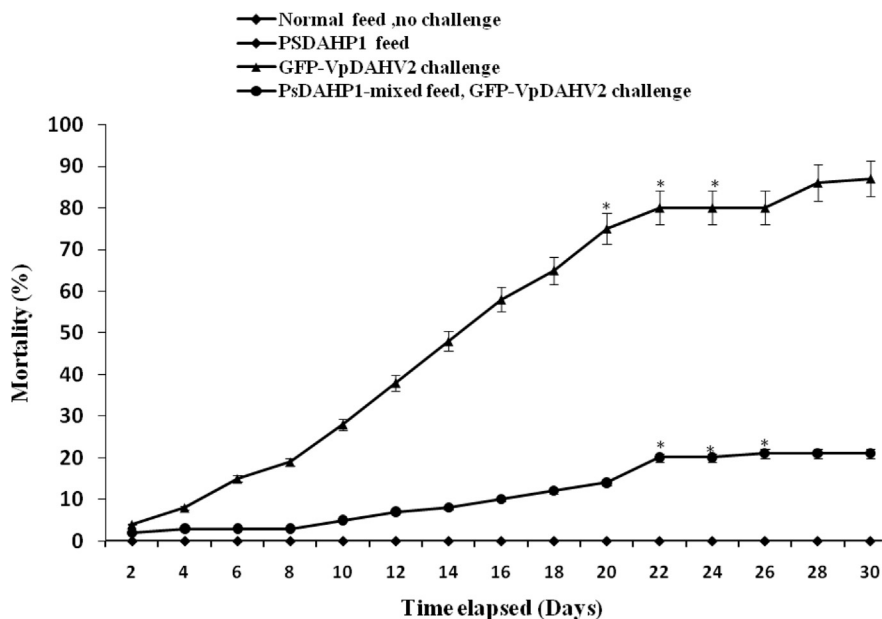


Fig. 5. Mortality rate of zebrafish in four groups of animals. Each point and error bar represents the mean of triplicate and standard deviation.

GFP signal was markedly reduced in the fish given PsDAHP1-mixed pellets during the challenge. The results were confirmed by plate counts of GFP-VpDAHV2 at day 30 post-challenge, in which the bacterial counts in the gills (4.1×10^4 CFU g^{-1}) and intestine (5.4×10^4 CFU g^{-1}) of the challenged fish were markedly reduced in both tissues in PsDAHP1-treated group (gills, 0.9×10^2 CFU g^{-1} ; intestine, 1.2×10^2 CFU g^{-1}).

3.7. Fish defence mechanisms following PsDAHP1 administration

Phagocytic cells of leucocytes of zebrafish fed with *Pseudomonas*-mixed feed for was evaluated for 30 days (Table 3). Initially without formalin-killed *V. parahaemolyticus* treated phagocyte cells in the fish receiving PsDAHP1 and control tank was $7 \pm 2\%$ and $13 \pm 3\%$ phagocytic cells respectively. Increased phagocytic cells were observed in leucocytes from all the fish that received the feed and highest rate was exhibited in the fish receiving PsDAHP1 ($51 \pm 0.5\%$ vs $40 \pm 1.5\%$ in control, $p < 0.05$).

Superoxide dismutase in the serum of the zebrafish receiving PsDAHP1-mixed feed was $23 \pm 0.1\%$, while the level 11.0 ± 0.5 in the control group. Similarly, the lysozyme activity in the serum of the fish receiving PsDAHP1-mixed feed was also significantly ($p < 0.05$) higher than that of the control ($40 \pm 0.5\%$ vs $18 \pm 0.2\%$ in control).

4. Discussion

The present study represents the first report on inhibition of GFP-tagged *V. parahaemolyticus* VpDAHV2 colonisation and

Table 3

Defence mechanisms in fish receiving *P. aeruginosa* PsDAHP1 cell-free extracts.

Treatment	Phagocytic cell	Superoxide dismutase activity	Lysozyme activity
Not receiving (control)	40 ± 1.5	11 ± 0.5	18 ± 0.2
PsDAHP1	51 ± 0.5	23 ± 0.1	40 ± 0.5

immune modulatory effects of zebrafish received AHL-degrading *P. aeruginosa* PsDAHP1 cell-free extract. The bacteria *V. parahaemolyticus* could persist for a long time in aquatic habitats, and replicate in the host [24]. Inhibition of bacterial biofilm has been proposed as one of the most potential strategies for increasing the sensitivity of pathogens in biofilm to antibiotics. In the present study, among the four *Pseudomonas* spp. studied, cell-free extract of *P. aeruginosa* PsDAHP1 was able to degrade C6-HSL, microbial adhesion to hydrocarbon, exopolysaccharide and biofilm metabolic activity, resulting in a decreased biofilm formation by the virulent *V. parahaemolyticus*. The reduction of biofilm by *Pseudomonas* extracts was also revealed morphologically by confocal light scanning microscope, showing disintegrated architecture and reduced thickness. This study is similar to a recent report that the *Bacillus licheniformis* cell-free extract disperses *Streptococcus aureus* biofilm [25]. Hydrophobicity index is an imperative factor for cell accumulation and targeting the hydrophobicity index is a novel way of inhibiting biofilm formation. Thus, reduction in the hydrophobicity index reduces the accumulation of bacteria that leads to the inhibition of biofilm formation. From microscopic observation, it was

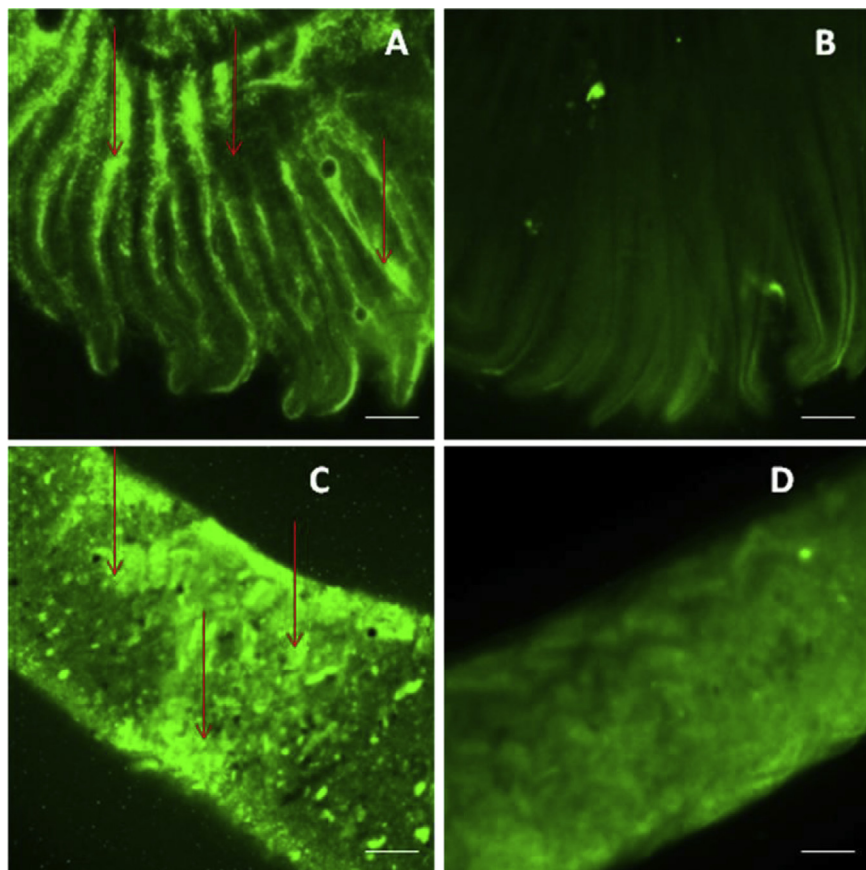


Fig. 6. Colonisation of GFP-VpDAHV2 in the gills (A) and intestine (C) of zebrafish challenged by GFP-VpDAHV2, observed under confocal light scanning microscope. The colonisation was markedly decreased in the gills (B) and intestine (D) of the fish that were given PsDAHP1-mixed pellets. Arrow represents the colonisation of GFP-VpDAHV2.

clear that the PsDAHP1 extract-treated biofilm architectures were looser than the control biofilm. Therefore, inhibition of hydrophobicity index and exopolysaccharide production by the PsDAHP1 cell-free extract slacked architecture of the GFP-VpDAHV2 micro-colonies. This would make *V. parahaemolyticus* more susceptible to antibiotics that in turn will facilitate the eradication of biofilm.

Exopolysaccharide and cell surface hydrophobicity play an important role in bacterium-host cell interactions [26]. Previous studies reported that exopolysaccharide play a vital role in biofilm architectures in *V. parahaemolyticus* [27]. High level of the substance leads to alterations in biofilm architecture that correlate with an increased resistance to biocides such as chlorine [28,29].

By applying the benefits of using *Pseudomonas* cell-free extracts found in the *in vitro* findings into the *in vivo* study, we found that zebrafish receiving feed mixed with PsDAHP1 survived better than the control group. The result was also supported by the findings that phagocytic cells of the leucocytes, serum SOD and lysozyme activities and colonisation of the challenged bacteria *V. parahaemolyticus* was markedly reduced in the gills and intestine of the fish receiving PsDAHP1-mixed feed. The findings were in agreement with others in different species of fish, pathogens and probiotics [30]. For instance, SOD activity of tilapia significantly increased by *Bacillus* spp.-mixed feed [31], and lysozyme activity increased in rainbow trout that received *Lactobacillus rhamnosus*, *Carnobacterium divergens*, and *Lactobacillus sakei* [32]. Likewise, in grouper, *Bacillus* enhanced growth and immune responses [33]. Since both phagocytic cells of zebrafish and leucocytes and SOD and lysozyme activities were increased by PsDAHP1 in this study, it therefore suggests that PsDAHP1 could increase resistance in the fish through the stimulation of both physiological (phagocytic cells) and biochemical responses (SOD and lysozyme activity). It remains to find out what substance(s) released from PsDAHP1 is responsible for the induction of these responses.

While the search for the active substance(s) is on-going, it is possible to apply PsDAHP1-mixed feed in zebrafish to protect the fish against *Vibrio* infections, as it was also found in this study that PsDAHP1 had no adverse effect on the fish.

In conclusion, cell-free extract of *P. aeruginosa* PsDAHP1 was found to antagonise the actions of *V. parahaemolyticus* by suppressing AHL synthesis, biofilm formation, production of exopolysaccharide, and by modifying the adhesion properties of the pathogen. Moreover, oral administration of PsDAHP1-containing feed reduced mortality caused by *V. parahaemolyticus* challenge in zebrafish, enhanced leucocyte phagocytic cells, and serum activities of SOD and lysozyme in the fish. Moreover, PsDAHP1 had no adverse effect to the fish. AHL-degrading *P. aeruginosa* PsDAHP1 could control biofilm-associated infections caused by *V. parahaemolyticus* and enhance defence mechanisms of zebrafish.

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