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Bioactive Prodigiosin Isolated from *Serratia marcescens* using Solid State Fermenter and its Bactericidal Activity Compared with Conventional Antibiotics

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Abstract

The prodigiosin was isolated from *Serratia marcescens* and the production process was optimized by response surface methodology-central composite design (CCD). The bactericidal efficiency of prodigiosin was analyzed against antibiotic resistant pathogens. The maximum yield of prodigiosin (70.4023 g) was achieved per kg of tannery fleshing at optimized condition time, 81.2 h; temperature, 29°C; pH, 6.8; moisture, 50%. The prodigiosin exhibited complete inhibition of the growth of *Pseudomonas aeruginosa* compared with other conventional antibiotics which is confirmed by disc diffusion method, LIVE/DEAD assay and fluorescence emission image. Prodigiosin, the bioactive compound was produced from tannery solid waste in solid state fermentation. The antimicrobial effect prodigiosin antibiotic resistant pathogens may make a possible platform in future for bacterial disinfection in clinical/ pharmaceutical wastewater.

Keywords: Serratia marcescens; Prodigiosin; Tannery fleshing; Solid state fermenter; Antibiotic resistant bacteria

Introduction

The tannery fleshing is proteinaceous solid waste generated from leather industries during leather manufacturing process. Tannery fleshing (TF) constitutes about 50 to 60 % of solid wastes generated from the leather industry [1]. They are dumped onto open land, digested in anaerobic digester or incinerated in thermal incinerators [2-4]. The inherent disadvantages associated with these disposal techniques are huge energy consumption, large retention period and objectionable odor emission [5]. The TF contains adequate carbon and nitrogen contents, besides the essential amino acids and favorable moisture content for the growth of microorganisms [6,7]. Despite the TF is the rich source of protein (50 - 60%) with moderate fat content (6-7%), it is underutilized due to the lack of viable technology. Prodigiosin is bioactive alkaloid compound, the secondary metabolite usually accumulates during the later stage of microbial growth in the process of submerged fermentation. Prodigiosin has pharmacological activity such as anticancer, antimicrobial, immunosuppressive, antimalarial and antiprotozoal. Lapenda JC et al has reported prodigiosin is the effective antimicrobial agent against gram negative and gram positive bacteria [8]. Despite wide pharmacological applications, the production of prodigioisin has not yet reached the economically viable process. Prodigiosin production depends on carbon and nitrogen sources of the substrate. The high carbon and nitrogen sources containing TF may support the production of prodigiosin. Sumathi et al. (2014) reported that TF was considered as the potential substrate for the production of prodigiosin in submerged fermentation [9]. The prodigiosin production was carried out in submerged fermentation and limited TF concentration (4%) was utilized with low yield of the prodigiosin. Moreover, separation of prodiogiosin through extraction from the submerged fermentation involves huge amount of solvents. Solid state fermentation (SSF) has been gaining renewed interest and focused attention from researchers owing to its importance in recent developments in biomass energy conservation. SSF has many advantages over submerged fermentation reactor including higher production yield, more effectiveness, more eco-friendly [10,11] and easy recovery of byproduct. Hence, the focal theme of the present investigation was to produce prodigiosin from proteinaceous TF through solid state fermentation using *Serratia marcescens* and to compare its bactericidal activity against *Pseudomonas aeruginosa* with conventional antibiotic compounds.

Materials and Methods

Isolation and characterization of microorganisms

The limed tannery fleshing was collected from a commercial tannery in Chennai, Tamilnadu, India. It was mixed with soil in the ratio of 1:1 and kept for 30 days. After 30th day 1g of composted material was collected and diluted with 10 ml of saline solution (0.85% NaCl), from this 1 ml was spread onto nutrient agar and incubated at 30 °C for 24 h. The red colour bacterial strain was isolated and cultivated in nutrient broth (peptic digest of animal tissue 5g/l; sodium chloride 5g/l; beef extract 1.5g/l; yeast extract 1.5g/l). This culture was used for the production of prodigiosin in Solid state fermenter (Figure 1). The basic biochemical tests were performed for the isolated red bacteria. DNA was extracted and sequencing was analyzed using the following method. 500 µl of 24 h old culture was added to 150 µl of TE buffer (10 mM TRIS-HCl, pH 8.0, EDTA1 mM) containing 8 µl proteinase K (20 mg/ ml) and 0.5% SDS and incubated at 56°C for 1 h. An equal volume of phenol: chloroform: isoamyl alcohol mixture was added and centrifuged at 15,600 g for 5 minutes. The aqueous phase was removed and DNA was precipitated with 0.1 volume of 3 M sodium acetate (pH 5.2) and absolute ethanol (2 volumes). It was cooled at -20°C for 30

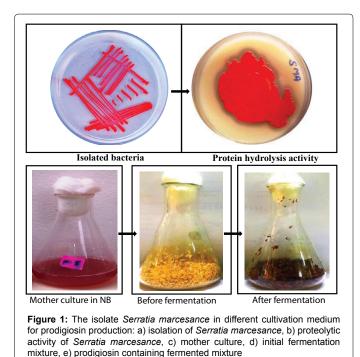
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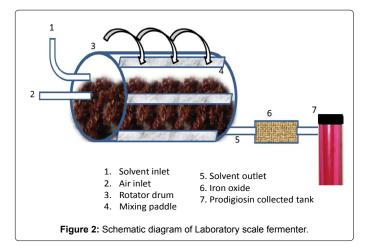
minutes, the precipitated DNA was washed again with 70% ethanol, dried, and reconstituted in 20 μ l TE buffer and stored at -20°C until use. The small subunit rRNA gene was amplified using primers 27F (AGAGTTTGATCMTGGCTCAG) and AGAGTTTGATCMTGGCT-CAG (TACGGYTACCTTGTTACGACTT) primers for bacteria, and then performed 35 amplification cycles at 94°C for 45 sec, 55°C for 60 sec, and 72°C for 60 sec. The PCR product was sequenced using the 518F (CCAGCAGCCGCGGTAATACG) and 800R (TACCAGGGTAATCC) primers [12]. The 16S rRNA sequence was analyzed for the similarity and homology with the existing sequences available in National Center for Biotechnology Information (NCBI) data base using BLAST search.

Preparation of substrate

Wheat bran was purchased from a commercial outlet and screened to remove impurities, washed with water to remove grit materials and dried. It was milled with an electric wearing blender and sieved through 20-40mm mesh sieve to get an uniform size fraction. Limed TF, the major solid waste generated from the tanneries during leather manufacture. It was collected from a commercial tannery in Chennai, Tamilnadu, India. It was treated with ammonia solution (25% v/v) for 3 to 4 h to remove the adhered calcium salts from TF. The TF was further washed in running water and cut into small pieces (1mm diameter) then stored at 4°C for further experiments.

Fermenter design and experimental setup

The fermenter consists of acrylic roller bottle of volume 1.61 L (length, 17 cm and width, 11 cm) connected to a filtered air supply source at the rate of 20 mL/min (Figure 2). The acrylic bottle was placed in a roller system, operated by a motor, to facilitate continuous rotation. The prodigiosin production media containing 1 Kg of fermented mixture (wheat bran, 700 g and TF, 300g) was supplemented with minimal medium of volume 500 mL (Na₂HPO₄7H₂O, 33.90g; KH₂PO4, 15g; NaCl, 2.5g; NH₄Cl, 5g/L) and 10% of inoculum was added to the



fermenter. The optimization of the prodigiosin production was carried out with different environmental factors such as time, temperature, pH, substrate concentration, moisture content and agitation. The optimization study was carried out by central composite design using design expert 8 software. The minimum and maximum values of the factors were entered into the software then the software automatically generated 50 experimental condition. The prodigiosin produced under the above 50 experimental condition were analyzed in laboratory scale fermenter and the values were enter into software. The significance of the experimental values were analyzed through ANOVA. Prodiogiosin was extracted from one gram of fermented SSF mixture by using various solvents (acetone, ethanol, methanol, isopropanol, benzene, hexane, diethyl ether, chloroform, dichloromethane, dimethyl sulfonate or petroleum ether) of volume 10 ml separately. The prodigiosin concentration was determined in each eluted solvent. The solvent which extracted high prodigiosin was used in further experiments to elute prodigiosin from fermented SSF mixture. After fermentation of TF, the solvent was added into the rotating fermenter and the solvent was collected from the fermenter outlet. The purified prodigiosin was characterized by UV-visible spectrum, high performance liquid chromatography and nuclear magnetic resonance spectrum. The presence of prodiogiosin in the SSF mixture was confirmed by diffuse reflectance spectrophotometer (Varian TCA Cary 300) at $\lambda_{200-800}$ nm.

Statistical analysis

The central composite design was used to optimize the most significant factors for improving prodigiosin production. The five independent factors were studied at three different levels and a set of 50 experiments were carried out (Table 1). Design expert, version 8.0.7 (statease Inc., Minneapolis, USA) was used for the experimental designs and regression analysis of the experimental data. Statistical analysis of the model was performed to evaluate the analysis of variance (ANOVA). The quality of the polynomial model equation was judged statistically by the coefficient of determination R² and its statistical significance was determined by an f-test.

Antimicrobial activity of prodigiosin

The antibacterial activity of prodigiosin was evaluated against *Pseudomonas aeruginosa* using the disc diffusion method. The pathogens, *Pseudomonas aeruginosa* was grown overnight on Muller Hinton agar plates. 24 h matured colonies were suspended into sterile saline (0.9%) solution of volume 5mL and the density of the suspension was adjusted to approximately 3×10⁸ colony forming units (CFU/ mL).

Factor	Parameter	Units	Minimum	Maximum	Coded	Values	Mean	Std. Dev.
Α	Time	hour	38.91806	153.08194	-1.000=72.00	1.000=120.0	96	22.33772
В	pН		2.24317	11.75683	-1.000=5.00	1.000=9.00	7	1.86148
С	Temperature	°C	18.10793	41.89207	-1.000=25.00	1.000=35.00	30	4.65369
D	TF concentration	%	6.21586	53.78414	-1.000=20.00	1.000=40.00	30	9.30739
E	Moisture	%	26.21586	73.78414	-1.000=40.00	1.000=60.00	50	9.30739

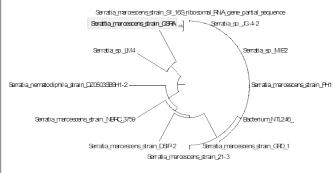
Table 1: Factors tested in the solid state fermentation

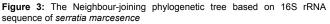
The swab was used to inoculate the dried surface of MH agar plate by streaking completely over the surface of the agar, rotating the plate approximately by 90° to ensure an even distribution of the inoculum. The medium was allowed to dry for about 3 min before adding a sterile paper disc of diameter 5 mm. The antimicrobial efficiency of prodigiosin was compared with conventional antibiotic compounds purchased from Himedia such as ampicillin (SD002), tetracycline (SD037), erythromycin (SD013), bacitracin (SD003) and chloramphenicol (SD006). The prodigiosin and conventional antibiotic compounds (1mg) were dissolved separately in methanol of volume 10 mL to get a homogenous solution. The discs were immersed in the prodigiosin solution and dried at 50°C for 2 h. Each disc was tapped gently down the agar to provide uniform contact and the plate was incubated at 37°C for 24 h. The zone of clearance of each antibiotics and prodigiosin were measured. The bacterial removal efficiency of prodigiosin was assessed by fluorescent dye uptake capacity of the bacterial cell. One milligram of ampicillin, tetracycline, erythromycin, bacitracin and chloramphenicol ampicillin, and prodigiosin were added separately in distilled water of volume 5 ml. The mature Pseudomonas aeruginosa culture solution of volume 10% was added into the antibiotic containing solution and incubated for 2 h at 37°C. The solutions were transferred into the illuminator and the fluorescence emission was recorded. Then 1 ml of solution was mixed with 0.05ml (1mg/mL) of propidium iodide and SYTO 9 then it was incubated at room temperature for 2 min. The cells were separated by centrifugation at 5000 rpm for 10min and pellets were re-suspended in 1 ml of phosphate buffer (pH, 7.0). Then, the suspension of volume 20µl was transferred onto microscopic slide and captured its fluorescence images using fluorescence microscope (Olympus BX-61) using red filter.

Results and Discussion

Isolation of microorganism and optimization of prodigiosin production

The 16S ribosomal RNA gene sequence of isolated red bacterial sp. was analyzed using similarity search tool blast. This indicated high degree of similarity (99%) with Serratia marcescens. The Phylogenetic tree constructed based on Neighbor joining method using ClustalW software (Figure 3). The present investigation, the process parameters of SSF were optimized for prodigiosin production using CCD. The CCD was employed to determine the optimal levels of the factors that significantly affected prodigiosin production. The high and low levels with the coded levels for the factors are as presented in Table 1. Based on the regression analysis of the data from the Table 2, the effects of factors on prodigiosin production were predicted by the second order polynomial function. The statistical significance of equation was checked by f-test and ANOVA for the second order polynomial model as shown in Table 2. The precision of a model could be checked by the regression coefficient (R²), 0.8708. Linear and quadratic terms were both significant at 1% level. Therefore, the quadratic model was selected in this present investigation. The P-values were used as the tool to check the significance of each coefficient, which was necessary to understand the pattern of the mutual interactions between the



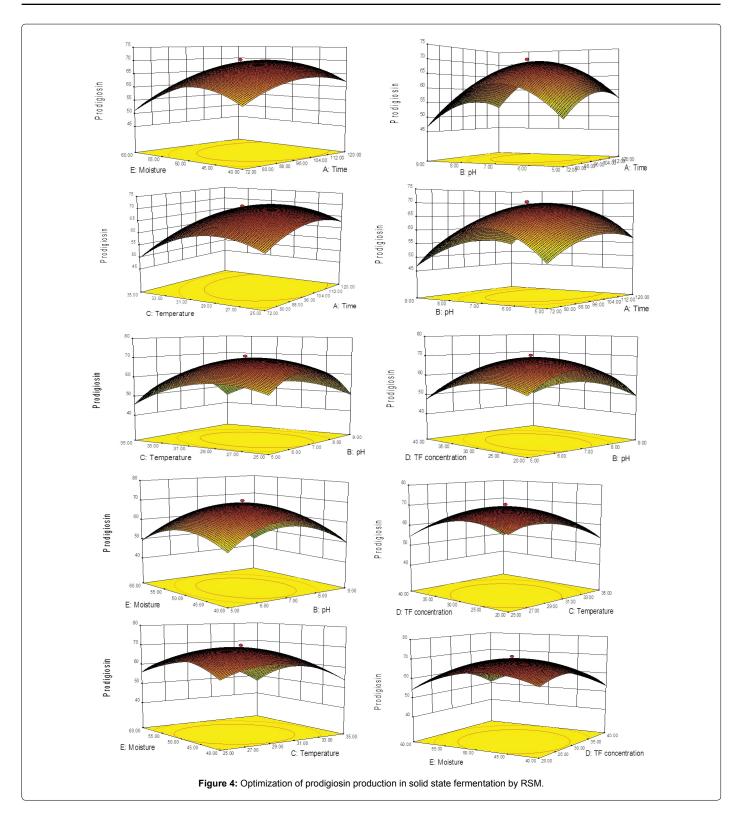


Source	Sum of Squares	df	Mean Square	F Value	p-value Prob > F	
Model	15964.52598	20	798.2263	9.77404	< 0.0001	Significant
A-Time	468.0205	1	468.0205	5.73077	0.0234	
В-рН	131.60701	1	131.60701	1.61149	0.2144	
C-Temperature	478.18802	1	478.18802	5.85527	0.022	
D-TF concentration	328.8718	1	328.8718	4.02694	0.0542	
E-Moisture	212.4835	1	212.4835	2.6018	0.1176	
Residual	2368.37136	29	81.66798			
Lack of Fit	2368.37136	22	107.65324			
Pure Error	0	7	0			
Cor Total	18332.89734	49				

Table 2: ANOVA for the second order polynomial model.

best selected factors. Smaller P-values indicate the significance of the corresponding coefficients. The Model F-value of 9.77 implies the model was significant. There was only a 0.01% chance that a "Model F-Value" this large could occur due to noise.Values of "Prob > F" less than 0.0500 indicate model terms are significant (Table 2). In this case A, C, A², B², C², D², E² are significant model terms. The Box-Cox plot is a tool to determine the most appropriate power transformation to apply to response data. Most data transformations could be described by the power function, $\sigma = fn (\mu^{\alpha})$, where sigma (σ) is the standard deviation, μ is the mean, and α is the power. Lambda (λ) is 1- σ in all cases. Power law transformations can only be performed on responses that are greater than zero. A recommended transformation is listed, based on the best lambda value, which was found to be + at the minimum point of the curve generated by the natural log of the sum of squares of the residuals. The standard deviation associated with an observation is proportional to the mean raised to the power, then transforming the observation by $\lambda = 0.5$ square root power gives a scale satisfying the equal variance requirement of the statistical model. The optimum conditions for the maximum prodigiosin yield (70.4g) at time, 96 h; pH, 7; temperature, 30°C and moisture content, 50% and mass of TF/ mass of SSF mixture 30% (w/w) (Figure 4).

Prodigiosin production = -651.6621.638189 * A* 42.2008 *

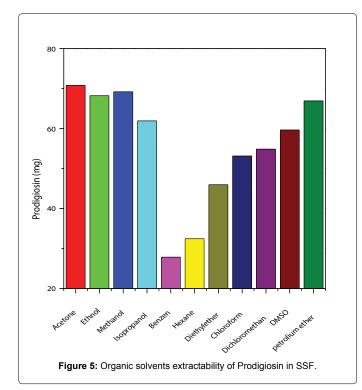


C17.96603 * B2.713859* D8.028217 * E-0.00421 * A * C0.00318 * A * B0.000281* A * D0.001216 * A * E0.118983 * C * B0.043777* C * D-0.04378 * C * E0.011225* B * D-0.022 * B * E0.010102 * D * E-0.00852 * A^2-3.24016* C^2-0.31676 * B^2-0.06941 * D^2-0.07703* E^2

The optimized conditions were maintained in SSF bioreactor for the production of prodigiosin. The agitation speed has no much

significant effect on production of prodigiosin (data not showed). The natural ventilation of the SSF reactor accessed enough oxygen for the metabolism of fermentative bacteria and in consequence a conducive condition for the production of prodigiosin was maintained. After 24 h, the colour of the support matrix of SSF reactor was changed from light yellow to red, this was an indication that prodigiosin production

was initiated, on further increase in incubation up to 96 h the intensity of the red colour was increased (Figure 1). The prodigiosin was extracted from the fermented matrix of SSF reactor using different solvents such as acetone, ethanol, methanol, isopropanol, benzene, hexane, diethyl ether, chloroform, dichloromethane, dimethyl sulfonate or petroleum ether. Among these solvents tested, the acetone extracted the maximum concentration of prodigiosin from the fermented matrix (Figure 5). Fermentation matrix was rotated continuously for providing aeration into the contents of SSF reactor which inturn to support microbial community attached to the surface of the matrix [13]. Prodigiosin production levels are greatly influenced by nutritional and physicochemical factors, such as nitrogen and carbon sources, inorganic salts, temperature, pH, agitation, and dissolved oxygen concentration [14-19]. Among them, the major factor influencing on prodigiosin production is medium composition. The production medium containing high protein content such as LB broth, nutrient broth, or peanut seed broth were found to support relatively high yields of prodigiosin and are usually better for pigment production in comparison to more defined media [16,19-21]. In general, carbohydrates were found to be poor nutrient sources for the production of prodigiosins, with glucose reported as a repressor of prodigiosin synthesis in Serratia marcescens [22,23]. Besides carbon source, the type of nitrogen source in the medium and the carbon-tonitrogen ratio also influenced the prodigiosin production. Overall, glycerol, glycine and fatty acids were found to be promising carbon sources or additives for increased production levels [24]. The influence of increased nitrogen sources on enhanced prodigiosin production. The maximum prodigiosin production was obtained with TF at 30% (w/w), this is comparatively higher than previously reported by researchers as shown in Table 3 [9,14-19,24-28]. The prodigiosin production reported with many bacterial strains isolated from different sources and with different carbon and nitrogen sources were compared with the present investigation (Table 3). An effective prodigiosin production could be achieved by increasing TF concentration from 10



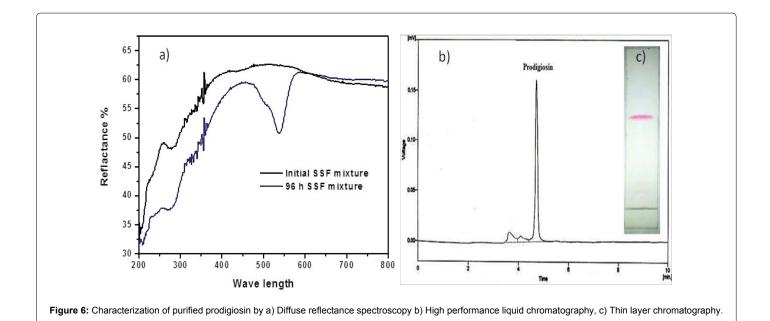
S. No	Prodigiosin Production media	Yield (g/l)	Reference			
1 2 3 4 5 6 7 8 9 10 12 13 14 15 16 17	Nutrient broth Peptone glycerol broth Sesame seed broth Sesame oil broth Peanut oil broth Peanut seed broth Copra seed broth Coconut oil broth Glucose Dextrose broth with casein Ethanol Ethanol and glucose Casein broth Sesame oil broth Corn steep mannitol medium Mannitol medium	1.9 1.35 16.68 0.767 2.89 38.75 1.94 1.42 0.005 1.3 3 4.28 0.66 37.5 34.00	[18] [18] [14] [14] [14] [14] [14] [27] [26] [24] [25] [28] [17] [16] [16] [16] [16]			
17 18 19 20	Mannitol medium Corn steep medium Cassava waste mannitol medium Cassava waste medium	34.00 25.60 49.50 27.00	[16] [16] [16]			
21 22 23	Luria Bertani glucose medium (NH ₄) ₂ PO ₄ + salt Glucose-glycerol medium	13.50 1.397 5.83	[16] [19] [28] [9]			
24 25 26	Animal fleshing solid waste Kitchen waste* Tannery animal fleshing solid waste*	8.3 4.155 (g/kg) 88.91 (g/kg)	[15] Present study			
	*solid state fermentation					

Table 3: Comparison of prodigiosin production with different growth medium

to 30 % (w/w) (Figure 4). The influence of above selected fermentation factors on prodigiosin production by this microbial strain, where minimum and maximum prodigiosin production obtained from 21.58 to 70.4 g/kg TF respectively. These results suggest that fermentation parameters greatly influenced the production of prodigiosin with this microbial strain and amino acids from TF played an important role in the biosynthesis of prodigiosin because of their regulatory role in the induction or repression of the prodigiosin production. Similarly, Sumathi et al 2014 also considered TF (animal protein) as the substrate in submerged fermentation for the production of prodigiosin. This present investigation focused on the solid state fermentation for the prodigiosin production. The solid state fermentation carried out in laboratory scale drum fermenter. The fermenter was designed with provision to extract prodigiosin after completion of fermentation. The prodigiosin production was favored by wheat bran (plant protein) with the TF (animal protein). The combination of plant (wheat bran) and animal (TF) proteins enhanced the prodigiosin production. The animal proteins helped the microbial growth and plant proteins provided the hydroxyl proline for the biosynthesis of prodigiosin. The hydroxyl proline mainly involved in the biosynthesis of prodigiosin production. The TF consists of proteins by 50 to 60 % of protein and lipid by 10 to 20%, the protein content provide amino acids for the formation of prodigiosin biosynthesis and the lipids provide carbon in the form of glycerol and fatty acid to enhance the prodigiosin production. This data helps in simulation of economic fermentation medium especially, with respect to selection of environmental factors and TF concentrations in scale-up studies to induce maximum prodigiosin production using Serratia marcescens.

Characterization of prodigiosin

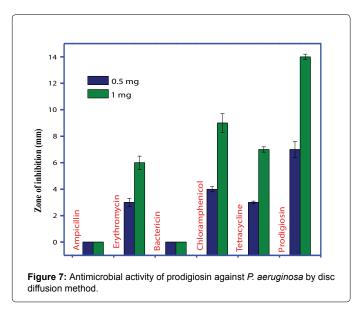
The fermented products before and after SSF were analyzed by diffuse reflectance spectroscopy. The change in colour was observed in SSF mixture from yellow to red denoted the presence of prodigiosin in solid state fermented medium (Figure 5). The Figure 6 shows a peak at λ_{ss2} nm in the final fermented mixture, this is considered to be the



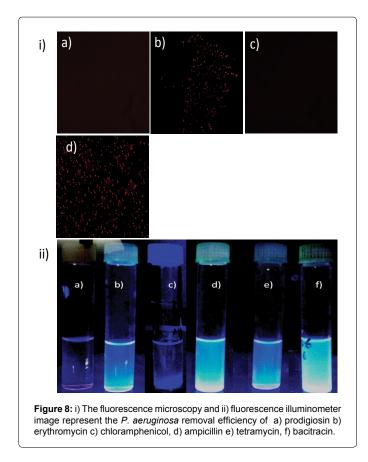
characteristic peak of prodigiosin [29]. The fermentation medium, before fermentation contained no peak at λ_{532} nm. The HPLC analysis was showed the >95% purity of the compound with single peak at retention time of 4.523 min as shown in Figure 6 [30].

Antimicrobial activity of prodigiosin

In the present investigation, the susceptibility of bacteria to inhibition towards the prodigiosin was assessed against P. aeruginosa and compared with conventional antibiotic compounds. The antimicrobial effect of prodigiosin was assessed on the basis of the zone of inhibition in the agar plate method. The prodigiosin impregnated disc showed the maximum zone of inhibition against P. aeruginosa (14 mm) as shown in Figure 7. The erythromycin, chloramphenicol and tetracycline were inhibited the P. aeruginosa growth with zone of influence 6, 9 and 7mm respectively. The prodigiosin exhibited more inhibition activity against P. aeruginosa compared with conventional antibiotic compounds. The ampicillin and bacitracin showed no inhibition of bacterial growth. The bacterial cell viability in antibiotic containing solution was investigated by fluorescence microscopic LIVE/DEAD assay. Fluorescent image of erythromycin and tetracycline containing bacterial solution showed 40 and 50% of bacterial cells with red colour. The prodigiosin and chloramphenicol solution showed 75 and 98% of bacterial cells with red colour. The red colored cell indicate, the propidium iodide stain was attached with the dead bacterial cell due to the presence of nucleic acid functional sites in the bacterial cell membrane [31]. The red colored cells indicated the dead cells because the addition of the propidium iodide was directly attached to the nucleic acid site. The nucleic acid leakage was occurred only in dead/damaged cells. The propidium iodide was attached with the nucleic acid site of the dead cells. So the dead cells only could emit the red fluorescent light under the fluorescent light due to the presence of propidium iodide. There were no red colored cells in ampicillin containing solution because the bacteria could survive in the presence of ampicillin and bacitracin. However, the prodigiosin containing bacterial cells were completely killed. It may due to the mechanism of action on the inhibition of cell wall biosynthesis, protein synthesis,



and DNA synthesis as described by peach et al 2013 [31,32]. Moreover, the same bacterial removal pattern was observed with illuminometer (Figure 7). The P. aeruginosa could emit the fluorescence light under the illuminometer. The ampicillin and bacitracin exposed solution were showed high fluorescence emission because the P. aeruginosa population was high in the ampicillin and bacitracin exposed solution. Erythromycin and tetracycline exposed P. aeruginosa solution showed less fluorescence emission and no fluorescence emission was observed in chloramphenicol and prodigiosin exposed P. aeruginosa solution. This indicate that the P. aeruginosa was completely eliminated in the prodigiosin and chloramphenicol containing solution. The prodigiosin is the effective antimicrobial compound compared to ampicillin, bacitracin, tetracycline and erythromycin. The fluorescence emission image also confirms the antimicrobial efficiency of prodigiosin on P. aeruginosa comparatively higher than conventional antibiotic compounds (Figure 8). The antimicrobial activity of prodigiosin



purchased from the Sigma Aldrich was similar with a conventional antibiotic as described by Jie Feng 2015 [33]. This present study shows prodigiosin has high antimicrobial activity compared than other antibiotic. The prodigiosin was effective enough to inhibit the activity of *P. aeruginosa* growth even at low concentration [9,34-38].

Conclusion

The present investigation provides the scope for effective production of pharmaceutical important secondary metabolite, prodigiosin from TF by solid state fermentation using *Serratia marcescens*. The Solid state fermenter shows high yield of prodigiosin production compared with submerged fermentation as reported earlier. The maximum prodigiosin (70.4 mg/kg of TF) yield was obtained at conditions such as fermentation time, 96 h; pH-7; temperature, 20 °C and moisture content, 50% and TF, 30%. The prodigiosin exhibited high antimicrobial activity against *P. aeruginosa* compared to other antibiotic compound. The antimicrobial potential of prodigiosin may make a possible platform in future for bacterial disinfection in clinical/ pharmaceutical wastewater.

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