

RESEARCH ARTICLE

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Isolation and Identification of Isolate, S1 with High Biotransformation Potential of Ferulic Acid to Vanillin

Nagaraju Bathini¹, Sai Krishna Esampally², Premsagar Korripally³, Vishnuvardhan Reddy Sultanpuram² and Thirumala Mothe^{4*}

Abstract

Nine actinobacterial isolates were purified from the sediment sample of Kogilvai village, Warangal, Telangana, based on their capability to grow on the minimal medium with Ferulic acid (FA) as only Carbon (C) source. FA to Vanillin conversion capacity of these isolates was identified by Thin-layer chromatography (TLC). Biotransformation of FA to Vanillin was high by four isolates, S1, S3, O3 and O4 when compared to other five isolates (O1, O2, S2, S4 and S5) with initial pH 7 in basal medium. Among these four isolates, optimal and rapid FA to Vanillin bioconversion of 140 mg/L was shown by isolate S1 with UV-spectrophotometry. Its conversion was confirmed by High Performance Liquid Chromatography (HPLC) analysis with retention time of 2.9 min after 28hrs of incubation at 37°C with 1g/L ferulic acid in the medium with 150 rpm. Isolate S1 could utilize Lactose, Maltose, Glycerol, Fructose, Galactose, Sucrose, Dextrose, L-Arabinose, ONPG, Esculin and not other carbohydrates present in the Himedia Hicarbo kit. Molecular characterization showed that 16S rDNA gene sequence of isolate S1 was 98.27% similar to Limosilactobacillus fermentum CECT 562 with completeness of 96.7% and was identified as Limosilactobacillus sp. 16S rDNA gene sequence of isolate S1 was submitted to NCBI GenBank and its accession number was OR136396. As this isolate has high potential of FA to Vanillin biotransformation capacity, it can be further explored to be used for industrial setups for commercial exploitation.

Keywords: Actinobacteria, Ferulic Acid, Molecular Characterization, High Performance Liquid Chromatography

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¹Department of Microbiology, Nagarjuna Government Degree college, Nalgonda, Telangana, India.

²Microbztech Labs Pvt. Limited, Sri Sai Dwarakapuri Colony, Cherlapally, Nalgonda, Telangana, India.

³Department of Biotechnology, University College of Science, Mahatma Gandhi University, Anneparthy, Yellareddygudem (PO), Nalgonda, Telangana, India.

⁴Microbial Ecology Laboratory, Department of Biochemistry, University College of Science, Mahatma Gandhi University, Anneparthy, Yellareddygudem (PO), Nalgonda, Telangana, India.

^{*}Correspondence: thirumala_21@yahoo.com

INTRODUCTION

Vanillin (4-hydroxy-3-methoxy benzaldehyde) & its related compounds have significant economic significance due to its rising demand and cost. It has numerous applications in the pharmaceutical, fragrance and food industries. Vanillin is frequently made using environmentally harmful methods and also involves no substrate selectivity. Even though direct extraction was used for natural vanillin production from vanilla costs more, it is nevertheless sought after on the global market.¹ There are numerous issues with vanilla extraction, including low concentration, which raises the cost of the extraction method.²,³

Microbial manufacturing is a potential and developing field for the creation of natural flavors. Recent advances have been explored in microbial Vanillin production through genetic and metabolic engineering methods.⁴

Consequently, using microbes which can bioconvert "cheap" substrate such as FA (trans-4-hydroxy-3-methoxycinnamic acid) to economically important Vanillin is an alternative cost-effective method.⁵

Studies on the microbial bioproduction of Vanillin have shown that several species of, *Streptomyces* and *Amycolatopsis* sp. ATCC 39116 are particularly effective at converting hydroxycinnamic acids specially FA into Vanillin.⁶

One of the most prevalent phenolic compounds is FA, which is found in the lignocellulose, plant cell walls, and agricultural plants' cell walls like grain, corn and sugar crops. According to some reports, a feruloyl-CoA synthetase first converts FA to feruloyl-CoA, and after that the CoA thioester is hydrated then cleaves into Vanillin & acetyl-CoA by an aldolase/enoyl-CoA hydratase. 99

Different microorganisms have varied capacity of biotransformation of FA to vanillic acid (VA) /Vanillin or both. Halomonas elognata strains quickly transformed FA to VA at 80% yield. However, in these studies, Vanillin was not formed which has considerably a greater value than VA. 16,17

Recombinant *Pseudomonas fluorescens* BF13 yielded of Vanillin and VA, 81% and 16%, respectively, in 3 L stirred tank reactor with incubation of 24 h. 18 Whereas, recombinant *Pseudomonas putida* strain KT2440 yield of Vanillin

was 86% in 3 h.¹⁹ *Halomonas* sp. B15 yielded FA biotransformation of 36.5% with production of VA, 365 mg/L after 31 h at 100 g NaCl/L.¹

Only a small number of microorganisms have so far been discovered which can bioconvert FA to Vanillin with increased yield. The main objective of this study was to isolate and identify the bacterial strain which can efficiently bioconvert FA to Vanillin.

MATERIALS AND METHODS

Chemicals

FA (99%) & Vanillin (99%) used in this research were bought from the Sigma-Aldrich company. Other chemicals purchased were Analytical Grade.

Enrichment culture & media

The sediment sample from Kogilvai Village, Warangal, Telangana was collected from the initial stages of paddy growing field in sterile polythene bags from around 5.0 cm below the soil surface and were stored in the laboratory until use. This sediment sample was then mixed with CaSO₄ and kept aside for about 20 days.²⁰ Isolation of microorganisms was performed using the serial dilution technique. Stock sediment solution was made by suspending 1 g of sediment sample in 100 ml sterile distilled water, then 1 ml of this suspension was inoculated in 9 mL sterile distilled water and mixed well. Then, 1 mL of this first dilution was used to prepare different dilutions using serial dilution method. Different dilutions (0.1 mL) of these were spread evenly on the sterilized Starch Casein Agar (SCA) and Oatmeal agar (OMA) plates. The SCA medium consists of (g/L): soluble starch (10), casein (0.3), K₂HPO₄ (2), NaCl (2), KNO₃ (2), CaCO₃ (0.02), MgSO₄.7H₂O (0.05), FeSO₄.7H₂O (0.01), agar (18) pH at 7.2.²¹ Oatmeal medium preparation: Oatmeal 20 g was cooked in 1000 ml distilled water for 20 min, filtered using cheese cloth and prepared the volume of filtrate to 1000 ml with distilled water. 20 g of agar was added, pH 7.2. Inoculated plates were incubated for 7 to 14 days at 30°C. The petri plates were noticed often for the growth of actinobacterial isolates. The isolates were randomly picked, purified & maintained on SCA and OMA plates at 4°C until further studies.

Screening of isolates converting ferulic acid to vanillin

The purified Actinobacterial isolates were plated on minimal medium supplemented with 0.2 μ m nylon filter sterilized Ferulic acid (500 mg/l) as only C source. After incubation of 48 h at 37°C, the colonies formed were later purified on the same media plates. Single bacterial colonies obtained were later maintained on the same media at 4°C until further research work.²² Minimal medium consists of (g/L): NH₄NO₃ (3.0) as a nitrogen source, KH₂PO₄ (1.0), Na₂PO₄ (1.0), NaCl (0.2), MgSO₄.7H₂O (0.2) CaCl₂ (0.05) and agar (18) pH was maintained at 7.

Effect of Ferulic acid on the growth of actinobacterial isolates

Purified isolates were grown in nutrient broth and incubated at 37°C for 24 h to prepare seed culture. Growth was measured at 600 nm in terms of OD. To determine the Ferulic acid biotransformation capacity of each isolate, seed culture (4% v/v) of each and every purified isolate was transferred to 30 ml of sterile basal media supplemented with 1g/L FA as C & energy source after 18 hrs of culture growth. The basal medium comprised of (g/L): Na₂HPO₄ (4), KH₂PO₄ (1), glucose (1), yeast extract (1), MgSO₄.7H₂O (0.2), NaCl (0.2), CaCl₂.2H₂O (0.05), pH at 7.2.²³ A control experiment was simultaneously performed by adding 1 g/L FA in the same medium without isolates. Ferulic acid was pre-dissolved in 37 g/L NaOH solution and sterilized with 0.2 µm syringe filters before use. Samples were withdrawn at random intervals (6, 24, 28, 30 and 48 hrs) to analyze the growth of culture. Cultures' growth was determined by taking spectrophotometric readings at 600 nm and were examined for FA bioconversion to Vanillin by TLC. The isolate with the quick and highest capacity of bioconversion of FA to Vanillin was later selected for further work.

FA to Vanillin bioconversion analysis

FA to Vanillin bioconversion by nine actinobacterial isolates was checked initially by TLC. Methylene chloride, methanol, and anhydrous acetic acid (98.5:1:0.5, v/v/v) were used as the mobile phase to create the chromatograms in a vertical glass chamber.²⁴ Ideal chamber

saturation time of 30 minutes at room temperature $(25 + 2^{\circ}C)$ was used for the mobile phase.

After chromatograms were developed, the TLC plates were dried & the components were observed under UV irradiation at 254 nm. Every analysis was performed in duplicate. After that, Vanillin concentration in the sample converted by isolate, S1 (with quick and high ferulic acid to vanillin transformation capacity) was analyzed by UV-spectrophotometry at 363 nm.²⁵ The Vanillin concentration present in the sample (isolate, S1 filtered culture broth) after 28 hrs of incubation period was extrapolated from Vanillin standard graph. Control experiment was performed by adding FA into the same media without isolate.

HPLC (Ultimate 3000, Thermo Scientific® with C18 column) analysis was also used to confirm the biocoversion of FA to Vanillin by isolate, S1. The analysis was done at 24 + 2°C or room temperature. The mobile phase was made up of 60% of Acetonitrile, 0.2% of Acetic acid and 38% water. Injection volume, 20 μl with flow rate of 1 mL/min. & detection wavelength, 280 nm.

Biochemical and Molecular characterization

The isolate with the quick and highest potential of bioconversion of FA to Vanillin was selected (isolate, S1) and characterized biochemically by testing different carbohydrates utilization using Hicarbo kit, Himedia.

The DNA of the isolate, S1 was extracted subjecting bacterial colony to incubation at high temperature (96°C). After DNA extraction, a PCR reaction was performed by employing two primers: 8F: 5'-AGA GTT TGA TCC TGG CTC AG 3' and 1525R: 5'-AAG GAG GTG ATC CAG CCG CA 3'. The PCR reaction was performed to amplify its 16S rDNA gene²⁶ according to the following protocol: Initial denaturation for 5 min at 95°C, then denaturation for 30 sec. at 95°C, primer annealing at 46.2°C for 1 min, and extension at 72°C for 1 min for 30 cycles. The last extension was performed at 72°C for 5 min and Amplicons were stored by keeping at 4°C. A PCR product with 1.5 kb size was eluted from agarose gel after PCR products underwent agarose gel electrophoresis, was purified and sequenced. 16S rDNA gene sequencing was outsourced. The EZ Biocloud public portal for data and analytics (https://www.ezbiocloud.net/apps) was utilized

to identify the isolate using its 16S rDNA gene sequence. The 16S rDNA gene sequence of the isolate, S1 was later submitted to NCBI GenBank to get its accession number. Phylogenetic analysis was done using MEGA 11 software.

RESULTS AND DISCUSSION

Isolates growing on FA as sole C source

Nine actinobacterial isolates (four on OMA (O1, O2, O3, O4) and five on SCA - (S1, S2, S3, S4, S5) were isolated and purified from the diluted sediment sample of Kogilvai Village, Warangal, Telangana on the minimal media with FA as sole C source. Growth of these nine actinobacterial isolates (optical density at 600 nm) before and after addition of FA in the media was shown in Figure 1.

Ferulic acid to vanillin bioconversion analysis

Ferulic acid biotransformation capacity of these nine isolates was determined using sterile basal medium supplemented with of 1 g/L Ferulic acid as substrate by drawing samples and analyzing for bioconversion capacity at different, random incubation times (6, 24, 28, 30 and 48 hrs) by TLC (Figure 2). Biotransformation of FA by some *Bacillus* sp. (*B. subtilis, B. coagulans* and *B. megaterium*) accumulated only minute concentrations of Vanillin and VA.²⁷⁻²⁹

Bioconversion process was shown by all the nine purified isolates with just incubation of 6 hrs after the addition of FA in the medium and continued till 28 hrs of incubation. After that, a substantial reduction in the quantity of Vanillin was seen as reaction time increased to 30 and 48 hrs, which was observed through analysis of TLC (Figure 2) and UV-spectrophotometry. HPLC confirmed the FA to Vanillin bioconversion by isolate, S1 (Figure 3). The current study has offered the first proof of *Limosilactobacillus* sp. strain, S1 with efficient FA biotransformation to Vanillin.

Research has been done on numerous bacteria, yeasts and filamentous fungi for the biotransformation of FA to Vanillin or VA. Low product concentrations were produced by the majority of the examined bioconversion processes. The yield of Vanillin formed by the biotransformation of FA has been extensively studied utilizing optimization techniques.

Table. Carbohydrates utilization test by Isolate, S1

No.	Carbohydrate	Result
	utilization	
1	Fructose	+
2	Xylose	-
3	Maltose	+
4	Lactose	+
5	Galactose	+
6	Dextrose	+
7	Trehalose	-
8	Raffinose	-
9	Melibiose	-
10	Glycerol	+
11	L-Arabinose	+
12	Mannose	-
13	Salicin	-
14	Sodium gluconate	-
15	Sucrose	+
16	Inulin	-
17	Inositol	-
18	Dulcitol	-
19	Sorbitol	-
20	Arabitol	-
21	Adonitol	-
22	Mannitol	-
23	Erythritol	-
24	α-Methyl-D-glucoside	-
25	Rhamnose	-
26	Xylitol	-
27	Melizitose	-
28	α-Methyl-D-Mannoside	<u> </u>
29	Cellobiose	-
30	Esculin	+
31	ONPG	+
32	D-Arabinose	-
33	Malonate	-
34	Citrate	-
35	Sorbose	_
55	22.232	

Note: - = negative test; + = positive test

Among the nine isolates tested in this report, four isolates, S1, S3, O3 and O4 showed increased FA bioconversion to Vanillin when compared to other five isolates with initial medium's pH 7. In a report, *Amycolatopsis* sp. began bioconverting FA present in broth after culturing the bacterial cells for 18 hours at various initial pH values between 7.0 and 8.5. It displayed an increased OD 600 value at pH 8.0 during the transformation than that at increased or decreased pH values.⁶

Among these four isolates (S1, S3, O3 and O4), optimal bioconversion was shown by isolate, S1 after the addition of FA in the medium. It was

converting 1 g/L FA in the medium to 140 mg/L Vanillin with just 28 hrs of incubation (Figure 4) with 150 rpm as analyzed by UV-spectrophometry and confirmed by HPLC (Figure 3).

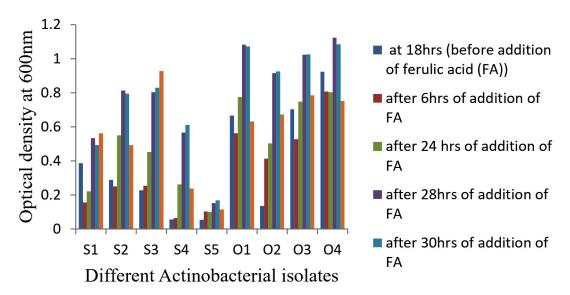


Figure 1. Growth (OD) of nine Actinobacterial isolates before and after addition of FA in the media

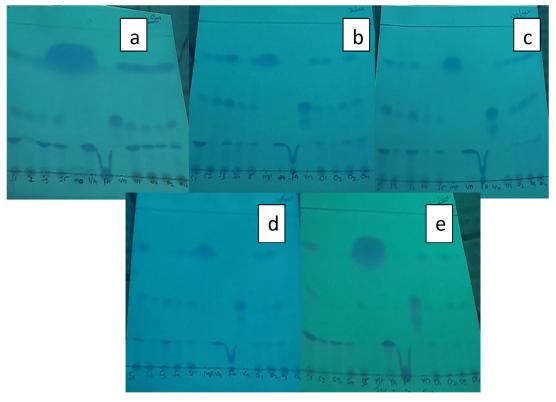


Figure 2. TLC of nine Actinobacteria with Bioconversion capacity of FA to Vanillin at different incubation periods

Biochemical and molecular characterization

Biochemical characterization (particularly carbohydrates utilization) of isolate S1 showed, it could utilize Lactose, Maltose, Glycerol, Fructose, Galactose, Sucrose, Dextrose, L-Arabinose, ONPG and Esculin. Whereas, other carbohydrates mentioned in Table were not utilized.

16S rDNA gene sequence of the isolate, S1 was amplified with PCR and the PCR product was subjected to sequencing. The 16S rDNA gene sequence of isolate, S1 was compared with other 16S rDNA gene sequences in EZ Biocloud public data and analytics portal. 16S rDNA sequence of isolate, S1 showed 98.27% similarity

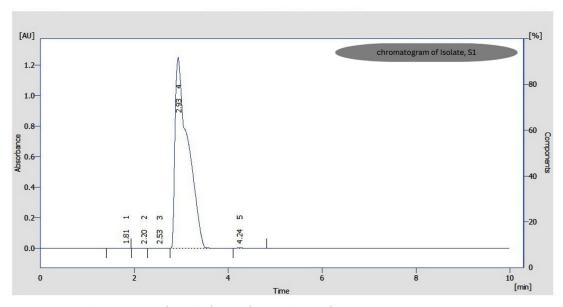


Figure 3. HPLC Chromatogram of Vanillin formed from FA biotransformation by Isolate, S1

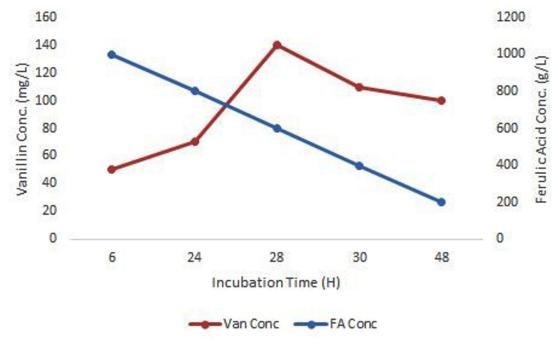


Figure 4. Biotransformation of FA to Vanillin by isolate, S1

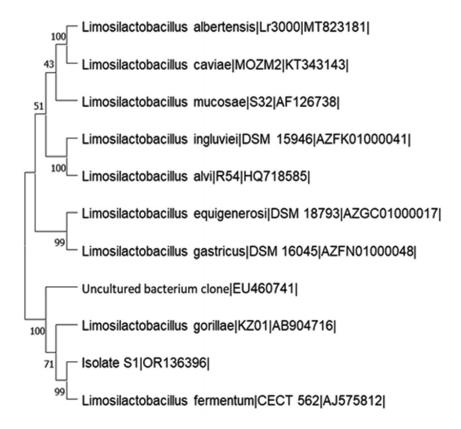


Figure 5. Neighbour-Joining Tree of the Isolate, S1 and its closely related type strains

to Limosilactobacillus fermentum CECT 562 with completeness of 96.7% and was identified as Limosilactobacillus sp. 16S rDNA gene sequence of this isolate, S1 was submitted to NCBI GenBank and its accession number was OR136396. The Neighbor-Joining tree was constructed to study phylogenetic analysis using MEGA 11 software (Figure 5).

CONCLUSION

All nine Actinobacteria isolated from the sediment sample of of Kogilvai Village, Warangal, Telangana, have the capacity of FA biotransformation to Vanillin. But, isolate S1 has the quick and potential conversion capacity of producing 140 mg/L (analyzed by HPLC) Vanillin with just 28 hours of incubation, after supplementation of 1 g/L FA in the medium at 37°C with 150 rpm. This study proved that the

Limosilactobacillus sp. strain, S1 has efficient FA biotransformation to Vanillin. Further research is required in this regard to determine the isolate's optimal conditions for the FA to Vanillin bioconversion process so that it can be applied in industrial setups for the commercial bioconversion process.

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None.

CONFLICT OF INTEREST

The authors declare that there is no conflict of interest.

AUTHORS' CONTRIBUTION

All authors listed have made a substantial, direct and intellectual contribution to the work, and approved it for publication.

FUNDING

None.

DATA AVAILABILITY

All datasets generated or analyzed during this study are included in the manuscript.

ETHICS STATEMENT

Not applicable.

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