

## Catalytic Zone, A Novel Screening Approach for the Production of Extracellular Tyrosinase by *Streptomyces vinaceusdrappus* DSV5

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An investigation was carried out to search potential actinobacteria for tyrosinase activity from different environmental samples. Detection of a typical catalytic zone developed on Tyrosine Gelatin Beef extract (TGB) agar indicating the tyrosinase activity by actinobacteria is a unique approach, which was never reported earlier. A total of five isolates of actinobacteria obtained from soils (DSV15 and DPG2) of Gulbarga India and marine sediments (DSV5) from Goa India were proved to be promising during primary and secondary screening for tyrosinase activity, based on the degree of catalytic zone and also intensity of color. Thus, observation of catalytic zone in addition to the intensity of color is a novel approach for the detection of tyrosinase activity. TGB medium has been proved as most suitable (high intensity of color and maximum catalytic zone) for tyrosinase activity, from among eight media, including two conventional and seven modified media, assessed during primary and secondary screening. An isolate of actinobacterium DSV5 has been proved as most efficient with the maximum production of tyrosinase in both Starch Tyrosine mineral ( $160 \pm 0.2$ U/ml) and TGB ( $175 \pm 0.2$ U/ml) media. The most potential isolate DSV5 was confirmed as *Streptomyces vinaceusdrappus* based on the molecular characterization by 16S rRNA analysis.

**Key words:** Streptomyces, Tyrosinase, Screening, Pigmentation, Catalytic zone, Production.

Tyrosinase (EC: 1.14.18.1) is an important copper containing high redox potential enzyme that catalyzes various phenolic substrates. It catalyzes the oxy hydroxylation of monophenols (Monophenolase activity) and the oxidation of oxy diphenols to reactive o-quinones (Diphenolase activity), using molecular oxygen<sup>1</sup>. The o-quinones undergo non enzymatic reactions with various nucleophiles producing an intermediate, which generates dark brown pigments<sup>2</sup>

Microbial tyrosinases are very prominent and attractive enzymes due to their several biotechnological and industrial applications. Tyrosinase has been employed as biosensor for the detection of various phenolic constituents from polluted soils and water samples<sup>3</sup>. It is also used as probe to sense the level of phenolic constituents in media during large scale/industrial fermentations<sup>4,5</sup>. Immobilization of antibody was achieved via tyrosinase catalyzed reactions<sup>6</sup>. Specific release of drug was targeted for melanoma by tyrosinase activity. Tyrosinase was used as an antigen to detect melanoma<sup>7,8,9</sup>. It is used as a biosensor for the detection of herbicides from agricultural soil<sup>10</sup>. It is also used as indicator for the potability of water by analyzing a phenolic compound released from colliforms in

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water<sup>11</sup>. Tyrosinases are found in both prokaryotes and eukaryotic microbes, invertebrates, plants and mammals<sup>12,13</sup>. The most extensively investigated tyrosinases are however reported from fungi, actinomycetes and bacteria. The predominant commercial tyrosinase was reported from *Agaricus bisporus*<sup>14,15</sup>, *Neurospora crassa*<sup>16</sup>, *Acremonium rutilum*<sup>17</sup>, *Agaricus oryzae*<sup>18</sup>, *Lentinus edodes*<sup>19</sup>, and *Pycnoporus sanguineus*<sup>20</sup>, *Streptomyces antibioticus*<sup>21</sup>, *Streptomyces glaucescens*<sup>22</sup>, *Streptomyces michiganensis* DSM 40015<sup>23</sup>, *Streptomyces griseus*<sup>24</sup>, *Streptomyces* sp. KY-453<sup>25</sup>, *Bacillus thuringiensis*<sup>26</sup> were also reported and known for the synthesis of tyrosinase.

The information on the tyrosinase structure and exact reaction mechanisms is limited. Some reasons for these deficits are difficulties in purification of sufficiently high amounts of tyrosinases from eukaryotic sources due to low enzyme concentrations, contamination with pigments, occurrence of isoenzymes or post translational modifications<sup>27</sup>. Despite of intensive screening investigations for the exploration of microorganisms for the synthesis of tyrosinase, no rapid and reliable methods are in practice. The development of dark brown color in the medium as visual observation, indicating the formation of melanin is the sole conventional method<sup>28</sup>, followed for the detection of tyrosinase activity by microorganisms. An investigation was carried out employing various media for the detection of tyrosinase activity by different approach, in addition to the visual observation of dark brown color, aiming at the exploration of actinobacteria for the enhanced production of tyrosinase.

## MATERIALS AND METHODS

### Collection of samples and isolation of actinobacteria

Soil samples from different ecological habitats such as agricultural fields, barren land, garden, graveyard, limestone quarry and tree bark scraping were collected<sup>29</sup>, in sterile polythene bags. Sediment from mangrove near Salim Ali Bird Sanctuary, Goa, India was also collected<sup>30</sup>. All the collected samples were processed aseptically and pretreated<sup>31</sup>. Actinobacteria were isolated from different samples on starch casein agar (SCA)<sup>32</sup>, and starch tyrosine mineral agar (STMA) by pour plate culture method<sup>33</sup>. The typical colonies of

actinobacteria were identified based on the standard colony and microscopic features<sup>34,35</sup>.

### Screening of actinobacteria

The identified prominent isolates of actinobacteria were subjected to primary, secondary and tertiary screening for tyrosinase activity. The selected isolates were screened for the synthesis of tyrosinase on STMA, where casein was replaced with tyrosine in SCA and Tyrosine-gelatin-beef extract agar (TGBA)<sup>36</sup>, by streak plate culture method<sup>37</sup>. Further, seven modified media including starch-tyrosine-mineral agar (STMA); tyrosine-beef extract agar (TBA); Tyrosine-gelatin agar (TGA); starch-tyrosine agar (STA); starch-tyrosine-beef extract agar (STBA); starch-tyrosine-gelatin agar (STGA) and starch-tyrosine-gelatin-beef extract agar (STGBA) were formulated based on the permutation combination of different ingredients of both conventional media (SCA and TGBA) and examined for the synthesis of tyrosinase. The potential isolates were employed to produce tyrosinase in selected media.

### Production and assay of tyrosinase

Production of tyrosinase in a batchwise bioprocess (100ml media in 250ml Erlenmeyer's flask) employing STM and TGB media was carried out as per the procedure prescribed by Konard lerch and Leopold ettlinger (1972)<sup>22</sup>. The quantity of tyrosinase produced was determined by measuring the dopachrome at 475 nm<sup>38</sup>, employing molar extinction co-efficient<sup>39</sup>. Standard reaction mixture containing 0.5 ml 4 mM L-dopa, 0.5 ml 0.1 M sodium phosphate buffer (pH 6.8) and enzyme extract in a total volume of 3 ml, mixture brought to 40°C in BOD incubator for of 10 min. The absorbance at 475 nm was monitored continuously for 3 min in (Systronics) 2201 UV-VIS spectrophotometer. One unit of tyrosinase activity was referred as the amount of enzyme required to catalyze 1 µmol of L-dopa per minute under the above conditions, which was calculated using the molar extinction coefficient of dopachrome 3600 M<sup>-1</sup> cm<sup>-1</sup><sup>39</sup>.

### Molecular characterization of the potential isolate

The potential isolate was identified and characterized by the standard colony characters<sup>34</sup>, microscopic features and biochemical/physiological properties<sup>35</sup>. In addition, the 16S rDNA sequence of strain was determined. The strain was cultivated on starch casein agar (SCA)

until sporulation and then the spores were collected. The genomic DNA of *Streptomyces* isolate DSV5 was extracted as prescribed by Rintala et al. (2001)<sup>40</sup> and further purified using Wizard DNA Clean-Up System A7280 (Promega, Madison, Wisconsin, USA) according to the manufacturer's instructions. The 16S rRNA gene was amplified by the polymerase chain reaction (PCR) using 50 µl reaction mixture with 1 µl of template DNA, 25 pmol of primers pA and pH (Oligomer, Helsinki, Finland)<sup>41</sup>, 10 nmol of each deoxynucleoside triphosphate (Finnzymes, Espoo, Finland), 5 µl of 10x Phusion HF buffer, 3 % dimethylsulphoxide, and 1.0 U of Phusion DNA Polymerase (Finnzymes, Espoo, Finland). The PCR conditions in DNA Engine DY AD Peltier Thermal Cycler (Bio-Rad, Hercules, CA, USA) were as follows: heat denaturation at 98 °C for 30 s, 30 amplification cycles of denaturation at 98 °C for 10 s, primer annealing at 61 °C for 30 s, and primer extension at 72 °C for 45 s. In the end of amplifications 72 °C was maintained for 10 min, followed by cooling to 4 °C. The 16S rRNA gene was sequenced in duplicate using the universal primers pE (forward and reverse), pB forward and pD forward<sup>41</sup>. The sequencings were done in the DNA Sequencing Laboratory (Institute of Biotechnology, University of Helsinki, Helsinki, Finland) as a purchased service. Sequence analysis was carried out at the Department of Biotechnology, University of Helsinki, Finland and submitted (Genbank ID: JX050266) to NCBI. BLAST search comparison was made against the Genbank databases and the related strains were selected for alignment by CLUSTAL X program<sup>42</sup>.

The evolutionary history was inferred using the Neighbor-Joining method and the evolutionary distances were computed using the Maximum Composite Likelihood method<sup>43</sup>. Phylogenetic analyses were carried out employing software MEGA4.

## RESULTS AND DISCUSSION

Exploration of microorganisms from various ecological habitats constituting unique niche is an ever continuing phenomenon, searching for the potential isolates targeting variety bioactive molecules. Few novel isolates of *Streptomyces* have been reported<sup>44,45</sup>, from the regional soil samples and were explored for different enzymes<sup>46,47</sup>. Though, various stages of isolation and screening of potential microorganisms have been well recorded<sup>48</sup>. Much more rapid and accurate methods are warranted, atleast for certain bioactive molecules. More accurate and different approach for the screening of actinobacteria from various habitats for the detection of tyrosinase activity was investigated. The *Streptomyces* sp. DSV5 was isolated from Tree bark sample, Chandarki, India.

### Primary screening

Two conventional media (SCA and TGBA) and seven modified media (starch-tyrosine-mineral agar (STMA); tyrosine-beef extract agar (TBA); Tyrosine-gelatin agar (TGA); starch-tyrosine agar (STA); starch-tyrosine-beef extract agar (STBA); starch-tyrosine-gelatin agar (STGA) and starch-tyrosine-gelatin-beef extract agar (STGBA) were employed (Table 1) in the primary

**Table 1.** Composition of different media employed

Composition (g)	Different media								
	SCA	STMA	TGBA	TBA	TGA	STA	STBA	STGA	STGBA
Starch	10.0	10.0	-	-	-	5.0	5.0	5.0	5.0
L-Tyrosine	-	4.0	5.0	5.0	5.0	3.0	5.0	5.0	5.0
Beef extract	-	-	3.0	3.0	-	-	3.0	-	3.0
Gelatin	-	-	5.0	-	5.0	-	-	5.0	5.0
Casein	0.03	-	-	-	-	-	-	-	-
K <sub>2</sub> HPO <sub>4</sub>	2.0	2.0	-	-	-	-	-	-	-
KNO <sub>3</sub>	2.0	-	-	-	-	-	-	-	-
NaCl	1.0	1.0	-	-	-	-	-	-	-
MgSO <sub>4</sub> ·7H <sub>2</sub> O	0.05	0.05	-	-	-	-	-	-	-
CaCO <sub>3</sub>	0.02	0.02	-	-	-	-	-	-	-
FeSO <sub>4</sub> ·7H <sub>2</sub> O	10.0	10.0	-	-	-	-	-	-	-

stage for the isolation and screening of actinobacteria aiming at the synthesis of tyrosinase.

Starch casein agar is an exclusive medium prescribed by Kuster and Williams (1964)<sup>32</sup> for the isolation of actinobacteria. Casein was replaced by tyrosine and used for the isolation and screening of actinobacteria for the synthesis of tyrosinase. In addition, TGB agar<sup>36</sup>, was also used for the said purpose. Degree of coloration or intensity of color (yellowish, brownish and blackish) was an exclusive conventional method prescribed<sup>28,49</sup>, and

followed for the detection of microorganisms synthesizing tyrosinase. The actinobacteria isolated and screened from different samples for the synthesis of tyrosinase based on intensity of color are as presented in Table 2. The degree of intensity of color was recorded as poor (+, yellowish), moderate (++, brownish) and higher (+++, blackish) and correlated for the level of synthesis of tyrosinase. TGB agar was never reported to be employed earlier, to the best of our knowledge, for the screening of actinobacteria to detect the tyrosinase activity. However, this

**Table 2.** Primary screening of soil actinobacteria for the synthesis of tyrosinase in different media based on intensity of color

Soil samples	Tyrosinase activity graded (+/++/+++)* in different media							
	STMA	TGBA	TBA	TGA	STA	STBA	STGA	STGBA
Agricultural field ( red)	++	++	+	+	++	++	+	+
Agricultural field (black)	++	++	+	++	+	++	+	+
Barren land	++	++	+	+	+	++	+	+
Fresh water	++	+++	+	++	++	+	+	+
Garden	++	++	+	+	+	+	+	+
Grass land	++	++	+	+	+	+	+	+
Grave yard	++	+++	+	++	++	+	+	++
Limestone quarry	+	+	+	+	+	+	+	+
Mangrove	+++	+++	+	++	++	++	+	++
Rhizosphere	++	++	+	+	+	+	+	+
Tree bark	++	+++	+	++	++	+	+	++
Wood ash	+	++	+	+	+	+	+	+

\* +: poor (yellowish); ++: moderate (brownish); +++: higher (blackish)

**Table 3.** Primary screening of soil actinobacteria for the synthesis of tyrosinase in different media based on catalytic zone

Soil samples	Tyrosinase activity measured (mm) in different media (Average $\pm$ SD)							
	STMA	TGBA	TBA	TGA	STA	STBA	STGA	STGBA
Agricultural field ( red)	0	8 $\pm$ 1.0	1 $\pm$ 0.2	3 $\pm$ 0.1	5 $\pm$ 2.0	4 $\pm$ 1.0	1 $\pm$ 0	2 $\pm$ 0.1
Agricultural field (black)	0	9 $\pm$ 0.4	2 $\pm$ 0.2	5 $\pm$ 0.2	5 $\pm$ 0.5	6 $\pm$ 0.1	4 $\pm$ 0.2	4 $\pm$ 0.2
Barren land	0	8 $\pm$ 1.0	2 $\pm$ 1.0	2 $\pm$ 0.1	3 $\pm$ 0.4	3 $\pm$ 0.1	1 $\pm$ 0.4	4 $\pm$ 1.0
Fresh water	0	9 $\pm$ 0.3	1 $\pm$ 0	4 $\pm$ 0.4	7 $\pm$ 0.2	5 $\pm$ 0.2	2 $\pm$ 0	1 $\pm$ 0.1
Garden	0	8 $\pm$ 1.1	2 $\pm$ 0	2 $\pm$ 0.4	2 $\pm$ 0.2	1 $\pm$ 0.4	2 $\pm$ 0.4	1 $\pm$ 0.2
Grass land	0	7 $\pm$ 0.6	1 $\pm$ 0.4	4 $\pm$ 0.1	3 $\pm$ 0.2	3 $\pm$ 0.2	3 $\pm$ 0.1	4 $\pm$ 0.4
Grave yard	0	15 $\pm$ 0.3	2 $\pm$ 0.2	2 $\pm$ 0.4	7 $\pm$ 1.0	5 $\pm$ 0.1	0 $\pm$ 0	1 $\pm$ 0.2
Limestone quarry	0	3 $\pm$ 1.0	1 $\pm$ 0.3	1 $\pm$ 0.2	1 $\pm$ 0	1 $\pm$ 2.0	1 $\pm$ 0	1 $\pm$ 0.1
Mangrove	0	25 $\pm$ 0.5	3 $\pm$ 0.2	7 $\pm$ 0.2	5 $\pm$ 0	4 $\pm$ 1.0	1 $\pm$ 0.1	3 $\pm$ 0.6
Rhizosphere	0	7 $\pm$ 0.1	1 $\pm$ 0.3	1 $\pm$ 0.4	3 $\pm$ 0.1	5 $\pm$ 2.0	2 $\pm$ 0	3 $\pm$ 0
Tree bark	0	20 $\pm$ 0.4	1 $\pm$ 0.2	4 $\pm$ 1.2	10 $\pm$ 0.1	5 $\pm$ 0.2	2 $\pm$ 0	7 $\pm$ 0.5
Wood ash	0	5 $\pm$ 1.0	2 $\pm$ 0.4	3 $\pm$ 0.4	2 $\pm$ 2.0	2 $\pm$ 0.4	1 $\pm$ 0.1	3 $\pm$ 0.1

medium<sup>36</sup> was recorded, to be used for the differentiation of *Streptomyces* isolates, but never reported the catalytic zone for the tyrosinase activity. Surprisingly, this medium exhibited catalytic zone (a clear zone around the colony catalyzing the tyrosine), indicating the synthesis of tyrosinase. Table 3 exhibit the degree of catalytic zone on different media (TGBA and modified media) by the isolates of actinobacteria obtained from various soil samples. The catalytic zone developed was measured (mm) and considered as novel criteria for the selection of potential isolate based on the degree of catalytic zone. Seven different modified media were also examined for the synthesis of tyrosinase by actinobacteria obtained from different soil samples. Tyrosinase is an induced enzyme<sup>50</sup>,

and tyrosine is an amino acid being the exclusive protein source, significantly revealing the tyrosinase activity. The typical colonies of actinobacteria isolated from different samples showing the maximum tyrosinase activity based on the intensity of color and catalytic zone were selected and examined further for the confirmation of tyrosinase synthesis.

#### Secondary screening

Three selected test isolates of actinobacteria (DSV5, DSV15 and DPG2) were subjected for secondary screening to confirm the synthesis of tyrosinase on STM agar based on intensity of color and catalytic zone on TGB agar (Table 4). It is interesting to note that, the graded intensity of color on STM agar can be matched

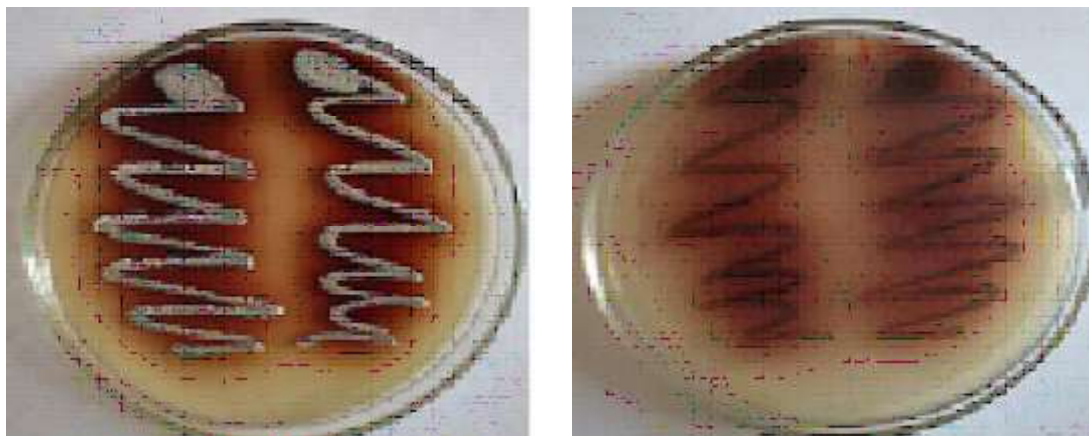
**Table 4.** Secondary screening of *Streptomyces* for the synthesis of tyrosinase on STM agar based on intensity of color and on TGB agar based on catalytic zone

Incubation Period (hrs)	Secondary Screening					
	Tyrosinase activity (intensity of colour) by selected isolates graded (+/++/+++)* on STM Agar			Tyrosinase activity (zone of catalysis) by selected isolates measured (mm) on TGB Agar(Average $\pm$ SD )		
	DSV5	DSV15	DPG2	DSV5	DSV15	DPG2
24	+	+	+	9 $\pm$ 0.57	7 $\pm$ 0.09	6 $\pm$ 0.2
48	++	++	+	15 $\pm$ 0.02	12 $\pm$ 0.5	7 $\pm$ 1.19
72	+++	++	+	20 $\pm$ 0.5	18 $\pm$ 1.0	12 $\pm$ 0.25
96	+++	++	++	25 $\pm$ 0.7	22 $\pm$ 1.0	15 $\pm$ 0.1
120	+++	+++	+++	30 $\pm$ 0.18	24 $\pm$ 0.57	18 $\pm$ 1.52
144	+++	+++	+++	32 $\pm$ 0.31	26 $\pm$ 1.0	20 $\pm$ 1.89

\* Yellowish = +; brownish = ++; blackish = +++

**Table 5.** Tertiary screening of the selected isolates of streptomyces for the production of tyrosinase in STM and TGB media

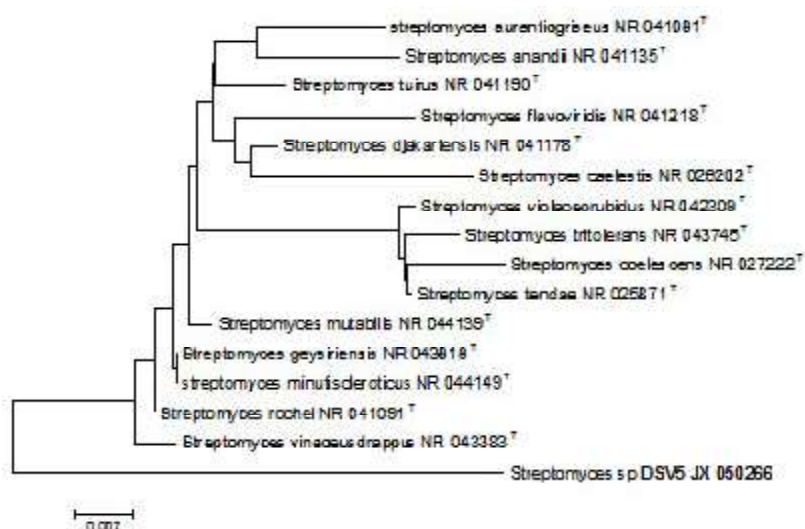
Incubation Period (hrs)	Tertiary Screening					
	Tyrosinase activity(U) in STM medium (Average $\pm$ SD )			Tyrosinase activity(U) in TGB medium (Average $\pm$ SD )		
	DSV5	DSV15	DPG2	DSV5	DSV15	DPG2
24	28 $\pm$ 1.0	23 $\pm$ 0.5	25 $\pm$ 0.2	25 $\pm$ 1.0	21 $\pm$ 0.4	24 $\pm$ 0
48	62 $\pm$ 0.2	47 $\pm$ 0.6	54 $\pm$ 0.4	53 $\pm$ 0.6	40 $\pm$ 0.6	58 $\pm$ 0
72	101 $\pm$ 0.1	93 $\pm$ 0.4	78 $\pm$ 0.4	105 $\pm$ 0.4	98 $\pm$ 0.2	102 $\pm$ 0.2
96	138 $\pm$ 0.4	124 $\pm$ 0.4	98 $\pm$ 0.2	140 $\pm$ 0.6	122 $\pm$ 0.2	125 $\pm$ 0.5
120	160 $\pm$ 0.2	155 $\pm$ 0.2	140 $\pm$ 0	175 $\pm$ 0.2	165 $\pm$ 0.4	160 $\pm$ 0.2
144	152 $\pm$ 0.2	140 $\pm$ 0.1	121 $\pm$ 0.2	172 $\pm$ 0.4	153 $\pm$ 0.4	158 $\pm$ 0.2



**Fig.1.** Synthesis of tyrosinase by the potential isolate of *Streptomyces* DSV5 on STM agar based on the intensity of color.



**Fig. 2.** Synthesis of tyrosinase by the potential isolate of *Streptomyces* DSV5 on TGB agar based on the intensity of color and catalytic zone.



**Fig.3.** Phylogenetic tree showing the systematic position of the potential isolate of *Streptomyces* - DSV5.

with the catalytic zone on TGB agar. However, incidentally TGB agar exhibits not only the catalytic zone but also the intensity of color similar to the STM agar. This ensures the accurate matching of the intensity of color and catalytic zone. Fig. 1 and 2 exhibiting the high intensity of color on STM agar and catalytic zone along with intensity of color on TGB agar by DSV5 significantly reveals it as the most potential isolate for the synthesis of tyrosinase.

### Tertiary screening

The qualitative attribute for the synthesis of tyrosinase shown during the primary and secondary screening was assessed and confirmed, by subjecting the selected isolates of actinobacteria to tertiary screening. Tyrosinase activity is generally measured by either determining the consumption of oxygen during the reaction or spectrophotometrically by following the increase of absorbance at 475 nm due to dopachrome formation<sup>51</sup>. The quantitative production of tyrosinase in STM and TGB media by all the three test isolates of actinobacteria are as shown in Table 5. In tertiary screening also, isolate DSV 5 has been proved as most potential isolate in both STM and TGB media. However TGB medium has resulted in more production ( $175 \pm 0.2$ U) of tyrosinase than STM ( $160 \pm 0.2$ U) medium. Thus, the intensity of color (a conventional approach) and the catalytic zone can be perfectly correlated for the synthesis of tyrosinase. This significantly reveals that, the catalytic zone can be an accurate and novel approach to screen actinobacteria for the synthesis of tyrosinase, which may be applied in general for screening of any microorganisms for the synthesis of tyrosinase. Recent available literature reveals 141 U/ml intracellular tyrosinase by *Acremonium rutilum*<sup>17</sup>, 94 U/ml extracellular tyrosinase by *Bacillus thuringiensis*<sup>26</sup>, and 72 U/ml extracellular tyrosinase by actinomycetes<sup>52</sup>. Highly varied range of tyrosinase activities from Streptomyces were reported by several researchers. Konard lerch and Leopold ettlinger (1972)<sup>22</sup>, reported 24400 units by *Streptomyces glaucescens*, following oxygen consumption method using oxygen electrode and Stephan Phillip *et al.* (1991)<sup>23</sup> reported 22700 nkat units by *Streptomyces michiganensis*, employing YSI Biological Oxygen Monitor model 5300. Various methods with different principles, type of substrates and their concentrations employed in

the process of enzyme assay would result such higher values. Molar extinction coefficient of Dopachrome is not an uncommon practice being followed by majority of researchers to determine tyrosinase activity. The potential isolate DSV5 detected in the present investigation was proved to be highly efficient with production of  $160 \pm 0.2$  U/ml (STM media) and  $175 \pm 0.2$  U/ml (TGB medium) extracellular tyrosinase.

Actinobacteria were being analyzed<sup>53</sup>, at various levels to gain information suitable for constructing databases and effecting identification. The highest level is the genome and its direct expression as RNA. Sequence analysis of various genes provides stable classification and accurate identification, which has become the cornerstone of modern phylogenetic taxonomy. The potential isolate in the present investigation was assigned to the genus *Streptomyces* based on colony characters and microscopic features. Further, the isolate was characterized by 16S rRNA analysis and has been confirmed (Fig. 3.) as *Streptomyces vinaceusdrappus* NR 043383<sup>T</sup> based on the phylogenetic illustration.

### CONCLUSIONS

Search for newer organisms with higher level production of tyrosinase is ever continuous phenomenon and essential for commercial viability. Screening of potential isolates from natural habitats is utmost important. Detection of catalytic zone, a novel approach can be a more reliable, accurate and simple method for rapid screening of microorganisms from natural sources for tyrosinase activity. The most potential isolate DSV5 *Streptomyces vinaceusdrappus* can be explored further for the large scale industrial production of extracellular tyrosinase with a greater commercial value. TGB medium may be developed further for the much enhanced production of extracellular tyrosinase.

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## REFERENCES

- Mayer, A.M. Polyphenol oxidases in plants – recent progress. *Phytochemistry*, 1987; **26**:11–20.
- Soler Rivas C., Jolivet S., Arpin N., Olivier, J.M., Wihers, H.J. Biochemical and physiological aspects of brown blotch disease of *Agaricus bisporus*. *FEMS Microbiol Rev.*, 1999; **23**:591–614.
- Streffer, K., Vijgenboom, E., Armand., Tepper, W.J.W., Makower, A., Frieder, W., Scheller., Gerard, W., Canters, Wollenberger, U. Determination of phenolic compounds using recombinant tyrosinase from *Streptomyces antibioticus*. *Analytica Chimica Acta*, 2001; **427**:201–210.
- Yaropolov, A.I., Kharybin, A.N., Emneus, J., Marko Varga, G., Gorton, L. Electrochemical properties of some copper-containing oxidases. *Bioelectrochem. Bioener.*, 1996; **40**:49–57.
- Mita, D.G., Attanasio, A., Arduini, F., Diano, N., Grano, V., Bencivenga, U., Rossi, S., Amine, A., Moscone, D. Enzymatic determination of BPA by means of tyrosinase immobilized on different carbon carriers. *Biosens. Bioelectron.*, 2007; **23**: 60–65.
- Ahmed, S.R., Lutes, A.T., Barbari, T.A. Specific capture of target proteins by oriented antibodies bound to tyrosinase-immobilized Protein A on a polyallylamine affinity membrane surface. *J Memb Sci.*, 2006; **282**(1-2), : 311–321.
- Morrison, M.E., Yagi, M.J., Cohen, G. In vitro studies of 2,4- dihydroxyphenylalanine, a prodrug targeted against malignant melanoma cells. *Proc Natl Acad Sci U S A.*, 1985; **82**:9, 2960–2964.
- Jordan. A.M., Khan, T.H., Malkin, H., Osborn, H.M., Photiou, A., Riley, P.A. Melanocyte-directed enzyme prodrug therapy (MDEPT): development of second generation prodrugs for targeted treatment of malignant melanoma. *Bioorg Med Chem.*, 2001; **9**(6); 1549–1558.
- Jordan, A.M., Khan, T.H., Osborn, H.M., Photiou, A., Riley, P.A. Melanocyte-directed enzyme prodrug therapy (MDEPT): development of a targeted treatment for malignant melanoma. *Bioorg Med Chem.*, 1999; **7**(9), 1775–1780.
- Xuejiang Wang., Ling Chen., Siqing Xia., Zhiliang Zhu., Jianfu Zhao., Jean-Marc Chovelon., Nicole Jaffrezic Renaul. Tyrosinase Biosensor Based on Interdigitated Electrodes for Herbicides Determination. *Int. J. Electrochem. Sci.*, 2006; **1**:55-61.
- Beatriz Serra, M., Dolores Morales., Jinbiao Zhang, A., Julio Reviejo., Elizabeth Hall, H., Jose Pingarron M. In-a-Day Electrochemical Detection of Coliforms in Drinking Water Using a Tyrosinase Composite Biosensor. *Anal. Chem.*, 2005; **77**(24): 8115–8121.
- Kwon, B.S., Haq, A.K., Pomerantz, S.H., Halaban, R. Isolation and sequence of a cDNA clone for human tyrosinase that map at the mouse c-albino locus. *Proc. Natl. Acad. Sci. U.S.A.*, 1987; **84**: 7473–7477.
- Kwon B.S., Wakulchik M., Haq AK., Halaban R., Kestler D. Sequence analysis of mouse tyrosinase cDNA and the effect of melanotropin on its gene expression. *Biochem. Biophys. Res. Commun.*, 1988; **153**:1301–1309.
- Seo, S.Y., Sharma, V.K., Sharma, N. Mushroom tyrosinase: Recent prospects. A review. *Journal of Agricultural and Food Chemistry.*, 2003; **51**(10), pp. 2837–2853.
- Selinheimo, E., Gasparetti, C., Mattinen, M., Steffensen, C.L., Buchert, J., Kruus, K. Comparison of substrate specificity of tyrosinases from *Trichoderma reesei* and *Agaricus bisporus*. *Enzyme and Microbial Technology*, 2009; **44**(1), pp. 1–10.
- Lerch, K. *Neurospora* tyrosinase: structural, spectroscopic and catalytic properties. *Mol Cell Biochem.*, 1983; **52**(2) :125–138.
- Krishnaveni. R., Vandana Rathod., Thakur, M.S., Neelgund, Y.F. Transformation of L-Tyrosine to L-Dopa by a Novel Fungus *Acremonium rutilum*, Under Submerged Fermentation. *Curr Microbiol.*, 2009; **58**:122–128.
- Wichers, H.J., Gerritse, Y.A., Chapelon, C.G.J. Tyrosinase isoforms from the fruitbodies of *Agaricus bisporus*. *Phytochemistry.*, 1996; **43**:333–337.
- Kanda, K., Sato, T., Ishii, S., Enei, H., Ejiri, S. Purification and properties of tyrosinase isozymes from gill of *Lentinus edodes* fruiting bodies. *Biosci Biotechnol Biochem.*, 1996; **60**:1273–1278.
- Halaouli, S., Asther, Mi., Kruus, K., Guo, L., Hamdi, M., Sigoillot, J.C., Asther, M., Lomascolo, A. Characterization of a new tyrosinase from *Pycnoporus* species with high potential for food technological applications. *J. Appl. Microbiol.*, 2005; **98**:332–343.
- Katz, E., Thompson, C.J., Hopwood, D.A. Cloning and expression of the tyrosinase gene from *Streptomyces antibioticus* in *Streptomyces lividans*. *J Gen Microbiol.*, 1983; **129**:2703–2714.
- Konrad, lerch., Leopold, Ettlinger. Purification and Characterization of a Tyrosinase from



- Streptomyces glaucescens*. *Eur. J. Biochem.*, 1972; **31**:427-437.
23. Stephan, Philipp., Thomas, Held., Hans., Kutzner, J. Purification and characterization of the tyrosinase of *Streptomyces michiganensis* DSM 40015. *J. Basic Microbiol.*, 1991; **31**, 4:293-300.
  24. Tadashi, Arai., Yuzuru, Mikami. Chromogenicity of *Streptomyces*. *Applied Microbiology.*, 1972; 402-406.
  25. Yoshimoto, T., Yamamoto, K., Tsuru, D. Extracellular Tyrosinase from *Streptomyces* Sp. KY-453: Purification and Some Enzymatic Properties. *J. Biochem.*, 1985; **97**:1747-1754.
  26. Liu, N., Zhang, T., Wang, Y.J., Huang, Y.P., Ou, J.H., Shen, P. A heat inducible tyrosinase with distinct properties from *Bacillus thuringiensis*. *Letters in Applied Microbiology.*, 2004; **39**:407-412.
  27. Harald, Claus., Heinz, Decker. Bacterial tyrosinases A review. *Systematic and Applied Microbiology.*, 2006; **29**:3-14.
  28. Taofeeq nurudeen, A., Donald Ahearn, G. Regulation of Melanin Production by *Cryptococcus Neoformans*. *Journal of Clinical Microbiology.*, 1979; 724-729.
  29. Grant, W.D., Mwatha, W.E., Jones, B.E. Alkaliphiles ecology, diversity and application. *FEMS Microbiol. Rev.*, 1990; **75**: 255-270.
  30. Kokare, C.R., Mahadik, K.R., Kadam, S.S., Chopade, B.A. Isolation of bioactive marine actinomycetes from sediments isolated from Goa and Maharashtra coastlines (west coast of India). *Indian Journal of Marine Sciences.*, 2004; **33**(3): 248-256.
  31. Hayakawa, M., Yoshida, Y., Iimura, Y. Selective isolation of bioactive soil actinomycetes belonging to the *Streptomyces violaceusniger* phenotypic cluster. *Journal of Applied Microbiology.*, 2004; **96**: 973-981.
  32. Küster, E., Williams, S.T. Starch-casein medium for isolation of *Streptomyces*. *Nature.*, 1964; **202**: 928-929.
  33. Treuhaft, M.W., Arden Jonest, M.P. Comparison of Methods for Isolation and Enumeration of Thermophilic Actinomycetes from Dust. *Journal of Clinical Microbiology.*, 1982; **16**(6): 995-999.
  34. Shirling, E.B., Gottlieb, D. Report of the International Streptomyces Project-five years of collaborative research, In H. Prauser(ed). The Actinomycetales. Gustav, Fischer. Verlag. Jena, 1970; 79-89.
  35. Gottlieb, D. An evaluation of criteria and procedures used in the description and characterization of Streptomyces: A cooperative study. *Appl. Microbiol.*, 1961; **9**:55-65.
  36. Ronald Atlas, M. Handbook of microbiological media. 4<sup>th</sup> Edition, CRC press, USA, 1947; 1862-1863.
  37. Lemos, M.L., Toranzo, A.E., Barja, J.L. Antibiotic activity of epiphytic bacteria isolated from intertidal seaweeds. *Journal of Microbial Ecology.*, 1985; **11**:149-163.
  38. Chiara Gasparetti., Emilia Nordlund., Janne Jänis., Johanna Buchert., Kristiina Kruus. Extracellular tyrosinase from the fungus *Trichoderma reesei* shows product inhibition and different inhibition mechanism from the intracellular tyrosinase from *Agaricus bisporus*. *Biochimica et Biophysica Acta.*, 2012; 1824: 598-607.
  39. Fling, M., Horowitz, N.H., Heinemann, S.F. The isolation and properties of crystalline tyrosinase from *Neurospora*. *J Biol Chem.*, 1963; **238**: 2045-2053.
  40. Rintala, H., Nevalainen, A., Suutari, M. PCR primers targeting the 16S rRNA gene for the specific detection of *Streptomyces*. *Molecular and Cellular Probes.*, 2001; **15**:337-347.
  41. Edwards, U., Rogall, T., Blöcker, H., Emde, M., Böttger E.C. Isolation and direct complete nucleotide determination of entire genes. Characterization of a gene coding for 16S ribosomal RNA. *Nucleic Acids Research.*, 1989; **17**, 7843-7853.
  42. Thompson, J.D., Gibson, T.J., Plewniak, F., Jeanmougin, F., Higgins, D.G. The Clustal X windows interface: flexible strategies for multiple sequence alignment aided by quality analysis tools. *Nucleic Acids Res.*, 1997; **24**: 4876-4882.
  43. Tamura, K., Nei, M., Kumar, S. Prospects for inferring very large phylogenies by using the neighbor-joining method. *Proc Natl Acad Sci.*, 2004; **101**:11030-11035.
  44. Dastager G Syed., Agsar Dayanand., Wen Jun Li., Li Hua Xu., Xin. Peng Tian., Chen Lin Jiang. *Streptomyces tritolerans* sp. nov, a novel actinomycetes isolated from soil in Karnataka, India. *Antonie Van Leeuwenhoek.*, 2007a; **92**: 391-397.
  45. Dastager G Syed., Wen Jun Li., Dayanand A., Mudgulkar B. Sulochana., Shu Ken Tang., Xin., Peng Tian., Xiao. Yang Zhi. *Streptomyces gulbargensis* sp. nov, isolated from soil in Karnataka, India. *Antonie Van Leeuwenhoek.*, 2007b; **91**: 99-104.
  46. Dastager G Syed., Jae Chan Lee., WenJun Li., ChangJin Kim., Dayanand Agsar. Production, characterization and application of keratinase from *Streptomyces gulbargensis*. *Bioresource Technology.*, 2008; **30**:1-4.

47. Dastager G Syed, Dayanand Agsar, Ashok Pandey. Production and partial purification of  $\alpha$  amylase from a novel isolate *Streptomyces gulbargensis*. *Journal of Industrial Microbial Biotechnology*, 2009; **36**:189-194.
48. Casida., Industrial Microbiology. Tata McGraw Hill., 1968; 55-63.
49. Livia Teixeira Duarte., Joyce Batista Tiba., Mariangela Fontes Santiago., Telma Alves Garcia., Maria Teresa Freitas Bara. Production and characterization of tyrosinase activity in *pyncoporus sanguineus* cct- 4518 crude extract. *Brazilian Journal of Microbiology*, 2012; 21-29.
50. Ikeda, K., Masujima, T., Sugiyama, M. Effects of methionine and  $\text{Cu}^{2+}$  on the expression of tyrosinase activity in *S. castaneoglobisporus*. *J. Biochem.*, 1996; **120**:1141–1145.
51. Greta Faccio., Kristiina Kruus., Markku Saloheimo., Linda Thony-Meyer. Bacterial tyrosinase and their applications, - a review. *Process Biochemistry.*, 2012; **47**:1749-1760.
52. Santosh Patil, Vandana Rathod, Ranganath E. Inducive effect of L-methionine in transformation of L-tyrosine to L-Dopa and tyrosinase production by *Streptomyces* sp. VRS9. *Indian Journal of Biotechnology*, 2012; **11**, 320-325.
53. Huo Hu., Hai-Peng Lin., Qingyi Xie., Lei Li., Xin-Qiang Xie., Ming Sun., Kui Hong. *Streptomyces shenzhenensis* sp. nov, a novel actinomycetes isolated from mangrove sediment. *Antonie van Leeuwenhoek*, 2011; **100**: 631-637.