

Isolation and Identification of Endophytic Bacteria from Day Flower of *Commelina benghalensis*

Archana S Injal^{1*}, Tejaswini C Mendke², Tara N Matnale³, Somnath Kshirsagar⁴

¹Department of Genetics, Walchand College of Art and Science, Solapur, India

²Department of Nanotechnology, Walchand College of Art and Science, Solapur, India

³Department of Biotechnology, Walchand College of Art and Science, Solapur, India

⁴Department of Zoology, K.N. Bhise College of arts and science, Solapur, India

Research Article

Received date: 02/01/2020

Accepted date: 13/01/2020

Published date: 20/01/2020

*For Correspondence

Archana S Injal, Department of Genetics,
Walchand college Of Art and Science,
Solapur, India.

E-mail: injalarchana29@gmail.com

Keywords: *Commelina benghalensis*,
Dayflower, Endophytic bacteria, Affinity
chromatography, *Micrococcus* species

ABSTRACT

Objective: Endophytes are nothing but microflora of plant. Each plant has its own endophytic flora. *C. benghalensis* is weed. Day flower is unique feature of *Commelina benghalensis*. Which is relatively difficult to control by herbicide? So, objective of this work is to isolate and identify endophytic bacteria from dayflower of *Commelina benghalensis* and finding role of entophyte in herbicide resistivity.

Methods: The plant of *C. benghalensis* was collected from nearby Solapur area and re-cultivated in college botanical garden. Dayflowers were collected from underground roots of *C. benghalensis*. Separated dayflowers were brought in laboratory and wash thoroughly by tap water. Surface sterilization was carried out by 70% alcohol, Tween 20, Sodium hypochloride and distilled water. Each dayflower was cut into two pieces from its central axis. These two cut pieces were separately inoculated on nutrient agar medium (cut portion facing toward medium) and incubate for 24 hours at 37 °C.

Result: After 24 hours of incubation, white bacterial growth was observed on nutrient agar medium exactly near surrounding area of cut portion of dayflower. Colonies were isolated and used for further biochemical tests. The Biochemical tests were performed on Vitek 2 biomérieux vitek 2 systems by using the identification software PIBWIN Version 2.0.0. On the basis of analysis of this system, it is finalized that the isolated bacterial strain was *Micrococcus* spp.

Conclusion: This resistivity to herbicides might be due to presence of *micrococcus* spp. in its dayflower. Because, *Micrococcus* actively participate in catabolic reactions by utilizing a wide range of unusual substrate like herbicides, pyridine, oil and chlorinated biphenyls.

INTRODUCTION

Bacteria and fungi which live inside plant cells without any harm to plant are called endophytes^[1]. Most plant species accommodate one, or more than one, kind of endophyte. Endophytes develop some associations with the host plant^[2-6]. These associations can be commensal or mutualistic but, sometime few bacteria and fungi live as parasite inside the plant cells^[7]. More work was done on endophytic fungi than bacterial endophytes^[8,9]. In this study, we have isolated endophytic bacteria from day flower of *Commelinabenghalensis*. The *Commelinabenghalensis* is a weed plant belongs to family *commelinaceae*^[10]. These weeds are normally found in crop field during monsoon season (June to September).

Commelinabenghalensis commonly known as the Benghal dayflower, tropical spiderwort, wandering jew, kanshira in Bengali language while in Hindi it is called as Kana and in Sanskrit: kanchata, kosapuspi, marishajalaja. Day flower is a unique character of *commelinaceae* family. This day flower is self-pollinating. It is a native to tropical Asia and Africa.

The study suggested that, there is a great diversity in reproductive systems of dayflower of *Commelina*^[11]. It produces

three types of flowers one male flower, chasmogamous and twohermaphroditeschasmogamous andcleistogamous. All three types develop on aerial branches. From this hermaphrodite chasmogamous flowers areon subaerial branches and cleistogamous flowers grow on underground branches. Also some plants may produce female flowers on aerial branches. The dayflowers have three ovules per ovary^[11].

As per the review of literature maximum work was done on different plant parts of *C. benghalensis* except dayflower and a bit of work was carried out on endophytes of this plant. So effort was made to isolated endophytic bacteria from day flower of *Commelina benghalensis* other than different plant portion.

MATERIALS AND METHODS

Collection of plant

The plant of *C. benghalensis* was collected from nearby Solapur areaand re-cultivated in college botanical garden (**Figures 1 and 2**).



Figure 1. *Commelina benghalensis* plant.



Figure 2. *Commelina benghalensis* plant.

Isolation of Endophytic bacteria

For isolation of endophytes, dayflowers were collected from underground roots of *C.benghalensis* (**Figure 3**). Separated dayflowers were brought in laboratory andwash thoroughly by tap water. Surface sterilization was carried out by 70% alcohol, Tween 20, Sodium hypochloride and distilled water. Each dayflower was cut into two pieces from its central axis. These two cut pieces were separately inoculated on nutrient agar medium (cut portion facing toward medium) and incubate for 24 hours at 37 °C.



Figure 3. Dayflowers of *commelina benghalensis*.

Biochemical characterization of Endophyte

The isolated endophytic bacterial colonies were identified on the basis of morphological, cultural and biochemical characteristics. The Biochemical tests were performed on Vitek 2 biomérieuxvitek 2 systems using the identification software PIBWIN Version 2.0.0 ^[12].

RESULTS

Isolation of Entophytic bacteria

In the present study, After 24 hours of incubation, white bacterial growth was observed on nutrient agar medium exactly near surrounding area of cut portion of dayflower. Four quadrant streaking method was performed to isolate single colony of respective bacteria. Well isolated colony was streak and maintain on separate nutrient agar slant. These isolated colonies were used for further biochemical tests (**Figure 4**).

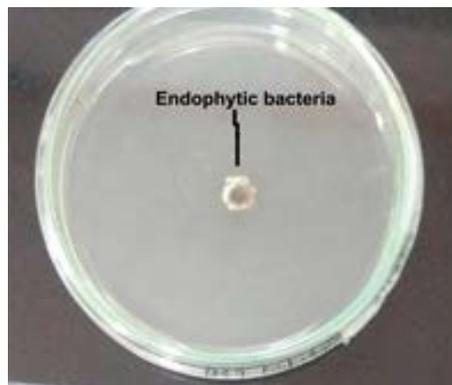


Figure 4. Isolation of endophytic bacteria of Dayflower of *commelina benghalensis*.

Biochemical test

The isolated endophytic bacteria were gram positive with white coloured colony morphology (**Figure 5**). These strain showed positive results for biochemical tests such as Catalase, Arginine, Hihydrolase 1, Beta galactosidase, Alpha glucosidase, Lascinarmylamidase, L-prolinearmylamidase, Alanine, Arylamidase, Tyrosine armylamidase, Urease, D-glucose, D-ribose, N-acetyl - D-glucosamine, D-maltose, Bactracin resistance, D-mannitol, D-mannose, Methyl -B-D-glucopyranoside, Saccharose \ Sucrose, D-trehalose and optochin resistance. The Biochemical tests were performed on Vitek 2 biomérieuxvitek 2 systems by using the identification software PIBWIN Version 2.0.0. On the basis of analysis of this system, it is finalized that the isolated bacterial strain was *Micrococcuspp*. The result of biochemical tests (**Table 1**).

Table 1. Biochemical tests for identification of isolated entophytic strains from Dayflower of *Commellina benghalensis*.

Biochemical Tests	Result
White Creame Colony	-
Yellow Orange colony	-
Pink Red colony	-
Catalase	+
Oxidase	-
Coagulase	-
Amygdalin	-

DISCUSSION

The planet having about 300,000 species of plants, from these, majority of plants contain endophytes^[13]. In nature, an endophyte-free plant is a rare exception^[14]. Endophytes help plant to cope up with infections developed by pathogen and various environmental stress^[15].

The most frequently found genera of bacterial endophytes are *Pseudomonas*, *Micrococcus*, *Bacillus*, *Stenotrophomonas*, *Burkholderia*, *Pantoea*, *Microbacterium*,^[16-22]. Genus micrococcus have been detected an endophytes in several other plants including the *Aloe vera*^[23], *zea*^[6], cotton^[24], rice seed^[5], the roots of *Polysporaaxillaris*^[25] and *Aquilariasinensis*^[26].

The result of this study indicates that the isolated strain of endophyte shows distinct features of different species of genus micrococcus, that it should be described as novel species in day flower of *C. benghalensis*. This *Micrococcus* spp. are likely involved in biodegradation of different environmental pollutant to and decrease their toxicity^[27-30]. Some *Micrococcus* isolates also can produce different useful products, such as long-chain aliphatic hydrocarbons for lubricating oils.

CONCLUSION

Medicinal plants are source of great economic value all over the World. On literature available it can be said that *C. benghalensis* plant, show unique characteristic of presence of day flower whose entophytic screening were studied. According to this study an endophytic bacteria of dayflower of said plant is nothing but *Micrococcus* spp.

C. benghalensis is relatively difficult to control by herbicide. This resistivity to herbicides might be due to presence of *micrococcus*spp. In its dayflower. Because, *Micrococcus* actively participate in catabolic reactions by utilizing a wide range of unusual substrate like herbicides, pyridine, oil and chlorinated biphenyls.

ACKNOWLEDGEMENTS

I would like to acknowledge Department of Genetics, Department of biotechnology Walchand College Solapur. Also I would to thank Dr. S. P. Gaikwad and Dr. R.D. Gore from Department of Botany, Walchand College Solapur for their help in identification of plant species. Also I acknowledge Dr. Dilip Karad, department of Microbiology, Shivaji college, Barshi for identification of endophyte.

REFERENCES

1. Wilson D. Endophyte: The evolution of a term, and clarification of its use and definition. *Oikos*. 1995; 73:274–276.
2. Adams PD, et al. Seed-borne bacterial endophytes in different cotton cultivars. *Phytopathology*. 1996;86: 597.
3. James EK, et al. Infection and colonization of rice seedlings by the plant growth-promoting bacterium *Herbaspirillum seropedicae* Z67. *Mol Plant Microbe Interact*. 2002; 15: 894–906.
4. Sessitsch A, et al. Cultivation independent population analysis of bacterial endophytes in three potato varieties based on eubacterial and Actinomycetes-specific PCROf 16S rRNA genes. *FEMS Microbiol Ecol* 2002; 39: 23–32.
5. Kaga H, et al. Rice seeds as sources of endophytic bacteria. *Microbes Environ*. 2009; 24: 154–162.
6. Johnston MD, et al. Conservation and diversity of seed associated endophytes in *Zea* across boundaries of evolution, ethnography and ecology. *PLoS ONE*. 2011; 6: e20396.
7. Rodriguez R, et al. More than 400 million years of evolution and some plants still can't make it on their own: plant stress tolerance via fungal symbiosis. *J Exp Bot*. 2008; 59: 1109–1114.
8. Verma S, et al. *Piriformosporoides* sp. nov., a new root Z-colonizing fungus. *J Mycologia*. 1998; 90: 896–903.
9. Waller F, et al. The endophytic fungus *Piriformosporoides* reprograms barley to salt-stress tolerance, disease resistance, and higher yield. *Proc Natl Acad Sci USA*. 2005; 102: 13386–13391.
10. Kharade A, et al. Photochemical investigation in *Commelinabenghalensis* and *Cyanotiscristata*, *Res J pharm App sci*. 2013; 3: 46-48.
11. Kaul V, et al. Reproductive effort and sex allocation strategy in *Commelinabenghalensis* L, a common monsoon weed. *Bot J Linn Soc*. 2002; 140: 403–413.
12. Karad D. Vitek 2 biomérieuxvitek 2 system using the identification software PIBWIN Version 2.0.0. 2018.
13. Smith SA, et al. Bioactive endophytes warrant intensified exploration and conservation. *PLoS One*. 2008; 3: e3052.
14. Partida-Martínez LP, et al. The microbe-free plant: fact or artifact? *Front Plant Sci*. 2011; 2: 100.
15. Timmusk S, et al. Bacterial distribution in the rhizosphere of wild barley under contrasting micro climates. *PLoS One*. 2011; 6: e17968.

16. Hallmann J, et al. Bacterial endophytes in agricultural crops. *Can J Microbiol.* 1997; 43: 895–914.
17. Sturz AV, Nowak J. Endophytic communities of rhizobacteria and the strategies required to create yield enhancing associations with crops. *Appl Soil Ecol.* 2000; 15: 183–190.
18. Sun Y, et al. The presence of a 1-aminocyclopropane-1-carboxylate (ACC) deaminase deletion mutation alters the physiology of the endophytic plant growth-promoting bacterium *Burkholderia phytofirmans* PsJN. *FEMS Microbiol. Lett.* 2009; 296: 131–136.
19. Rosenblueth M, et al. Bacterial endophytes and their interactions with hosts. *Molec. Plant Microbe Interact.* 2006; 19: 827–837.
20. Marquez-Santacruz HA, et al. Diversity of bacterial endophytes in roots of Mexican husk tomato plants (*Physalis ixocarpa*) and their detection in the rhizosphere. *Gen Mol Res.* 2010; 9: 2372–2380.
21. Shi Y, et al. Illumina-based analysis of endophytic bacterial diversity and space-time dynamics in sugar beet on the north slope of Tianshan mountain. *Appl Microbiol Biotechnol.* 2014; 98: 6375–6385.
22. Romero FM, et al. The communities of tomato (*Solanum lycopersicum* L) leaf endophytic bacteria, analyzed by 16S-ribosomal RNA gene pyrosequencing. *FEMS Microbiol Lett.* 2014; 351: 187–194.
23. Om Prakash, et al. Description of *Micrococcus aloevera* sp. nov., an endophytic actinobacterium isolated from *Aloe vera*, *Int J Syst Evol Microbiol.* 2014; 64: 3427–3433.
24. McInroy JA, et al. Survey of indigenous bacterial endophytes from cotton and sweet corn. *Plant Soil.* 1995; 173: 337–342.
25. Zhao GZ, et al. *Micrococcus yunnanensis* sp. nov., a novel actinobacterium isolated from surface-sterilized *Polyspora axillarum* roots. *Int J Syst Evol Microbiol.* 2009; 59: 2383–2387.
26. Chen HH, et al. *Micrococcus endophyticus* sp. nov., isolated from surface-sterilized *Aquilaria sinensis* roots. *Int J Syst Evol Microbiol.* 2009; 59: 1070–1075.
27. Doddamani H, et al. Biodegradation of carbaryl by a *Micrococcus* species. *Curr Microbiol.* 2001; 43: 69–73.
28. Sims GK, et al. Riboflavin Production during Growth of *Micrococcus luteus* on Pyridine. *Appl Environ Microbiol.* 1992; 58: 3423–3425.
29. Wilson AK. Commelinaceae - a review of the distribution, biology and control of the important weeds belonging to this family. *Tropical Pest Management.* 1981; 27: 405-418.
30. Huang W, et al. Importance of Gram-positive naphthalene-degrading bacteria in oil-contaminated tropical marine sediments. *Lett Appl Microbiol.* 2003; 36: 251–257.