

Original Research Article

Fungal diversity associated with the hive stored pollen of stingless bees *Tetragonulatravancorica* Shanas and Faseeh

ABSTRACT

The process of collecting and storing pollen by stingless bees involves a complex fermentation process, enriching it with nutrients and probiotics, making it valuable as a dietary supplement. However, the presence of mycotoxins-producing fungi in bee pollen poses health risks to humans. Thus, the study aims to characterize the fungi associated with hive-stored pollen of stingless bees. The results revealed the presence of various fungal species, including *Penicillium* and *Aspergillus*, which can contaminate pollen with mycotoxins. This necessitates post-harvest processing to reduce microbial contamination, ensuring the safety and quality of bee-derived food products for consumer health

Keywords: Stingless bees, *Tetragonulatravancorica*, Microbiota, hive-stored pollen, and Fungi.

1. INTRODUCTION

Pollen is often regarded as “the world's best food product” [1]. Stingless bees gather pollen during foraging, transport it to the hive in their specialized pollen baskets, and store it in designated pollen pots. While packing the pollen in pots, bee pollen is enriched with honey, as well as digestive enzymes and organic acids from the salivary glands secretions of bees [2]. This process initiates spontaneous lactic fermentation of the pollen by *Lactobacillus* bacteria within the pollen pots. Pollen sheaths are dissolved in the process of pollen transformation. Fermentation not only protects the pollen against the loss of properties but also gives rise to new components as a result of enzymatic transformations. Proteins in the pollen degrade into peptides and amino acids during this fermentation

process. Studies revealed that pollen contains higher protein concentrations than bee bread, although amino acid levels are generally lower. Elevated levels of free amino acids may result from specific proteolytic enzyme activity breaking down polypeptide chains. The compound content in bee bread can be influenced not only by the pollen source but also by the genotype of the bees converting it. Furthermore, lactic acid concentration in bee bread is approximately six times higher than in pollen [3]. The presence of lactic acid helps preserve bee bread, thereby extending its shelf life.

The combination of protein content and added probiotics renders bee bread a valuable dietary supplement. While the recommended daily intake for adults stands at approximately 20 grams. Due to its health-promoting attributes, it's crucial to monitor bee pollen for contamination with harmful substances from a food safety perspective [4,5]. The presence of mycotoxin-producing fungi in bee pollen was documented by [6]. Many fungi produce mycotoxins, which can cause acute or chronic intoxication and pose risks to human health upon consumption of contaminated food [7]. Thus, the present study was carried out to characterize the fungi associated with the hive-stored pollen of stingless bees.

2. MATERIAL AND METHODS

2.1. Collection of hive-stored pollen

The hive-stored pollen of stingless bees was collected from commercial stingless beekeepers of Kerala. Pollen was collected by taking out the pollen pots and removing the caps of the pollen pots and the extracted pollen was placed inside a plastic container. The procedure was repeated at least 4 times to gather a minimum of 5 grams of pollen. Later the pollen samples were labelled with a sample code for easy identification and stored in a refrigerator (-20° C).

2.2. Isolation and purification of fungi associated with hive-stored pollen

The fungi associated with hive-stored pollen of stingless bees were isolated using serial dilution and plate count technique using Potato dextrose agar (PDA) [8]. The isolated plates were incubated at room temperature for 5 days. After this incubation period, the plates

were examined for colony growth, and the cultural and morphological characteristics of the isolates obtained in the isolated plates were recorded. The distinct fungal colonies from each plate were purified separately using the disc method and stored in a refrigerator for identification.

2.3. Molecular characterization of fungal isolates obtained from the hive-stored pollen

The fungal isolates were identified through molecular techniques by sequencing the ITS region. The pure fungal cultures were sent to Biokart, Bangalore for genomic-level sequencing and identification. BLASTn was utilized to search for homologous sequences using nucleotide data obtained from Biokart. Subsequently, all isolate sequences were deposited into the NCBI GENBANK database.

3. RESULTS

The fungal composition present in the hive-stored pollen of stingless bees was analyzed through serial dilution and plate count techniques. The morphological and cultural characteristics of fungal isolates were used for the identification of fungi. Further confirmation is through molecular characterization.

A total of 21 fungal isolates were identified in the hive-stored samples collected from Kerala. Among them, four distinct fungal isolates were identified from all the samples. The morphological and cultural characteristics of fungal isolates aid in the identification of fungi such as *Penicillium* spp., *Penicillium chrysogenum*, *Aspergillus aculeatus*, and *Aspergillus flavus*. The colony characteristics of the *Penicillium* spp. were filamentous, medium-sized, flat with entire margins, rough, and green colonies with white borders. While *Penicillium chrysogenum* colonies were irregular, small-sized, raised elevation, filiform borders, rough surface, and yellow colour with green spores. The fungal genus, well known as laboratory weed *Aspergillus* is also recorded in the hive-stored pollen samples. The various species of *Aspergillus* recorded in the hive stored pollen including *Aspergillus flavus* and *Aspergillus aculeatus*. The colony morphology of *Aspergillus flavus* was circular-shaped, small-sized,

and raised colonies, with entire margins, wrinkled surfaces, and white margins with brown to black spores. Whereas, the colony morphology of *Aspergillus aculeatus* was irregular, small-sized, raised elevation, filiform borders, rough surface, and brown to black spores (Plate 1).

Molecular characterization of fungal isolates associated with the hive stored pollen of *T. travancorica* was carried out using ITS gene sequencing from Biokart India Pvt Ltd. The PCR product amplified on 1.2 per cent agarose showed a band between 600-00bp (Fig. 1). The sequence analysis was used to confirm the fungal isolates as *Penicillium sp.* (Accession number: PP296549), *Aspergillus aculeatus* (Accession number: PP291951), *Aspergillus flavus* (Accession number: PP291949), and *Penicillium chrysogenum* (Accession number: PP301322). This confirmed the cultural, and morphological characterization.

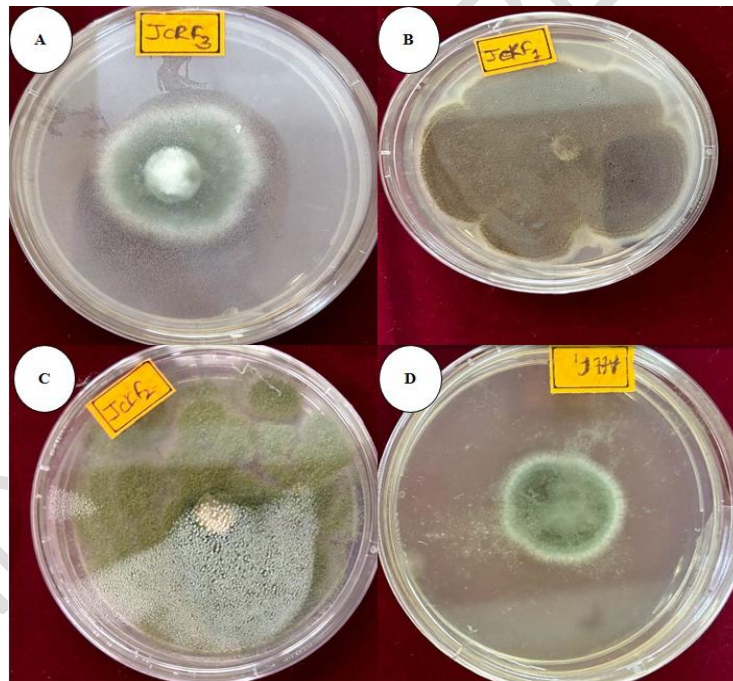


Plate 1. Fungi obtained from the hive stored pollen of stingless bee *T. travancorica*, A-*Penicillium chrysogenum*, B-*Aspergillus aculeatus*, C-*Aspergillus flavus*, and D-*Penicillium spp.*

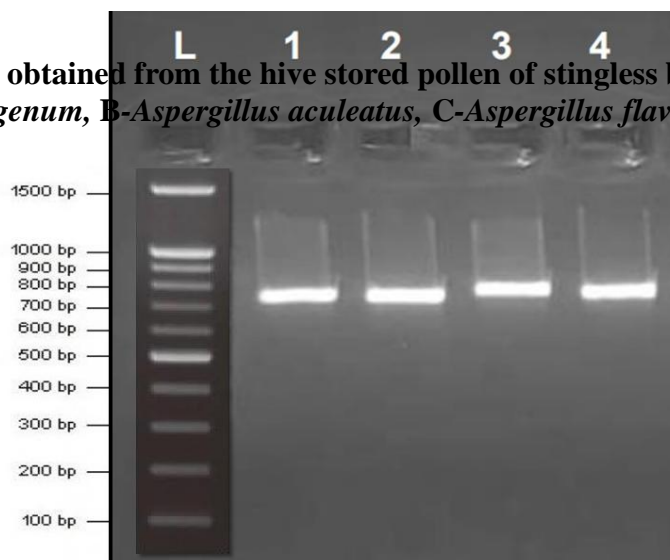


Figure 1. Agarose gel electrophoresis of Genomic DNA isolated from the fungi associated with hive stored pollen. 1-*Penicillium chrysogenum*, 2-*Aspergillus aculeatus*, 3-*Aspergillus flavus*, and 4-*Penicillium spp.*

4. Discussion

The fungi associated with the hive stored pollen of stingless bees revealed the presence of *Penicillium spp.*, *Penicillium chrysogenum*, *Aspergillus flavus*, and *Aspergillus aculeatus*. Similar fungal compositions have been observed in the previous studies of bee pollen and bee bread [9,10,11]. The major factors such as plant source, geographical origin, and bee-keeping practices significantly influence the microbial composition in the bee pollen [12,13]. *Aspergillus spp.* can thrive on decomposing plant matter and are adapted for the degradation of complex plant polymers [14]. These species are often found in association with bees and bee products, particularly pollen. Pollen serves as a point of entry for fungal pathogens into beehives. Studies have shown that spores of *Aspergillus spp.* can contaminate pollen while it is still on plants [15]. Bees collect pollen, store, and consume pollen, allowing the spores to reach their gut, which is the primary site of infection for bee pathogens [16]. *Aspergillus spp.* are opportunistic pathogens capable of infecting bees at all developmental stages [17]. *Aspergillus flavus* was the causal agent of the stone brood disease of honey bees and is poorly studied in the honey bee pathogens [16,18]. Bee pollen acts as a substrate for the production of mycotoxins [19]. The commercial pollen may still

contain *A. flavus* spores [2,6,20]. Thus, posing a risk to human health due to high mould contamination levels and mycotoxins. Therefore, it is important to implement post-harvest processing to reduce microbial contamination.

The association of *Penicillium* spp. with honey and bee bread has been reported earlier, but their specific functions within the hives of stingless bees have been lacking. Studies showed that *Penicillium* spp. is capable of secreting organic acids that aid in the preservation of pollen. The enzymes secreted by the *Penicillium* contributed to the nutritional value of the pollen [21], and protection against pathogens [22]. Some species of *Penicillium* are producing mycotoxins that impair the health of humans, hence the actual role and specific mechanisms need to be explored.

5. Conclusion

The present work highlights the intricate relationship between fungi and bee products like pollen. The fungi enter the hive from flowers through bees during foraging, proliferate within stored pollen, and produce mycotoxins. These mycotoxins can pose health risks to humans. Hence, it's crucial to implement measures to minimize microbial contamination in food products derived from bee pollen through careful post-handling processes. This emphasizes the importance of ensuring the safety and quality of bee-derived food products for consumer health.

REFERENCES

1. Bobiș O, Mărghitaș LA, Dezmirean D, Morar O, Bonta V, and Chirilă F. Quality parameters and nutritional value of different commercial bee products. Bull Univ Agric Sci Vet Med Cluj Napoca. 2010; 67(2).
2. Deveza MV, Keller KM, Lorenzon MCA, Nunes LMT, Sales ÉO, and Barth OM..Mycotoxicological and palynological profiles of commercial brands of dried bee pollen. Braz. J. Microbiol. 2015;46:1171-1176.

3. Nagai T, Nagashima T, Myoda T, and Inoue R. Preparation and functional properties of extracts from bee bread. *Food/nahrung*, 2004;48(3), 226-229.
4. Kostić AŽ, Milinčić DD, Petrović TS, Krnjaja VS, Stanojević SP, Barać MB, Tešić ŽL, and Pešić MB. Mycotoxins and mycotoxin-producing fungi in pollen. *Toxins*, 2019;11(2):64.
5. Végh R, Csóka M, Sörös C, and Sipos L. Food safety hazards of bee pollen—A review. *Trends Food Sci. Technol.* 2021;114:490-509.
6. González G, Hinojo MJ, Mateo R, Medina A, and Jiménez M. Occurrence of mycotoxin producing fungi in bee pollen. *Int. J. Food Microbiol.* 2005;105(1):1-9.
7. Moss MO. Centenary review: mycotoxins. *Mycological Res.* 1996;100(5):513-523.
8. Kačániová M, Pavličová S, Haščík P, Kociubinski G, Křázovická V, Sudzina M, Sudzinová J, and Fikselová M. Microbial communities in bees, pollen and honey from Slovakia. *Acta Microbiol. Imm. H.* 2009;56(3):285-295.
9. Nardoni S, D'Ascenzi C, Rocchigiani G, Moretti V, and Mancianti F. Occurrence of moulds from bee pollen in Central Italy-A preliminary study. *Ann. Agr. Env. Med.* 2016;23(1).
10. Barbosa RN, Bezerra JD, Souza-Motta CM, Frisvad JC, Samson RA, Oliveira NT and Houbraken J. New *Penicillium* and *Talaromyces* species from honey, pollen and nests of stingless bees. *Antonie Van Leeuwenhoek*, 2018; 111:1883-1912.
11. Bush DS, Calla B, and Berenbaum MR. An *Aspergillus flavus* strain from bee bread of the Western honey bee (*Apis mellifera*) displays adaptations to distinctive features of the hive environment. *Ecology and Evolution*, 2024;14(2):e10918.
12. Nogueira C, Iglesias A, Feás X, and Estevinho LM. Commercial bee pollen with different geographical origins: a comprehensive approach. *Int. J. Mol. Sci.* 2012;13(9):11173-11187.
13. De-Melo AAM, Estevinho MLMF, Sattler JAG, Souza BR, da Silva Freitas A, Barth OM, and Almeida-Muradian LB. Effect of processing conditions on characteristics of

- dehydrated bee-pollen and correlation between quality parameters. *LWT-Food Science and Technology*, 2016;65:808-815.
14. Bennett JW. An overview of the genus *Aspergillus*. *Aspergillus molecular biology and genomics*. Caister Academic Press, Norfolk, United Kingdom. 2010; 1–17.
 15. Gilliam M, Prest DB, and Lorenz BJ. Microbiology of pollen and bee bread: taxonomy and enzymology of molds. *Apidologie*, 1989;20(1):53-68.
 16. Foley K, Fazio G, Jensen AB, and Hughes WO. The distribution of *Aspergillus* spp. opportunistic parasites in hives and their pathogenicity to honey bees. *Vet. Microbiol.* 2014;169(3-4):203-210.
 17. Foley K, Fazio G, Jensen AB, and Hughes WO. Nutritional limitation and resistance to opportunistic *Aspergillus* parasites in honey bee larvae. *J. Invertebr. Pathol.* 2012;111(1):68-73.
 18. Schwarz RS, Huang Q, and Evans JD. Hologenome theory and the honey bee pathosphere. *Curr. Opin Insect Sci.* 2015;10:1-7.
 19. Medina Á, González G, Sáez JM, Mateo R, and Jiménez M. Bee pollen, a substrate that stimulates ochratoxin A production by *Aspergillus ochraceus* Wilh. *Syst. Appl. Microbiol.* 2004;27(2):261-267.
 20. Bucio Villalobos CM, López Preciado G, Martínez Jaime OA, and Torres Morales JJ. Mycoflora is associated to bee pollen collected by domesticated bees (*Apis mellifera* L). *Nova scientia*, 2010;2(4):93-103.
 21. Hsu CK, Wang DY, and Wu MC. A potential fungal probiotic *Aureobasidium melanogenum* CK-CsC for the western honey bee, *Apis mellifera*. *J. Fungi*, 2021;7(7):508.
 22. Disayathanoowat T, Li H, Supapimon N, Suwannarach N, Lumyong S, Chantawannakul P, and Guo J, Different dynamics of bacterial and fungal communities in hive-stored bee bread and their possible roles: a case study from two commercial honey bees in China. *Microorganisms*, 2020;8(2):264.