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# Remarkable fungal biodiversity on *Arachis hypogea* L. field at Dhule, Maharashtra, India.

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## **ABSTRACT**

Groundnut (Arachis hypogea L.) accounts for almost half of India's oil seed area, making it the country's one of most important oil seed crop. Plants of the genus A. hypogea L. are susceptible to suffering from fungal diseases. A purposeful and random sampling was done to record the course of two consecutive kharif seasons in 2022 and 2023, the current aerobiological study was conducted over the A. hypogea L. Var. TAG-24 fields in four talukas of the Dhule district. These talukas are Dhule, Sakri, Shirpur, and Shindkheda. When doing the survey, a volumetric Tilak air sampler is utilized to ascertain the fungal concentration that is present in the crop. It always maintains a height of 0.75 meters and is positioned in the middle of the groundnut field. Over sixty distinct fungal spore species, belonging to five different divisions such as Basidiomycota, Ascomycota, Oomycota, Mucoromycota, and Myxomycota, eleven classes, and more than fifty-five families, were found to be concentrated in a single region under the observation of aerobiological monitoring. Over the course of the kharif seasons, meteorological information was maintained at a current level.

Keywords: Air spora; Aerobiology; Arachis hypogaea; Groundnut; Fungal Spores, Basidiomycota, Ascomycota.

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## INTRODUCTION

The present study is all about the qualitative analysis of air spore found in one of the most important oil seed crops Arachis hypogea L, this analysis is unique in its own way because, it is found that only one aerobiological survey is carried by Dr. M. T. Bhadane in over 34 years back at Dhule city [1]. Microorganism shows important relationship with crops where they are found on, characteristics of Gram-positive and filamentous bacteria were seen in actinomycetes that were isolated from the rhizospheric areas of *Arachis* hypogea L. and Gossypium herbaceum L. The isolates demonstrated the capacity to effectively utilize a wide range of nutritional sources present in the rhizosphere [2]. The research revealed that the predominant species of arbuscular mycorrhizal fungi (AMF) indigenous spores in the rhizosphere of peanuts were Glomus sp., Acaulospora sp., and Glomus sp. The elevation level, population size, and percentage of infection were found to be linked with the type of infection [3]. Uncultivated varieties of the *Arachis* genus exhibited higher resistance to fungal infections compared to farmed peanuts. Some wild accessions exhibited comparable or superior resistance to A. cardenasii. [4]. The etiological agent responsible for leaf spot in peanut plants in China is Cladosporium tenuissimum. The presence of C. tenuissimum infection presents a significant risk to both the quantity and quality of peanut production [5]. Hydrolysis of peanut protein concentrates by fungal crude protease extract affects structural, functional, and in-vitro protein digestibility properties. Hydrolysis with crude extract from Aspergillus oryzae showed the highest solubility, water and oil binding capacity, foaming capacity, foam stability, and in-vitro protein digestibility [6]. The experimental treatments NLEFS, DLEFS, DebLEFS, and BBSFS exhibited superior efficacy in managing leaf spot and enhancing pod yield. Based on the data, BBSFS were having the highest overall profit and Benefit-Cost Ratio (Hasan et al., 2016). Pseudomonas fluorescens and Trichoderma viride can be used to biologically control Fusarium wilt disease in peanuts. The efficacy of Pseudomonas fluorescens in

regulating the growth of Fusarium oxysporum was found to be superior, leading to elevated amounts of chlorophyll and carotenoids [7]. Aspergillus niger, Fusarium solani, Rhizopus stolonifer, and Aspergillus flavus are common pathogenic fungi causing rot in groundnut seeds. Ethanolic Azadirachta indica extracts have more inhibitory compounds than aqueous extracts [8]. Arsule and Pande studied fungal airspora over the groundnut fields at Newasa (M.S.) for two consecutive Kharif seasons by employing Tilak volumetric air sampler [1, 2]. Fungal spores are prevalent in Arachis hypogaea L. fields and can have an impact on crop yield. Various fungal pathogens, such as Passalora arachidicola, Aspergillus flavus, A. niger, A. fumigatus, A. nidulans, Fusarium oxysporum, Sclerotium rolfsii, Rhizoctonia bataticola, Curvularia lunata, Alternaria alternata, and Rhizopus stolonifer, have been detected in groundnut fields[9]. These fungal pathogens can cause diseases like early leaf spot and aflatoxin contamination, leading to reduced yield and quality of the crop [10, 11]. However, the prevalence and impact of fungal spores on crop yield can be mitigated through various strategies. For example, the colonization of the endophytic fungus *Phomopsis liquidambaris* has been found to promote nodulation and increase rhizosphere nitrogen availability, resulting in improved overall nitrogen utilization and yield stabilization [12]. Additionally, the use of improved peanut varieties, such as ICGV-SM 90704, has shown lower disease rates, lower levels of aflatoxins, and higher yields compared to locally used varieties.

### **MATERIAL AND METHODS**

For current study air sampling is done using Tilak Air sampler for Plants kharif and summer seasons from the year 2022 and 2023 at Dhule, Sakri, Shirpur and Shindkheda Taluka regions (**Figure 1**). Preparation and scanning of slides is done with the help of PVLG and Motic (Compound Microscope) respectively. Airspora were identified based on spore morphology (the physical characteristics of spores). While spore morphology involves examining the shape, size, colour, and other features of spores produced during asexual or sexual reproduction. For comparison the crop land characters were examine for some values such as pH, Av. Nitrogen, Av. Phosphorus, Av. Potassium (Table-1).



Figure 1 Area of Study: Dhule District, Maharashtra (India)

Table 1 Varieties used for respective seasons with sampling days

Sr No.	Place	Season	Varieties	Total No. of
				Sampling Days
1	Dhule	*K 1	SB-11	129
2		*K 2	SB-11	126
3		*S 1	TAG-24	119
4		*S 2	TAG-24	123
5	Sakari	*K 1	SB-11	129
6		*K 2	JL-24	126
7		*S 1	TAG-24	123
8		*S 2	TAG-24	119
9	Shindkheda	*K 1	JL-24	125
10		*K 2	JL-24	126
11		*S 1	TAG-24	119
12		*S 2	TAG-24	127
13	Shirpur	*K 1	JL-24	125
14		*K 2	JL-24	126
15		*S 1	TAG-24	119
16		*S 2	TAG-24	128

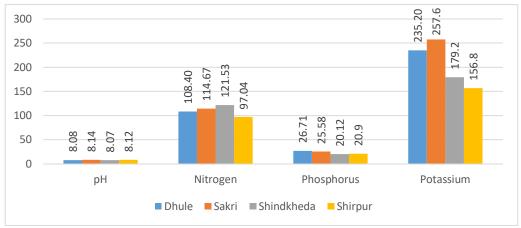


Figure 2 Average pH and NPK Lavels in study area

## RESULTS AND DISCUSSION

The presence of fungal spores in Arachis hypogaea L. fields can have a negative impact on the quality of the peanuts produced. Fungal diseases, such as groundnut seed rot caused by organisms like Aspergillus niger, Fusarium solani, Rhizopus stolonifer, and Aspergillus flavus, can lead to rot incidences and reduce the yield of peanuts [13]. These fungi are not only a constraint to peanut production but also pose health hazards for human consumption [14]. Aflatoxin contamination, primarily caused by Aspergillus flavus, is a major concern in groundnut cultivation as it affects the quality and safety of the peanuts [15]. To mitigate the impact of fungal diseases, various approaches have been explored, including cultural management, diseaseresistant cultivar development, and the use of plant extracts like neem to control fungal growth [3, 5, 12]. Additionally, the use of arbuscular mycorrhizal fungi (AMF) has shown potential in improving yield and yield components in peanuts. Overall, managing fungal spores in peanut fields is crucial for ensuring the quality and safety of the peanuts produced. The study demonstrates a wide range of fungal spores. The divisions encompassed within this taxonomic classification are Deuteromycota, Zygomycota, Ascomycota, Basidomycota, and Myxomycota. The investigation of groundnut cultivation in the talukas like Sakari, Shirpur, and Shindkheda of Dhule District is totally new. Air spora from groundnut field ware studied in 1990 at Dhule taluka of Dhule district. The aerobiological diversity in ground nut fields in areas such as Sakri, Shindkheda, and Shirpur of Dhule district is being examined and reported for the first time. The study revealed that there was a greater variation in the number of spores or fungal hyphae under high humidity circumstances. However, modest differences for presence of spores were noted in all four locations due to variations in climatic parameters. The kharif season months (July-November) had the highest spore count compared to the summer season. Four different types of groundnut seeds, namely Gujrat, Western, Ghungroo, and Local, were obtained from several markets in the state of Madhya Pradesh. The mycoflora present in these seeds was extracted using both the usual blotter paper method and the agar plate method. The seeds in all four kinds demonstrated a higher amount of fungus, accompanied by a greater proportion of incidence. The major fungi identified in the study were Aspergillus flavus, A. niger, Fusarium oxysporum, Macrophomina phaseolina, and Penicillium sp. A greater quantity of fungus was obtained through the utilization of the agar plate approach in comparison to the conventional blotter paper method. The use of HgCl2 for surface sterilizing decreases the occurrence of Aspergillus flavus and Aspergillus niger [16]. The prevailing strain of peanut rhizobia in Henan Province, China is Bradyrhizobium guangdongense. The

occurrence of various strains is correlated with soil pH and the availability of phosphorus [17].

Table 2 Fungal spores were observed in study area.

	rable 2 Fullgal spore					
Sr. No.	Spore Name	MycoBank #				
Asco	Ascomycota					
1.	Aspergillus P. Micheli Ex Haller	7248				
2.	Alternaria Nees	7106				
3.	Arthrinium Kunze	7214				
4.	Beltrania Penz.	7355				
5.	Bispora Corda	7377				
6.	Botryodiplodia Sacc.	7420				
7.	Cephaliophora Thaxt.	7518				
8.	Ceratophorum Sacc.	7534				
9.	Cercospora Fresen. Ex Fuckel	7545				
10.	Chaetomella Fuck.	7575				
11.	Cladosporium Link.	7681				
12.	Cordana Preuss.	7777				
13.	Corynespora Güssow.	92201				
14.	Curvularia Boed.	7847				
15.	Deightoniella S. Hughes.	7934				
16.	Dictyoarthrinium Hughes.	7993				
17.	Dictyosporium Corda.	8001				
18.	Diplodia Fr.	8047				
19.	Epicoccum Link.	8188				
20.	Fusariella Sacc.	8282				
21.	Fusarium Link.	8284				
22.	Haplosporella Speg.	8441				
23.	Harknessia Cook.	8449				
24.	Helminthosporium Link.	8495				
25.	Heterosporium Klotzsch.	8529				
26.	Lacellina Sacc.	8693				
27.	Memnoniella Höhnel.	8900				
28.	Nigrospora Zimm.	9124				
29.	Periconia Tode.Ex Schw.	9263				
30.	Pestalotia De Not.	9271				
31.	Pithomyces Berk.	9412				
32.	Pseudotorula Subram.	9620				
33.	Pyricularia Sacc.	9670				
34.	Spegazzinia Sacc.	9963				
35.	Sporidesmium Link.	10024				
36.	Tetraploa Berk. & Br.	10199				

vere observed in study area.						
Sr. No.	Spore Name	MycoBank #				
37.	Torula (Pers) Link. This	10248				
38.	Trichothecium Link.	10303				
39.	Amphisphaerella (Sacc.) Kirsch.	261				
40.	Ascotricha Berk	384				
41.	Bitrimonospora Sivanesan, Talde & Tilak	587				
42.	Chaetomium Kunz. Ex. Fr.	953				
43.	Claviceps Tul.	1092				
44.	Cucurbitaria Gray.	1348				
45.	Didymosphaeria Fuckel.	1562				
46.	Hypoxylon Bull. Ex Fr.	2456				
47.	Hysterium Tode. Ex Fr.	2464				
48.	Lophiostoma Ces. & De Not.	2933				
49.	Massarina Sacc.	3016				
50.	Melanospora Corda.	3085				
51.	Nodulosphaeria Rabh.	3517				
52.	Otthia Nke.	3656				
53.	Pleomassaria Speg.	4214				
54.	Pleospora Rabh.	4233				
55.	Sordaria Ces. & De Not.	5061				
56.	Valsaria Ces. & De Not	5704				
Basi	diomycota					
57.	Basidiospores					
58.	Ganoderma Karst.	17639				
59.	Smut Spores					
Muc	oromycota					
60.	Cunninghamella Matr.	20150				
61.	Mucor Micheli Ex. Fr.	20348				
Myx	Myxomycota					
62.	Physarum Pers.	12178				
63.	Stemonitis Roth, Mag.	1787				
Oom	ycota					
64.	Albugo Pers. Ex. S. F. Gray.	20015				
65.	Phytophthora De Bary.	20418				
66.	Scelerospora (Sacc.) Schroet.	20514				

Effective management strategies for controlling fungal spores in *Arachis hypogaea* L. fields include the use of recommended fungicides [18]. Planting moderately resistant varieties can also reduce the need for fungicide application and associated expenses [19]. Integrated management approaches, which consider all available pest control techniques, are preferred for managing agricultural pests [20]. Biological control, such as the use of native bacterial and fungal bio agents, has shown to be beneficial in reducing the reliance on agricultural chemicals [21]. Additionally, the application of plant extracts, such as neem extracts, has demonstrated inhibitory effects on fungal mycelial growth [4]. Soil application of mineral nutrients, particularly copper and potassium, has also been effective in reducing disease incidence and increasing pod yield. Therefore, a combination of these management strategies, including the use of fungicides, resistant varieties, biological control agents, plant extracts, and soil amendments, can help control fungal spores in *Arachis hypogaea L.* fields.

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