

RESEARCH ARTICLE

MYCOLOGICAL ISOLATION AND IDENTIFICATION IN THE CAFETERIA OF HAWLER CITY KURDISTAN REGION, IRAQ

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ABSTRACT

In order to forecast the effects of the fungi on humans, this study was conducted to isolate and identify the fungi in three distinct samples in the college of science cafeteria. In order to obtain well-isolated colonies, the power plate method with Potato Dextrose Agar (PDA) was used to isolate and identify various fungal species. The results show the identity and the total colony forming units (CFU) for fungi. The most frequently isolated fungi were *Cladosporium*, *Rhizopus* sp., *Aspergillus* Sp, *Alternaria*, *Yeast*, *Penicillium*, . While the less frequently detected fungi species were *Trichocladium*, *Drecheslera* sp. and *Botrytis* sp. According to the fungi the higher numbers of fungi were isolated on menu and containers.

KEYWORDS

Cafeteria, Fungi, *Aspergillus niger*, tables, containers

1. INTRODUCTION

Due to daily handling, cafeteria and restaurant items (menus, tables, chairs, and containers) may pose a risk of cross-contamination between patrons' hands and food and, if not cleaned and disinfected, may serve as a vector for certain food-borne illnesses. (Martin et al., 2016)

The kingdom Myceteae includes fungi. This group's unique traits include being eukaryotic, not photosynthetic, lacking tissue differentiation, having a chitin or other polysaccharide-based cell wall, and spreading through spores (either sexual or asexual) (Benson, 2002). The most prevalent fungal infections in humans are opportunistic fungal infections that occur in individuals with immune system disorders. (Kavanagh, 2007). The majority of bacteria in humans and the majority of other animals are found in the gut and on the skin. Though many are helpful, especially in the gut flora, the immune system's defenses render the great majority of bacteria in the body harmless. On the other hand, a number of bacterial species are harmful and can result in infectious illnesses such as leprosy, cholera, syphilis, anthrax, and bubonic plague. Respiratory infections are the most common and deadly bacterial diseases. (Sears, 2005). In many infectious diseases, microorganisms are the pathogens that cause the illness. The organisms involved include fungi that cause diseases like candidiasis, ringworm, or histoplasmosis, as well as pathogenic bacteria that cause illnesses like anthrax, TB, and plague (Lepp et al., 2004). The aim of the research is to identify and separate the bacteria and fungi found in restaurants, as well as to ascertain their existence and potential health impacts on people

2. MATERIALS AND METHODS

2.1 Sampling

Sample swapping was done in the cafeteria at various locations (chairs, tables, and containers). The samples were then brought to the lab the same day of the sample collection to be cultured on the growth medium for identification and isolation.

2.2 Identification Of The Fungal Genera

Transferring the fungal isolates to sterile plates allowed for their identification and purification. grown using the power plate technique on PDA medium and incubated for three to five days at 25°C. Media utilized in the identification and isolation of fungi Potato Dextrose Agar (PDA) (Onuorah, 1982).

After the fungi were grown, they were put on a slide, stained with lactophenol-cotton blue to look for fungal structures, covered with a cover slip, looked at under a microscope, and identified using atlas books and the morphology of their colonies and spores (Ronhede et al., 2005 and Rajankar et al., 2007).

Chemical compound Quantity of PDA

Dextrose 20 g.

Potato... .200 g.

Nutrient Agar.....500g.

Distill water (D.W.).....1 L

3. RESULT AND DISCUSSION

The total colony forming units (CFU) of the fungi in sample 1 on PDA medium are displayed in the results table (table 1) in each of the three locations where samples were collected (table, chair, and container). Tables held the highest quantity of isolated fungi, with 62 CFU, followed by chairs with 47 CFU. In this sample, *Aspergillus Niger*, *Penecillium* sp., *Vorticella* sp., and *Cladosporium* sp. were the most often isolated fungi.

The total colony forming units (CFU) of the fungi in sample 2 on PDA medium and SDA medium are displayed in table 2 along with the three locations where samples were collected (table, chair, and container). There were 308 CFU more isolated fungi on the chair than there were on

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the tables (96 CFU). In this sample, *Penecillium* sp., *Aspergillus* sp., and *Cladosporium* sp. were the most often isolated fungi.

The findings displayed in Table 3 demonstrate the total colony forming units (CFU) of fungi in Sample 3 on PDA and SDA medium in the three locations where samples were collected (the table, chair, and container). With 148 CFU on a container and 96 CFU on chairs, respectively, the higher concentrations of isolated fungi were found. In this sample, *Alternaria* sp., *Penecillium* sp., *Cladosporium* sp., and *Mycelia Sterilia* were the most often isolated fungi.

Figure (1) and Table (4) display the proportion of fungal genera in the initial sample. *Aspergillus niger* had a higher percentage of isolated fungi (77%), compared to 30% for *Penecillium* sp. *Scopula* and *Alternaria* sp., with a combined 0.8% of the total, have the lowest percentage of isolated fungi. Figure (2) and Table (5) display the proportion of fungal genera in the second sample. *Aspergillus niger* had a higher percentage of isolated fungi (71%), compared to 14% for *Cladosporium* sp. *Candida*, on the other hand, made up only 0.2% of the isolated fungi.

The third sample's percentage of fungal genera is displayed in Table (6) and Figure (3). *Aspergillus niger* accounted for 45% of the isolated fungal percentages. *Cladosporium* sp., on the other hand, had a lower percentage of isolated fungi—2%. The results across all samples show that there were more microorganisms on the chairs and containers (sugar, salt, pepper, etc.) figure (4,5,6), which may be related to the infrequent cleaning and disinfection of these areas. Aerobic bacteria might be proliferating in your essential salt and pepper shakers. It is well known that these thrive in

environments with oxygen. *E. coli* and certain Coliform bacteria, which are found in the gut and in feces, are among the bacteria that could be there (Barbara et al 2000; Joanne and others, 2008). Menus are a breeding ground for contamination, just like almost anything that is passed around and touched regularly. Bacteria such as *E. Coli* and *Staph* were detected in one study. When exposed to rhizopus, most healthy individuals are not at risk for serious health issues. The illness known as mucormycosis is an uncommon but dangerous risk. When this mold gets into the air and gets into the sinuses or lungs, it can cause illness (Bjorklund et al 2005). Open wounds are another way for it to get into the bloodstream. Because their skin is so susceptible to infection, people who have burns are more likely to get this illness. Fever, cough, eye swelling, and black discharge from the nose are a few of the symptoms of mucormycosis (Bjorklund et al 2005; Kirk et al 2008). Numerous molds, particularly those belonging to the *Cladosporium* and *Aspergillus* genera, can grow indoors where they can enter through dust particles that are attached to spores or active fungi (Ainsworth, 2009). The *Aspergillus* and *Penicillium* species are typically found isolated in indoor settings, typically originating from indoor sources. In spite of the fact that Hindy and Awad noted that *Cladosporium* species typically originate in outdoor environments, we were able to identify them in indoor air during our study. Because *Aspergillus niger* is a common species of the genus *Aspergillus*, it is a fungus and the most commonly isolated type of *Aspergillus*. It contaminates food frequently and causes a disease known as "black mold" on some fruits and vegetables, including grapes, apricots, onions, and peanuts. It grows widely in soil and is frequently observed in indoor environments, where its black colonies are mistaken for *Stachybotrys* colonies (a species also known as "black mold") (Kirk et al., 2008).

Table 1: Variation Of Fungi Count (CFU.m⁻¹) in first sample.

chairs (Fungi)	CFU	Table (Fungi)	CFU	Containers(fungi)	CFU
<i>Penicillium</i> sp.	30	<i>Cladosporium</i> sp.	3	<i>Vorticella</i> sp.	4
<i>Cladosporium</i> sp.	7	<i>Rhizopus</i> sp	2	<i>Alternaria</i> sp.	1
<i>Aspergillus niger</i>	100	<i>Drechslera</i> sp.	1	<i>Aspergillus niger</i>	30
<i>Yeast</i> .	3	<i>Yeast</i> .	1	<i>Cladosporium</i> sp.	1
<i>Trichoderma</i> sp.	1	<i>Aspergillus niger</i>	55	<i>Torula</i> sp.	2
<i>Drechslera</i> sp.	1			<i>Humicola</i> sp.	3
<i>Torula</i> sp.	2			<i>Trichocladium</i>	1
<i>Penicillium</i> sp.	2			<i>Mycelia sterilia</i>	1
<i>Alternaria</i> sp	1				
<i>Total</i>	47		62		43

Table 2: Variation Of Fungi Count (CFU.m⁻¹) in second sample.

chairs (fungi)	CFU	Table (Fungi)	CFU	Containers(fungi)	CFU
<i>Cladosporium</i> sp.	60	<i>Cladosporium</i> sp.	11	<i>Aspergillus</i> sp.	30
<i>Altrenaria</i> sp.	0	<i>Altrenaria</i> sp.	30	<i>Cladosporium</i> sp.	2
<i>Aspergillus</i> sp.	300	<i>Aspergillus</i> sp.	54		
		<i>Candida</i> sp.	1	<i>Penicillium</i> sp.	40
<i>Total</i>	360		96		72

Table 3: Variation Of Fungi Count (CFU.m⁻¹) in Third sample.

chairs (fungi)	CFU	Table (Fungi)	CFU	Containers(fungi)	CFU
<i>Yeast</i>	5	<i>Yeast</i>	10	<i>Yeast</i>	12
<i>Mycelia Sterilia</i>	2	<i>Alternaria</i> sp.	0	<i>Cladosporium</i> sp.	4
<i>Penicillium</i> sp.	20	<i>Penicillium</i> sp.	6	<i>Aspergella niger</i>	91
<i>Alternaria</i> sp.	40	<i>Cladosporium</i> sp.	2	<i>Alternaria</i> sp.	1
<i>Aspergellus nigar</i>	29	<i>Mycelia sterilia</i>	1	<i>Penicillium</i> sp.	40
<i>Total</i>	96		19		148

Table 4: Total fungi count (CFU.ml⁻¹) in first sample

Fungi	Count	Percentage%
<i>Penicillium</i> sp.	30	12
<i>Cladosporium</i> sp.	11	4
<i>Aspergillus niger</i>	185	77
<i>Yeast</i>	4	1.6

Table 4: Total fungi count (CFU.ml-1) in first sample

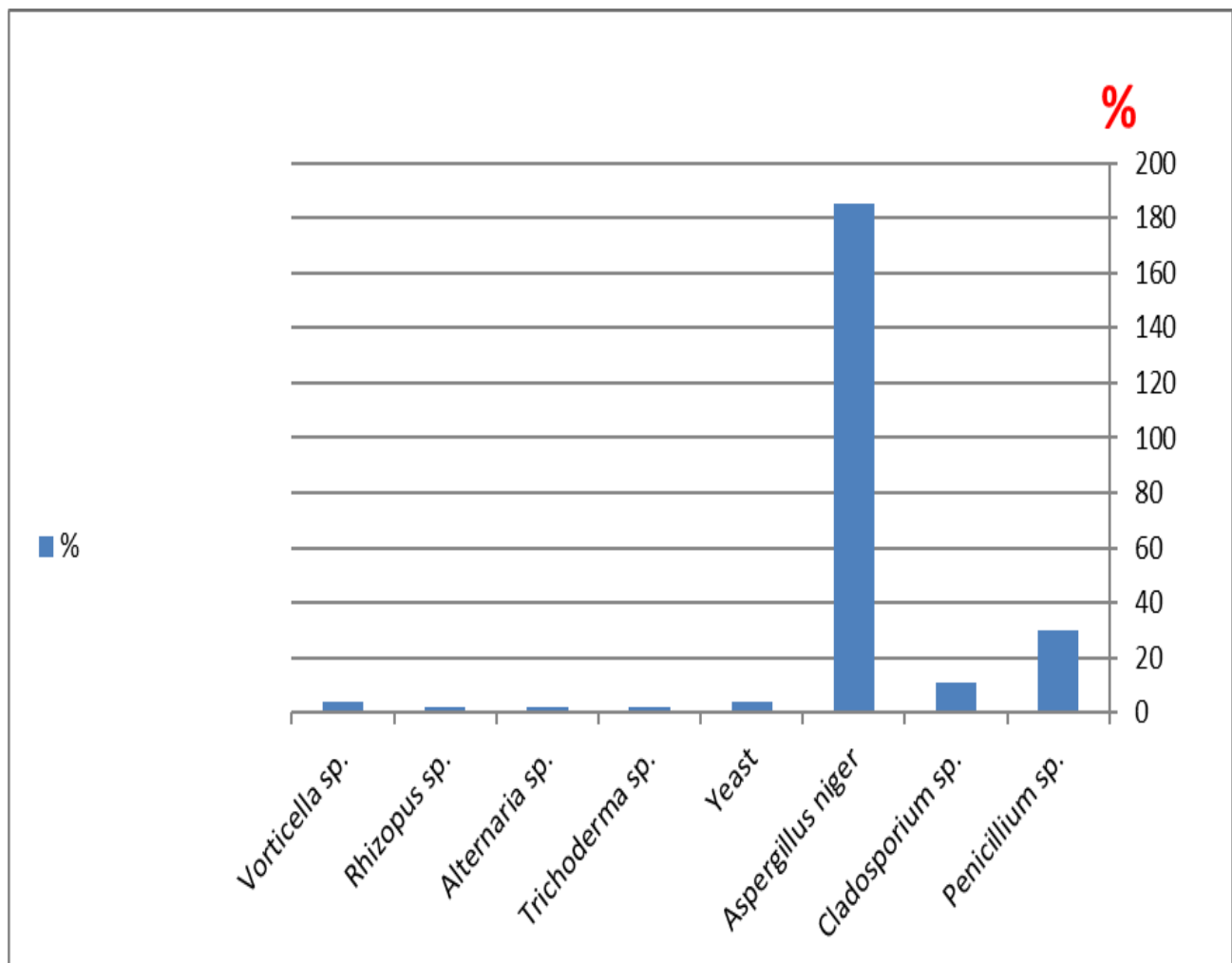
<i>Trichoderma sp.</i>	2	0.8
<i>Alternaria sp.</i>	2	0.8
<i>Rhizopus sp.</i>	2	0.8
<i>Vorticella sp.</i>	4	1.6
<i>Total</i>	240	100

Table 5: Total Fungi Count (CFU.Ml-1) In Second Sample

Fungi	Count	Percentage %
Cladosporium spp.	73	14
Alternaria	30	6
Aspergillus sp.	354	71
Candida sp.	1	0.2
Penicillium spp.	40	8
Total	498	100

Table 6: Total fungi count (CFU.ml-1) in third sample

Fungi	Count	Percentage%
Yeast	27	10
Penicillium sp.	66	25
Alternaria sp.	41	15
Cladosporium sp.	6	2
Aspergella Niger	120	45
Total	260	100

**Figure 1:** Shows fungi percentage in first sample

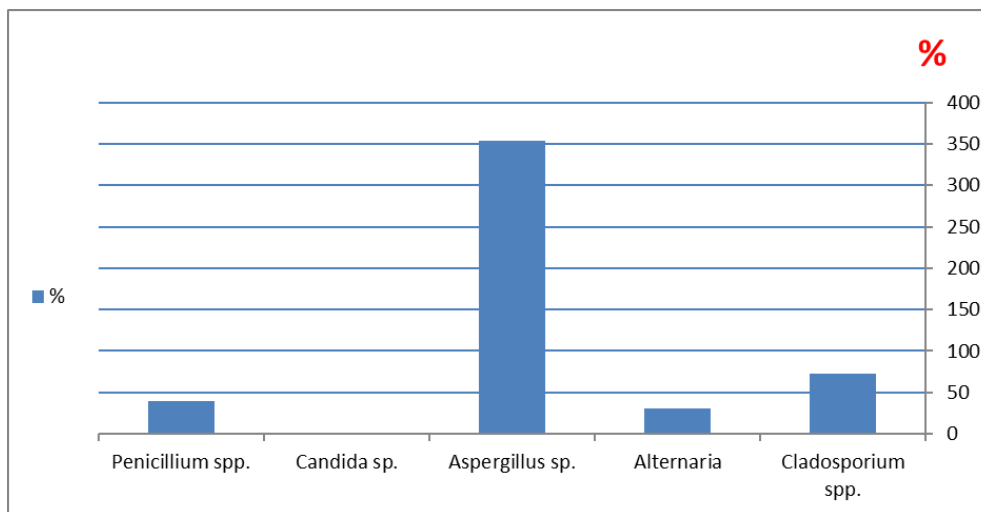


Figure 2: Shows fungi percentage in second sample

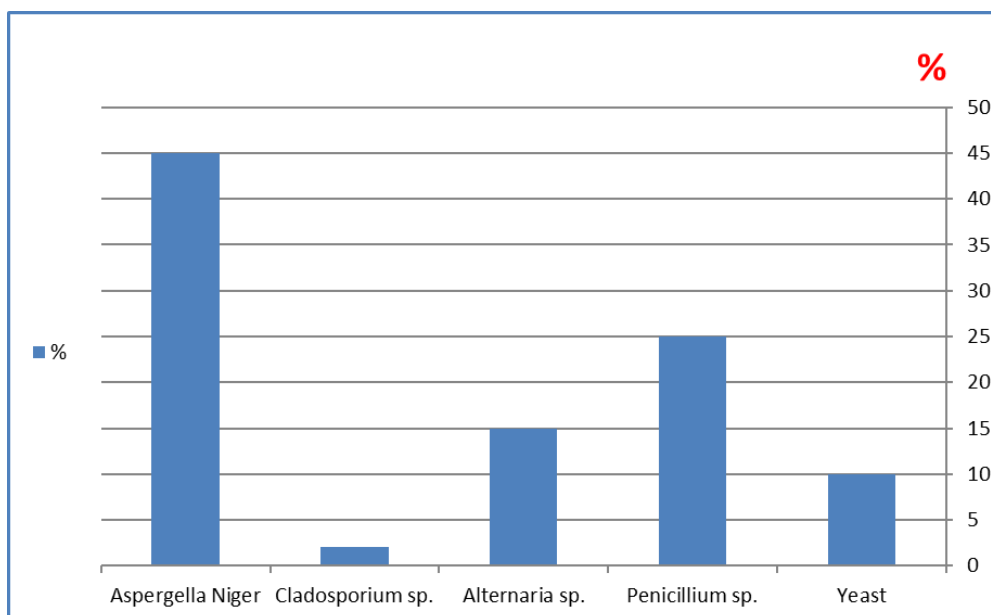
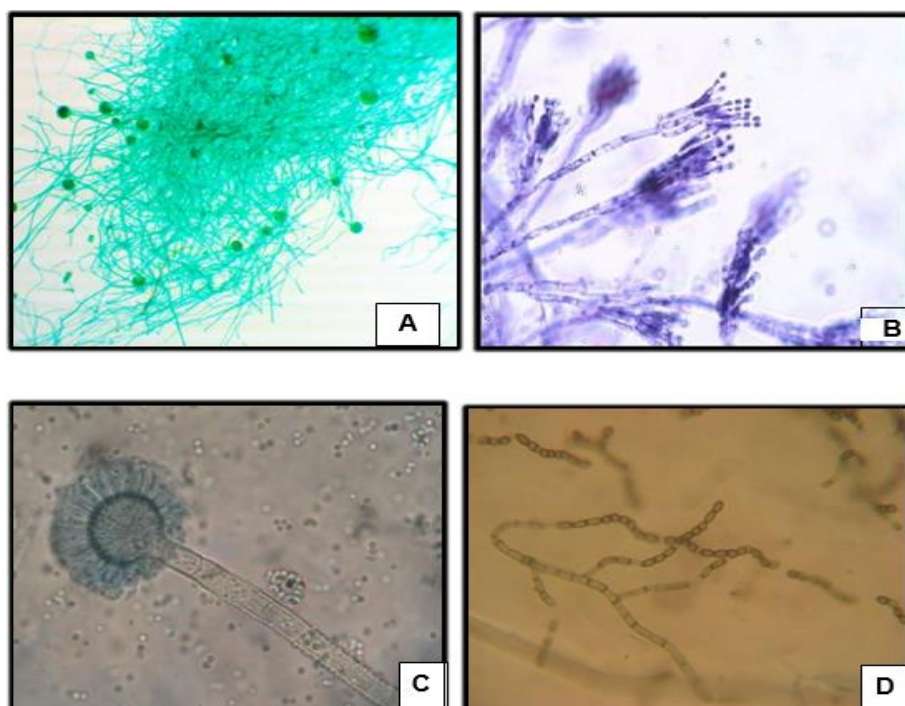


Figure 3: Shows Fungi Percentage In Third Sample



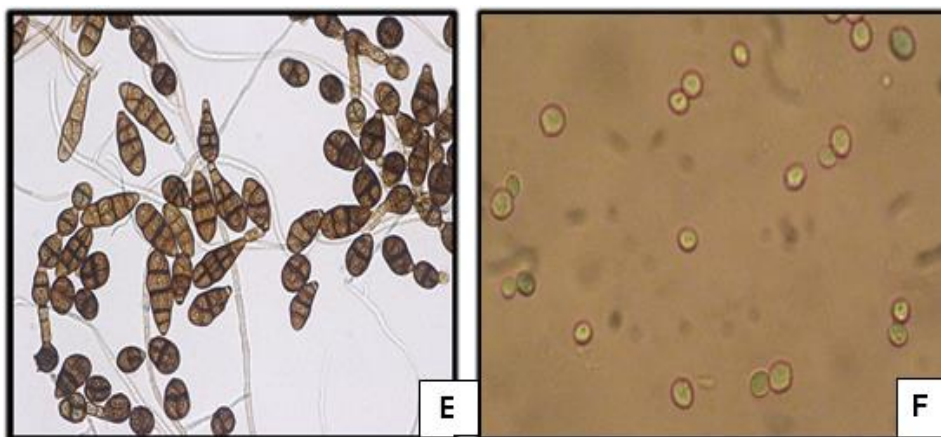


Figure 4: Shows Some Genera of Fungi Isolated On PDA Medium

A- *Rhizopus* sp. B- *Penicillium* sp. C- *Aspergillus* sp.

D- *Cladosporium* sp. E- yeasts F- *Alternaria* sp.

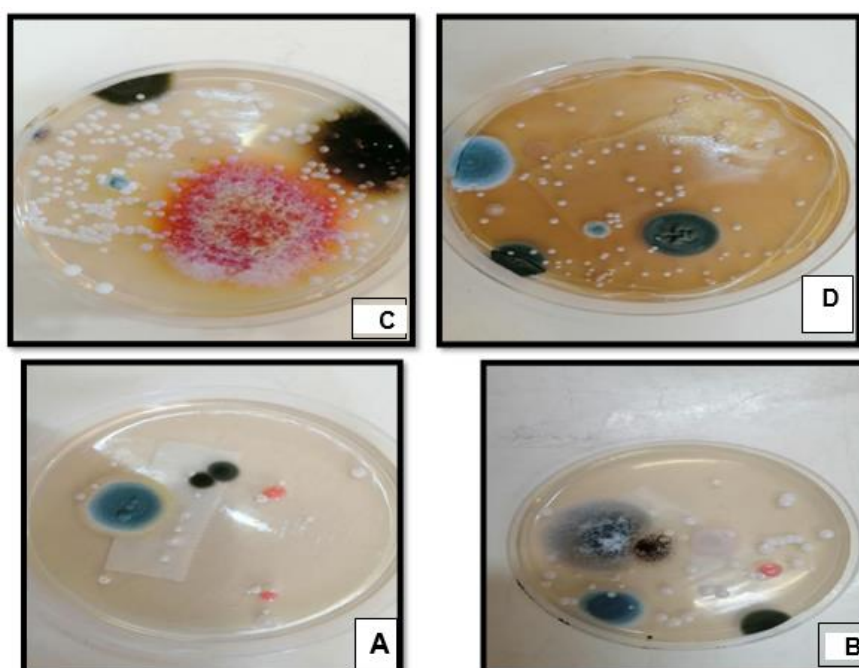


Figure 5 : Show different fungal colonies growth on PDA

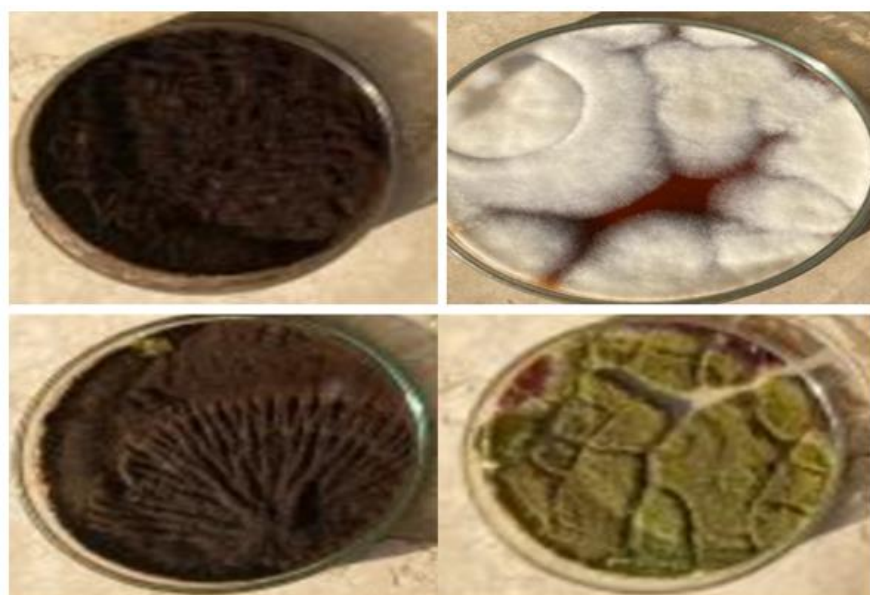


Figure 6: Show different fungal colonies (pure culture) growth on PDA

4. CONCLUSION

Several kinds of fungi were isolated from Salahaddin University's College of Science cafeteria. *Penicillium*, *Cladosporium*, *Alternaria*, and *Aspergillus* Sp were the most often isolated fungi. *Trichocladium*, *Drecheslera* sp., and *Botrytis* sp. were the less commonly found fungal species. Of all the *Aspergillus* species, *Aspergillus Niger* was the most prevalent. The fungi indicated that larger concentrations of fungi were isolated on chairs and containers.

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