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Preliminary Study on the Determination of Heat Resistant Fungi in Agricultural Soils

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Abstract: Heat resistant fungi are capable of surviving temperature at short heat applications. These organisms continue to growing and metabolic activities. Therefore, they cause spoilage of food products and effect on health because of mycotoxin production. This study focused on determination of density and biodiversity of heat resistant fungi in agricultural soil samples. For this aim, two different soil samples were collected from agricultural lands at June 2018. Isolation process was performed by using main soil dilution method after heat treatment at 75°C for 30 minutes of diluted soil samples. After incubation at 25°C for 7-14 days, all colonies were purified. All of the isolates were diagnosed by using conventional methods according to the macromorphological and micromorphological properties. We determined 11 species belong to Alternaria, Aspergillus, Cladosporium, Epicoccum, Humicola, Penicillium and Taloromyces genera. As a result of this study, we showed that agricultural soil areas have a potential for heat resistance fungi biodiversity and exhibited that we should concentrate on this subject with investigation of different areas and sources.

Key words: Agricultural Soils, Heat Resistant Fungi, Conventional Method

Tarım Topraklarındaki Isıya Dirençli Fungusların Belirlenmesi Üzerine Ön Çalışma

Öz: Isıya dayanıklı mantarlar, kısa ısı uygulamalarına dayanabilir. Bu organizmalar büyümeye ve metabolik aktivitelere devam ederler. Bu nedenle, gıda ürünlerinde bozulmaya ve mikotoksin üretme yetenekleri sayesinde sağlık üzerinde etkiye neden olurlar. Bu çalışma, tarımsal toprak örneklerinde ısıya dirençli mantarların yoğunluğu ve biyolojik çeşitliliğinin belirlenmesi üzerine odaklanmıştır. Bu amaçla, Haziran 2018'de tarım alanlarından iki farklı toprak örneği toplanmıştır. İzolasyon işlemi dilüe edilmiş toprak örneklerinin 75 °C'de 30 dakika boyunca ısı uygulaması yapıldıktan sonra toprak dilüsyon yöntemi ile gerçekleştirilmiştir. 25 °C'de 7-14 günlük inkübasyondan sonra tüm koloniler saflaştırılmıştır. İzolatların tamamı makromorfolojik ve mikromorfolojik özelliklere göre geleneksel yöntemler kullanılarak teşhis edilmiştir. Alternaria, Aspergillus, Cladosporium, Epicoccum, Humicola, Penicillium ve Taloromyces cinslerine ait 11 tür tespit edilmiştir. Bu çalışma sonucunda, tarımsal toprak alanlarının ısıya dirençli mantar biyolojik çeşitliliği potansiyeline sahip olduğunu belirledik ve farklı alanların ve kaynakların araştırılmasıyla bu konuya odaklanmamız gerektiğini gösterdik.

Anahtar kelimeler: Tarım Toprağı, İsiya Dirençli Fungus, Geleneksel Yöntem



Introduction

Soil; is a structure which physical and chemical properties including organic matter, minerals and large numbers of macro and microorganisms. This complex structure of the soil provides a suitable habitat for many microorganisms. The physicochemical structure of the soil, climate changes and agricultural applications have a significant impact on the biodiversity of soil (Chandrashekar et al., 2014).

Heat resistant fungi are microorganisms that resist short heat applications and whose main source is soil. These fungi are capable of surviving temperatures at or above 75°C for 30 or more minutes and can grow and cause spoilage during storage (Houbraken and Samson, 2006; Ali et al., 2009; Ibrahim et al., 2014). In addition to cause food spoilage, heat resistant fungi have direct



negative impact on human health because of mycotoxins, which are produced by these fungi (Demirci and Arıci, 2006).

Material and Method Site Description and Soil Sampling

The sampling area is an agricultural land located within the latitude 39°59'43.14"N to 32°20'39.09"E coordinates and 3 km away from Ayaş district of Ankara (Figure 1). Typical continental climate characteristics are observed in the region. During the year, various agricultural crops such as barley are cultivated. Two soil samples were collected with a sterilized trowel from 5 profiles in 0–10 cm depth at June 2018. Samples were transported in sterile plastic bags and stored at +4°C until analysed.



B)

Figure 1. Photograph of the work area (A) and the location on the map (B)

Isolation

A)

For isolation of the heat resistant fungi, 100 g of each soil sample was diluted in 150 ml sterilized distillate water in stomacher bag and homogenized by shaking. After homogenization, heat treat this stomacher bags at 75°C for 30 minutes in a water bath for inactivation of vegetative cells of fungi and bacteria. After the heat treatment, samples were cool to about 55°C. The contents of stomacher bag were transferred a sterile bottle (500 ml) and added 250 ml melted double strength Dichloran Glycerol Agar (DG18) medium. Sheerly mixed and poured into seven large petri dishes (diameter 12.0 mm). the petri dishes were incubated at 30°C for 7-14 days (Houbraken and Samson, 2006).

After incubation, colonies were counted and fungal number was calculated as CFU/g soil. Come into existence fungal colonies were subcultivated on Potato Dextrose Agar (PDA). Microscopic examination was

performed for the diagnosis of the genus level at the pure isolates that developed at 25°C incubation for 7 days.

Identification of isolates

Taxonomic identification of heat resistant fungi was evaluated according to their cultural characteristics and morphological structures. For identification of members of *Aspergillus* genus, inoculation process were performed on Malt Extract Agar (MEA), Czapek Yeast Extract Agar (CYA) (for 25 and 37°C), Creatine Sucrose Agar (CREA) media, incubated at 25°C for 7 days and investigated by using conventional method according to Klich (2002) and Samson et al. (2010). For members of *Penicillium* genus, MEA, CYA (for 25 and 30°C) and CREA media were used by incubation at 25°C for 7 days and investigated according to Samson et al. (2010). The other genera members were inoculated on PDA and MEA, incubated at



25°C for 7 days and investigated according to Barnet and Hunter (1999).

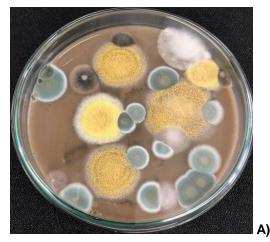
Result and discussion

The main objectives of this study were to determine the concentration and identification of heat resistant fungi in agricultural soil. Fungal counts and biodiversity in soil samples determined as 56 CFU/100 g soil (91.80 %) for sample A and 5 CFU/ 100 g soil (8.2 %) for sample B (Figure 2). The heat resistant fungi obtained in this study composed of 16 isolates belong to 7 genera that were encountered including Alternaria (1 species), Aspergillus (3), Cladosporium (2), Epicoccum (1), Humicola (1), Penicillium (2) and Talaromyces (1) according to their colony characteristics (Figure 3, 4). The highest fungal biodiversity was obtained from soil sample A as 75 % of total isolates (Table 1). The most common species belonged to the Aspergillus (3 species) genus and followed by Penicillium (2 species) and Cladosporium (2 species), respectively.

In this study, *Aspergillus* genus that attracts attention with widespread encounter is one of the most popular endophytic fungi. Endophytes live symbiotically in plants without causing significant diseases, involved in the biosynthesis of plant products and known as producer of many substances with potential use in the modern medicine, agriculture and pharmaceutical industry (Tawfik et al., 2017). Identified some members of *Aspergillus* genus as *A. glaucus* L. Link and *A. ochraceus* G. Wilh.commonly known as their mycotoxigenic properties

especially ochratoxin A producer (Valente et al., 2015; Moazam and Denning, 2017; Gil-Serna et al., 2018; Lim et al., 2019). In addition to some important members of Aspergillus genus, we determined 2 significant Penicillium genus members. One of them is P. aurantiogriseum Dierckx that causes contamination in foodstuffs such as olives, grains and especially on cereals (Bouhoudan et al., 2018; Neto et al., 2018). The other is P. griseofulvum Dierckx produces a mycotoxin named patulin which causes neurotoxicity, nephrotoxicity and hepatotoxicity (Benkerroum, 2016). Investigated soil samples were exhibited Cladosporium genus members rich profile. In previous studies, this genus was recorded important plant pathogen such as C. cladosporioides (Fresen.) G.A. de Vries (Nam et al., 2015) and mycotoxin producer and allergens such as C. herbarum (Pers.) Link (Hurrass et al., 2017, Sahib, 2019; Shuryak, 2019).

In addition to these common species, we obtained some isolates as *Alternaria* sp. that is pathogen on plant and human (Woudenberg et al., 2015; Patriarca, 2016; Tralamazza et al., 2018), *Epicoccum* sp. and *Humicola* sp. that known as important secondary metabolite (antioxidants, anticancers, and antimicrobial compounds) and bioactive compounds beside the saprophytic activities (Oliveira et al., 2018; Braga et al., 2018; Chadha et al., 2019; Wang et al., 2019), and *Talaromyces* sp. that is revised as new genus and include significant human pathogens (Chen et al., 2016).



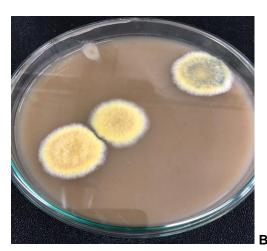


Figure 2. Colony perspective on DG18 (A) soil sample A, (B) soil samples B

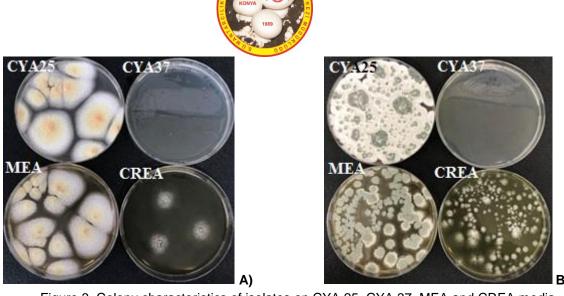


Figure 3. Colony characteristics of isolates on CYA 25, CYA 37, MEA and CREA media (A) isolate A5 (B) isolate B3

Table 1. List of the isolates and identification result

| Identification Result | Number of the Isolates | Soil Sample |
|--|---------------------------|-------------|
| Alternaria Nees | 1 | Α |
| Aspergillus chevalieri Thom & Church | 1 | Α |
| Aspergillus glaucus L. Link | 2 | В |
| Aspergillus ochraceus G. Wilh. | 2 | Α |
| Cladosporium cladosporioides (Fresen.) G.A. de Vries | 1 | Α |
| Cladosporium herbarum (Pers.) Link | 1 | Α |
| Epicoccum Link | 1 | Α |
| Humicola Traaen | 1 | В |
| Penicillium aurantiogriseum Dierckx | 4 | А |
| Penicillium griseofulvum Dierckx | 1 | А |
| Talaromyces C.R. Benj. | 1 | В |
| Total | 16 | |

A)

A)

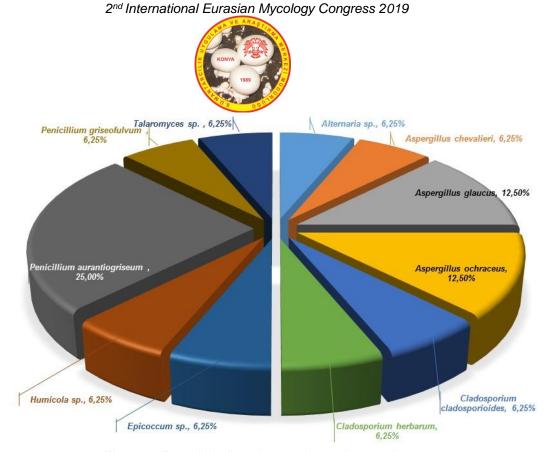


Figure.4. Fungal biodiversity together with annual average

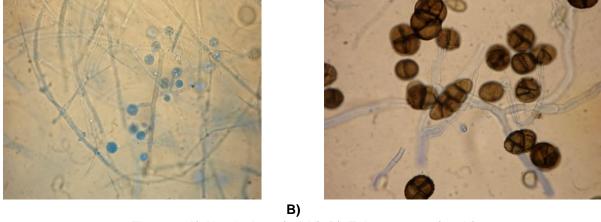


Figure 5. A) Humicola sp (100X), B) Epicoccum sp. (100X)

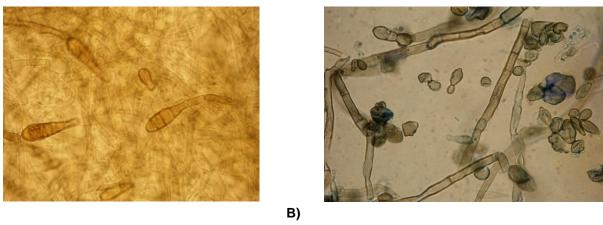


Figure 6. A) Alternaria sp (100X), B) Cladosporium herbarum (100X)



Conclusion

As results of this preliminary study, species belonging to the *Aspergillus* and *Penicillium* genus were commonly found to be heat resistant fungi in the investigated farmland. In particular, the presence of saprophytic, pathogenic and mycotoxigenic members were evaluated as potential threat to agricultural products

and consumers. Our results shown that agricultural soils contain heat-resistant fungi on significant level and their metabolic effects may have significant effects on the life cycle. Therefore, the density and diversity of heat-resistant fungi in different sources should be detailed with new studies.

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