# IDENTIFICATION AND QUANTIFICATION OF FUNGI ASSOCIATED WITH SEEDS OF BARLEY, IN TERMS OF 2014

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#### Abstract

Seeds are considered very effective means of plant pathogens transport over long distances. The use of seed material, free from pathogens, is one of the basic conditions for obtaining healthy crops. If infested seed is used for sowing, then the seed will cause diseases in crop and crop yields will be reduced considerably. The study aimed to identify and quantify the fungi associated with barley (Hordeum vulgare L.) seeds. We have analysed samples of seeds of barley varieties grown in the area of South-Eastern Romania. The mycoflora spectrum determined from analyzed samples consists of a variety of parasitic and saprophytic fungi. Species of Alternaria and Fusarium genera were the most frequently isolated with an incidence of 93.3% and 80% respectively. We also identified Curvularia lunata and Pyrenophora species with an incidence of 20% and 13.3% respectively. The group of saprophytic fungi like Oedocephalum sp., Penicillium sp., Stemphylium sp., Trichoderma viride and Epicoccum purpurascens, was also isolated.

Key words: mycoflora, seeds, barley, varieties.

# INTRODUCTION

Barley (Hordeum vulgare L.) and wheat (Triticum aestivum L.) are two of the most important cereal crops in the entire world. Generally the most important factor in disease transmission from one year to another is the seed (Beratlief and Oprea, 1994). Infected seeds are the most frequent and richest source of transmission of pathogens, especially fungi (Alexandri et al., 1967; Hulea, 1973). The quality as vegetal material of seeds reproduction has a great influence on the crops' vield (Clear and Patrick, 1993). Barley seeds are rich in proteins, carbohydrates and mineral substances which represent a good medium for the development of microorganisms. Knowing the pathology of the seed makes it possible to prevent the introduction of new pathogen agents in the barley crop as well as to avoid the apparition of massive spread of infection in the field (Chong and Sheridan, 1982; Bateman and Kwasna, 1999; Kubiak and Korbas, 1999; Gheorghies and Cristea, 2001). Pathogens transmitted by seed, and those who like a vector seed, are a great problem which needs to be dealt with taking into consideration various management strategies, including aspects concerning their biology (Bărbulescu and Popov, 1995). A clear identification and quantification of the pathogen transmitted through seeds is necessary in order to determine the means of control and the active substances which are to be applied (Cană et al., 2010). Association of several fungal, in some cases, can induce the black-point phenomenon characterized by blackening of the embryo (Cristea and Berca, 2003).

## MATERIALS AND METHODS

The study aimed to identify and quantify the associated fungi of barley (*Hordeum vulgare* L.) seeds. We have analized samples of barley seeds varieties grown in the area of South-Eastern Romania.

The samples which were examined were collected from various lots from local producers at the moment of harvest, in the climatic conditions of 2014. After harvesting, the samples were packed for transportation and storing them in order to maintain the physico -

chemical and microbiological analyzes up to the time of the laboratory in order to obtain more accurate results (Petcu, 2014). The biological material was represented by 15 barley samples such as: Alora baza, Amical, Andreea, Andrei, Cardinal, Compact, Friderichus, Hanzi baza, Maresal, Madalin, Orizont, Regent, Scarpia, Univers, Vanessa.

The barley seeds were placed in Petri dishes with a diameter of 8 cm on PDA culture medium (potato-dextrose-agar), sterilized at 121°C/20 min). The culture medium was prepared after the classic recipe (Hulea, 1969; Constantinescu, 1974). For each variety was analyzed a total of 100 seeds arranged in three variants each with three repetitions. The seeds have been not disinfected before this stage because we want to estimate the presence of pathogens on seeds surface not into seeds tissues. It had been determined the frequency of affected seeds by the identified micromycetes. The dishes were incubated at a temperature of 22°C for 10 days. The identification of micromycetes was performed with a Zeiss Primo Star microscope based on scientific literature (Raicu and Baciu, 1978; Hulea and Iliescu, 1986).

# **RESULTS AND DISCUSSIONS**

The spectrum of identified pathogens includes fungi belonging to the Ascomycotina and Deuteromycotina, consisting of species of the genus \_ Alternaria. Curvularia lunata. purpurascens, Fusarium, Epicoccum Oedocephalum, Penicillium, Pyrenophora, Rhizopus, Stemphylium and Trichoderma viride (Table 1). The micromycetes Alternaria sp., Penicillium sp., Rhizopus sp., and Fusarium sp. were also detected on the seeds of other plant species in Romania (Cozea et al., 2011; Zala et al., 2011; Mardare et al., 2014; Pana et al., 2014).

It is noted that on the surface of the seeds, the *Alternaria* species were the most frequently 93.3% (Figure 1), the analyzed varieties developing colonies, except the Scarpia variety. Also, there was a high incidence of *Fusarium* species, which considered to be higly important pathogens for the seeds of barley; 80% of the species studied presented spores that developed

colonies of the genus (*F. graminearum*, *F. moniliforme*), except the Andreea, Maresal and Regent varieties.

*Epicoccum purpurascens*, was detected with a 60% incidence, *Rhizopus* sp. and *Stemphylium* sp. with 40% incidence, are also part of the saprophytic determined.

The *Pyrenophora* species were detected in the Andreea, Orizont and Univers varieties, with 20% incidence. *Penicillium* species were identified in 33.3% of the analyzed varieties (Compact, Maresal, Madalin, Scarpia, Univers) and the *Oedocephalum* sp. in 20% (Amical, Friderichus and Madalin). In Andreea and Cardinal varieties, *Curvularia lunata* and *Trichoderma viride* saprophytes were determined in 13.3% of the cases.

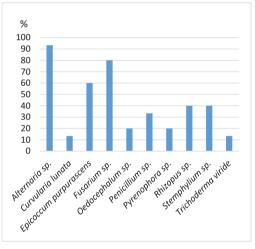


Figure 1. Incidence of pathogens spectrum on the seeds of barley analysed

Figure 2 presents aspects of the fungal colonization on barley seeds, in laboratory conditions on PDA culture medium.

Similar results were reported by other authors in the field of study of wheat seeds pathology (Cristea et al., 2008) as well as barley seeds pathology (Valceva, 2006; Karadjova, 1979 and Ivanov, 1980, cited by Georgieva-Andreeva al, 2011, Cristea-Manole et al, 2015), who have found infections similar to those of the fungi in the lots of seeds from the north of Bulgaria and southern Romania.

Table 1. Identified mycoflora in barley seeds

	Variety														
Microscopic fungi	Alora baza	Amical	Andreea	Andrei	Cardinal	Compact	Friderichus	Hanzi baza	Maresal	Madalin	Orizont	Regent	Scarpia	Univers	Vanessa
Alternaria spp.	х	х	х	х	х	х	х	х	х	х	х	х	-	х	х
Curvularia lunata	-	-	х	-	х	-	-	-	-	-	-	-	-	-	-
Epicoccum purpurascens	х	-	-	-	х	-	Х	х	х	х	-	х	Х	-	Х
Fusarium spp.	х	х	-	х	х	х	х	х	-	х	х	-	х	х	Х
Oedocephalum sp.	-	Х	-	-	-	-	х	-	-	х	-	-	-	-	-
Penicillium sp.	-	-	-	-	-	х	-	-	х	х	-	-	х	х	-
Pyrenophora sp.	-	-	Х	-	-	-	-	-	-	-	х	-	-	х	-
Rhizopus sp.	-	-	х	х	х	-	-	-	-	-	х	х	Х	-	-
Stemphylium sp.	-	-	-	-	х	х	-	х	-	х	-	-	Х	х	-
Trichoderma viride	-	-	х	-	х	-	-	-	-	-	-	-	-	-	-

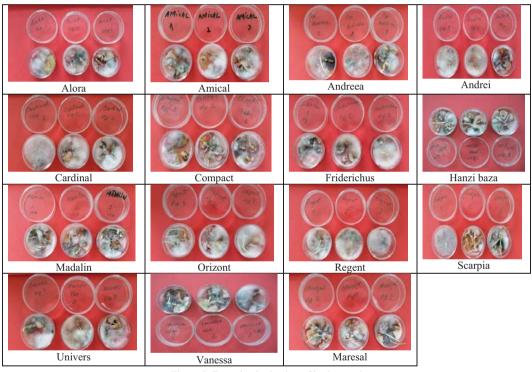


Figure 2. Fungal colonization of barley seeds

The varieties examined showed a variable number of fungi, which is between 3 and 7 pathogens. The lowest number of pathogens was identified Alora baza, Amical, Andrei, Maresal, Orizont and Vanessa varieties, and most pathogens were present on the seeds of the variety Cardinal (Figure 3).

# CONCLUSIONS

Our research confirmed the presence of a large number of fungi associated barley seeds.

The fungi isolated from the seeds barley were included in the genera: *Alternaria, Curvularia, Epicoccum, Fusarium, Oedocephalum, Penicillium, Pyrenophora, Rhizopus, Stemphylium* and *Trichoderma.* 

The most frequent pathogenic species isolated belong to the genus *Alternaria* and *Fusarium* and they colonized the barley seeds in most of the varieties studied, with an incidence of 93.3% and 80% respectively.

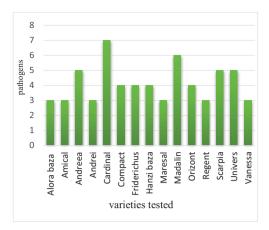


Figure 3. Fungi at seed level

### ACKNOWLEDGEMENTS

This paper was published under the frame of European Social Fund, Human Resources Development Operational Programme 2007-2013, project no. POSDRU/159/1.5/S/132765.

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