## Fungal root endophytes of Northern Kazakhstan isolated from agricultural and non-agricultural crops

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Endophytes are microorganisms that colonize plants without any visible symptoms of disease and might produce a plethora of secondary metabolites to prevent plants from various phytopathogens [4]. Root colonizing fungal endophytes, the so called dark septate endophytes (DSE) are world-wide common in both agronomical areas and natural habitats and can enhance the growth of host plants and provide resistance to plant diseases, drought stress and enhance plant performance [3, 5]. Hence, they may play important role in ecosystem functioning and can have major importance in agricultural applications. However, our knowledge on DSE communities is limited in Kazakhstan.

In this study, we aimed to identify the DSE community of different plant species agricultural and non-agricultural areas in the steppe zone of Northern Kazakhstan.

Roots were collected from the gramineous species barley ( $Hordeum\ vulgare$ ), wheat ( $Hordeum\ jubatum$ ), oat ( $Avena\ sativa$ ),  $Stipa\ capillata$ , and also lentil ( $Lens\ culinaris$ , Fabaceae) and flax ( $Linum\ usitatissimum$ , Linaceae). The sampling was carried out in 2018-2019. The surface sterilization of plant roots was conducted by placing them in  $H_2O_2$  for 90 sec, then 90 sec in ethanol 70% and twice washed in distilled  $H_2O$  [2]. Then, root pieces were placed onto PDA media and the hyphae growing out were subcultured [1]. For molecular identification, DNA isolation was carried out by NucleoSpin Plant II Kit and the internal transcribed spacer (ITS) region of the nrDNA was amplified and sequenced. In case of Fusarium species, the translation elongation factor 1-alpha (TEF). The sequences gained were compared with those of public databases, and phylogenetic analyses were used for identification of DSE species [6].

Altogether 605 isolates were collected from roots of different agricultural and non-agricultural plants in Northern Kazakhstan. The isolates represented Fusarium spp., Alternaria sp., Bipolaris sp., Aspergillus sp., Nigrospora oryzae, Cladosporium sp., Talaromyces purpureogenus, Periconia macrospinosa, Acremonium sp., Crinipellis scabella, Microdochium bolleyi, Cadophora sp., Phaeosphaeria sp., Rhizoctonia sp., Epicoccum nigrum, Clonastachus rosea, Sarocladium strictum, Arthrinium arundinis, Slopeiomyces cylindrosporus, Chaetomium sp. and Darksidea sp. The most common and widespread genus was Fusarium, which was represented by at least 15 species. Therefore, Fusarium species such as F. redolens, F. culmorum and F. clavum dominate the DSE community of the investigated plant species in Northern Kazakhstan. We found diverse fungal community in the sites and the major presence of DSEs in this area might be indicating an important function for the plants in defence mechanisms due to the severe climate of Northern Kazakhstan.

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