

Studies on occurrence of Rhizospheremycoflora of Jatropha curcas L.

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I. Introduction

Jatropha curcas L is the most important biofuel plant belonging to Euphorbiaceae family. Among plant evaluated for the extraction of biodiesel from its seeds, Jatropha curcas has been found to be most promising. Several parts of Jatropha plant have medicinal and cosmetic uses also. In India, Jatropha curcas is gaining importance commercially as biodiesel plant. It was found that performance of Jatropha seed oil based biodiesel was superior among the other biodiesel derived from different vegetable oils (Lutz, 1992; Pak and Alexi, 1994). Jatropha seeds contain 30-35% viscous non edible oil that has attracted the attention of the world as an alternate fuel (Takeda, 1982; Banerji et al., 1985; Martin and Mayenx, 1985; Openshaw, 2000).

The original definition of the rhizosphere means the zone of soil in which the microflora is influenced by plant roots (Rovira, 1965). A series of reports have suggested that microbial community structure in the rhizosphere are dependent upon the type of plant species (Germida et al., 1998) including the genotypes within a species.

II. Materials & Methods

The rhizospheremycoflora of Jatropha curcas was studied during March 07 to Feb 08. The rhizosphere soil samples were collected from the Jatropha plantation site monthly for a year during March 2007 to February 2008. Soil particles adhering to the roots of Jatropha curcas were used as the study materials. Enumerations of rhizospheremycoflora were done by serial dilution technique (Harley and Waid, 1955). Martin's agar medium supplemented with Rose Bengal was used to isolate the rhizosphere fungi of Jatropha curcas. The petri plates were incubated at 27±2°C for a week.

III. Results & Discussion

During investigation period 22 species belonging to 11 genera of fungi were recorded and identified as Aspergillus cervinus; Aspergillus flavus; Aspergillus niger; Aspergillus terreus; Chaetomella raphigera; Cladosporium cladosporioides; Cladosporium oxysporum; Curvularia lunata; Curvularia pallescens; Emericella nidulans; Fusarium equiseti; Fusarium solani; Monodictyscastraneae; Monodictys fluctuata; Paecilomyces lilacinus; Penicillium aurantiogriseum; Penicillium citrinum; Penicillium thomii; Rhizopus oryzae; Rhizopus stolonifer; Trichoderma hamatum and Trichoderma reesei (Table-1). A total of 1220 colonies of rhizosphere fungi were isolated. Maximum number of colonies were isolated during Rainy Season (757) followed by summer (323) and winter season (140). Colonies of Aspergillus niger was isolated throughout the study periods. Maximum number of colonies of A. terreus was isolated (317) during investigation period followed by Paecilomyces lilacinus (309) and Penicillium aurantiogriseum (207). Monthly analysis of the rhizosphere fungi showed that highest 109 colonies of Paecilomyces lilacinus were isolated in the month of August followed by Aspergillus terreus (108) in Sept. Cladosporium oxysporum was present only in the month of Feb 08 with its highest (45) number of colonies. Thus many pathogenic, non-pathogenic, saprophytic and antagonistic fungi were isolated from the Rhizosphere of Jatropha curcas. Two species of the most important antagonistic fungus Trichoderma were also present.

Several fungal and bacterial biocontrol agents have been used for achieving plant disease control, amongst them Trichoderma group has been found effective against aerial, root and soil pathogens (Weller, 1988; Kumar and Mukerji, 1996; Van Loon et al., 1998; Whipsett et al., 1993; Elad et al., 1998 a, b and 2000; Chaube et al., 2002; Harman et al., 2004). Trichoderma is a potent fungal biocontrol agent against a range of plant pathogen has attracted considerable scientific attention (Rini and Sulochana, 2007). Along with other factors, presence of Trichoderma species in the rhizosphere of Jatropha curcas may be one of the cause for its disease resistant nature.

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