



Degradation of Pyrazosulfuron Ethyl in Rice Ecosystem by *Aspergillus terreus*

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Authors' contributions

This work was carried out in collaboration among all authors. Author SSM designed the study, performed the statistical analysis, wrote the protocol and wrote the first draft of the manuscript. Authors PPC and SRM managed the analyses of the study. Author AR managed the literature searches and edited for publication. All authors read and approved the final manuscript.

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ABSTRACT

Investigation on the degradation of pyrazosulfuron ethyl by *Aspergillus terreus* in the rice ecosystem was conducted at the Department of Biochemistry, UBKV, Pundibari Coochbehar, West Bengal, India. The soil was collected from the paddy field to isolate fungal inoculums. Results revealed that *Aspergillus terreus* can survive in minimal broth containing pyrazosulfuron ethyl at 1000 ppm and degrades through two major pathways, first involves the cleaves of sulfonylurea bridge resulting in the formation of two major metabolites viz., 2-amino-4, 6 –dimethoxyprimidine, 5-aminosulfonyl-1-methyl-pyrazole-4-carboxylic ethyl ester and, second was the cleavage of sulfonyl amide linkage which forms the metabolite viz; 1 methyl pyrazole-4-carboxylic acid ethyl ester, 1 methyl pyrazole -4-carboxylic acid, 5-carbamoyl -1-methyl pyrazole -4-carboxylic acid ethyl-ester. The enzyme involved in these transformations can be utilized to decontaminate soil and water from Pyrazosulfuron ethyl residue. Even, the gene responsible for the production of these useful enzymes can be exploited for future research.

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1. INTRODUCTION

Sulfonylureas are new class of herbicides characterized by low application rate, having high herbicidal activity, broad action spectrum, low toxic, used on wide range of crops, such as rice, wheat, barley, corn, cotton, potato, and soybean [1,2,3]. The application of these herbicides kill the weeds in 3-5 week [4]. Pyrazosulfuron ethyl is a popular recent origin Sulfonylureas class of herbicides, widely used in rice for pre-and post-emergent control [5] of weeds. Due to frequent use of these herbicides, a majority of these chemicals entered in to soil to form short or long term residues and resulted in phytotoxicity to sensitive crops [6]. There are few studies which report adverse effect of sulfonylureas to soil microbial population. [7]. Like other herbicides sulfonylurea also degrades in soil through microbial degradation [8,9,10,11]. Although pyrazosulfuron are reported to be fast degraded in the soil [6] but its frequent use showed adverse effect on paddy production and enhanced risk to environment and human health [6]. There are few reports on bacterial degradation of Sulfonylureas in to soil [7,12]. Degradation of chlorsulfuron and metasulfuron methyl by *Streptomyces griseolus* bacteria [13], [14] and Trisulfuron by *S. griseolus* [15] have been well established. Like bacteria, fungal species are also found to be efficient to degrade phenylurea herbicides, but, a little study has been done on Pyrazosulfuron -ethyl degradation by fungi in the context of the ecological risk of its long term application. *Aspergillus terreus* is considered as an important pyrazosulfuron ethyl degrading fungi [16]. In this article, we described the isolation of the new Pyrazosulfuron ethyl degrading fungi, *Aspergillus terreus* and studied the Pyrazosulfuron ethyl degradation in contaminated soil. The metabolites formed during the biodegradation of Pyrazosulfuron ethyl were also analyzed.

2. MATERIALS AND METHODS

2.1 Chemical

A sample of technical grade Pyrazosulfuron ethyl (90.5%) and analytical Pyrazosulfuron ethyl (99.9%) were obtained from United Phosphorus Limited Company, Mumbai. Technical grade Pyrazosulfuron was further purified by recrystallisation. The purity was checked by Thin

layer chromatography (TLC) (solvent system - chloroform: acetonitrile = 2: 3, v/v, $R_f = 0.35$). It is a white amorphous solid. Laboratory grade reagents and solvents were locally procured, purified and used. All the solvents were distilled and dried before use.

2.2 Equipments

Autoclave, BOD incubator, laminar air flow, mechanical shaker, pH meter, vortex mixture, compound microscope, thin layer chromatography (TLC) applicator, Liquid chromatography with tandem mass spectrometry (LC-MS/MS), gas chromatography-mass spectrometry (GC-MS/MS) and Vacuum drier.

2.3 Preparation of Soil Suspension

A sample of 1.0 g of soil collected from rice field of research farm, Uttar Banga Krishi Viswavidyalaya, Pundibari, Coochbehar, North Bengal, was suspended in 100 ml distilled water in a conical flask. The flask was thoroughly shaken to make uniform soil suspension to use as inoculums for soil fungi.

2.4 Isolation of Pyrazosulfuron Degradative Microorganisms from Soil Samples

Minimal media (100 ml) of fungi was taken in 250 ml sterilized conical flask. The minimal media (a culture medium for microorganisms that contains the minimal necessities for growth) is prepared with one fifth of the strength of original fungal media. Required quantity of Pyrazosulfuron ethyl was added to minimal broth media in order to get 100 ppm, 250 ppm, 500 ppm, 1000 ppm of active ingredient. Similarly, distilled water containing mineral solution was also taken in conical flask and pyrazosulfuron ethyl was added to get final concentration of 1000 ppm. From each category, 1.0 ml of soil suspension was added to each flask as inoculums. All the flasks were incubated at room temperature for a long period (65 days) with a view to enrich the herbicide tolerant microorganisms. For isolation of Pyrazosulfuron ethyl tolerant/transforming microorganisms, enriched culture from each flask was streaked on media of fungi amended with 100 ppm pyrazosulfuron ethyl [17]. The colony grown in 1000 ppm conc. of pyrazosulfuron ethyl in water

plus minerals media was considered as resistant one for fungi. It was taken out after 65 days, purified by serial dilution and stored for taxonomic identification.

2.5 Identification of Isolated Fungus from Soil

For fungal identification used stain such as lacto phenol/cotton blue mounting procedure: A drop of lacto phenol or cotton blue was placed on a clean, dust free slide. A small tuft of the fungus preferably with spores and spore bearing structures was transferred onto the drop under aseptically environment and was gently teased by the sterile mounted needle. After proper mixing of the fungal material a cover-glass was placed over the preparation taking care to avoid trapping air bubble in the stain.

2.6 Degradation of Pyrazosulfuron ethyl by Isolated Fungus and Isolation of Metabolites in Broth Media

1ml broth media of the isolated soil fungi was transferred to 100 ppm. Pyrazosulfuron ethyl conc. in a conical flask of fungi media for incubation at room temperature. The growths of the soil fungus in the above flask were checked by drowning 1 ml from it and pour plating in petriplate. In order to assess further tolerant of the fungus to Pyrazosulfuron ethyl 1ml of broth was drown from the 1st flask (100ppm PSE conc.) and transferred to flask of 250 ppm pyrazosulfuron ethyl concentration. It gives growth visualization in 30 days. Similarly the flasks which carry 500 & 1000 ppm pyrazosulfuron ethyl were also inoculated. These flasks were incubated up to 50 days and 21 days, respectively. The growth of the fungus bacteria inoculated was periodically checked by visual observation and pour plating. A fungus grown in 1000 ppm conc. of pyrazosulfuron ethyl in water plus minerals media was considered as resistant one. It was taken out after 21 days. During these phenomena pyrazosulfuron ethyl was degraded in to several components. This component was studied by chromatographic technique.

2.7 Component Separation of Fungal Degraded

Growth minimal media of 100 ppm, 250 ppm, 500 ppm and 1000 ppm water medium in fungus of each soil samples were filtered separately.

The bunch of microbe was drowning out. Balance solution was extracted first with chloroform solvent then ethyl acetate solvent by gravity separator funnel. Pyrazosulfuron ethyl component were soluble in the solvents. Similarly ethyl acetate extract was also extracted by gravity separation method to finding the degraded product of metabolites.

2.8 Extraction and Characterization of Metabolites from Broth and Soil Product

Degraded products were extracted from broth in different time interval by partitioning in chloroform. The incubated soil was also extracted in ethyl acetate by continuous shaking for 4 hours and filtered. Solvent was then evaporated under low pressure in rotary vacuum evaporator to obtain a crude mixture of products. Products were purified and characterized by the following chromatographic and spectroscopic techniques.

2.9 Chromatography

2.9.1 Preparative thin layer chromatography

Silica gel based TLC plates were prepared by spreading a slurry of silica gel G containing 10 per cent binder (gypsum) and 1g fluorescence in water on 20cm X 20cm glass plates, uniformly maintaining a thickness of 0.75mm using a Thin layer chromatography (TLC) applicator. Prepared plates were air dried first and then activated at 120°C for 2 h. The sample solutions were spotted on the TLC plates using capillary tubes. Spots were visualized under UV light, marked and scrapped. The scrapped silica gel was extracted with mobile phase solution (40% acetone in Hexane).

2.9.2 Liquid chromatography mass spectroscopy

An API 3200 Qtrap mass spectrometer hyphenated to Shimadzu UFLC was used to mass characterization of degraded products. Mass spectrometric analysis was performed with electro spray ionization (ESI) in positive (5500 eV.) mode for each sample. The nebulizer gas and heater gases were adjusted at 30 psi and 55 psi, respectively. The ion source temperature was set at 500°C. Each sample was injected by infusion technique at the rate of 10 μLs^{-1} .

2.9.3 Gas chromatography mass-mass spectroscopy

Varian Saturn 2200 GC/MS with the CP-3800 and 2000 Series Ion Trap MS was used for the separation and characterization of degraded products.

3. RESULTS AND DISCUSSION

3.1 Isolation and Characterization of Soil Fungus

The media incubated with Pyrazosulfuron ethyl showed a rapid growth in colonies of fungus, appeared as a brown cinnamon with various shades. Microscopic observations (X 100) showed columnar like structure having various conidiophores, with smooth and colorless hyalines. Conodia was measured 1.5-2.3 μm diameter by ocular microscope and appeared as slightly elliptical and strait. A special property was seen in this genera that globose was stitch on hyphae. Thus from above characters it was concluded that the isolates of farm soil, contained the fungus *Aspergillus terreus* [18,19].

3.2 Degradation of Pyrazosulfuron-Ethyl by *Aspergillus terreus*

Isolated fungus *Aspergillus terreus* degrades the Pyrazosulfuron ethyl in incubated media, formed five metabolites, designated as compound F₁, F₂, F₃, F₄ & F₅ (analyzed by GC-MS/MS and LC-MS/MS and characterized by their mass ions and fragmentation patterns). Pyrazosulfuron ethyl degrades through two major pathways, first involves the cleavage of sulfonyleurea bridge resulting in the formation of two major metabolites viz; 2-amino-4,6-dimethoxy-pyrimidine (F₅), 5-aminosulfonyl-1-methyl-pyrazole-4-carboxylic ethyl ester (F₁) and, second was the cleavage of sulfonamide linkage which forms the metabolite viz; 1-methyl pyrazole-4-carboxylic acid ethyl ester (F₂), 1-methyl pyrazole-4-carboxylic acid (F₃) and 5-carbamoyl-1-methyl pyrazole-4-carboxylate (F₄). The metabolites formed was separated and characterized by GC-MS/MS and mass characterization was performed by LC-MS/MS through mass ions and fragmentation patterns which are given below:

Compound F₁: m/Z 206 with the Compound F₂ showed a molecular ion peak (MH⁺) at m/Z 234. The loss of -C₂H₄ radicals from the molecular ion peak gave a peak at m/Z 206. A fragmentation at m/Z 188 was generated from elimination of molecular H₂O. Hence from the mass fragmentation pattern the structure of F₁ may be elucidated as 5-aminosulfonyl-1-methyl pyrazole-4-carboxylic acid ethyl ester. This suggested that the primary step of the degradative route is the cleavage of sulfonyleurea bridge (Fig. 1). According to previous reports, the cleavage of sulfonyleurea bridge is the common degrading route of sulfonyleurea herbicides as reported for the degradation of sulfonyleureas by bacteria and fungi [20,21].

Compound F₂: F₂ showed a molecular ion peak (M⁺) at m/Z 154 together with diagnostic peak at m/Z 139, 125, 109 and 81 by the elimination of radicals -CH₃, C₂H₅, -OC₂H₅, and CO₂C₂H₅ respectively from (M⁺). Hence considering the above mass fragmentation pattern compound F₂ to be assigned as 1-methyl pyrazole-4-carboxylic acid ethyl ester (Fig. 2).

Compound F₃: Molecular ion peak of F₃ (M⁺) at m/Z 126 and other two fragments m/Z 109 (M⁺ - OH), m/Z 81 (M⁺ - COOH) revealed that the structure of F₃ may be elucidated as 1-methyl pyrazole-4-carboxylic acid (Fig. 3).

Compound F₄: This compound showed a molecular ion peak (M⁺) at m/Z 197. The ionic fragment at m/Z 182 could be resulted through the loss of NH₂ radical and addition of hydrogen radical from M⁺. A peak at m/Z 153 was formed by the elimination of -COH radical from m/Z 182. The peak at m/Z 81 was formed from m/Z 153 through the simultaneous elimination of -CH₃, (m/Z 138), CH₂ (m/Z 124), -O (m/Z 108) and -CO (m/Z 81). Finally at peak at m/Z 68 was formed by the elimination of CH₃ radicals from m/Z 81. Analysis of all the fragments the structure of F₄ was assigned as ethyl 5-carbamoyl-1-methyl-1-pyrazole-4-carboxylate (Fig. 4).

Compound F₅: The molecular peak at m/Z 156 (MH⁺) of F₅ may be assigned as 2-amino-4,6-dimethoxy-pyrimidine by the analysis of mass spectral data m/Z 140 (MH⁺ - NH₂), m/Z 124 (m/Z 140 - CH₄ radical) and m/Z 100 (elimination of 24 atomic mass unit), Fig. 5.

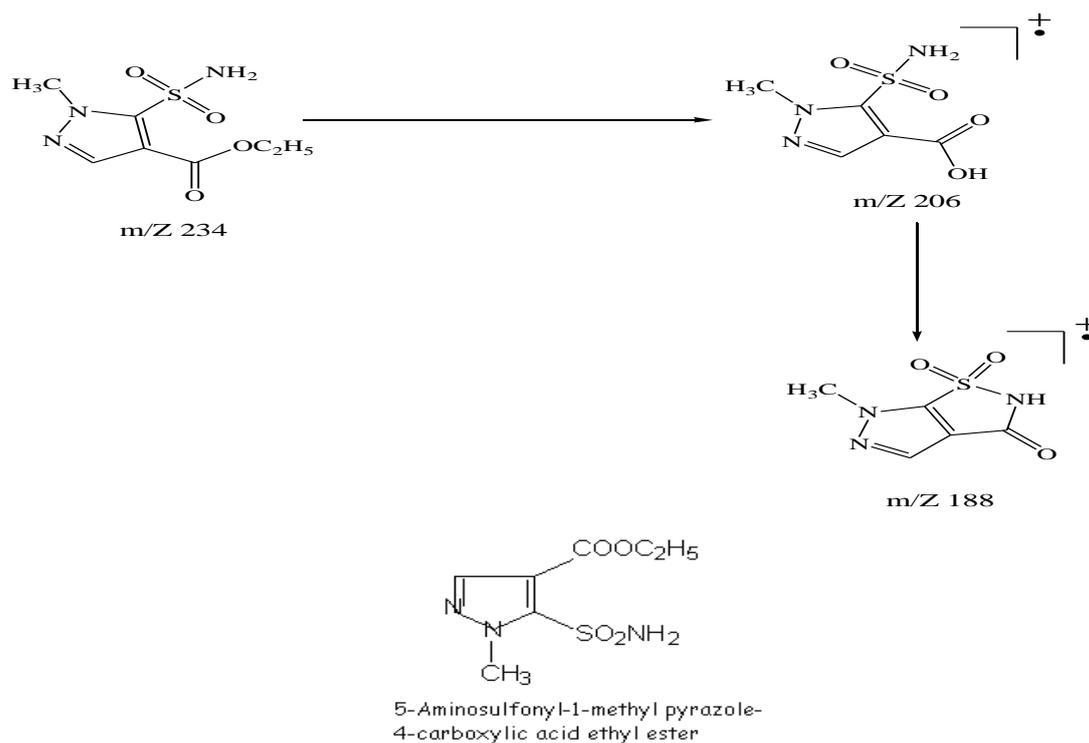


Fig. 1. Mass fragmentation pattern of compound F₁ with elucidated name

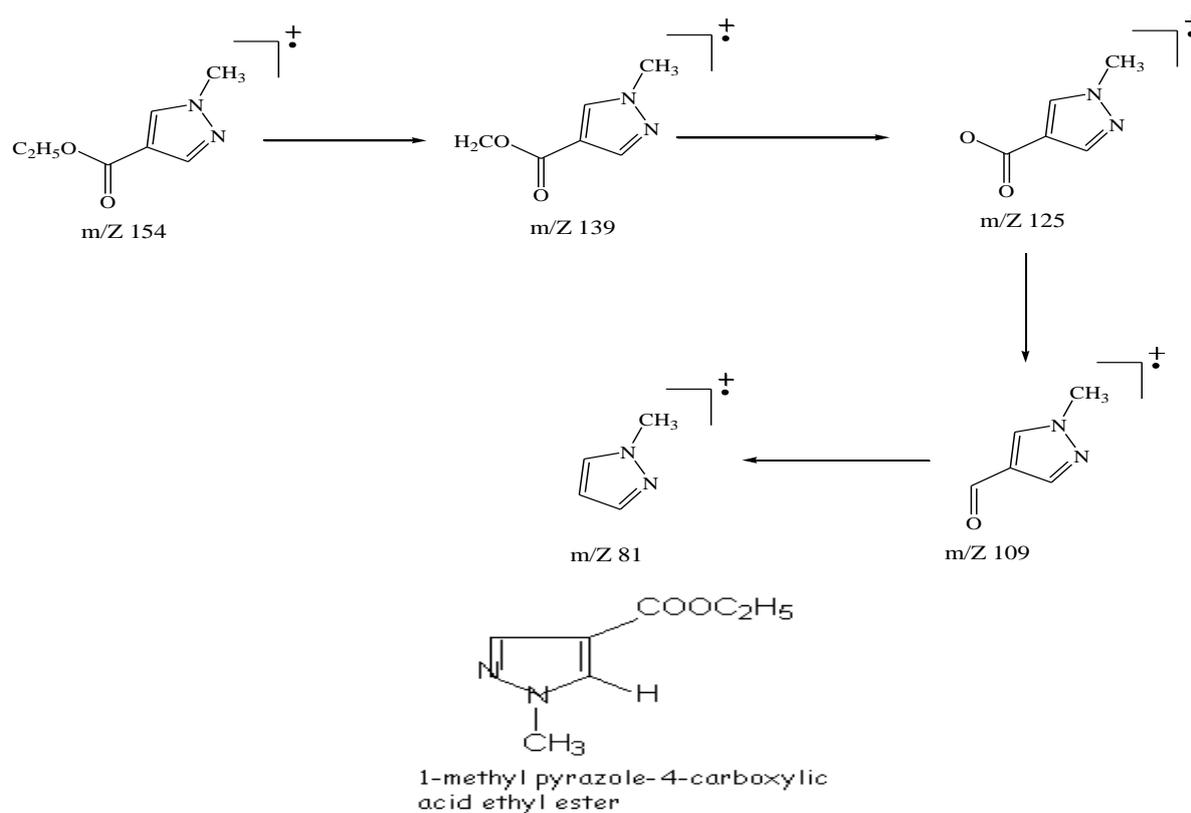


Fig. 2. Mass fragmentation pattern of compound F₂ with elucidated name

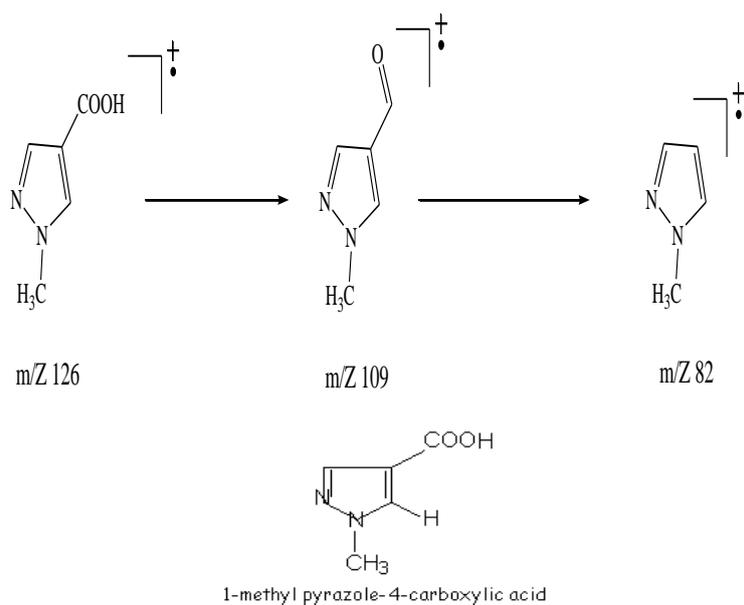


Fig. 3. Mass fragmentation pattern of compound F₃ with elucidated name

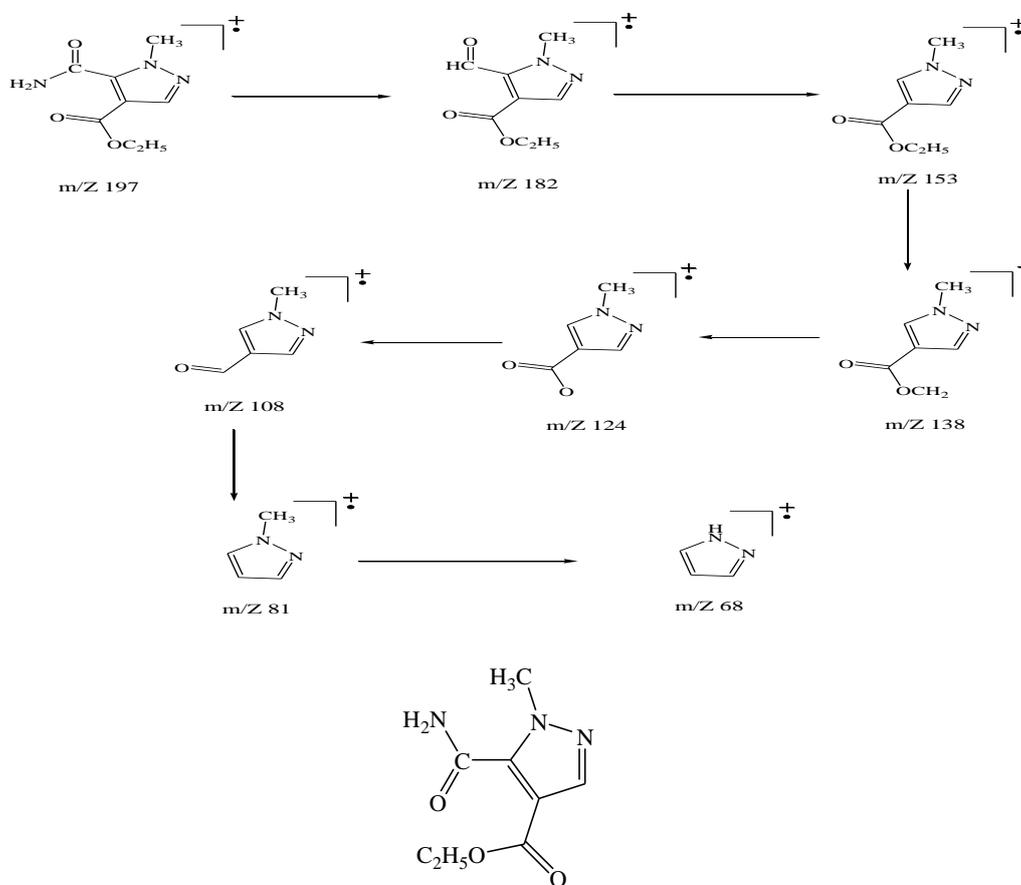


Fig. 4. Mass fragmentation pattern of compound F₄ with elucidated name

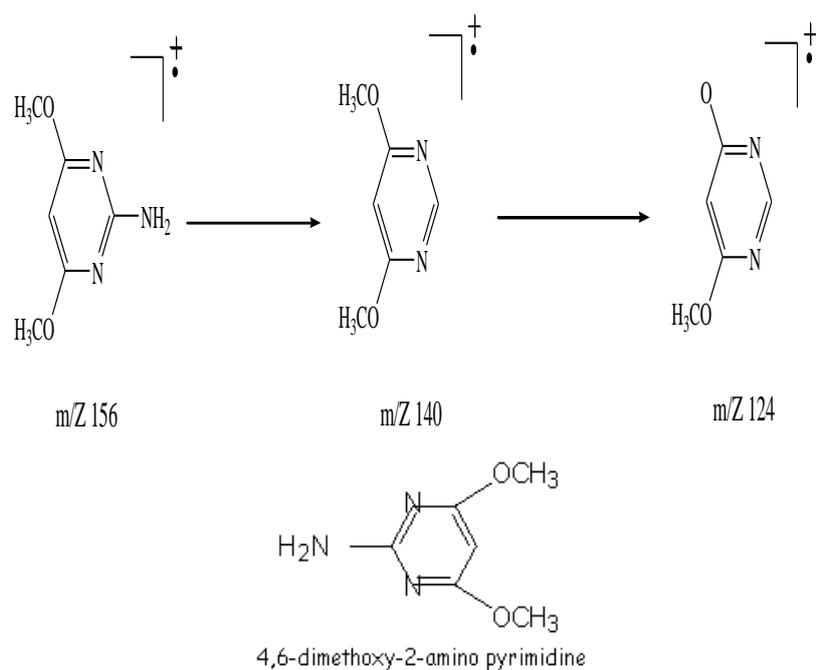


Fig. 5. Mass fragmentation pattern of compound F₅ with elucidated name

4. CONCLUSION

The isolated Pyrazosulfuron ethyl degrading fungi (*Aspergillus terreus*) can survive in minimal broth containing pyrazosulfuron ethyl at 1000 ppm and degrades the herbicide by harvesting energy through two major pathways. One path involves the cleavage of sulfonylurea bridge resulting in the formation of two major metabolites, first involves the cleavage of sulfonylurea bridge resulting in the formation of two major metabolites viz., 2- amino-4,6-dimethoxyprimidine (F₅) , 5-aminosulfonyl-1-methyl-pyrazole-4-carboxylic ethyl ester (F₁) and, second was the cleavage of sulfonyl amide linkage which forms the metabolite viz; 1 methyl pyrazole-4-carboxylic acid ethyl ester (F₂), 1 methyl pyrazole -4-carboxylic acid (F₃) and 5-carbamoyl-1-methyl pyrazole-4-carboxylate (F₄). On the basis of results obtained, it can be concluded that the enzyme involved in these transformations can be utilized to decontaminate soil and water from Pyrazosulfuron ethyl contamination. Even, the gene responsible for the production of these useful enzymes can be exploited for future research.

COMPETING INTERESTS

Authors have declared that no competing interests exist.

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