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Distribution of vesicular-arbuscular mycorrhizal fungi in soils polluted with industrial and sewage effluents

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ABSTRACT

The microbial communities including VAM fungi get affected by the sewage and industrial effluent. About this there is not much information. Soil polluted with sewage effluents supported less VAM population than non-polluted. 44 VAM fungal species were collected and indentified.

Tables: 03 Figure: 00 References: 06

KEY WORDS: Effluents, Industrial, Pollution, Sewage, Soil, VAM fungi.

Introduction

Indiscriminate use of pesticides, fertilizers and other pollutants cause undesirable changes in the soil ecosystem thus disturbing its physico-chemical set up and biological spectrum. Sewage effluents are a rich source of organic wastes, while industrial effluents contain many chemicals, including many heavy metals. The microbial communities, including vesicular-arbuscular mycorrhizal (VAM) fungi, may get affected by the sewage and industrial effluents. About this there is not much information. Hence, this study has been taken up.

Materials and Methods

In the present study, the soil which was distant from the effluent site and supported many weeds, grasses and other plant communities was considered as control. The soil site which received industrial effluents rich in caustic soda, acetic acid, soap, sodium silicate, ammonium sulphate, print colour and other effluents has been considered as the second sampling site. The third sampling soil site received only sewage effluents and supported only on type of grass. The soils were analysed for VAM fungi quantitatively following the wet sieving and decanting technique¹. The soils were analysed for soil type, pH, soil temperature, moisture, chlorides, available nitrogen, available phosphorus, available potassium, organic carbon, ferric, manganese, zinc, copper, cobalt, nickel, lead and cadmium following standard

TABLE-1: Number of VAM fungal propogules in 100 g of three sampling sites

Months /Year 2015-2016	S ₁	S_2	S ₃	
June 2015	740	300	116	
July	664	328	180	
August	632	500	204	
September	328	320	208	
October	532	312	172	
November	572	192	168	
December	352	220	132	
January 2016	250	129	44	
February	460	128	64	
March	580	480	80	
April	784	440	96	
Мау	642	472	108	

a = absent: b = present:

Notes- S₁ non-polluted soil, S₂ polluted soil with industrial effluents; S₃ polluted soil with sewage.

methods²⁻⁶.

Results and Conclusion

Soil polluted with the sewage effluent supported less VAM population than non-polluted soils. Altogether,

TABLE-2 : Distribution of VAM species in nonpolluted and polluted soils

VAM Species	S ₁	S ₂	S ₃
Acaulospora appendicula	_a	1	+ b
A. delicata	-	+	-
A. denticulata	+	-	-
A. foveata	-	-	+
A. lacunosa	+	-	+
A. laevis	-	+	-
A. mellea	+	-	+
A. nicoisonii	-	+	-
A. spinosa	-	+	+
A. tuberculata	-	+	-
Gigaspora albida	+	-	-
G. decipiensps	+	-	-
G. gigantea	+	-	-
G. margarita	+	-	-
Glomus aggregatum	-	+	-
G. botryoides	-	+	-
G. citricolum	-	-	+
G. claroideum	-	+	-
G. delhiense	-	+	-
G. dimorphicum	-	+	-
G. fascicalatum	-	+	-
G. fecundisporum	-	+	-

44 VAM fungal species were collected and indentified. The non-polluted soil contained 15 VAM species, whereas the soil polluted with industrial effluents harboured 26; only 9 species were found associated with the soil receiving sewage effluents. *Gigaspora* and *Scutellospora*,

VAM Species	S ₁	S ₂	S ₃
G. flavisporum	-	+	-
G. geosporum	-	+	-
G. hoi	-	-	+
G. intraradices	-	+	-
G. leptotichum	-	+	-
G. macrocarpum	-	+	-
G. maculosum	-	+	-
G. manihotis	-	+	-
G. mosseae	-	+	+
G. pustulatum	-	+	-
G. reticulutum	-	+	-
G. tortuosum	-	-	+
G. versiforme	-	+	-
Sclerocystis clavispora	+	-	-
S. coccogena	-	+	-
S. pakistanica	+	+	-
S. sinuosa	+	+	-
Scutellospora albarosea	+	-	-
S. calospora	+	-	-
S. gregaria	+	-	-
S. nigra	+	-	-
S. reticulata	+	-	-

which were found in non-polluted soils were absent in the soil polluted with industrial and sewage effluents. *Glomus* was completely absent in the non-polluted soil. The soils below neutral pH favoured more VAM fungal population while those with the pH range 7.02 to 8.1 had supported more VAM fungal species. Soil moisture was inversely related to VAM fungi. Chlorides, bicarbonates and nitrogen

did not affect VAN fungal dynamics. The soils with moderate to less phosphorus content supported more VAM fungal population. Soils with less potassium, 1-3% organic carbon, 53 to 330 ppm of iron and 4.5 to 17.5 ppm of manganese favoured more VAM fungi. Nickel and lead were found toxic at higher levels. The soil type therefore affects the VAM fungi dynamics.

TABLE-3: Physico-chemical factors and statistical data

Factors	S ₁ (Sandy loam)			S ₂ (Sandy silt loam)		S ₃ (Sandy clay loam)			
	1	2	3	1	2	3	1	2	3
рН	6.06	0.32005	1.06830	7.3	0.20362	0.65770	7.2	0.06088	0.19289
Moisture (%)	5.53	0.13123	0.41862	14.725	0.10022	0.31853	11.08	0.1722	0.55283
Temperature	37ºC	0.32505	1.08692	31ºC	0.24940	0.81442	29.7ºC	0.31252	1.04039
Bicarbonates	26.440	0.01534	0.04853	37.118	0.50049	1.82817	34.659	0.04123	0.13050
Chloride	11.700	0.50629	1.85656	114.74	0.36839	1.25311	22.072	0.17253	0.55390
Nitrogen (ppm)	70.58	0.03724	0.11786	124.32	0.10984	0.34947	247.21	0.45744	1.62674
Phosphorus (ppm)	29.52	0.40025	1.38117	32.9	0.23642	0.76944	125.2	0.038612	1.32370
Potassium (ppm)	107.8	0.37492	1.27891	125.3	0.182592	0.58729	200.5	0.18701	0.60201
Organic Carbon (%)	1.24	0.14364	0.45899	3.40	0.28379	0.93591	4.05	0.20566	0.66456
Iron (ppm)	108	0.19574	0.63121	163.1	0.19141	0.61672	218	0.70235	3.12020
Manganese (ppm)	9.63	0.51733	1.91166	6.70	0.20671	0.66811	16.7	0.34014	1.14385
Zinc (ppm)	2.023	0.22436	0.72808	6.37	0.14652	0.46840	2.99	0.05679	0.17990
Copper (ppm)	1.216	0.16028	0.51351	3.54	0.15952	0.51100	4.08	0.34572	1.16512
Cobalt (ppm)	8.091	0.02593	0.08204	10.17	0.32022	1.06892	12.67	0.4288	1.50101
Nickel (ppm)	0.629	0.48404	1.74928	30.39	0.10857	0.34537	2.132	0.3398.3	1.14266
Lead (ppm)	3.25	0.07940	0.25191	2.945	0.55385	2.10354	2.744	0.55385	2.10354
Cadmium (ppm)	0.103	0.15963	0.51135	0.119	0.28646	0.94552	0.153	0.50441	1.84734

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