

## INHIBITION OF *IN VITRO* GROWTH OF SOIL-BORNE PATHOGENS BY COMPOST-INHABITING INDIGENOUS BACTERIA AND FUNGI

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### Abstract

During the present studies, compost-inhabiting microorganisms including 44 fungi and 15 bacteria isolated from different compost samples were evaluated for their *in vitro* efficacy against soil-borne pathogens viz., *Fusarium solani*, *Macrophomina phaseolina*, *Pythium aphanidermatum*, *Rhizoctonia solani*, and *Sclerotium rolfsii*. Compost inhabiting microbes like *Trichoderma harzianum*, *T. virens*, *Bacillus cereus*, *B. pumilus*, *B. subtilis*, *Micrococcus varians* and *Pseudomonas fluorescens* were found to inhibit all the test pathogens. *Acrophialophora fuispora* and *Penicillium citrinum* reduced the mycelial growth of all the test pathogens except *Sclerotium rolfsii*. *Bacillus licheniformis* and *Bacillus megaterium* showed biocontrol activity against all the pathogens except *Rhizoctonia solani*. *Trichoderma harzianum* parasitized mycelia of all the tested pathogens and produced coiling around the mycelium.

### Introduction

Compost is a solid mature product obtained by the controlled biological decomposition of organic matter by different microorganisms like bacteria and fungi. These microorganisms decompose the organic matter in to a stable product that contains digested organic matter and nutrients, which can be easily utilize by plants (Rebollido *et al.*, 2008). Soils generally have complex compounds and low level of nutrients. Compost amendments generally enhance the nutrient activity of soil and stimulate plant growth as well as the development of microflora (Bailey & Lazarovits, 2003; Muhammad & Amusa, 2003).

Soil-borne pathogens like *Fusarium solani*, *Macrophomina phaseolina*, *Pythium aphanidermatum*, *Rhizoctonia solani* and *Sclerotium rolfsii* are known to produce serious losses to crop plants (Ghaffar, 1998; Agrios, 2005). Chemical control is often not sufficiently helpful, and hazardous for the environment. Use of biological control provides an alternative safe method of control instead of chemicals.

Use of composts for disease suppression is economical and eco-friendly. Soil amendment by compost suppressed the diseases caused by soil-borne pathogens in different crops (Hoitink *et al.*, 2001; Reuveni *et al.*, 2002; Noble & Coventry, 2005). Lewis *et al.*, (1992) found that cotton stand improved by compost treatment that also appreciably reduced the inoculum of *R. solani* in soil. Rhizospheric and root associated bacteria isolated from suppressive compost-based plant growth media provided protection to tomato plants against *Fusarium oxysporum* (Kavroulakis *et al.*, 2010). *Phytophthora* and *Pythium* root rots in many plants were suppressed by beneficial microbes found in compost (Rijckeboer *et al.*, 2002; Postma *et al.*, 2003).

Generally, the main protection mechanism against plant pathogens depends on microbial activity in the compost. Certain microbial inhabitants of composts viz., *Aspergillus*, *Bacillus*, *Penicillium*, *Pseudomonas*, *Streptomyces*, and *Trichoderma* species are potential biocontrol agents (Hoitink *et al.*, 1997; De Ceuster & Hoitink, 1999; Tilston *et al.*, 2002; Fuchs & Larbi 2004).

Control of pathogens by compost application may involve different mechanisms such as competition for nutrients, antibiotic production by beneficial microorganisms or activation of disease-resistance genes in plants (Hoitink & Boehm, 1999). The aim of the present study was to investigate the *in vitro* inhibition and suppression of soil-borne pathogens viz., *F. solani*, *M. phaseolina*, *P. aphanidermatum*, *R. solani*, and *S. rolfsii* by indigenous fungi and bacteria isolated from compost samples collected from Karachi.

### Materials and Methods

**Isolation of pathogens:** *F. solani*, *R. solani*, and *M. phaseolina* were isolated from roots and seeds of mung bean (*Vigna radiata* (L.) Wilczek). Cultures of *P. aphanidermatum* and *S. rolfsii* were obtained from Karachi University Culture Collection (KUCC).

**Isolation of compost inhabiting microbes:** Bacteria and fungi were isolated from different compost samples using serial dilution (Waksman & Fred, 1922) and soil plate (Warcup, 1950) methods. Five days old colonies used to identify fungi after reference to Ellis (1971), Barnette & Hunter (1972), and Domsch *et al.*, (1980). Bacterial identification carried out after 2-3 days incubation on the bases of morphological as well as biochemical tests following the standard procedure describe in Bergey's manual (Holt *et al.*, 1994).

***In vitro* interactions of soil-borne pathogens with compost-inhabiting microbes:** Interactions of soil-borne pathogens with isolated microorganisms were evaluated by dual culture plate assays. A 5mm inoculum disc of a pathogen was cut from an actively growing colony on potato sucrose agar (PSA) medium and placed near the edge of a Petri plate containing PSA medium. A similar inoculum disc of a test fungus was placed at the opposite end of the Petri plate, whereas, bacteria were streaked at the opposite end. There were three replicates of each treatment. The plates were incubated at 28±2°C and colonies diameter of pathogens and test microorganisms

were measured at 24 hours intervals and the type of interaction was recorded using the following key:

- A: Test microorganism inhibited growth of the pathogen and produced coiling around mycelium of the pathogen.  
 B: Colonies of test microorganism and pathogen met each other; No further growth of either the pathogen or the test microorganism observed.  
 C: Test microorganism inhibited growth of the pathogen and produced a zone of inhibition.  
 D: Pathogen inhibited and overgrew the colony of test microorganism.  
 E: Colonies of pathogen and test microorganism intermingled.

## Results

During the present studies, 20 fungi and seven bacteria showed biocontrol potential in dual culture plate assay against the soil-borne pathogens (Tables 1-5, Fig. 1).

***Fusarium solani*:** Six fungi viz., *Acrophialophora fuscispora*, *Penicillium chrysogenum*, *P. citrinum*, *Stachybotrys atra*, *Trichoderma harzianum* and *T. virens*, and seven bacteria viz., *Bacillus cereus*, *B. licheniformis*, *B. megaterium*, *B. pumilus*, *B. subtilis*, *Micrococcus varians* and *Pseudomonas fluorescens* inhibited the growth of *F. solani* (Table 1).

***Macrophomina phaseolina*:** Growth of *M. phaseolina* was inhibited by 11 fungi viz., *A. fuscispora*, *Aspergillus flavus*, *A. fumigatus*, *A. niger*, *Drechslera hawaiiensis*, *Emericella nidulans*, *Penicillium chrysogenum*, *P. citrinum*, *S. atra*, *T. harzianum* and *T. virens*, and seven bacteria viz., *B. cereus*, *B. licheniformis*, *B. megaterium*, *B. pumilus*, *B. subtilis*, *M. varians* and *P. fluorescens* (Table 2).

***Pythium aphanidermatum*:** 20 fungi viz., *A. fuscispora*, *A. flavus*, *A. fumigatus*, *A. niger*, *A. nidulans*, *A. terreus*, *A. terreus* var. *flococcus*, *Botrytis cinerea*, *Chaetomium thermophilum* var. *coprophile*, *Conidiobolus thermophilus*, *Corynascus sepedonium*, *E. nidulans*, *E. nivea*, *E. rugulosa*, *Paecilomyces variotii*, *P. chrysogenum*, *P. citrinum*, *S. atra*, *T. harzianum* and *T. virens*, and seven bacteria viz., *B. cereus*, *B. licheniformis*, *B. megaterium*, *B. pumilus*, *B. subtilis*, *M. varians* and *P. fluorescens* inhibited the growth of *P. aphanidermatum* (Table 3).

***Rhizoctonia solani*:** Five fungi viz., *A. fuscispora*, *A. niger*, *P. citrinum*, *T. harzianum* and *T. virens*, and five bacteria viz., *Bacillus cereus*, *B. pumilus*, *B. subtilis*, *M. varians* and *P. fluorescens* were antagonistic to *R. solani* (Table 4).

***Sclerotium rolfsii*:** Growth of *S. rolfsii* was inhibited by only 2 fungi viz., *T. harzianum* and *T. virens*, and seven bacteria viz., *B. cereus*, *B. licheniformis*, *B. megaterium*, *B. pumilus*, *B. subtilis*, *M. varians* and *P. fluorescens* (Table 5).

**Table 1. In vitro interactions of *Fusarium solani* with fungi and bacteria isolated from compost.**

Test microorganisms	Days of incubation	Colony diameter (mm)		Type of interaction
		Pathogen	Test microorganism	
<b>Fungi</b>				
<i>Acronium thermophilum</i>	5	35	56	D
<i>Acrophialophora fuscispora</i>	6	60	22	C
<i>Alleschleria crocea</i>	4	35	61	D
<i>Alternaria alternata</i>	6	59	33	D
<i>Annelophora africana</i>	5	42	50	D
<i>Aspergillus flavus</i>	6	38	50	D
<i>Aspergillus fumigatus</i>	5	35	54	D
<i>Aspergillus nidulans</i>	5	38	44	D
<i>Aspergillus niger</i>	6	46	34	D
<i>Aspergillus terreus</i>	7	59	30	D
<i>Aspergillus terreus</i> var. <i>flococcus</i>	7	50	38	D
<i>Aureobasidium pullulans</i>	4	31	60	D
<i>Botryotrichum</i> sp.	6	48	43	D
<i>Botrytis cinerea</i>	8	65	29	D
<i>Chaetomium globosum</i>	5	55	35	E
<i>Chaetomium thermophilum</i> var. <i>coprophile</i>	7	48	42	E
<i>Chaetomium thermophilum</i> var. <i>dissitum</i>	8	45	45	E
<i>Conidiobolus thermophilus</i>	7	56	37	D
<i>Coniothyrium</i> sp.	5	32	58	D
<i>Corynascus sepedonium</i>	7	56	35	D
<i>Drechslera hawaiiensis</i>	6	47	44	D
<i>Emericella nidulans</i>	7	65	26	D
<i>Emericella nivea</i>	8	45	44	D
<i>Emericella rugulosa</i>	6	57	35	D
<i>Gilmaniella humicola</i>	5	29	61	D
<i>Humicola fuscoatra</i>	5	32	60	D
<i>Humicola grisea</i>	5	31	60	D
<i>Mucor fragalis</i>	3	23	59	E
<i>Mucor</i> sp.	3	19	73	E
<i>Myrothecium</i> sp.	9	61	30	D
<i>Paecilomyces fumosoroseus</i>	6	46	45	D
<i>Paecilomyces variotii</i>	5	36	54	E
<i>Papulaspora thermophila</i>	5	36	56	D
<i>Penicillium chrysogenum</i>	7	46	42	C
<i>Penicillium citrinum</i>	7	53	23	C
<i>Rhizomucor pusillus</i>	5	30	60	E
<i>Scytalidium</i> sp.	6	45	45	E
<i>Scytalidium thermophilum</i>	8	54	37	D
<i>Stachybotrys atra</i>	10	70	18	C
<i>Syncephalastrum racemosum</i>	6	35	55	E
<i>Talaromyces thermophilus</i>	6	46	46	D
<i>Thermomyces lanuginosus</i>	5	47	44	D
<i>Trichoderma harzianum</i>	4	29	64	A
<i>Trichoderma virens</i>	4	28	60	B
<b>Bacteria</b>				
<i>Aeromonas</i> sp.	7	72	21	D
<i>Bacillus cereus</i>	7	60	22	C
<i>Bacillus licheniformis</i>	7	63	20	C
<i>Bacillus megaterium</i>	7	62	25	C
<i>Bacillus pumilus</i>	6	63	24	C
<i>Bacillus stearothermophilus</i>	7	65	25	E
<i>Bacillus subtilis</i>	7	66	19	C
<i>Enterococcus</i> sp.	7	69	23	D
<i>Escherichia coli</i>	7	66	25	D
<i>Geobacillus toebii</i>	7	68	23	D
<i>Klebsiella</i> sp.	7	69	22	D
<i>Micrococcus varians</i>	7	65	20	C
<i>Pseudomonas fluorescens</i>	7	68	22	B
<i>Staphylococcus aureus</i>	7	67	23	E
<i>Thermus thermophilus</i>	6	68	22	D

Table 2. *In vitro* interactions of *Macrophomina phaseolina* with fungi and bacteria isolated from compost.

Test microorganisms	Days of incubation	Colony diameter (mm)		Type of interaction
		Pathogen	Test micro-organism	
Fungi				
<i>Acremonium thermophilum</i>	4	52	40	D
<i>Acrophialophora fusispora</i>	4	68	17	C
<i>Alleschleria crocea</i>	4	58	36	D
<i>Alternaria alternata</i>	4	67	24	D
<i>Annelophora africana</i>	4	47	45	D
<i>Aspergillus flavus</i>	4	59	30	C
<i>Aspergillus fumigatus</i>	3	54	33	C
<i>Aspergillus nidulans</i>	5	51	39	E
<i>Aspergillus niger</i>	4	53	54	B
<i>Aspergillus terreus</i>	4	68	23	D
<i>Aspergillus terreus</i> var. <i>flococcus</i>	5	68	22	E
<i>Aureobasidium pullulans</i>	4	26	65	D
<i>Botryotrichum</i> sp.	4	61	29	D
<i>Botrytis cinerea</i>	4	76	15	D
<i>Chaetomium globosum</i>	4	64	26	E
<i>Chaetomium thermophilum</i> var. <i>coprophile</i>	4	67	23	E
<i>Chaetomium thermophilum</i> var. <i>dissitum</i>	4	70	20	E
<i>Conidiobolus thermophilus</i>	4	57	40	E
<i>Coniothyrium</i> sp.	4	32	58	D
<i>Corynascus sepedonium</i>	4	66	24	E
<i>Drechslera hawaiiensis</i>	5	63	26	B
<i>Emericella nidulans</i>	5	66	18	C
<i>Emericella nivea</i>	4	67	23	E
<i>Emericella rugulosa</i>	4	66	24	E
<i>Gilmaniella humicola</i>	4	36	54	D
<i>Humicola fuscoatra</i>	4	41	49	E
<i>Humicola grisea</i>	4	35	55	E
<i>Mucor fragalis</i>	3	46	46	E
<i>Mucor</i> sp.	3	34	57	E
<i>Myrothecium</i> sp.	5	73	17	E
<i>Paecilomyces fumosoroseus</i>	4	47	44	D
<i>Paecilomyces variotii</i>	4	56	34	E
<i>Papulaspora thermophila</i>	4	34	56	D
<i>Penicillium chrysogenum</i>	4	67	24	B
<i>Penicillium citrinum</i>	4	64	20	C
<i>Rhizomucor pusillus</i>	3	44	46	E
<i>Scytalidium</i> sp.	4	63	27	E
<i>Scytalidium thermophilum</i>	4	65	25	D
<i>Stachybotrys atra</i>	5	69	17	C
<i>Syncephalastrum racemosum</i>	4	58	38	E
<i>Talaromyces thermophilus</i>	4	51	39	D
<i>Thermomyces lanuginosus</i>	4	49	42	D
<i>Trichoderma harzianum</i>	3	35	53	A
<i>Trichoderma virens</i>	4	52	45	B
<b>Bacteria</b>				
<i>Aeromonas</i> sp.	4	67	24	D
<i>Bacillus cereus</i>	4	66	18	C
<i>Bacillus licheniformis</i>	4	64	20	C
<i>Bacillus megaterium</i>	4	65	22	C
<i>Bacillus pumilus</i>	4	63	25	C
<i>Bacillus stearothermophilus</i>	5	65	25	E
<i>Bacillus subtilis</i>	4	65	20	C
<i>Enterococcus</i> sp.	4	70	23	D
<i>Escherichia coli</i>	4	68	25	D
<i>Geobacillus toebii</i>	4	68	24	D
<i>Klebsiella</i> sp.	4	69	22	D
<i>Micrococcus varians</i>	4	60	26	C
<i>Pseudomonas fluorescens</i>	4	66	22	C
<i>Staphylococcus aureus</i>	5	67	23	E
<i>Thermus thermophilus</i>	5	69	21	D

Table 3. *In vitro* interactions of *Pythium aphanidermatum* with fungi and bacteria isolated from compost.

Test microorganisms	Days of incubation	Colony diameter (mm)		Type of interaction
		Pathogen	Test micro-organism	
Fungi				
<i>Acremonium thermophilum</i>	3	56	27	D
<i>Acrophialophora fusispora</i>	4	70	17	C
<i>Alleschleria crocea</i>	4	57	46	D
<i>Alternaria alternata</i>	3	69	21	E
<i>Annelophora africana</i>	4	57	45	D
<i>Aspergillus flavus</i>	3	63	25	C
<i>Aspergillus fumigatus</i>	3	60	31	B
<i>Aspergillus nidulans</i>	3	62	30	B
<i>Aspergillus niger</i>	3	57	29	C
<i>Aspergillus terreus</i>	3	67	20	C
<i>Aspergillus terreus</i> var. <i>flococcus</i>	4	64	21	C
<i>Aureobasidium pullulans</i>	3	32	57	D
<i>Botryotrichum</i> sp.	3	62	28	D
<i>Botrytis cinerea</i>	3	62	17	C
<i>Chaetomium globosum</i>	5	64	26	E
<i>Chaetomium thermophilum</i> var. <i>coprophile</i>	4	59	25	C
<i>Chaetomium thermophilum</i> var. <i>dissitum</i>	4	72	19	D
<i>Conidiobolus thermophilus</i>	3	48	37	C
<i>Coniothyrium</i> sp.	2	36	55	D
<i>Corynascus sepedonium</i>	4	66	20	C
<i>Drechslera hawaiiensis</i>	4	65	26	D
<i>Emericella nidulans</i>	4	73	16	C
<i>Emericella nivea</i>	4	62	23	C
<i>Emericella rugulosa</i>	4	64	24	C
<i>Gilmaniella humicola</i>	3	36	55	D
<i>Humicola fuscoatra</i>	3	29	61	D
<i>Humicola grisea</i>	3	38	52	D
<i>Mucor fragalis</i>	3	54	44	E
<i>Mucor</i> sp.	3	54	42	E
<i>Myrothecium</i> sp.	4	77	14	D
<i>Paecilomyces fumosoroseus</i>	4	44	46	D
<i>Paecilomyces variotii</i>	3	55	35	B
<i>Papulaspora thermophila</i>	4	35	56	D
<i>Penicillium chrysogenum</i>	4	68	20	C
<i>Penicillium citrinum</i>	3	66	19	C
<i>Rhizomucor pusillus</i>	3	50	40	E
<i>Scytalidium</i> sp.	4	63	30	D
<i>Scytalidium thermophilum</i>	3	63	28	D
<i>Stachybotrys atra</i>	4	74	13	C
<i>Syncephalastrum racemosum</i>	4	60	30	E
<i>Talaromyces thermophilus</i>	4	49	45	D
<i>Thermomyces lanuginosus</i>	4	48	42	D
<i>Trichoderma harzianum</i>	3	46	56	A
<i>Trichoderma virens</i>	3	47	55	B
<b>Bacteria</b>				
<i>Aeromonas</i> sp.	4	68	23	D
<i>Bacillus cereus</i>	4	60	18	C
<i>Bacillus licheniformis</i>	4	65	15	C
<i>Bacillus megaterium</i>	3	60	22	C
<i>Bacillus pumilus</i>	4	55	25	C
<i>Bacillus stearothermophilus</i>	4	67	23	E
<i>Bacillus subtilis</i>	3	60	25	C
<i>Enterococcus</i> sp.	4	78	18	D
<i>Escherichia coli</i>	3	79	17	D
<i>Geobacillus toebii</i>	4	67	24	D
<i>Klebsiella</i> sp.	3	71	20	D
<i>Micrococcus varians</i>	3	54	24	C
<i>Pseudomonas fluorescens</i>	3	60	21	C
<i>Staphylococcus aureus</i>	4	70	22	D
<i>Thermus thermophilus</i>	3	69	22	D

Table 4. *In vitro* interactions of *Rhizoctonia solani* with fungi and bacteria isolated from compost.

Test microorganisms	Days of incubation	Colony diameter (mm)		Type of interaction
		Pathogen	Test micro-organism	
<b>Fungi</b>				
<i>Acremonium thermophilum</i>	3	53	42	D
<i>Acrophialophora fusispora</i>	4	69	17	C
<i>Alleschleria crocea</i>	3	40	57	D
<i>Alternaria alternata</i>	4	71	21	D
<i>Annelophora africana</i>	3	48	46	D
<i>Aspergillus flavus</i>	4	58	32	E
<i>Aspergillus fumigatus</i>	3	61	32	D
<i>Aspergillus nidulans</i>	3	68	25	D
<i>Aspergillus niger</i>	4	52	36	C
<i>Aspergillus terreus</i>	4	67	25	D
<i>Aspergillus terreus</i> var. <i>flococcus</i>	4	70	23	D
<i>Aureobasidium pullulans</i>	3	29	62	D
<i>Botryotrichum</i> sp.	3	63	28	D
<i>Botrytis cinerea</i>	4	76	16	D
<i>Chaetomium globosum</i>	4	70	20	D
<i>Chaetomium thermophilum</i> var. <i>coprophile</i>	4	65	26	D
<i>Chaetomium thermophilum</i> var. <i>dissitum</i>	4	74	17	D
<i>Conidiobolus thermophilus</i>	3	63	30	D
<i>Coniothyrium</i> sp.	3	36	55	D
<i>Corynascus sepedonium</i>	4	73	21	D
<i>Drechslera hawaiiensis</i>	3	67	27	D
<i>Emericella nidulans</i>	4	78	13	D
<i>Emericella nivea</i>	4	71	23	D
<i>Emericella rugulosa</i>	4	68	22	E
<i>Gilmaniella humicola</i>	3	35	55	D
<i>Humicola fuscoatra</i>	3	44	48	D
<i>Humicola grisea</i>	3	39	52	D
<i>Mucor fragalis</i>	3	48	45	D
<i>Mucor</i> sp.	3	49	49	D
<i>Myrothecium</i> sp.	4	75	16	D
<i>Paecilomyces fumosoroseus</i>	3	47	44	D
<i>Paecilomyces variotii</i>	3	61	30	D
<i>Papulaspora thermophila</i>	4	41	53	D
<i>Penicillium chrysogenum</i>	3	72	19	D
<i>Penicillium citrinum</i>	4	67	20	C
<i>Rhizomucor pusillus</i>	3	61	41	D
<i>Scytalidium</i> sp.	3	63	28	D
<i>Scytalidium thermophilum</i>	3	64	27	D
<i>Stachybotrys atra</i>	4	78	12	D
<i>Syncephalastrum racemosum</i>	3	62	34	D
<i>Talaromyces thermophilus</i>	4	58	33	D
<i>Thermomyces lanuginosus</i>	4	46	47	D
<i>Trichoderma harzianum</i>	4	51	50	A
<i>Trichoderma virens</i>	4	50	55	B
<b>Bacteria</b>				
<i>Aeromonas</i> sp.	4	70	22	D
<i>Bacillus cereus</i>	3	64	25	C
<i>Bacillus licheniformis</i>	3	65	25	E
<i>Bacillus megaterium</i>	3	70	20	D
<i>Bacillus pumilus</i>	3	65	21	C
<i>Bacillus stearothermophilus</i>	4	65	25	D
<i>Bacillus subtilis</i>	3	69	20	B
<i>Enterococcus</i> sp.	4	79	15	D
<i>Escherichia coli</i>	4	76	15	D
<i>Geobacillus toebii</i>	4	74	18	D
<i>Klebsiella</i> sp.	4	71	20	D
<i>Micrococcus varians</i>	3	64	20	C
<i>Pseudomonas fluorescens</i>	3	68	21	C
<i>Staphylococcus aureus</i>	3	73	20	D
<i>Thermus thermophilus</i>	4	68	23	D

Table 5. *In vitro* interactions of *Sclerotium rolfsii* with fungi and bacteria isolated from compost.

Test microorganisms	Days of incubation	Colony diameter (mm)		Type of interaction
		Pathogen	Test micro-organism	
<b>Fungi</b>				
<i>Acremonium thermophilum</i>	4	55	40	D
<i>Acrophialophora fusispora</i>	4	71	19	D
<i>Alleschleria crocea</i>	3	61	31	D
<i>Alternaria alternata</i>	4	68	24	D
<i>Annelophora africana</i>	4	47	46	D
<i>Aspergillus flavus</i>	4	64	30	D
<i>Aspergillus fumigatus</i>	4	56	36	D
<i>Aspergillus nidulans</i>	4	61	32	D
<i>Aspergillus niger</i>	4	64	32	D
<i>Aspergillus terreus</i>	4	66	25	D
<i>Aspergillus terreus</i> var. <i>flococcus</i>	4	73	24	D
<i>Aureobasidium pullulans</i>	4	32	58	D
<i>Botryotrichum</i> sp.	4	68	26	D
<i>Botrytis cinerea</i>	5	80	15	D
<i>Chaetomium globosum</i>	5	70	21	D
<i>Chaetomium thermophilum</i> var. <i>coprophile</i>	4	65	31	D
<i>Chaetomium thermophilum</i> var. <i>dissitum</i>	4	71	24	D
<i>Conidiobolus thermophilus</i>	3	54	39	D
<i>Coniothyrium</i> sp.	3	41	51	D
<i>Corynascus sepedonium</i>	4	78	15	D
<i>Drechslera hawaiiensis</i>	4	67	27	D
<i>Emericella nidulans</i>	4	80	12	D
<i>Emericella nivea</i>	4	70	23	D
<i>Emericella rugulosa</i>	4	67	25	D
<i>Gilmaniella humicola</i>	3	35	57	D
<i>Humicola fuscoatra</i>	3	44	47	D
<i>Humicola grisea</i>	3	36	56	D
<i>Mucor fragalis</i>	4	51	44	D
<i>Mucor</i> sp.	4	49	53	D
<i>Myrothecium</i> sp.	5	74	17	D
<i>Paecilomyces fumosoroseus</i>	4	47	44	D
<i>Paecilomyces variotii</i>	4	61	32	D
<i>Papulaspora thermophila</i>	4	55	36	D
<i>Penicillium chrysogenum</i>	4	74	23	D
<i>Penicillium citrinum</i>	4	68	25	D
<i>Rhizomucor pusillus</i>	3	47	45	D
<i>Scytalidium</i> sp.	4	59	32	D
<i>Scytalidium thermophilum</i>	4	61	29	D
<i>Stachybotrys atra</i>	5	85	15	D
<i>Syncephalastrum racemosum</i>	3	61	33	D
<i>Talaromyces thermophilus</i>	4	56	39	D
<i>Thermomyces lanuginosus</i>	3	50	42	D
<i>Trichoderma harzianum</i>	3	37	54	A
<i>Trichoderma virens</i>	4	53	56	B
<b>Bacteria</b>				
<i>Aeromonas</i> sp.	3	78	14	D
<i>Bacillus cereus</i>	4	55	20	C
<i>Bacillus licheniformis</i>	4	53	23	C
<i>Bacillus megaterium</i>	4	56	24	C
<i>Bacillus pumilus</i>	4	54	28	C
<i>Bacillus stearothermophilus</i>	4	78	12	D
<i>Bacillus subtilis</i>	4	55	25	C
<i>Enterococcus</i> sp.	4	66	25	D
<i>Escherichia coli</i>	3	67	22	D
<i>Geobacillus toebii</i>	4	75	26	D
<i>Klebsiella</i> sp.	4	70	22	D
<i>Micrococcus varians</i>	4	50	20	C
<i>Pseudomonas fluorescens</i>	4	74	12	C
<i>Staphylococcus aureus</i>	4	72	22	D
<i>Thermus thermophilus</i>	4	68	23	E

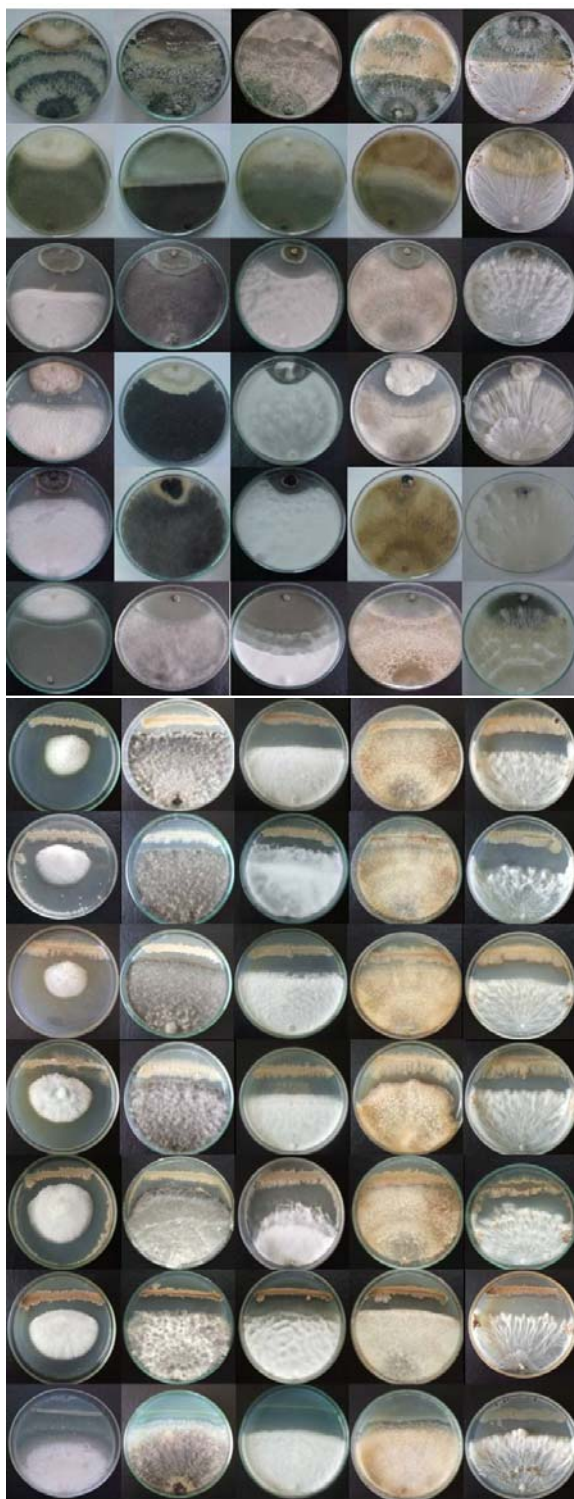


Fig. 1. Interactions of soil-borne pathogens with microorganisms isolated from compost (Pathogens at the bottom and antagonists at the top side of the plates).

**Columns (from left):** *Fusarium solani*, *Macrophomina phaseolina*, *Pythium aphanidermatum*, *Rhizoctonia solani*, and *Sclerotium rolfsii*.

**Rows (from top):** *Trichoderma harzianum*, *T. virens*, *Penicillium citrinum*, *Acrophialophora fusispora*, *Stachybotrys atra*, *Aspergillus fumigatus*, *Bacillus cereus*, *B. licheniformis*, *B. megaterium*, *B. pumilus*, *B. subtilis*, *Micrococcus varians* and *Pseudomonas fluorescens*.

## Discussion

Of the 44 fungi and 15 bacteria isolated from different compost samples, 20 species of fungi belonging to 12 genera viz., *Acrophialophora*, *Aspergillus*, *Botrytis*, *Chaetomium*, *Conidiobolus*, *Corynascus*, *Drechslera*, *Emericella*, *Paecilomyces*, *Penicillium*, *Stachybotrys* and *Trichoderma*, and seven species of bacteria belonging to three genera viz., *Bacillus*, *Micrococcus* and *Pseudomonas* showed antagonism against different soil-borne pathogens used during the present studies. No previous report on antagonistic activity of *M. varians* is available but in our studies, it successfully inhibited the growth of all the pathogens. *Trichoderma harzianum* parasitized all the pathogens and produced coiling around their mycelia, whereas, *Trichoderma virens* did not produce any coiling around the mycelium of the pathogens. Bacteria viz., *B. cereus*, *B. pumilus*, *B. subtilis*, *M. varians* and *P. fluorescens* inhibited all the test pathogens and produced zones of inhibition. *B. licheniformis* and *B. megaterium* showed biocontrol activity against all the pathogens except *R. solani*. *A. fusispora* and *P. citrinum* also produced zones of inhibition against all the pathogens except *S. rolfsii*. The maximum zones of inhibition were produced by *P. citrinum* against *F. solani* and *M. varians* against *M. phaseolina*, *P. aphanidermatum*, *R. solani* and *S. rolfsii*.

*A. thermophilum*, *A. crocea*, *A. alternata*, *A. africana*, *A. pullulans*, *Botryotrichum* sp., *C. thermophilum* var. *dissitum*, *Coniothyrium* sp., *G. humicola*, *H. fuscoatra*, *H. grisea*, *Mucor* sp., *Myrothecium* sp., *P. fumosoroseus*, *P. thermophila*, *R. pusillus*, *Scytilidium* sp., *S. racemosum*, *T. thermophilum* and *T. lanuginosus* showed 'D' type of interaction with all the pathogens, where the pathogens inhibited and overgrew the colonies of the test microorganisms. Similarly, bacteria viz., *Aeromonas* sp., *B. stearothermophilus*, *Enterococcus* sp., *E. coli*, *Klebsiella* sp., *G. toebii* and *T. thermophilum* did not show antagonistic activity against any pathogen.

Muhammad & Amusa (2003) found that compost-inhabiting microbes inhibited the growth of pathogens and thus suppressed the diseases. During the present studies, growth of *Fusarium solani* was inhibited by *T. harzianum*, *T. virens*, *P. chrysogenum*, *P. citrinum* and *S. atra* that corroborates well with the results reported by Akrami *et al.*, (2012) and Dwivedi & Dwivedi (2012). Suppression in growth of *F. solani* by *B. cereus*, *B. licheniformis*, *B. megaterium*, *B. pumilus*, *B. subtilis* and *P. fluorescens* during the present studies supports the findings of Amalraj *et al.*, (2012) and Haggag & El-Gamal (2012).

Growth of *Macrophomina phaseolina* was found to be inhibited by *T. harzianum*, *T. virens*, *A. niger*, *A. fumigatus*, *Penicillium* sp., *B. cereus*, *B. licheniformis*, *B. megaterium*, *B. pumilus*, *B. subtilis* and *P. fluorescens*. Similar results have been reported by Shivakumar (2007), El-Mougy & Abdel-Kader (2008), Hajjegrari *et al.*, (2008), Bandopadhyay *et al.*, (2011), Mahalingam, *et al.*, (2011), Amalraj *et al.*, (2012), and Gajera *et al.*, (2012).

*Rhizoctonia solani* was inhibited by *A. fusispora*, *A. niger*, *P. citrinum*, *T. harzianum*, *T. vires*, *B. cereus*, *B. pumilus*, *B. subtilis*, *M. varians* and *P. fluorescens*. There are several reports where growth of *R. solani* was inhibited by *T. harzianum* (Demirci *et al.*, 2011), *T. vires* and *A. niger* (Vibha, 2010), *B. subtilis* and *P. fluorescens* (Haggag & El-Gamal, 2012). Ganesan & Sekar (2011) reported that *B. pumilus* and *B. licheniformis* inhibited the mycelial growth of *R. solani* *in vitro*, but in present studies, *B. licheniformis* did not find to inhibit the mycelial growth of *R. solani*.

*Pythium aphanidermatum* was the most sensitive among all the pathogens tested since 20 fungi and seven bacteria inhibited its growth. Similar results have been reported by Kamil *et al.*, (2007), Mishra (2010), Gomathi & Ambikapathy (2011), El-Mohamedy *et al.*, (2011), Basurto-Cadena *et al.*, (2012), and El-Bramawy *et al.*, (2012). On the other hand, *Sclerotium rolfsii* showed highest tolerance to the antagonistic effects of microorganisms since only *T. harzianum* and *T. vires* were able to inhibit its growth, whereas, all other fungi and bacteria were over grown by *S. rolfsii*. Inhibition in growth of *S. rolfsii* has also been reported by Yaqub & Shahzad (2005) and Bhuiyan *et al.*, (2012). Bosah *et al.*, (2010) reported that *A. niger* reduced the mycelial growth of *S. rolfsii* but *Penicillium* did not. In our studies, *A. niger* and *Penicillium* did not inhibit the mycelial growth of *S. rolfsii*. *B. cereus*, *B. licheniformis*, *B. megaterium*, *B. pumilus*, *B. subtilis* and *P. fluorescens* reduced the mycelial growth of *S. rolfsii*. Similar results were observed by Nalisha *et al.*, (2006), Shivakumar (2007), and Amalraj *et al.*, (2012).

The growth inhibition of pathogen may be due to hyperparasitism, antibiosis (We *et al.*, 1986) or production of chitinase and B-1,3-glucanase enzymes which degrade the cell wall of the pathogens (Ahmed & Baker, 1987). *Trichoderma* generate many enzymes that are used against cell walls of fungi to utilize the fragment of pathogens (Grosch *et al.*, 2006). *B. subtilis* produces five antibiotics viz., bacillin, bacitracin, bacilomycin, subtilin and subtenolin. Muhammad & Amusa (2003) reported that two mechanisms were involved during the interactions between the antagonist and the pathogen: First one, the production of biologically active metabolites, and the other one, the rapid growth of biocontrol agent on moist surface of agar, which inhibited the growth of pathogens. Various *Trichoderma* and *Bacillus* species have the potential to control plant pathogens (Utkhede *et al.*, 1999; Begum & Begum, 2010). Inhibition of *Rhizoctonia solani* growth by *Bacillus subtilis* is attributed to the production of lipopeptides such as iturin and surfactin (Yu *et al.*, 2002). The results of the present study confirm that compost contains a large number of indigenous beneficial microbes capable of suppressing soil-borne pathogens. This character of the compost can be utilized to overcome the difficulties in mass production of biocontrol agents for field application and may lead to suppression of diseases leading to increased crop productivity.

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