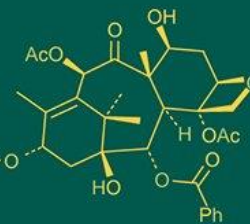
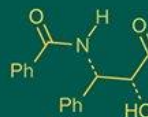


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Trichoderma as a possible biocontrol agent and its use in agriculture

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Abstract

Increased agricultural food production is required to feed the growing global population. However, risk of pest infestation hugely hampers the agricultural production every year. The main factor reducing this loss is the usage of chemical insecticides. Even though these compounds are highly effective, they have a severe impact on both human life and the environment. Therefore, it is crucial to create innovative, secure alternatives that are both effective and safe. *Trichoderma* species are opportunistic, avirulent plant symbionts. Their symbiotic relationship with plants increases plant resilience to diseases, enhances growth and productivity, encourages nutrient uptake and fertilizer use efficiency. Antibiosis, competition, and mycoparasitism are a few of the key ways that *Trichoderma* responds to the presence of other competing pathogenic organisms by inhibiting or blocking their growth, among other biocontrol mechanisms. Indirectly and directly, the use of *Trichoderma* as a biocontrol agent for insect pest has been contemplated recently. *Trichoderma* is capable of directly reducing insect pests through parasitism and the formation of insecticidal secondary metabolites, antifeedant chemicals, and repellent metabolites, according to studies conducted so far. Additionally, indirectly, through the induction of innate plant defence mechanisms, the recruitment of natural enemies, or the parasitism of microorganisms that lives in symbiotic relationship with insects. The existing data on *Trichoderma* spp. and its biocontrol activity in long-term disease management programmes are reviewed in this work.

Keywords: *Trichoderma*, Biocontrol agent, antibiosis, mycoparasitism

Introduction

Biological control, as defined by Cook and Baker 1983, is the process that reduces the number of microbes or pathogens without the assistance of outside humans. Tubef first used the term "biocontrol" in 1914, and Smith first connected it to insects and plant pathogens in 1909. By examining the antagonistic potential of some microorganisms, this methodological scheme uses natural predators that have the capacity to eradicate and control the growth of pests as well as pathogens, making it an environmentally friendly method for controlling crop diseases (Gawai, 2018) ^[14]. Chemical compounds have been used to control plant diseases (chemical control), but overuse of these compounds has facilitated the evolution of fungicide resistance in pathogens. In contrast, using antagonistic microbes to control plant pathogens (biological control) poses no risk when it strengthens the local antagonist population. Many microbial antagonists have been reported to possess antagonistic activities against plant fungal pathogens, such as *Pseudomonas fluorescens*, *Agrobacterium radiobacter*, *Bacillus subtilis*, *B. cereus*, *B. amyloliquefaciens*, *Trichoderma virens*, *Burkholderia cepacia*, *Saccharomyces* sp., *Gliocadium* sp. etc. The specificity of the biological control agent released in a new ecosystem is one of the major issues with this approach, so the viability of such introductions must be carefully considered.

The genus *Trichoderma* belongs to the Kingdom-Fungi, Phylum-Deuteromycotina, Class-Sodariomycetes, Order-Hypocreales, Family-Hypocreaceae. However, *Trichoderma* fungi, which are common in most ecosystems, can eliminate this significant concern (Brotman *et al.*, 2010) ^[15] of chemical control. In recent years, the use of pesticides in economically significant crops has decreased as a result of integrated pest management strategies, reduced or controlled pesticide use, and increased use of fungal biocontrol agents, particularly *Trichoderma* spp. The most widely and frequently used fungal biocontrol agents against fungi are *Trichoderma* spp.

They are used in different crops as bio control agents which include pulses, grapes, cotton, onions, carrots, peas, plums, maize, and apples. Due to their capacity to produce enzymes that break down polysaccharides, *Trichoderma* spp. can grow on a wide range of substrates and has a very rapid growth rate. In addition, they can withstand a variety of environmental conditions (Papavizas 1985; Elad *et al.*, 1993) ^[97, 38]. Through the use of antagonists, phytopathogenic fungi have been used to control plant diseases, *Trichoderma* has been utilized in 90% of these applications. *Trichoderma* spp. are also used to treat plant diseases brought on by nematodes and insects that rob crops in a variety of settings, including greenhouses, fields, and post-harvest (Ferreira and Musumeci 2021) ^[40]. Additionally, *Trichoderma* species provide benefits to agriculture such as increased photosynthetic capacity and yields, effective nutrient uptake, and resistance to abiotic stress (Sood *et al.*, 2020) ^[117]. Effectiveness of *Trichoderma* as a biocontrol agent, the mechanisms of its biocontrol, and the variables that affect its biocontrol is discussed in this brief review article.

Morphology and features of *Trichoderma*

Rifai 1969 and Domsch *et al.*, 1980 explained that colonies of *Trichoderma* spp. grown in culture appeared floccose, tufted green, rapidly multiplying, and sporulating well under incandescent light or produced spores in bands under normal conditions. The sterile, creeping, septate tufts of conidiophores that rise upright from a small branch are known as phialides in *Trichoderma* spp. When *T. viride* was observed, it was discovered that conidia have double-layered walls made up of an inner layer with a moderate electron density and an outer layer with a coarse outer layer known as an epipore. According to Hashioka *et al.*, 1996 ^[50], the conidia and hyphae have a distinct mucilaginous substance surrounding them. Chlamydospores are formed in the cultural media as well, but they do so later, according to Majumdar 1993 ^[73] and Sengupta 1995 ^[113]. These round, double-walled, intercalary or, rarely, terminal chlamydospores serve as resting spores under unfavorable circumstances. In order to see the branching pattern of the fungus that produces cellulolytic enzymes, *T. reesei*'s micromorphology was studied by confocal laser scanning microscopy a few years ago (Novy *et al.*, 2016) ^[95]. According to Kubicek *et al.*, 2009 ^[63] and Jaklitsch 2009 ^[57], *Trichoderma* spp. are typically found in areas with decomposing plant materials, primarily cellulosic materials. The presence of branched conidiophores with bright green conidia is the distinguishing characteristic of *Trichoderma* spp. The strain's phialides and phialospores are found in the conidiophores. *T. viride* bears globose conidia, while *T. harzianum* has light green conidia that are globose to sub-globose, according to Shah *et al.*, 2012 ^[16], *T. pseudokoningii* showed small, light green conidia.

Trichoderma as a biocontrol agent

Early in the 1930s, researchers began to notice *Trichoderma*'s efficacy as a biocontrol agent. *Trichoderma* is thought to have a biocontrol mechanism that involves mycoparasitism, competition, antibiosis, or a combination of all of these (Elad 1996) ^[36]. In 1933, Weindling claimed that a *T. lignorum* strain produced a "Lethal principle" that was secreted in the environment and caused the biocontrol agent to engage in parasitic activity. Lumsden *et al.*, 1992 ^[71]

found that *T. virens* can suppress the damping-off of zinnias brought on by *R. solani* and *Pythium ultimum*. Utilizing microorganisms as biocontrol agents to combat plant pathogens is very environmentally friendly. About 20 different species of *Trichoderma* have the potential to function as biocontrol agents for soil-borne and foliar plant pathogens. They include species with significant antagonistic characteristics such as *T. harzianum*, *T. viride*, *T. atroviride*, *T. pseudokoningii*, *T. longibrachiatum*, *T. hamatum*, *T. polysporum*, and *T. reesei* (Monaco *et al.*, 1991) ^[82]. Mukherjee and Raghu 1997 observed *Glicocladium virens* and *Trichoderma* sp. suppressing *S. rolsfii* on ginger rhizomes. Through the production of specific volatile compounds, four *Trichoderma* species were able to inhibit the growth of *Fusarium oxysporum* (Li *et al.*, 2018) ^[67].

Bicontrol Mechanisms of *Trichoderma*

The biocontrol action of *Trichoderma* is through various mechanisms. The major mechanisms include antibiosis, competition and mycoparasitism. These mechanisms are described below.

Antibiosis action of *Trichoderma*

One of the primary elements determining the fungus' saprophytic capacity is antibiosis. A group of researchers (Manibhusanrao *et al.*, 1989) ^[74] revealed that antibiotics produced by *T. harzianum* such as trichodermin, suzukacillin, and alamethicin impact the morphological or physiological sequences that lead to its effective penetration. Volatile and non-volatile toxic metabolites that inhibit colonization by antagonized microorganisms are produced by most of the *Trichoderma* strains; among all these metabolites, Vey *et al.*, 2001 described the production of harzianic acid, alamethicins, tricholin, peptaibols, antibiotics, 6-penthy- α -pyrone, massoilactone, viridin, gliovirin, glisoprenins, heptelidic acid and others. Antibiotic studies on both volatile and non-volatile antibiotics revealed that *T. harzianum* and *T. viride* were particularly effective in reducing the radial growth of *S. rolsfii* (Rao and Kulkarni, 2003) ^[102]. *Trichoderma* spp. has been shown to produce volatile and nonvolatile antibiotics that are antagonistic to a variety of pathogenic fungi (Mukhopadhyay and Kaur, 1990) ^[92]. *T. harzianum*, *T. viride*, *T. aureoviride*, and *G. virens* were utilized to isolate their five local isolates from ginger rhizosphere, and their method of antagonism against *R. solani*, harming *Capsicum annum* and *C. frutescens*, was evaluated *in vitro*. Non-volatile antibiotics were found to be more efficiently effective than volatile antibiotics (Bunker and Mathur, 2001) ^[17]. There have also been reports of overproduction of antibiotics by strains, such as *T. virens* mutants producing too much gliovirin which offer similar biocontrol to the wildtype, and gliovirin deficient mutants that did not safeguard cotton seedlings from *Pythium ultimum*, while the parental strain did (Chet *et al.*, 1997) ^[22]. *T. virens* strains that are most effective as biocontrol agents may generally produce gliovirin (Howell 1998) ^[53]. Furthermore, the most successful *T. harzianum* strains generate pyrone antibiotics against *Gaeumannomyces graminis* var. *tritici*, which clearly contributed to its success. Bhagat and Pan, 2010 ^[11] used dual culture tests and antibiotic production (volatile and nonvolatile) to test 12 strains of *Trichoderma* spp. *in vitro* against *R. solani* Kuhn., which causes root and collar rot of French bean (*Phaseolus*

vulgaris L.), and found that all isolates significantly reduced *R. solani* mycelial growth. When antibiotics and various types of hydrolytic enzymes were combined and applied to *B. cinerea* and *F. oxysporum*, synergism occurred, but it was lesser when the antibiotics were incorporated first, indicating that cell wall deterioration was required to establish the relationship (Howell 2003) [54]. The synergistic effects of a *T. harzianum* endochitinase and the antibiotic gliotoxin, as well as hydrolytic enzymes and peptaibols, on *B. cinerea* conidial germination are well known (Howell 2003) [54].

Mycoparasitic action of *Trichoderma*

Through activation of plant defense systems

Trichoderma settles only the root's external layers, acting as a root endophyte as a result of a salicylic acid mediated plant reaction that prevents the fungus from becoming a systemic pathogen by preventing it from reaching vascular bundles (Alonso Ramirez *et al.*, 2014; Poveda *et al.*, 2020a) [1, 97]. *Trichoderma* is thus capable of activating systemic plant resistances against pests and/or pathogens (Poveda *et al.*, 2020b) [98]. *Trichoderma* species have been shown to parasitize not only fungus, but additionally eggs and larvae of early stages of nematodes (Zhang *et al.*, 2014; Ibrahim *et al.*, 2020) [129] and insects (Hatvani *et al.*, 2019) [51]. *T. longibrachiatum* and *T. harzianum* parasitize mature hemipteran insects of the silverleaf whitefly (*Bemisia tabaci*) and the tropical bedbug (*Cimex hemipterus*), resulting in 40% death for the whitefly in 5 days and 90% death for the bedbug in 14 days (Anwar *et al.*, 2016) [15]. When sprayed to wheat grains, *T. album* destroys 94% of the lesser grain borer (*Rhyzopertha dominica*) in 7 days (Mohamed and Taha, 2017) [2]. Inoculating the leaves of *Cirsium arvense* and *Arabidopsis thaliana* with *T. viride* and *T. gamsii*, respectively, activates plant systemic defenses and reduces eating of the thistle tortoise beetle (*Cassida rubiginosa*) (Gange *et al.*, 2012) [43] and cabbage looper (*Trichoplusia ni*) (Zhou *et al.*, 2018) [129]. *T. harzianum*, *T. longibrachiatum*, and *T. atroviride* may stimulate S A mediated systemic defense against the potato aphid (*Macrosiphum euphorbiae*) by settling in tomato plant roots, resulting in 100% aphid mortality in 25 days. This occurs as a result of plant synthesis of VOCs such as methyl salicylate, which draws the parasitoid wasp (*Aphidius ervi*) (Coppola *et al.*, 2017, 2019a, 2019b) [25, 26, 27] as well as the predatory aphid (*Macrolophus pygmaeus*) (Battaglia *et al.*, 2013) [8]. When *T. harzianum*, *T. asperellum*, and *T. atroviride* spores are inoculated directly with hazelnut branches, they greatly diminish the beetle number by simply mycoparasitizing the beetle-symbiotic fungus. (Kushiyev *et al.*, 2020) [64].

Through cell wall degrading lytic enzymes

Chitinases derived from bacteria and fungi are thought to behaving greater potential antifungal agents than plant chitinases. Chitinases are classified as 1, 4 acetylglucosaminidases (GlcNAcases), endochitinases, and exochitinases. *T. harzianum* TM and *T. asperellum* strains both had the glcNAcases CHIT73 and CHIT102 (Haran *et al.*, 1996) [47]. Fungal chitinases have an antifungal impact due to their chitinolytic activity, particularly in *Trichoderma*, with an ED50 value equivalent to several commercial fungicides (Lorito *et al.*, 1993) [69]. *Trichoderma* derived chitinase enzymes have the ability to cause

distortions in ascomycetes and basidiomycetes' cell walls (Monte 2001) [83]. Chit42 is an endochitinase secreted by *T. harzianum* that may hydrolyze the cell walls of Botrytis cinerea and inhibit spore germination and pollen tube elongation in a variety of fungi (Marcovich and Konova, 2002) [77]. Two *T. harzianum*-derived chitinases with a similar molecular weight but separate PI values may be employed to inhibit spore germination and germ tube development in fungi from various genera such as *Fusarium*, *Gliocladium*, *Trichoderma*, *Rhizoctonia*, *Ustilago*, *Botrytis*, *Sclerotium*, and *Alternaria* (Marcovich and Konova 2002) [77]. *Trichoderma longibrachiatum*'s chitinase action may be useful in controlling Aphis gossypii infestations in cotton plants (Anwar *et al.*, 2023) [15]. The antifungal efficacy of *Trichoderma asperellum*, as well as its involvement in inducing a defensive reaction against leaf spot disease in lettuce via chitinase action, has already been established (Baiyee *et al.*, 2019) [7]. Just a few of the many identified 1, 3 glucanases have been successfully cloned, including bgn 13.1.1 am 1.3 (Cohen Kupiec 1999) [23] from *T. harzianum*, glu 78 (Donzelli *et al.*, 2001) [33] from *T. atroviride*, and Tv-bgn1 and Tv-bgn2 from *T. virens* (Kim *et al.*, 2002) [58]. Glucanases appear to be responsible for *Trichoderma*'s antagonism of plant pathogenic oomycetes such as *Pythium*. When chitinase and glucanase were expressed together in a transgenic tobacco plant, immunity towards bacterial infection was increased (Jach *et al.*, 1995) [56]. Several *Trichoderma* spp. 1, 6 glucanases have exhibited antagonistic action, either alone or in conjunction with chitinases (da Silva Aires *et al.*, 2012) [29]. Prb1, an alkaline protease isolated from *T. harzianum* IMI 206040, has been shown to have an important role in biological regulation. *T. harzianum* protease Pral has been shown to exhibit attraction for fungus cell walls, which could be effective in biocontrol (Elad *et al.*, 2000) [35]. When the *T. virens* extracellular serine protease gene (tvsp1) was cloned, it was discovered that overexpression significantly increased cotton seedling protection against *R. solani* (Pozo *et al.*, 2004) [99].

Competition mechanism of Biocontrol

Trichoderma strains, like any other beneficial antagonist, can withstand the fungistatic effects of soil caused by the presence of plant metabolites and can grow quickly in soil while being resilient to toxic substances such as herbicides, fungicides, phenolic compounds, and pesticides such as DDT (Chet *et al.* 1997) [22]. *Trichoderma* strains are very effective in managing multiple phytopathogens such as *R. solani*, *P. ultimum*, or *S. rolfii* as an alternative to chemicals such as capatan, benomyl, and others because they can recover very quickly after the addition of sublethal doses of some of these toxic compounds (Vyas and Vyas 1995) [122]. Starvation is the most prevalent cause of mortality in microorganisms. As a result, Chet *et al.*, 1997 [22] argued that competing for nutrition availability might result in fungal pathogen biological control. Iron absorption, for example, is required for most filamentous fungus to survive (Eisendle *et al.*, 2004) [34]. A variety of *Trichoderma* BCAs can produce siderophores (low-molecular-weight ferric-iron specific chelators) that are highly effective at chelating ambient iron and thereby inhibiting the growth of other fungus (Chet and Inbar 1994) [21]. As a result, the soil composition influences the efficiency of *Pythium* biocontrol by *Trichoderma* in relation to iron availability.

T. harzianum T35 management of *Fusarium oxysporum* through competition for both Rhizosphere colonization and nutrients is becoming increasingly useful when nutrient concentrations decline progressively (Tjamos 1992) [199]. Competition has become extremely significant in the biocontrol of phytopathogens such as *B. cinerea*, which is the most important pre- and post-harvest pathogenic agent in numerous countries (Latorre *et al.*, 2001) [65]. The main advantage of using *Trichoderma* to control *B. cinerea* is that *Trichoderma* uses an array of biocontrol mechanisms, making it nearly impossible for *B. cinerea* to generate a resistant strain, despite the fact that *B. cinerea* has outstanding genetic variability (Latorre *et al.* 2001) [65]. Among all methods of competition, nutritional competition is the most important because *B. cinerea* is especially vulnerable to nutrient deficit. *Trichoderma* spp. and *R. solani* may compete for nutrition, rhizosphere, and root colonization (Yu *et al.*, 2022) [127] and for seed exudates, which stimulate the growth of *R. solani* propagules in soil (Nawrocka *et al.*, 2018) [93]. *Trichoderma* genes can be used to make transgenic plants resistant to fungal pathogens.

Factors influencing biocontrol potential of *Trichoderma* species

Environmental factors

The optimal growth temperature in *Trichoderma* species varies, however the majority of them are mesophilic (Samuels 1996) [110]. Several researchers determined that water conditions have a substantial impact on *Trichoderma* activities such as spore germination and germ tube growth (Magan, 1998) [7], contact with other fungus (Badham 1991) [5], and the generation of enzymes (Grajek and Gervais 1987) [45]. Cellobiohydrolase and NAGase enzyme secretion was found to be optimal at higher water potentials, whereas secreted glucosidase, xylosidase, and chymotrypsin like enzyme activities were found to be optimal at lower water potentials (Mukhopadhyay and Kumar 2020) [92]. A group of scientists namely, Zehra *et al.*, 2017 [128] investigated the biocontrol ability of several *Trichoderma* species against *Alternaria alternata* and *Fusarium oxysporum* under various environmental circumstances such as salt, pH, and temperature, and discovered that *T. harzianum* was particularly most effective.

Practical factors influencing fungicide characters of *Trichoderma*

Preventive treatments are found effective when *Trichoderma* species are inoculated before the pathogen or when the time gap between antagonist and pathogen inoculation is concised. This was demonstrated by Diaz-Gutiérrez *et al.*, 2021 [30] through their experiment. They assessed the obstructive and curative capabilities of *T. asperellum* UDEAGIEM-H01 against *Fusarium oxysporum* in stevia plants. The *Fusarium* wilt infestation was only 10% when the antagonist was inoculated 6 days before the pathogen whereas the infestation was about 70% when the inoculation was done 6 days after the pathogen attack. This indicated the greater effectiveness of biocontrol in early inoculation of antagonist. However, protective effect was seen in a therapeutic treatment against *Phytophthora cactorum* in graft wound of pear plant when the application of *Trichoderma* strains was done 24 h after pathogen inoculation (Sánchez *et al.*, 2019) [111]. Advanced symptoms of diseases on infected plants and tissues have not been yet

reported that have shown curative effects on application of *Trichoderma* spp. In another proposition Harman, 2000 stated that *Trichoderma* spp. should be used as a integrated management strategy along with the application of systemic fungicides when the disease pressure is high as because some *Trichoderma* spp. resist amalgamation with fungicides. Another group of scientists demonstrated that the mutual application of difenoconazole-propiconazole and *T. harzianum* was observed to control 60% of the Southern corn leaf blight in maize under natural field conditions. The singular treatment of the fungicides was seen to give no better results (Wang *et al.*, 2019) [123].

The frequency of application too affects the biocontrol efficiency of *Trichoderma* spp. It was seen by Harman, 2011 that in orderly applications, 500 mg/Ha of commercial preparation 1×10^{10} cfu/g is recommendable for treatments with seed of tested crops; whereas 1×10^4 - 1×10^5 cfu/mL are applied for potting soils in greenhouse. However, upon heavy disease infestation situations, more frequent inoculations and higher concentrations of *Trichoderma* spp. are essential to reduce any more spread of the pathogens (Hanada *et al.*, 2009) [46].

The application procedure of *Trichoderma* spp. to the ecosystem effects the biocontrol efficacy (Rojo *et al.*, 2007) [104]. Application of *T. harzianum* spores by pollinators like honey bees (*Apis mellifera*) and bumblebees (*Bombus terrestris*) were seen to be more effective than spray applications to control *B. cinerea* in strawberry (Kovach *et al.*, 2000) [62]. The method of application also depends on the nature of the pathogen to be controlled, the site where it attacks and the tissues where the disease occurs. Applications in soil are useful to suppress sclerotia and other survival structures of soil borne pathogens that decreases the levels of primary inoculum of pathogens (Amira *et al.*, 2017) [2]. However, control of foliar phases of diseases caused by polycyclic pathogens would not be effectively controlled by soil application of inoculum (Elad 2000) [35].

Development of transgenics

T. virens genes encoding hydrolytic enzymes have been extracted to investigate their function in mycoparasitism and antifungal activity (Baek *et al.*, 1999; Kim *et al.*, 2002) [6, 58]. Emani *et al.*, 2003 [39] conducted research on the efficiency of the 42 kDa endochitinase genes from *T. virens* against fungal infections in cotton. Cotton plants altered with one of the 42 kDa endochitinase genes obtained from *T. virens* demonstrated a high level of resistance to *R. solani* and *A. alternata* infestation. Though previous research had shown that overexpression of plant chitinase genes in transgenics had increased disease resistance (Brogue *et al.*, 1991) [14], it was verified when *Trichoderma* chitinase (chit42) expression in tobacco and potato raised resistance to both foliar (*B. cinerea* and *Alternaria alternata*) and soilborne (*Rhizoctonia solani*) pathogens (Lorito *et al.*, 1998) [70]. Transgenic apple plants that overexpressed *T. atroviride* CHIT42 exhibited increased resistance to *Venturia inaequalis* (Bolar *et al.*, 2000) [12]. The same research group (Brants *et al.*, 2001) [13] discovered that cell cultures of transgenic tobacco expressing the chit 42 gene together with a *T. atroviride* secretion signal peptide might prevent the germination of *Penicillium digitatum* conidia. Chit42 expression in broccoli by (Mora *et al.*, 2001) [85] resulted in a considerable reduction in the severity of disease caused by

A. brassicicola in leaves, which was shown to be comparable to the efficiency of fungicides on non-transgenic plants.

When the Gluc78 gene, which encodes the exo1,3glucanase a from *T. atroviride*, was inserted into the pearl millet genome, varied levels of resistance to the downy mildew disease (*Sclerospora graminicola*) were seen in different transgenic types (O'Kennedy *et al.*, 2011)^[95]. The expression of the endochitinase chit 36 gene obtained from *T. asperelloides* T203 inhibited the growth of *Alternaria radicina* and *B. cinerea* on carrot leaves (Baranski *et al.*, 2008)^[100]. When investigated by Shah *et al.* 2010^[15], the same chit42 of another strain of *T. virens* was shown to have higher endo chitinase activity in tobacco and tomato leaf and stem tissue compared to root tissue. The first glucanase of this genus to be expressed in plants, the alpha-1, 3-glucanase gene agn 13.1 isolated from *T. harzianum*, showed antifungal and antioomycete activity, and the expression of this enzyme in *Arabidopsis trichomes* resulted in significant resistance to infection by *B. cinerea* (Calo *et al.*, 2006)^[18]. *T. virens* endochitinase gene (cht 42) identified and inserted in transgenic rice shown increased resistance to sheath blight (Shah *et al.*, 2009)^[14]. The heat shock protein (HSP) hsp70 gene was isolated from *T. harzianum* and introduced to *Arabidopsis* to induce stress tolerance (Montero-Barrientos *et al.*, 2010)^[85]. When *T. harzianum* strain genes encoding a 42 kDa endochitinase (harchit) and a chitosanase (harcho) were inserted in sorghum plants, seedling tolerant to the anthracnose disease was reported (Kosambo-Ayoo *et al.*, 2011)^[59]. A list

of genes used as bio control agents isolated from *Trichoderma* enlisted in table no 1.

Conclusion

Trichoderma can play an important part in integrated pest management which is becoming popular day by day. The combination of *Trichoderma* with GRAS drugs, cultural practices, physical techniques, and other antagonists in an integrated treatment system can be used to increase food producer output and income in a sustainable manner. Different *Trichoderma* have been isolated and studied for their biocontrol mechanisms. The principal activity noticed was the infections' development and proliferation being limited by parasitism, competition, and antibiosis. The mechanisms of biocontrol exerted by *Trichoderma* strains on various phytopathogens and pests under study have provided us with a better understanding of those mechanisms, as well as the signaling pathways and associated components involved in procedures such as host acknowledgment by *Trichoderma* spp. These approaches will aid in the efficient isolation of better enhanced strains, allowing for the development of more effective formulations for disease management before and after harvest periods. Despite the fact that *Trichoderma* has demonstrated similar or even greater efficacy than fungicides and pesticides in specific conditions (Ferreira and Musumeci 2021)^[40], complete replacement of fungicides with *Trichoderma* is still far from practical, as much work remains to be done to make this possibility a reality.

Table 1: List of some biocontrol genes isolated from *Trichoderma* strains

Sl. No.	Name of the strain	Isolated gene	Function	Reference
1.	<i>T. harzianum</i> strain IMI206040)	proteinase prb1 and endochitinase (ech42 genes	This gene expression helps to regulate hydrolytic enzymes.	Cortes <i>et al.</i> 1998 ^[28]
2.	<i>T. harzianum</i> strain P1 74058	ech 42 gene.	Disruption of this gene affects the biocontrol activity	Woo <i>et al.</i> 1999 ^[127]
3.	<i>Trichoderma</i> strain SY	Xylanase gene Xyl	Assists to breakdown hemicellulose.	Min <i>et al.</i> 2002 ^[80]
4.	<i>T. longibrachiatum</i> wild type strain CECT2606	B-1,4-endoglucanase gene, egl1	Shows better biocontrol activity against <i>Pythium ultimum</i> on cucumber	Migheli <i>et al.</i> 1998 ^[79]
5.	<i>T. atroviride</i> strain P1 (ATCC 74058)	1,3-βglucosidase gene, gluc78	Helps to degrade cell wall of <i>Pythium</i> and <i>Phytophthora</i> pathogens.	Donzelli <i>et al.</i> 2001 ^[33]
6.	<i>T. harzianum</i> strain ATCC 90237	trichodiene synthase tri5 gene	Enhances the virulence against <i>Fusarium</i> spp.	Gallo <i>et al.</i> 2004 ^[41]
7.	<i>T. virens</i> strain IMI 304061	TgaA, TgaB genes	Enhances virulence against the plant pathogenic interactions.	Mukherjee <i>et al.</i> 2004 ^[87]
8.	<i>T. hamatum</i> strain LU593	chitinase chit42 and proteinase prb1 gene	Exhibits moderate biocontrol activity against <i>Sclerotinia sclerotiorum</i> .	Steyaert <i>et al.</i> 2004
9.	<i>T. virens</i> strain IMI 304061	TmkA Mitogen Activated Protein kinase gene	This gene inhibits the formation of conidia in <i>R. solani</i> .	Mukherjee <i>et al.</i> 2003 ^[90]
10.	<i>T. virens</i> wildtype strain Gv298 and an arginine auxotrophic strain, Tv10.4	tvsp1 serine protease encoding gene	Involved in pathogenesis or biocontrol process of <i>R. solani</i> .	Pozo <i>et al.</i> 2004 ^[99]
11.	<i>T. harzianum</i> T88	beta tubulin gene	Expresses biocontrol mechanisms like mycoparasitism, and antifungal activity	Li <i>et al.</i> 2007 ^[66]
12.	<i>T. hamatum</i> LU593	monooxygenase gene	Shows enhanced antagonist activity against <i>S. sclerotiorum</i> , <i>S. minor</i> and <i>S. cepivorum</i> .	Carpenter <i>et al.</i> 2008 ^[20]
13.	<i>T. virens</i> Gv298	TvBgn2 and TvBgn3 genes	These genes help to encode cell wall degrading enzymes.	Dzonovic <i>et al.</i> 2007 ^[31]
14.	<i>T. harzianum</i> CECT 2413	erg1 gene	erg1 gene silencing increases resistance towards terbinafine that shows antifungal activity.	Cardoza <i>et al.</i> 2006 ^[19]
15.	<i>T. harzianum</i> Rifai CECT 2413	qid74 gene	This gene is involved in protection of cell and adherence to hydrophobic surfaces that aids in antagonism against <i>R. solani</i> .	Rosado <i>et al.</i> 2007 ^[106]
16.	<i>T. viride</i> IFO31137	endo β-1,6-galactanase gene	Expression of this gene influences the production of proteins.	Kotake <i>et al.</i> 2004 ^[60]

17.	<i>T. atroviride</i> strain PIATCC 74058	tga1 gene	Enhances the antifungal activity through formation of chitinase and production of antifungal metabolites.	Reithner <i>et al.</i> 2005 [103]
18.	<i>T. harzianum</i> CECT 2413	ThPTR2 gene	Induces locomotion of peptide that increases mycoparasitism	Vizcaino <i>et al.</i> 2006 [19]
19.	<i>T. virens</i> IMI 304061	tac1, adenylate cyclase gene	This gene expression leads to Mycoparasitism against <i>R. solani</i> , <i>S. rolfii</i> , <i>Pythium</i> Spp. and production of secondary metabolites.	Mukherjee <i>et al.</i> 2007 [87]
20.	<i>T. harzianum</i>	ThChit gene	This gene shows antifungal activity in transgenic tobacco.	Saiprasad <i>et al.</i> 2009 [110]
21.	<i>T. harzianum</i> CECT 2413	T34 hsp70	Enhancement of fungal resistance to heat and abiotic stresses.	MonteroBarrientos <i>et al.</i> 2008 [75]
22.	<i>T. harzianum</i>	serine protease gene SL41	This gene expresses biocontrol activity against pathogens.	Liu <i>et al.</i> 2009 [68]
23.	<i>T. atroviride</i> P1 (ATCC 74058)	Taabc2 gene	Plays important role in antagonistic role against <i>R. solani</i> , <i>P. ultimum</i> , and <i>B. cinerea</i> .	Ruocco <i>et al.</i> 2009 [108]
24.	<i>T. harzianum</i> CECT 2413	Thctf1 transcription factor gene	This gene shows antifungal action against <i>R. solani</i> , <i>Fusarium oxysporum</i> and <i>B. cinerea</i>	Rubio <i>et al.</i> 2009 [107]
25.	<i>T. harzianum</i> T34 CECT 2413	endopolygalacturonase ThPG1 gene	This gene expression helps in secretion of plant cell wall degrading enzymes against <i>R. solani</i> and <i>P. ultimum</i> .	MoranDiez <i>et al.</i> 2009 [86]
26.	<i>T. asperellum</i> (Enzymology Group collection, UFGICB)	tag 3 gene	Induces production of cell wall degrading enzyme glucanase.	Marcello <i>et al.</i> 2010 [76]
27.	<i>T. virens</i> strain TvSMOE38	Sm1 gene, cysteine rich protein	This gene codes for a small cysteine rich protein that induces defense responses in dicot and monocot plants and in protection of crop diseases.	Buensanteai <i>et al.</i> 2010 [16]
28.	<i>T. harzianum</i> E58	CRE1 gene	This gene helps to produce cellulase and hemicellulase enzymes. Accession number not available Shows enhanced biocontrol activity.	Saadia <i>et al.</i> 2008 [109]
29.	<i>T. brevicompactum</i> IBT40841	tri5 gene	Production of trichodermin and antifungal activity against <i>C. albicans</i> , <i>C. glabrata</i> and <i>A. fumigatus</i> .	Tijerino <i>et al.</i> 2011 [119]
30.	<i>T. harzianum</i> CECT 2413	Thke11 gene	This gene expression modulates glucosidase activity, and increases salt and osmotic stress tolerance in <i>A. thaliana</i>	Hermosa <i>et al.</i> 2011 [52]

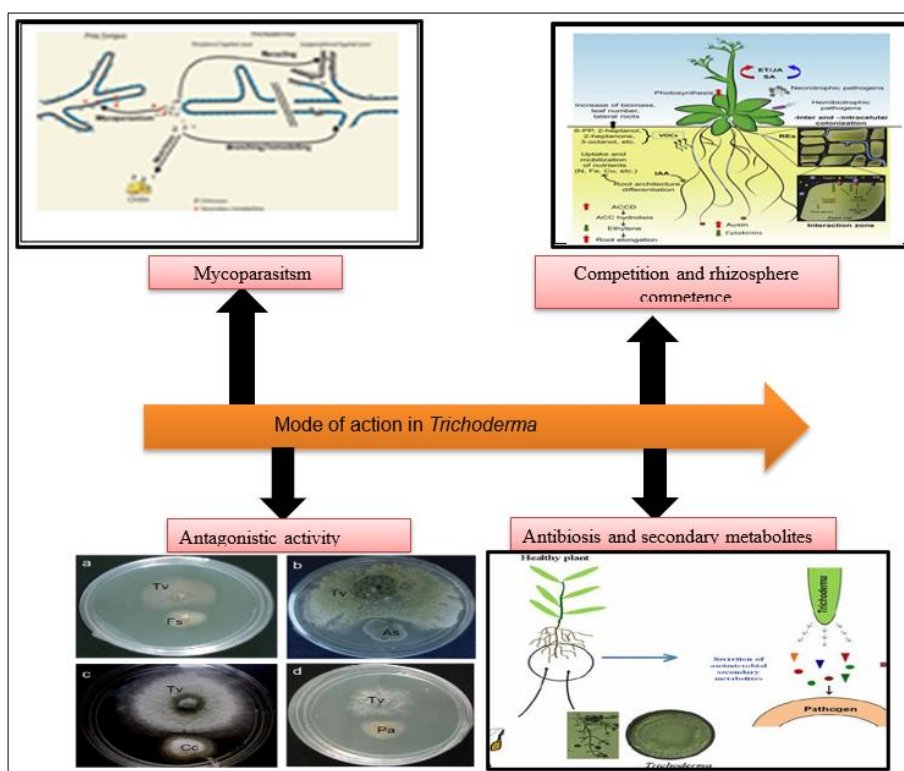


Fig 1: Different biocontrol mechanism of *Trichoderma*

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