Antifungal activity of soil actinomycetes to control chilli anthracnose caused by *Colletotrichum gloeosporioides*

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One-hundred and ninety one isolates of actinomycetes were screened for antifungal activity against anthracnose of long cayenne chilli pepper causing *C. gloeosporioides*. With this, six effective isolates of soil actinomycetes shwoed their potent antagonistic efficiency against *C. gloeosporioides*. Those isolates were belong to genus *Streptomyces* based on morphology and chemotaxonomy and 16S rDNA sequences. The non filtrates of isolates NSP1 and NSP2 were significantly highest to inhibit the colony growth of *C. gloeosporioides* and followed by isolate NSP 3. The filtrates of isolate NSP 1 gave significantly highest colony inhibition and followed by isolate NPS2. The application of NF-NSP1 before inoculation of *C. gloeosporioides* was significantly lowest number of anthracnose lesions and highest in disease reduction of 81.66 %, followed by NF-NSP1 after inoculation of *C. gloeosporioides* was significantly lowest number of anthracnose lesions and highest in disease reduction of 81.66 % the inoculated control. These effective antagonistic actinomycetes are identified as candidate to be a new *Streptomyces* spp and new candidate as biological control agents against chilli anthracnose.

Key words: Colletotrichum gloeosporioides, soil actinomycetes, antifungal activity

Introduction

Long cayenne chilli pepper (*C. annuum* var *acuminatum*) is one of the economically crop in Thailand and severely infected by *Colletotrichum* spp. (*C. capsici* and *C. gloeosporioides*) causing anthracnose disease (Than *et al.*, 2008, Waller *et al.*, 2002) due to yield loss and reduced marketability (Mananadhar *et al.*, 1995). Anthracnose disease can infect on every part of chilli and typical symptoms on stem such as leaf tip die-back, stem die-back, foliar blight, leaf

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spot, leaf lesion and fruit also include sunken necrotic tissues, with concentric rings of acervuli that are often wet (Mananadhar *et al.*, 1995; Than *et al.*, 2008).

The management and control of anthracnose disease are still under extensive research. In some case eradicative sanitation methods are not feasible, systemic fungicides were substituted. Due to the reliance on fungicides, pathogen resistance to some systemic fungicides repeatedly used throughout the year, and the deleterious to the environment and consumer were concerned. The benzimidazole fungicides, particularly carbendazim, have been reported to be effective against higher fungi. However, continuous using showed the frequency of resistant isolates in some regions of China increased gradually (Zhang and Huang, 2007). Antimicrobial organisms are interested over the decade and has been extensively studied (Shimizu et al., 2009), the actinomycetes are important producers of bioactive compounds such as chitinase and β -1, 3-glucanase (El-Tarabily *et al.*, 2000) and constitute a potential as biological control agents because of its intense antagonistic activity via the production of various antifungal substances (El-Tarabily and Sivasithamparam, 2006; Rothrock and Gottlieb, 1984; Xiao et al., 2002). The aims of this study were to isolate, identify and investigate antimicrobial activities of soil actinomycetes to control C. gloeosporioides causing chilli anthracnose disease.

Materials and methods

Isolation soil actinomycetes

Soil samples were collected from Suthep-Pui National Park and soil planted to chilli pepper brought to laboratory. Soil samples were air dried at room temperature for 7 days and to isolate thermotolerant actinomycetes, air dried soil samples were ground and pretreated at 120°C for 1 h (Xu *et al.*, 1996) become fine soil particles ready to use for isolation of actinomycetes. Soil extract agar (SEA) medium was used for isolation, modified from Takefumi et al. (2005) was prepared as soil solution by using the other organic soil from Suthep-Pui Nattional Park. It consisted of 500 g of soil sample mixed to 50 mM NaOH/L and incubated overnight at room temperature, then the soil solution was filtrated and the soil filtrate was then centrifuged for 60 min at 15,000 rpm. The 500 ml of supernatant was sterilized under 121°C at 15 psi for 15 min, then mixed into agar 20 g/L of water. The soil particles were taken in random and spread over SEA medium and incubated at room temperature. The colonies of actinomycetes were observed and transferred to yeast extract-malt extract (YEME) agar plates which consisted yeast extract 4 g, malt 10 g, glucose 4 g, agar 20g/L (Shirling and Gottlieb, 1966) for isolation into pure culture. Pure

cultures were transferred to Emerson's agar slant (Gottlieb *et al.*, 1948) which consisted of 4.0 g beef extract, 4.0 g peptone, 2.6 g NaCl, 1.0 g yeast extract, 10.0 g glucose and 20 g agar/L of water as stock cultures.

Isolation of pathogen and pathogenicity test

C. gloeosporioides causing chilli anthracnose was isolated from fruits of long cayenne chilli pepper (*C. annuum* var *acuminatum*). The fruiting bodies of pathogen were taken from anthracnose lesion and placed onto potato dextrose agar (PDA), incubated at room temperature to get pure culture. The pathogen isolates were identified and tested for pathogenicity on chilli fruits.

Screening of antifungal activities of actinomycetes

Dual culture method was used to screen effective antagonistic actinomycetes against *C. gloeosporioides*. Each actinomycete isolate was cultured on glucose yeast malt agar (GYM) which consisted of glucose 4 g, yeast extract 4 g, malt extract 10 g, CaCO₃ 2 g, agar 20 g/L (Shirling and Gottlieb, 1966). Culture of actinomycete was transferred by a single streak on one side of GYM plate at a distance of 1.5 cm and incubated at 27° C for 7 days (modified from Rothrock and Gottlieb, 1981). Thereafter, the pathogenic isolate of *C. gloeosporioides* growing on PDA for 7-days was cut using sterilized cork borer at peripheral colony to get mycelial agar plug, then transferred to the middle of dual culture plate. Data were collected as percent inhibition of colony growth (PICG) (modified from Soytong, 1989; Lokesha and Benagi, 2007). The most effective isolates of actinomycetes were used for the next experiment.

Identification of the effective soil actinomycetes

Morphological and chemotaxonomy identification: The antagonistic effective actinomycete isolates were identified into genera based on morphological characteristics and chemotaxonomy of cell wall (Williams *et al.*, 1989; Miyadoh, 1997). The morphological characters were observed as colony shape and color, sporulation, pigment diffusion etc. on GYM and IMA2 (Inhibitory mold agar consisted of glucose 5 g, soluble starch 5 g, beef extract 1 g, yeast extract 1 g, NaCl 2 g, CaCO₃ 1 g and agar 20 g/L) (Shimizu *et al.*, 2000). Chemotaxonomy of cell wall component was determined by thin layer chromatography (TLC) to detect the diaminopimelic acid (DAP or A2 pm) isomeric forms either *LL* or *meso* forms or both by using 0.01 M *LL-meso* A2 pm as a standard (Lechevalier and Lechevalier, 1970). All actinomycetes

isolates were grown on YEME agar plates for 14-days. The culture of each isolate was harvested and used as whole cells, 100 mg of biomass cells were placed in 1.5 ml microcentrifuge tube contained with 0.1 ml of 6 N HC1 and homogenized for 6 min vigorously with vortex and sterilized by autoclaving 20 min at 121°C. After cooling, the lyses cells were centrifuged for 5 min at 10000 rpm. Three μ l of supernatants were spotted directly on a thin cellulose sheet. Three μ l of 0.01 M *LL- meso* A2 pm was spotted on the same sheet as a standard. Ascending chromatography was performed on the solvent system, methanol:distilled water:6 N HC1:pyridine (80: 26: 4:10, v/v) for 3 hr. After the sheet was dried, it was sprayed with 0.2% ninhydrin reagent and heated at 100° for 5 min to reveal the spots (modified from Hasegawa *et al.*, 1983).

16s rDNA sequencing and phylogenetics analysis

Total genomic DNA of effective actinomycetes were extracted and amplified for the16S rDNA fragments from the crude lysates. The products were amplified by PCR by using universal primers 27f (5'-AGA GTT TGA TCC TGG CTCAG -3') and 1492r (5'-GGC TAC CTT GTT ACG ACTT-3') (Lane, 1991) under the following condition: 96°C for 5 min, 35 cycles of 94°C for 60 sec, 55°C for 60 sec, 72°C for 120 sec and final extension at 72°C for 3min. The amplicons were separated by 1% gel electrophoresis analysis and purified by using a Nucleospin® Extraction Kit (Biogen Ltd.) and 16S rDNA were sequenced. The resultant of 16S rDNA sequences were aligned against corresponding sequences of the type strains of *Streptomyces* species, retrieved from the GenBank and RDP databases, using the PHYDIT and TREECON programs.

Test of culture filtrates from effective actinomycetes against C. gloeosporioides

The effective actinomycete isolates were separately cultured in enzyme producing medium (EPM) modified from Rattanakit *et al.* (2002); consist of 3.0 g/L glucose, 1.0 g/L bacto peptone, 0.3 g/L urea, 1.4 g/L (NH₄)₂SO₄, 0.3 g/L MgSO₄·7H₂O, 0.3 g/L CaCl₂·6H₂O, 200 mL/L colloidal chitin (described previously on Hsu and Lockwood, 1975), trace salt solution 1.0 ml/L (5.0 g/L FeSO₄·7H₂O, 1.6 g/L MnCl₂·7H₂O, 1.4 g/L ZnSO₄·7H₂O, 2.0 g/L CoCl₂) and incubated at 35°C for 15 days with continuous shaking at150 rpm. The cultures were centrifuged for 20 min at 6,000 rpm (4°C) and the supernatants were collected as non culture filtrate (NF). The supernatant was then filtrated through membrane filter pore size 0.2-µm (Minisart[®]) to get culture filtrate (F) (Chareunrat, 1999). Agar well method was done by following the technique of

Bauer *et al.* (1996) which performed on double-layer PDA plates to test the culture filtrates (F) and non culture filtrates (NF) of each actinomycete isolate. Wells were made on upper layer of PDA plates after solidification, the upper layer of agar was punched by cork borer (5 mm diameter) about 3 cm away from center for 3 wells in each plate. One agar disc (5 mm diameter) of *C. gloeosporioides* culture was cut from peripheral colony of 5 day-old fungal culture grown in PDA at 28°C and transferred to the center of the plate. Thirty microlitre of F, NF and EPM were put to each well, respectively and the plates were incubated at 28°C for 7 days. Data were collected as percent inhibition of colony growth (PICG) modified from Soytong (1989), Lokesha and Benagi (2007).

Effect of the actinomycetes culture filtrates on chilli seeds germination

Seeds of long cayenne chilli pepper, local variety and Chomthong 2 (F1), were soaked with sterilized water overnight at room temperature. The experiment was performed using two factors in Completely Randomized Design (CRD) with three replications. Factor A was actinomycetes isolates and factor B was NF, F, EMP and distilled water. The soaked chilli seeds were placed on water-soaked blotting paper and incubated for 14 days at room temperature. Data were collected as seed germination.

Effect of actinomycetes culture filtrates as biocontrol of chilli anthracnose

antagonistic actinomycetes The effective isolates against С. gloeosporioides causing chilli anthracnose were screened and used in this experiment. The experiment was done using CRD with three replications. First experiment was done as the application before inoculation of the pathogen and treatments were NF, F from each isolate of actinomycetes and pathogen alone served as control. The application before inoculation was done by spraying five milliliter of NF and F from each isolate of actinomycetes and inoculated control on wounded leaves, 12 hours before inoculation with 10^6 conidia/mL of C. gloeosporioides. The second experiment was done as the application after inoculation of the pathogen and treatments were NF, F from each isolate of actinomycetes and pathogen alone served as control. The application after inoculation of the pathogen was done by inoculated 10^6 conidia/mL of C. gloeosporioides on wounded leaves for 12 hours, then applied NF and F from each isolate of actinomycetes and inoculated control. Both experiments, leaves of 45-days chilli seedling were used to inoculated, 4 leaves/plant were wounded by using needle pricked gently (20 spots/leaf). Data were collected as the number of anthracnose lesions per plant and computed to disease reduction as

following formula:- number of lesions in control –number of lesions in treatment/ number of lesion in control X 100.

Statistical analysis

Data were computed analysis of variance (ANOVA). Treatments mean were compared using Duncan's Multiple Range Test (DMRT) at P=0.05 and 0.01.

Results

Isolation soil actinomycetes and screening of antifungal activities of actinomycetes

One-hundred and ninety one isolates of actinomycetes were encountered from Suthep-Pui National Park and soil planted to chilli. Anthracnose on fruits of long cayenne chilli pepper was isolated and identified as *C. gloeosporioides* which was virulent isolate to the same host plant in pathogenicity test. Dual culture test showed that six isolates of soil actinomycetes revealed the highest antifungal activity against *C. gloeosporioides* as 100 % inhibition of colony growth as follows:- NSP1, NSP2, NSP3, NSP4, NSP5 and NSP6.

Identification of the effective soil actinomycetes

Morphological and chemotaxonomy identification: The actinomycetes isolates were incubated for 7-day on GYM and IMA-2 plates for morphological studies. The isolates NSP4, NSP1, NSP2, NSP6 and NSP3 are whitish mass color, rough and convex colonies whereas NSP5 showed whitish mass color and flat colony. Under microscopic observation of the six actinomycetes form coiled and flexible spore chain with more than 10 spores in chain; no diffusible soluble pigment produced on agar and showed it is consisted of *LL*-diaminopimelic acid isomer type of cell wall component (Fig. 1). Both taxonomic criteria showed that these screened isolates are similar to *Streptomyces* spp.



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Fig. 1. Separation of A2pm isomers from NSP4 (1), NSP1 (2), NSP2 (3), NSP6 (4), NSP3 (5), NSP5 (6) and 0.01 M *LL*, *meso* A_2 pm (S) by thin-layer chromatography.

16s rDNA sequencing and phylogenetics analysis

Six actinomycete isolates were subjected to a 16S rDNA sequencing to reveal its phylogenetic relationship with representative *Streptomyces*. The fragment of 16S rDNA sequences were determined (0.5 kb). The six actinomycetes isolates were clearly separated from a clade containing 27 *Streptomyces* species and closely related to *Kitasatospora* species. Their similarity to nucleotide sequences of thirty representatives of the genus, the similarity among *Kitasatospora griseola* (Syn. *Streptomyces griseolosporeus*) (Hamano *et al.*, 2002) and the isolates showed that NSP3 and NSP4 exhibited comparable levels of similarity 93.08 and 91.33%, respectively whereas NSP1, NSP2, NSP5 and NSP6 exhibited comparable levels of similarity 89.60, 89.31, 85.84 and 89.02%, respectively (Fig. 2). The evident from the 16S rDNA similarity values of the 6 representative actinomycetes indicated that the six actinomycetes are candidate to be a new *Streptomyces* spp.



Fig. 2. Phylogenetic tree of the 16S rDNA sequences from NSP4, NSP1, NSP2, NSP6, NSP3 and NSP5 isolates. The sequence data for several closely related *Kitasatospora griseola* (Syn. *Streptomyces griseolosporeus*) were recovered from GenBank and included in the tree. The percentage of replicate tree in which the associated taxa clustered together in the bootstrap test is shown above the branches. The tree is drawn to scale, with branch lengths in the same units as those of the evolutionary distances. *Nocardiopsis dassonvillei* are designated outgroup.

Test of culture filtrates from effective actinomycetes against C. gloeosporioides

The non filtrate (NF) of NSP1 and NSP2 was significantly highest to inhibit the colony growth of *C. gloeosporioides* of 66.67 and 70.00 %, respectively, and followed by NSP3 (58.33%). NSP1 and NSP5 were not significantly differed which inhibited the colony growth of 55.00 %. The filtrates (F) of NSP1 gave significantly highest colony inhibition (55.00 %) and followed by NSP2 (50.00%). The filtrates from other isolates showed colony inhibition lower than 50 %. (Table1).

Table 1. Culture and non-culture filtrates of effective Streptomyces sp. againstC. gloeosporioides

Isolates of Streptomyces sp.	Inhibition of colony growth (%)	
NSD4	NF	53.33cd ¹
11314	F	15.00f
NGD1	NF	66.67a
INSF I	F	55.00bc
NGD2	NF	70.00a
INSF2	F	50.00d
NSD6	NF	53.33cd
1131.0	F	0.00g
NGD2	NF	58.33b
NSF 3	F	1.67g
NGD5	NF	55.00bc
11013	F	25.00c

¹Means of three replications. Means followed by a common letter in each column are not significantly different by DMRT at P < 0.05.

Effect of the actinomycetes culture filtrates on chilli seeds germination

The actinomycetes isolates NSP1, NSP2 and NSP3 which gave very good results of colony inhibition were then selected to test the effect of filtrates (F) and non filtrates (NF) from those isolates for seed germination of two varieties of chilli. Result showed that F and NF of isolates NSP1, NSP2 and NSP3 were significantly differed in seed germination in local variety, which lower seed germination than non-treated control. While, in Chomthong 2 variety showed that F of NSP1, NSP2 and NF of NSP2 and NSP3 were not significantly differed in seed germination when compared to the non-treated one (Table 2).

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Treatments	Isolates	Chilli seed germina	Chilli seed germination (%)				
		local variety	Chomthong 2				
Distillated water		99.00a ¹	95.00abc				
EPM		93.00b	98.00a				
	NSP1	96 00ab	97 00ab				

95.00ab

97.00ab

97.00ab

99.00a

98.00ab

94.00abc

94.00abc

92.00bc

91.00c

96.00ab

NSP2

NSP3

NSP1

NSP2 NSP3

Table 2. Culture and non-culture filtrates of effective *Streptomyces* sp. affected to chilli seed germination for 12 hours

¹Means of three replications. Means followed by a common letter in each column are not significantly different by DMRT at P < 0.05.

Effect of actinomycetes culture filtrates as biocontrol of chilli anthracnose

The application of NF-NSP1 before inoculation of pathogen was significantly lowest number of anthracnose lesions and highest in disease reduction was 81.66 %, and followed by NF-NSP2, F-NSP1, F-NSP3, NF-NSP3 and F-NSP2, respectively. On the other hand, the application of NF-NSP1 after inoculation of pathogen was also significantly lowest number of anthracnose lesions and highest in disease reduction was 66.66 %, and followed by NF-NSP2, NF-NSP3, F-NSP1, F-NSP3, respectively when compared to the inoculated control (Table 3).

Table 3. Number of anthracnose lesions and disease reduction after application of actinomycete culture filtrate and non-filtrate for 10 days inoculation

Treatments	applied before inoculated pathogen		applied after inoculated pathogen	
	Lesion no.	Disease reduction	Lesion no.	Disease reduction
Pathogen	20.00 a ¹	0.00 d	20.00 a	0.00 e
NF-NSP1	3.67 d	81.66 a	6.67 e	66.66 a
NF-NSP2	7.67 cd	61.66 ab	9.33 d	53.33 b
NF-NSP3	9.33 bc	53.33 bc	12.00 c	40.00 c
F-NSP1	8.00 c	60.00 b	13.33 bc	33.33 cd
F-NSP2	13.00 b	35.00 c	14.00 bc	30.00 d
F-NSP3	7.00 cd	65.00 ab	20.00 a	0.00 e
C.V.(%)	24.97	24.04	6.21	13.24

¹Means of three replications. Means followed by a common letter in each column are not significantly different by DMRT at P < 0.05.

NF

F

Discussion

Six isolates of soil actinomycetes of NSP1, NSP2, NSP3, NSP4, NSP5 and NSP6 exhibited high antagonistic efficiency against *C. gloeosporioides* causing chilli anthracnose which were identified as *Streptomyces* based on morphological and chemotaxonomic of cell wall component (Becker *et al.*, 1964; Lechevalier and Lechevalier, 1970; Cummins and Harris, 1985; Williams *et al.*, 1989; Miyadoh, 1997) and 16S rDNA sequence (Kumar *et al.*, 2007). With this, these isolates of *Streptomyces* were confirmed based on *LL*-A2pm isomer of DAP in cell wall component, on the contrary *meso*-A2pm isomer of DAP were found in rare actinomycetes which stated by Lechevalier and Lechevalier (1970).

However, those actinomycetes isolates were clearly separated from a clade that comparing 27 *Streptomyces* species and closely related to *Kitasatospora* species. Their similarity to nucleotide sequences of representatives showed 16S rDNA similarity to *Kitasatospora griseola* (Syn. *Streptomyces griseolosporeus*). These effective antagonistic actinomycetes are then identified as candidates to be a new *Streptomyces* spp.

The potential of antifungal activity of the soil actinomycetes culture as non-filtrate (NF) and filtrate (F) to inhibit growth of *C. gloeosporioides* showed that NF and F of isolates NSP1, NSP2 and NSP3 gave the highest per cent inhibition of colony growth and no adverse effect on seed germination of chlli. The inhibition of fungal colony of *C. gloeosporioides* in dual culture test in this experiment may be due to the presence of extracellular metabolites both hydrolytic enzyme and secondary antifungal compound might be produced by six isolates of *Streptomyces* (Prapagdee, 2008).

The application of NF-NSP1 before inoculation of C. gloeosporioides was significantly lowest number of anthracnose lesions of long cayenne chilli pepper (C. annuum var acuminatum) and highest in disease reduction was 81.66 %. In this study the non filtrates showed better inhibition than filtrates. The non filtrates are usually contained living cells of actinomycetes. Bibb (2005) and Prapagdee (2008) stated that the non filtrate of actinomycetes shown higher potent than filtrate treatment due to active actinomycetes cells remained in culture and may be produced antibiosis and secondary metabolites against the pathogen cells. In this study, the tested actinomycetes were cultured for 15 days to obtain filtrates and it showed efficacy to control chili anthracnose. Abyad et al. (1993) and Chang (2007) reported that degree of growth inhibition were decreased in association with incubation period of the fungal culture were increased (El-Abyad et al., 1993, Chang, 2007). It is suggested that further study would be concerned on incubation period of culture to get the highest potential filtrates to control the disease. It is recommended 735

that the six actinomycetes isolates from soil in this study are successfully candidates as potentially biological control agent against chilli anthracnose caused by *C. gloeosporioides*

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