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35	Abstract	<p>White root disease causing by <i>Rigidoporus</i> sp. is a severe problem that decreases latex productivity and can even cause mortality of rubber trees. With the aim to control biologically this disease, antifungal actinobacteria were isolated from rhizospheric soils of some medicinal plants cultivated in Thailand. Among all isolated actinobacteria, an isolate TM32 exhibited distinctive antagonistic activity against the fungus. Cell-free culture broth of the isolate showed median effective dose (ED<sub>50</sub>) of 2.61 ml l<sup>-1</sup> (equal to 1.19 g l<sup>-1</sup> of metalaxyl). The isolate was also able to solubilize phosphate and to produce chitinase (enzyme activity = 0.093 ± 0.004 U ml<sup>-1</sup>), siderophore (average clear zone, 11.75 ± 0.96 mm) and indole-3-acetic acid (54.00 ± 1.00 µg ml<sup>-1</sup>). Application of biocontrol starters produced by this isolate in nursery stage of rubber trees farming showed greater suppression of the disease than direct use of its biocontrol agents. The biocontrol starters also enhanced growth of the rubber trees by increasing their heights. This might be due to the persistent growth of the isolate by using the organic substrate remaining in soil, which could later antagonize fungal pathogens through colonization at the rhizosphere and immunization of the rubber trees. The isolate revealed phylogenetically related to <i>Streptomyces sioyaensis</i> supported by 99 % similarity of 16S rRNA gene sequences. We concluded that application of the biocontrol starters produced by this <i>Streptomyces</i> isolate would be an alternative approach for sustainable control of soil-borne fungal invasion in long-term rubber tree farming.</p>
36	Keywords separated by ' - '	<i>Hevea brasiliensis</i> - Soil-borne fungal pathogen - Biocontrol - Actinobacteria - Antifungal activity
37	Foot note information	

# Utilization of rhizospheric *Streptomyces* for biological control of *Rigidoporus* sp. causing white root disease in rubber tree

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**Abstract** White root disease causing by *Rigidoporus* sp. is a severe problem that decreases latex productivity and can even cause mortality of rubber trees. With the aim to control biologically this disease, antifungal actinobacteria were isolated from rhizospheric soils of some medicinal plants cultivated in Thailand. Among all isolated actinobacteria, an isolate TM32 exhibited distinctive antagonistic activity against the fungus. Cell-free culture broth of the isolate showed median effective dose (ED<sub>50</sub>) of 2.61 ml l<sup>-1</sup> (equal to 1.19 g l<sup>-1</sup> of metalaxyl). The isolate was also able to solubilize phosphate and to produce chitinase (enzyme activity = 0.093±0.004 U ml<sup>-1</sup>), siderophore (average clear zone, 11.75±0.96 mm) and indole-3-acetic acid (54.00±1.00 µg ml<sup>-1</sup>). Application of biocontrol starters produced by this isolate in nursery stage of

rubber trees farming showed greater suppression of the disease than direct use of its biocontrol agents. The biocontrol starters also enhanced growth of the rubber trees by increasing their heights. This might be due to the persistent growth of the isolate by using the organic substrate remaining in soil, which could later antagonize fungal pathogens through colonization at the rhizosphere and immunization of the rubber trees. The isolate revealed phylogenetically related to *Streptomyces siوياensis* supported by 99 % similarity of 16S rRNA gene sequences. We concluded that application of the biocontrol starters produced by this *Streptomyces* isolate would be an alternative approach for sustainable control of soil-borne fungal invasion in long-term rubber tree farming.

**Keywords** *Hevea brasiliensis* · Soil-borne fungal pathogen · Biocontrol · Actinobacteria · Antifungal activity

## Introduction

Rubber trees, known also as para rubber trees (*Hevea brasiliensis* (Willd. ex Adr. de Juss) Muell. Arg.), are an important economic crop in southern Thailand (Kaewchai et al. 2009). There are a number of plants that can produce natural latex but *H. brasiliensis* is a major contributor. It contributes 99 % of the natural latex in the world market for the manufacture of natural rubber products. These products include automobile tires, gloves, condoms and clothes (Nandris et al.

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1987a; Rose and Steinbüchel 2005). More than 70 % of the natural latex was produced from Southeast Asia mainly in Thailand, Indonesia and Malaysia (Rose and Steinbüchel 2005). In 2010, the export market of this natural latex from Thailand was increased 78 % over the previous year with the gross income of USD8.1 billion, and it seems to be increasing year by year (Business Report Thailand 2011). Productivity of the natural latex depends significantly on the health of the rubber trees and their planting season. The rubber trees take approximately 5–6 years to grow to be ready to harvest their latex (Evueh et al. 2011). During this long term cultivation together with preference of wet weather, there is a high risk of invasion by soil borne fungal pathogens (Kaewchai et al. 2009). Fungal infection especially at the root system of the rubber trees can cause a severe reduction in latex productivity and may lead to their mortality (Wilhelm 1973; Nandris et al. 1987a, b; Evueh and Ogbebor 2008). This infection problem critically influences the investment for rubber trees farming.

Among the soil borne fungal pathogens, a Basidiomycete, *Rigidoporus* sp. (formerly called as *Fomes lignosus*) causing white root disease is a worldwide serious problem that can deplete the productivity of the natural latex and lead to massive death of the rubber trees (Wilhelm 1973; Nandris et al. 1987a, b; Jayasuriya and Thennakoon 2007; Kaewchai et al. 2009; Ogbebor et al. 2010). The fungal pathogen lives and remains in soil for a long period of time by production of white mycelia (1–2 mm thickness) that can adhere to the surface of the root bark where their rhizomorphs develop to several metres length in the soil nearby. Healthy rubber trees around can be infected directly by this fungal network or even by root contact between the trees themselves. Infection mechanisms of this fungus involve release of extracellular enzymes that are able to decay the wood and collar of host plants, while other parts of the plants' root system are invaded by its hyphae (Nandris et al. 1987a; Nicole and Benhamou 1991; Kaewchai et al. 2009). Where the infection network develops, it causes much damage over a wide area. Therefore, healthy and immunized rubber tree seedlings are required prior to cultivation.

Chemical treatment is a choice to reduce severity of the disease and to prevent invasion of pathogenic fungi. However, this protection may be expensive. Moreover, awareness of chemical use in agriculture is an important issue due to both toxic accumulation in food and effects on biodiversity of useful soil organisms (Wilhelm 1973;

Ogbebor et al. 2010; Evueh et al. 2011). Biological control is an attempt to reduce chemical use and pollution in agriculture. It can be applied to control white root disease in rubber trees but it is limited especially in large scale plantations when the infection happened while the trees are mature. Only antagonistic fungi like *Trichoderma* spp. (Jayasuriya and Thennakoon 2007; Kaewchai and Soyong 2010), *Lentinus squarrosulus* (Idwan et al. 1992), *Chaetomium* spp. and *Aspergillus niger* (Kaewchai and Soyong 2010) have been investigated as biological agents for suppression of the white root disease in rubber trees. However, the field application of these fungi has yet been unclear. Use of potential rhizospheric actinobacteria for the biological control of this disease has not yet been studied. In rhizosphere, these actinobacteria are some of the dominant soil bacteria; they are fast-growing, and play an important role in protecting their associated plants from soil borne pathogens. In addition, most of the actinobacteria can produce plant growth hormones and exchange soil mineral nutrients, which further enhance growth and improve plant health (Tokala et al. 2002; Vasconcellos and Cardoso 2009; Loqman et al. 2009).

In order to reduce the use of chemical fungicides and to sustain rubber tree farming with a biological approach, the main objective of this work is to isolate potential actinobacteria from rhizospheric soils of medicinal plants for biological control of *Rigidoporus* sp. causing white root disease in rubber trees. Both in vitro modes of action for inhibiting the fungal growth and plant growth promoting abilities of some distinct actinobacterial isolates obtained were evaluated. Either direct application of biocontrol agents or biocontrol starters produced by the actinobacteria were tested as alternative approaches for suppression of the disease at the nursery stage of rubber tree farming, and the optimal approach is proposed and discussed in this article.

## Materials and methods

### White root disease causing fungus

A fungus, *Rigidoporus* sp. causing white root disease was kindly provided by the Rubber Research Center, Surat Thani province, Thailand. Pathogenicity of the fungus was confirmed by inoculation and re-isolation from living roots of rubber trees, where the disease symptoms were observed. The inoculation was done

152	after growing the fungus on 100 g of sterilized wheat	phenotypic data (morphological characteristics of	197
153	grains at 30 °C for 15 days, and the whole fungal culture	mycelia and spores, and cell wall diaminopimelic acid	198
154	was inoculated at the bottom of a pot before planting	(DAP)) into three groups including <i>Streptomyces</i> ,	199
155	rubber trees. Budded rubber trees (cultivar RRIM600) at	<i>Micromonospora</i> and unknown genus. The	200
156	an age of 8–10 months were purchased from Wangthong	actinobacterial isolates were grown on International	201
157	district, Phitsanulok, Thailand and used throughout this	<i>Streptomyces</i> project medium II (ISP2) agar (Atlas	202
158	work. The fungus was also isolated from infected roots	1946) at 30 °C for a number of days, allowing them to	203
159	of diseased rubber trees grown at the Rubber Research	form mature spores. Morphological observations were	204
160	Center. Briefly, the infected roots were washed by run-	carried out microscopically under a light microscope	205
161	ning tap water to remove soil particles for 10 min and	(Olympus BH-2, Japan). DAP isomeric analysis was	206
162	washed twice with sterilized distilled water. Small	done following a procedure described by Hasegawa	207
163	pieces of the roots were cut and put on Potato dextrose	et al. (1983).	208
164	agar (PDA) (Difco™, MD, USA) plus chloramphenicol		
165	and streptomycin at the final concentration of 50 mg l <sup>-1</sup>	Screening of antifungal actinobacteria	209
166	each. The agar plates were then incubated at 25 °C until		
167	the fungal mycelia were growing out of the roots. Iso-	Antifungal activity of all rhizospheric actinobacteria	210
168	lated fungi were sub-cultured until they were pure cul-	was assayed against <i>Rigidoporus</i> sp. by dual culture	211
169	tures and identified preliminarily based on morpholog-	technique. Each actinobacterial isolate was grown by	212
170	ical observations compared to known <i>Rigidoporus</i> sp.	streak-plate method on ISP2 agar at 30 °C for 7 days,	213
171	The observations were done microscopically and mac-	where an agar plug (5 mm Ø) of the fungus grown	214
172	roscopically. All isolates of <i>Rigidoporus</i> were allowed	previously on PDA at 30 °C for 7 days was placed	215
173	to grow on PDA and maintained at 4 °C for further use.	4 cm from the actinobacterial streak line. The dual	216
		culture plate was incubated at 30 °C, where the antifun-	217
174	Isolation of actinobacteria from rhizospheric soils	gal activity was observed everyday. Size of any apparent	218
		inhibition zone at the interaction area of both microor-	219
175	Rhizospheric soils were collected from the field of me-	ganisms was measured (approximately a week after	220
176	dicinal plants: turmeric ( <i>Curcuma longa</i> ) and ginger	incubation). Antagonistic effects of the actinobacteria	221
177	( <i>Zingiber officinale</i> ) in Chiang Mai province, Thailand.	on the physiology of the fungal mycelia at the inhibition	222
178	The soils were air-dried at ambient temperature prior to	zone were observed microscopically. Different antifun-	223
179	oven-drying at 120 °C for 1 h (Tamura et al. 1997).	gal levels were determined by different sizes of the	224
180	Dried soil samples were then pretreated with 1.5 % (w/v)	inhibition zones (Table 1).	225
181	phenol solution (Hayakawa et al. 2004). The superna-		
182	tant of this solution was serially diluted by sterilized	Phylogenetic analysis of the most potent actinobacterial	226
183	distilled water before spreading on Humic acid-vitamin	isolate	227
184	agar (Hayakawa and Nonomura 1987) plus nystatin and		
185	cycloheximide at the final concentration of 25 and	The most potent actinobacterial isolate that exhibited the	228
186	10 µg ml <sup>-1</sup> , respectively. The agar plates were incubated	largest inhibition zone against <i>Rigidoporus</i> sp. was se-	229
187	at 30 °C for a month. All visible actinobacterial isolates	lected. Its genomic DNA was extracted following the	230
188	were selected, sub-cultured and purified individually on	method described by Hopwood et al. (1985). The 16S	231
189	Hickey-Tresner (HT) agar (Atlas 1946), which were	rRNA gene was amplified by polymerase chain reaction	232
190	further maintained at 4 °C as a working stock and in	(PCR) using a pair of universal primers: 27 F (5'-AGAG	233
191	20 % (v/v) glycerol at -20 °C for a long term storage.	TTTGATCMTGGCTCAG-3') and 1525R (5'-AAGG	234
		AGGTGWTCCARCC-3') (Lane 1991). PCR product	235
192	Determination of generic abundance of rhizospheric	was purified using QIAquick PCR Purification Kit	236
193	actinobacteria	(Qiagen) and sequenced by Macrogen Inc. (Seoul, Ko-	237
		rea). The gene sequence obtained was compared with all	238
194	Total actinobacterial isolates obtained from the same	accessible sequences in GenBank database using	239
195	amount of each rhizospheric soil were counted. These	BLASTN software. Multiple sequence alignments and	240
196	rhizospheric actinobacteria were classified using some	phylogenetic tree were carried out using the neighbour-	241

**Table 1** Actinobacteria isolated from rhizospheric soils of medicinal plants

	Rhizospheric soil	Actinobacteria (no. of isolate)	Total	No. of bioactive isolate showing different sizes of inhibition zone				No. of inactive isolate
				>20 mm	16–20 mm	11–15 mm	<11 mm	
t1.4	Ginger	<i>Streptomyces</i> (82)	127	0	1	5	3	73
t1.5	( <i>Zingiber officinale</i> )	<i>Micromonospora</i> (8)		0	0	2	1	5
t1.6		Other genera (37)	0	1	1	3	6	27
t1.7	Turmeric	<i>Streptomyces</i> (48)	82	1	0	4	9	34
t1.8	( <i>Curcuma longa</i> )	<i>Micromonospora</i> (8)		0	1	1	2	4
t1.9		Other genera (26)	0	0	0	3	3	20
t1.10	Total		209	1	3	18	24	163

242 joining method (Saitou and Nei 1987) in the Molecular  
243 Evolutionary Genetics Analysis (MEGA) program ver-  
244 sion 4 (Tamura et al. 2007), where the bootstrap confi-  
245 dence levels were computed based on 1000 replications.

#### 246 Evaluation of in vitro antifungal activity

247 Some antifungal actinobacteria (giving relatively large  
248 ( $\geq 16$  mm) inhibition zones) were evaluated in their  
249 modes of antifungal action including productions of  
250 antifungal agents, siderophores and cell wall degrading  
251 enzymes. Briefly, the antifungal actinobacteria were  
252 grown individually in 100 ml of ISP2 broth at 30 °C  
253 with shaking at 120 rpm for 7 days, while their cell-free  
254 culture broths (CCBs) were prepared by filtering  
255 through 0.20- $\mu$ m filter paper (Millipore Corporation,  
256 Bedford, MA). Antifungal activity of the CCBs against  
257 *Rigidoporus* sp. was confirmed by agar well diffusion  
258 assay. Inhibition capacity (% inhibition) of the CCBs  
259 was evaluated using dry weight of fungal biomass (dry-  
260 ing at 55 °C for 3 days) after growth at 30 °C for 20 days  
261 on PDA supplemented with each CCB (at variant con-  
262 centrations of 10, 20, 30, 40 and 50 % (v/v)) that was  
263 covered with cellophane. The dry weight of fungal  
264 biomass after growth in the same conditions on PDA  
265 without CCB supplement was used as a control. The  
266 CCB giving the highest % inhibition was evaluated by  
267 its median effective dose (ED<sub>50</sub>) for inhibition of fungal  
268 growth compared with known concentrations of a chemi-  
269 cal fungicide, Metalaxyl (Local trademark, Samut  
270 Prakan, Thailand).

271 The production of siderophores was assayed using  
272 Chrome azurol S (CAS) agar (Schwyn and Neilands  
273 1987). The CCBs were individually applied to both

274 iron-restricted and iron-rich CAS agars. Appearance of  
275 a yellow to orange halo on the blue CAS agar according  
276 to the production of siderophores was observed, where  
277 the diameter of the halo was measured.

278 Production of some cell wall degrading enzymes  
279 ( $\beta$ -1,3-glucanase, cellulase, chitinase and protease)  
280 was assayed by growing the antifungal  
281 actinobacteria at 30 °C for 7 days on a set of agar  
282 media supplemented with respective substrates of  
283 the enzymes tested. The agar media used were  
284 Minimal agar (% (w/v) of yeast extract, 0.2;  
285 KH<sub>2</sub>PO<sub>4</sub>, 0.1; MgSO<sub>4</sub>, 0.5; agar powder, 1.5) plus  
286 0.5 % (w/v) of carboxymethyl cellulose (CMC) for  
287 glucanolytic and cellulolytic activities, Chitin agar  
288 (Malviya et al. 2009) for chitinolytic activity, and  
289 Skim milk agar (Atlas 1946) for proteolytic activ-  
290 ity. Staining with an aqueous solution of  
291 1 mg ml<sup>-1</sup> Congo red for 15 min followed by  
292 washing with 1 M NaCl solution was applied for  
293 observations of glucanolytic and cellulolytic activ-  
294 ities (Teather and Wood 1982). The enzyme activ-  
295 ities of CCBs derived from the antifungal  
296 actinobacteria were also determined, while these  
297 CCBs were prepared after the actinobacterial  
298 growth at 30 °C by shaking at 120 rpm, for 7 days  
299 in 100 ml of the liquid media mentioned previous-  
300 ly, except for the chitinase activity that Chitin-  
301 peptone medium (pH 6.8) (Malviya et al. 2009)  
302 was used instead. The  $\beta$ -1,3-glucanase and cellu-  
303 lase activities of the CCBs were measured by  
304 dinitrosalicylic acid method (DNS) Cattelan et al.  
305 (1999), while the chitinase activity was measured  
306 following the protocol described by Malviya et al.  
307 (2009). The protease activity was measured

- 308 according to the protocol described by Yang and  
309 Huang (1994).
- 310 Evaluation of plant growth promoting potentials
- 311 Some plant growth promoting potentials of the antifun-  
312 gal actinobacteria such as productions of indole-3-acetic  
313 acid (IAA) and ammonia together with capable of solu-  
314 bilization of phosphate, were assessed. The ability to  
315 produce IAA was evaluated following the method de-  
316 scribed by Bano and Musarrat (2003). Briefly, one ml of  
317 the same CCB as used for antifungal activity test was  
318 mixed with 2 ml of Salkowski's reagent, where the  
319 appearance of a pink colour indicates IAA production.  
320 Optical density at 530 nm was used to estimate the level  
321 of IAA produced compared with an IAA standard curve.
- 322 A full loop of each isolate of the actinobacteria was  
323 inoculated into 10 ml Peptone water (Atlas 1946) and  
324 incubated at 30 °C for 48 to 72 h. Nessler's reagent  
325 (0.5 ml) was added into the culture broth, where the  
326 colour brown turning to yellow was determined as posi-  
327 tive ammonia production.
- 328 The ability of the actinobacteria to solubilize inor-  
329 ganic phosphate was assessed using Pikovskaya's agar  
330 (pH 7) (Nautiyal 1999). The actinobacteria were inocu-  
331 lated on the agar by spot technique, and incubated at  
332 30 °C for 7 days. Appearance of a clear zone reflecting  
333 the removal of phosphate was determined as a positive  
334 phosphate solubilization. Size of the clear zone was  
335 measured by subtracting the diameter of the  
336 actinobacterial colony.
- 337 Suppression of white root disease in nursery stage  
338 of rubber trees
- 339 The budded rubber trees, cultivar RRIM600 as men-  
340 tioned previously, were screened based on similarities of  
341 fertility and size. Agricultural loamy soil was used nat-  
342 urally for planting the rubber trees in all treatments. Five  
343 treatments were carried out: 1) a control of planted  
344 rubber trees untreated (R), 2) a pathogenic treatment of  
345 R invaded by the fungus (RF), 3) a chemical treatment  
346 of RF with metalaxyl (RFM), 4) a supplemental treat-  
347 ment of R with the biocontrol agents (RBa), and 5) a  
348 treatment of RF with the biocontrol agents (RFBa). The  
349 soil invaded with the fungus was prepared by inoculat-  
350 ing the whole volume of *Rigidoporus* sp. growing on  
351 100 g of sterilized wheat grains at 30 °C for 15 days,  
352 onto the bottom of a pot before planting the rubber trees.
- The chemical treatment was prepared by mixing the soil  
with metalaxyl at the final concentration 1 mg ml<sup>-1</sup> per  
1 kg of soil. The biocontrol agents were prepared by  
growing the most potent actinobacterial isolate in  
200 ml of ISP2 broth at 30 °C with shaking at  
120 rpm for 7 days. Whole culture broth was used as  
the biocontrol agents, which was then mixed well with  
the soil at the final concentration of 6.67 % (v/v). Ten  
rubber trees were used individually for each treatment.  
All treatments were carried out for 5 months under the  
same weather condition with approximate 10 h of day-  
light period, and were watered every 3 days using  
100 ml of tap water per pot per time. Survival rate (%)  
of the rubber trees in each condition was calculated.  
Healthy score of the survival trees was observed and  
determined in 3 levels; weak = 1, medium = 2, firm = 3  
(for stem and branch), and yellow or burned = 1, yel-  
lowish green = 2, green = 3 (for leaf). The total score  
was summarized and calculated as a percentage.
- The biocontrol agents were also developed in the  
forms of biocontrol starters. Four different starters were  
prepared using different holding organic materials, in-  
cluding: starter Bs1) paddy: rice bran (3:1 v/v), starter  
Bs2) sorghum grain: agricultural loamy soil (3:1 v/v),  
starter Bs3) solely sorghum grain, and starter Bs4) solely  
agricultural loamy soil. Most prepared materials were  
sterilized once at 121 °C for 20 min before use, except  
for the soil that was sterilized twice at the same condi-  
tion. The sterilized materials (160 cm<sup>3</sup>) were inoculated  
with 50 ml of the biocontrol agents and incubated at  
30 °C for 30 days. These prepared biocontrol starters  
were individually mixed with the soil at the final con-  
centration of 10 % (v/v). These soils were used to plant  
the rubber trees under the same planting conditions  
described previously for the evaluation of the biocontrol  
agents. Five rubber trees were used individually for each  
treatment. Cumulative increased-height of survival rub-  
ber trees was recorded by subtracting initial height at the  
start of the experiments, and used to determine potential  
of each biocontrol starter.
- Statistical analysis
- Comparisons of multiple means with standard deviation  
(SDs) obtained from any treatments were performed  
using the SPSS 16.0 computer program (SPSS, Chicago  
IL, USA), with one-way analysis of variance (ANOVA)  
and Tukey's *post hoc* tests at different significant levels  
(*P*). *F*-distribution (*F*) value and its appropriate

400 significant level derived from each comparison were  
401 indicated elsewhere of this article.

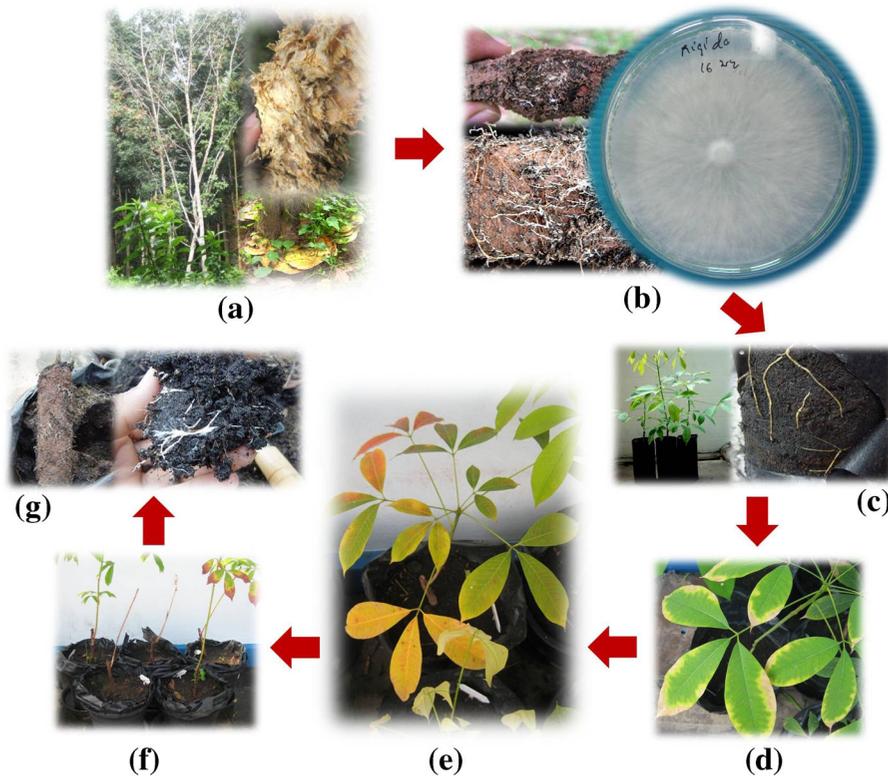
412 all areas of fibrous and lateral roots of the trees.  
413 The macroscale observation of the disease symp-  
414 toms is shown in Fig. 1.

402 **Results**

415 Actinobacteria from rhizospheric soils of medicinal  
416 plants

403 Phytopathogenicity of *Rigidoporus* sp. causing white  
404 root disease in rubber trees

417 A total of actinobacteria (209 isolates) was isolated  
418 from rhizospheric soils of medicinal plants  
419 (Table 1). Most of the actinobacteria (61 %) were  
420 found in the rhizospheric soils of ginger. On the  
421 basis of preliminary phenotypic classification,  
422 *Streptomyces* was the dominant group of  
423 actinobacteria found in rhizospheric soils of both  
424 ginger (65 %) and turmeric (58 %). Most of the  
425 actinobacteria (78 %) were inactive against



**Fig. 1** The macroscale observation of the white root disease caused by *Rigidoporus* sp. Dead rubber tree after infection by the fungus for several months standing in farm without leaf (a-left), while the fungus formed fruit body at the base of the rubber tree (a-right down) where the white fungal mycelia was observed in the inner wood of this infected zone (a-right top). The root of this rubber tree was colonized with the white fungal mycelia, which was later isolated as a pure culture of the fungus (b) (see

also **Materials and Methods**). Greenish leaves with firm root system were determined as healthy rubber trees (c). After inoculating the fungus into the soil for cultivation of rubber trees, the leaf color turned to yellow at week 7–8 (d) and be burned at week 10–12 (e). Most leaves had fallen at week 12–13, determined as the dead rubber trees (f). At the base and root of the dead rubber trees is a covering with white mycelia of the fungus (g)

t2.1 **Table 2** Modes of action against *Rigidoporus* sp. and plant growth promoting potentials of distinct antifungal actinobacteria isolated from rhizospheric soils of medicinal plants

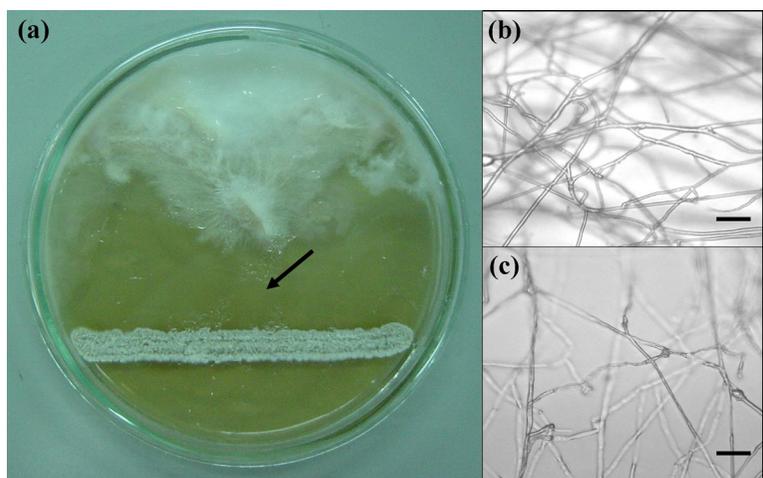
t2.2	Modes of action and plant growth promoting potentials	Actinobacterial isolate			
t2.3		TM32	GN12	GN15	GN20
t2.4	Antagonistic activity [inhibition zone (mm)]	24.00±1.00	18.33±0.58	16.00±1.00	16.67±0.58
t2.5	Siderophore production [halo zone (mm)]	11.75±0.96	8.00±1.41	18.50±1.29	5.25±0.50
t2.6	β-1,3-glucanase	0 <sup>a</sup> , 0 <sup>b</sup>	20.33±0.58 <sup>a</sup> , 0.83±0.05 <sup>b</sup>	42.50±1.32 <sup>a</sup> , 1.16±0.12 <sup>b</sup>	0 <sup>a</sup> , 0 <sup>b</sup>
t2.7	Cellulase	0 <sup>a</sup> , 0 <sup>b</sup>	20.33±0.58 <sup>a</sup> , 0.0097±0.0021 <sup>b</sup>	42.50±1.32 <sup>a</sup> , 0.71±0.13 <sup>b</sup>	0 <sup>a</sup> , 0 <sup>b</sup>
t2.8	Chitinase	22.67±0.58 <sup>a</sup> , 0.093±0.004 <sup>b</sup>	0 <sup>a</sup> , 0 <sup>b</sup>	17.17±0.76 <sup>a</sup> , 0.060±0.002 <sup>b</sup>	17.50±0.50 <sup>a</sup> , 0.050±0.004 <sup>b</sup>
t2.9	Protease	0 <sup>a</sup> , 0 <sup>b</sup>	0 <sup>a</sup> , 0 <sup>b</sup>	0 <sup>a</sup> , 0 <sup>b</sup>	0 <sup>a</sup> , 0 <sup>b</sup>
t2.10	Production of indole-3-acetic acid (μg ml <sup>-1</sup> )	54.00±1.00	34.33±1.53	51.67±1.53	72.50±2.18
t2.11	Production of ammonia	–	–	+	+
t2.12	Phosphate solubilization [solubilizing zone (mm)]	8.83±0.76	0	5.50±0.50	15.17±0.76

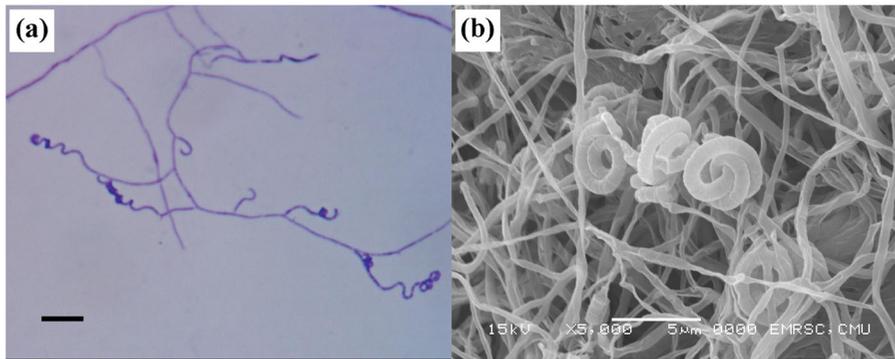
Actinobacteria isolated from rhizospheric soils of turmeric and ginger, were coded TM and GN, respectively. All bioassay results except for production of ammonia are presented as mean±standard deviation (SD), while the alphabetic indices refer to <sup>a</sup>; clear zone of enzyme assay on solid agar (mm) and <sup>b</sup>; enzyme activity of CCB (U mg<sup>-1</sup>). The production of ammonia was reported with positive (+) and negative (–) tests. All tests were done in triplicate

426 *Rigidoporus* sp., while four actinobacterial isolates  
 427 (TM32, GN12, GN15 and GN20) showed distinct  
 428 antifungal activity by giving relatively large inhi-  
 429 bition zone (≥16 mm). Only two (TM32 and  
 430 GN12) out of these four isolates belonged to the  
 431 *Streptomyces* group. Isolate TM32 originating from  
 432 the rhizospheric soil of turmeric was determined as  
 433 the most potent actinobacterial isolate, giving the  
 434 largest inhibition zone (24.00±1.00 mm) against  
 435 the fungus (Table 2). The antifungal activity of  
 436 this isolate at the inhibition zone revealed that

437 the isolates produced antifungal substances causing  
 438 the fungal mycelial wilt (Fig. 2). The isolate was  
 439 classified into the genus *Streptomyces* on the basis  
 440 of its morphology to form long spiral chain spores  
 441 (Fig. 3) and its cell wall comprised of LL-DAP  
 442 and glycine, which is the cell wall chemotype I of the  
 443 genus *Streptomyces* (Lechevalier and Lechevalier  
 444 1970). In addition, a total 1452 nucleotides sequence  
 445 of its 16S rRNA gene revealed it was closely related to  
 446 *Streptomyces sioyaensis* (DQ026654) supported by  
 447 99 % sequence similarity (Fig. 4).

**Fig. 2** Dual culture assay of *Rigidoporus* sp. and *Streptomyces* sp. TM32. The assay was done on ISP2 agar incubated at 30 °C for 7 days (a), while the arrow points at the interaction zone. Morphology of the fungal mycelia grown on the agar without isolate TM32 (b) was compared with the wilt fungal mycelia taken from the interaction zone of the dual culture (c). Bars = 10 μm



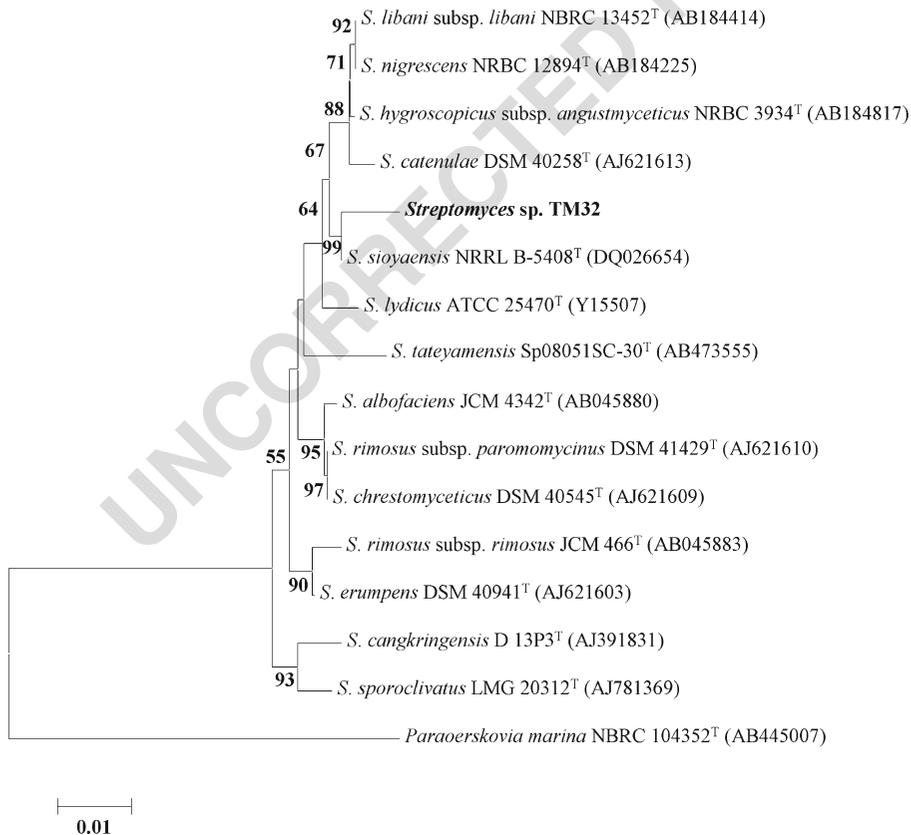


**Fig. 3** Morphology of *Streptomyces* sp. TM32. Isolate TM32 was grown on HT agar at 30 °C for 10 days. Then, the aerial mycelia and spores of isolate TM32 were observed and photographed under light (a) and scanning electron (b) microscopes. Bars=5 µm

448 Antifungal activity and plant growth promoting  
 449 potentials of the rhizospheric actinobacteria

450 Modes of action and some plant growth promoting  
 451 potentials of the four distinct actinobacterial isolates  
 452 are shown in Table 2. They inhibited the growth of

*Rigidoporus* sp. with different antifungal levels deter- 453  
 mined by the sizes of inhibition zones. They could also 454  
 produce siderophores and at least one cellulolytic en- 455  
 zyme, but no production of protease was observed. 456  
 Isolate TM32 showed the highest chitinase activity 457  
 (0.093±0.004 U mg<sup>-1</sup>) followed by isolates GN15 and 458



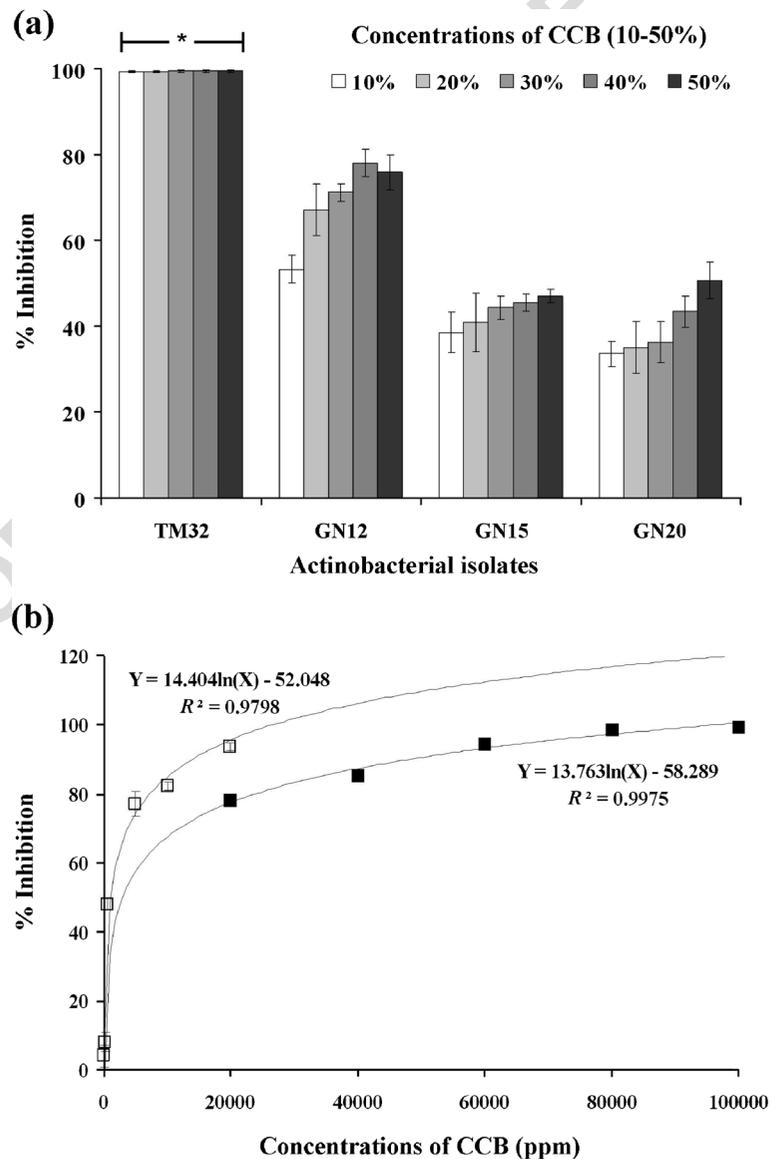
**Fig. 4** Phylogenetic tree based on the 16S rRNA gene sequence of *Streptomyces* sp. TM32. The gene sequence (1452 nucleotides without gap) of isolate TM32 (in bold) was aligned and compared to the available sequences in GenBank database (accession

numbers are given in parentheses). The out-group used was *Paraoerskovia marina*. The bootstrap values greater than 50 % of 1000 replications are shown as percentages at branching points of the tree, where the bar is 0.01 *K<sub>nuc</sub>*

459 GN20. In vitro antifungal activities of the CCBs  
 460 derived from the four isolates are shown in  
 461 Fig. 5a. The CCB derived from isolate TM32  
 462 could inhibit the fungal growth nearly 100 %,  
 463 while its inhibition values were not different sig-  
 464 nificantly across the different concentrations tested  
 465 ( $F_{(4, 15)}=0.62$ ,  $P=0.0001$ ). At the same concentra-  
 466 tion ( $10^{F1}$ ,  $20^{F2}$ ,  $30^{F3}$ ,  $40^{F4}$  and  $50\%^{F5}$  (v/v))  
 467 compared, the CCB of isolate TM32 showed sig-  
 468 nificantly greater % inhibition than those of other  
 469 CCBs derived from other isolates ( $F_{1-5(3, 12)}=$   
 470 346.92, 117.29, 361.88, 426.98 and 256.16,  $P=$

0.0001). Median effective dose ( $ED_{50}$ ) of this  
 471 CCB calculated by logistic regression equation  
 472 was  $2.61 \text{ ml l}^{-1}$  or  $0.26 \%$  (v/v) compared with a  
 473 standard curve of the known concentrations of  
 474 metalaxyl (Fig 5b). Some plant growth promoting  
 475 potentials of the four isolates are shown in Table 2.  
 476 All isolates could produce IAA, while the highest pro-  
 477 duction was found by isolate GN20 ( $72.50 \pm$   
 478  $2.18 \mu\text{g ml}^{-1}$ ). The isolate GN20 was also the greatest  
 479 phosphate solubilizer, giving the largest solubilizing  
 480 zone ( $15.17 \pm 0.76 \text{ mm}$ ). Only the isolates GN15 and  
 481 GN20 were able to produce ammonia.  
 482

**Fig. 5** In vitro antifungal activity of actinobacteria isolated from rhizosphere soils of turmeric (TM) and ginger (GN). Percent inhibition (mean $\pm$ SD) was determined by dry weight of the fungal biomass grown on PDA plus different concentrations of CCB obtained from the actinobacteria (see also [Materials and Methods](#)) (a), while the asterisk refers to no difference of means compared by ANOVA at  $P=0.0001$ . Logarithmic regression equation of the linear relation between the different concentrations of CCB obtained from *Streptomyces* sp. TM32 (black square) was compared to the chemical treatment using the known-concentration of metalaxyl (white square) (b). All experiments were done in four replications

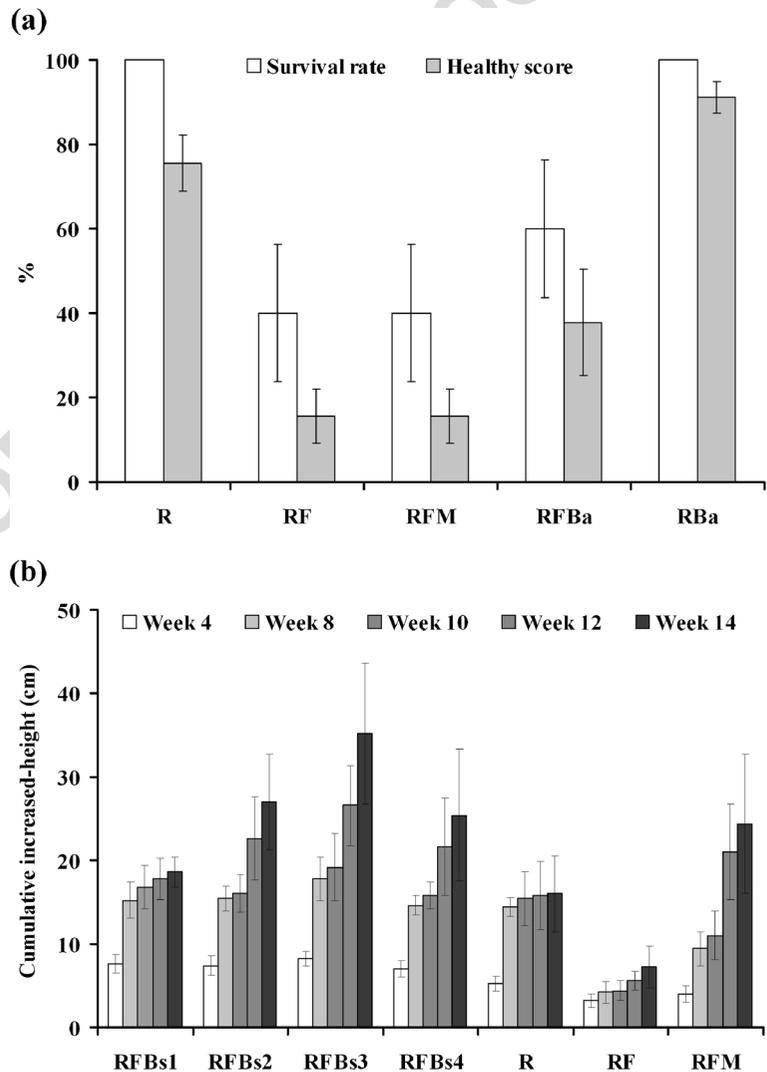


483 Utilization of *Streptomyces* sp. TM32 for suppression  
484 of white root disease in rubber trees

485 Uses of both biocontrol agents and biocontrol starters in  
486 nursery stage could suppress the disease causing by  
487 *Rigidoporus* sp. (Fig. 6). The highest survival rate  
488 (100 %) and healthy score (91 %) of rubber trees were  
489 observed in the supplemental treatment (RBa) (Fig. 6a).  
490 The chemical treatment (RFM) showed equal both sur-  
491 vival rate (40 %) and healthy score (16 %) to the  
492 pathogenic treatment (RF). The biocontrol agents could  
493 suppress the disease caused by the fungus (RFBa) to  
494 approximately 20 % greater extent than the chemical  
495 treatment. Modification of biocontrol agents in the  
496 forms of biocontrol starters (Bs1-4) was carried out

497 using different holding materials for sustaining the  
498 growth of isolate TM32. No dead rubber tree was ob-  
499 served when any biocontrol starters were applied. Potent-  
500 ial of each biocontrol starter was determined by cumu-  
501 lative increased height of the rubber trees grown in fungal  
502 invaded soil (RF) treated with each biocontrol starter  
503 (Fig. 6b). Among all biocontrol starters, starter Bs3 pre-  
504 pared with sorghum grain was the optimal one for sup-  
505 pression of the disease, revealing the highest cumulative  
506 increased-height at week 14 of the experiments. Howev-  
507 er, the statistical comparison demonstrated that the po-  
508 tential of this starter was not significantly different from  
509 starter Bs2, starter Bs4 and chemical treatment but sig-  
510 nificantly different from starter Bs1, the control and the  
511 pathogenic treatment ( $F_{(6, 28)}=10.66, P=0.01$ ).

**Fig. 6** Applications of biocontrol agent (Ba) and biocontrol starter (Bs) for suppression of white root disease caused by *Rigidoporus* sp. in nursery stage of rubber tree farming. Survival rate and healthy score of rubber trees growing in agricultural soil (R), were compared to other treatments (see also Materials and Methods) (a). Ten rubber trees were used in each treatment (a). Means of cumulative increased-height of rubber trees growing in each treatment mentioned previously together with the application of biocontrol starters (Bs1-4) were compared (see also Materials and Methods for the composition of each starter) (b). Five rubber trees were used in each treatment (b)



512 **Discussion**

513 Our results indicate that the genus *Streptomyces* is one  
 514 of the predominant bacteria found in soil habitats. Its  
 515 number isolated from the rhizospheric soils was higher  
 516 than other actinobacterial genera. The different level of  
 517 actinobacterial abundance might due to either the vari-  
 518 ations of physicochemical and biological properties of  
 519 the rhizospheric soils or the selective association be-  
 520 tween plant and microorganisms (Jayasinghe and  
 521 Parkinson 2007; Pandey and Palni 2007; Tewtrakul  
 522 and Subhadhirasakul 2007). Mostly, *Streptomyces* is a  
 523 representative producer of diverse bioactive compounds  
 524 that are used in various disciplines. Although the potent  
 525 antifungal isolate TM32 was phylogenetically related to  
 526 *Streptomyces siوياensis*, it showed some dissimilar phe-  
 527 notypes to its closest species. It was found that the  
 528 isolate possesses chitinase activity but no  $\beta$ -1,3-  
 529 glucanase activity. Otherwise,  $\beta$ -1,3-glucanase was re-  
 530 ported previously as a key tool of *S. siوياensis* for its  
 531 antifungal activity (Hong et al. 2002, 2008). We sup-  
 532 posed that the isolate TM32 might be a novel candidate  
 533 of the genus *Streptomyces*, while its complete taxonom-  
 534 ic evaluation using polyphasic approaches would be  
 535 further required to support its novelty.

536 Abilities to produce cellulolytic enzymes and  
 537 antifungal activity were observed in all distinct  
 538 antifungal actinobacteria. The isolate GN15 could  
 539 produce various enzymes but showed lowest inhibitory  
 540 level against the fungus. On the other hand, a small  
 541 amount of CCB obtained from the isolate TM32 could  
 542 inhibit the growth of the fungus equal to a gram of  
 543 metalaxyl. Although it was the most potent antifungal  
 544 actinobacterium, it was able to produce only chitinase as  
 545 a cellulolytic enzyme. This might due to either the cell  
 546 wall structure of the pathogenic fungus, comprising  
 547 chitin, or different modes of action constituted by the  
 548 antifungal isolate. Yu et al. (2008) mentioned that  
 549 chitinolytic microorganisms are considered to be more  
 550 effective antagonists of fungal pathogens because of the  
 551 direct action of chitinase alone. Based on our observa-  
 552 tion, the antifungal activity of the isolate TM32 might  
 553 not only due to the action of chitinase but also the  
 554 mixture of diverse bioactive compounds that remain in  
 555 its crude culture fluid such as antibiotics, antifungal  
 556 agents, siderophore, etc.

557 Application of *Streptomyces* sp. TM32 as a biocon-  
 558 trol agent at the nursery stage of rubber tree cultivation  
 559 did not harm to the trees, and could suppress the disease

caused by the fungus. In general, rhizospheric 560  
 actinobacteria are not only a defence against soil borne 561  
 pathogens but they can also colonize on plant root and 562  
 promote the plant growth. The capacity to suppress the 563  
 disease in nursery farming of rubber trees by our bio- 564  
 control agents showed similar results to chemical treat- 565  
 ment. However, it was found that the suppression po- 566  
 tential was greater when the biocontrol starters were 567  
 applied. Uses of both chemical and biocontrol agent 568  
 treatment seemed to be influenced by technical dilution 569  
 through watering process. This process may limit the 570  
 plant root colonization of the rhizospheric 571  
 actinobacteria, while the solid state cultivation of them 572  
 in the form of biocontrol starters could allow persistent 573  
 growth of their mycelia and later promote their mutual 574  
 coexistence in the rhizospheric habitat. 575

576 Widespread use of synthetic chemicals to control 577  
 various fungal pathogens in agriculture can induce re- 578  
 sistance to those chemicals (Para and Ristaino 1998). 579  
 With the aim to reduce chemical use in agriculture, 580  
 biological control could be an alternative approach for 581  
 the farmers. Utilization of plant-microbe association 582  
 might be a sustainable method of control of plant path- 583  
 ogens. Here, we conclude that use of biocontrol starters 584  
 from *Streptomyces* sp. TM32 could suppress the white 585  
 root disease caused by *Rigidoporus* sp. in nursery farm- 586  
 ing of rubber trees. It could also promote and immunize 587  
 the health of rubber tree seedlings, which would be a 588  
 great start for further long-term plantation of rubber 589  
 trees in the farm-scale. 590

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 597

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