

# Staurosporine from the endophytic *Streptomyces* sp. strain CNS-42 acts as a potential biocontrol agent and growth elicitor in cucumber

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**Abstract** Chinese medicinal plants and their surrounding rhizospheric soil serve as promising sources of actinobacteria. A total of 180 actinobacteria strains were isolated from the rhizosphere soil, leaves, stems, and roots of nine selected plants and have been identified as potential biocontrol agents against *Fusarium oxysporum* f. sp. *cucumerinum*. An endophytic strain CNS-42 isolated from *Alisma orientale* showed the largest zone of inhibition demonstrating a potent effect against *F. oxysporum* f. sp. *cucumerinum*

and a broad antimicrobial activity against bacteria, yeasts, and other pathogenic fungi. The in vivo biocontrol assays showed that the disease severity index was significantly reduced ( $P < 0.05$ ), and plant shoot fresh weight and height increased greatly ( $P < 0.05$ ) in plantlets treated with strain CNS-42 compared to the negative control. This isolate was identified as *Streptomyces* sp. based on cultural, physiological, morphological characteristics, and 16S rRNA gene analysis. Further bioassay-guided isolation and purification revealed that staurosporine was responsible for its antifungal and plant growth promoting activities and the latter property of staurosporine is reported for the first time. The in vivo assay was further performed and indicated that staurosporine showed good growth promoting effect on the plant shoot

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biomass of cucumber. This is the first critical evidence identifying CNS-42 as a biocontrol agent for the soil borne pathogen, *F. oxysporum* f. sp. *cucumerinum*.

**Keywords** Chinese medicinal plants · Endophytic actinobacteria · Antimicrobial activity · Plant growth promoter · Biocontrol · Cucumber · Staurosporine

## Introduction

Fusarium wilt of cucumber (*Cucumis sativus* L.) caused by *Fusarium oxysporum* f. sp. *cucumerinum* is one of the major limitations for the stable production of cucumber, especially in greenhouse vegetable production (Vakalounakis et al. 2004). A few traditional chemical methods have been used for the control of this disease. However, environmental problems could arise due to the chemical dispersion. Thus, previous studies focusing on environmentally friendly biocontrol agents have been conducted and several potential candidates have been reported (Pavlou and Vakalounakis 2005; Postma et al. 2008; Wei et al. 2008; Xian and Peng 2008; Xue et al. 2008).

Actinobacteria play a significant role in the control of plant diseases and their biocontrol mechanisms have been clearly demonstrated. Diversified secondary metabolites have been identified from actinobacteria including various antibiotics, antitumor and immunosuppressive agents, and plant growth hormones (Fiedler et al. 2008; Schulz et al. 2009; Strobel et al. 1999). Several studies have reported the antimicrobial activities of actinobacteria in the protection of host plants against pathogens thus promoting the growth of such plants (de Araújo et al. 2000; Schippers et al. 1987; Taechowisan et al. 2005; Tahvonen 1982). Nishimura et al. (2002) isolated an endophytic *Streptomyces* strain AOK-30, which showed antagonistic effect against pathogenic fungi from the cuckoo families of flowering plants. Shimizu et al. (2006) identified that disease resistance could be induced by some non-antagonistic endophytic *Streptomyces* sp. on tissue-cultured seedlings of rhododendron. Cao et al. (2004) indicated that endophytic actinobacteria could reduce the impact of ‘take-all’ on wheat.

Actinobacteria are widespread in the environment and most of the species are chemo-organotrophic, aerobic, mesophilic, and grow optimally at a pH near

neutral (Goodfellow and Williams 1983; Williams and Wellington 1982). They are affected directly by the presence of available carbon sources and their numbers are especially high in land rich in organic matters (Alexander 1977). A large number of actinobacteria have been isolated and described from traditional sources and many researchers have started to select and isolate untapped microbes from special environments. Kitouni et al. (2005) identified some strains with antimicrobial activities from water, soil, and tree barks and an endophytic *Streptomyces* from the medicinal plant *Hibbertia scandens* capable of producing antibiotics with a broad antimicrobial spectrum was isolated in Australia by Castillo et al. (2003). Two novel strains were recently isolated from the rhizosphere soil of TCMs (Wei et al. 2009). One of them showed strong inhibitory activity against the growth of *Mycobacterium smegmatis* and the other isolate could inhibit the growth of *Pseudomonas aeruginosa*. Yuan et al. (2009) isolated a total of 959 strains from 62 rhizosphere soil samples of medicinal plants from Yunnan province and Chongqing area, China, among which 31 % of these isolates showed positive bioactivities.

In this study, we attempted to investigate endophytic or rhizospheric actinobacteria from Chinese medicinal plants as potential biocontrol agents against *F. oxysporum* f. sp. *cucumerinum* and a plant growth promoter of cucumber. A total of 180 actinobacteria were isolated from the rhizosphere soils, leaves, stems, and roots of selected plants. An endophytic strain CNS-42, identified as *Streptomyces* sp., isolated from *Alisma orientale* showed the strongest antagonistic effect against *F. oxysporum* f. sp. *cucumerinum* in vivo. Further bioassay-guided purification and in vivo evaluation revealed that staurosporine was responsible for its antifungal and plant growth promotion activities.

## Materials and methods

### General experimental procedures

UV spectra were obtained on a Cary 50 spectrophotometer. NMR spectra were obtained on a Bruker Avance DRX600 spectrometer. HRESIMS measurements were obtained on a Bruker micro TOF mass spectrometer. Resin HP-20 (Diaion, Japan), and ODS-

A (YMC, Japan) were used for purification. RP-HPLC was carried out using an Eclipse XDB-C18 column (5  $\mu$ m) semi-preparative column (9.4  $\times$  250 mm) on a Shimadzu 20A series HPLC.

Isolation of the actinobacteria from Chinese medicinal plants and rhizosphere soils

Nine Chinese medicinal plants including *A. orientale*, *Platycodon grandiflorum*, *Curcuma longa*, *Platycodon grandiflorum*, *Polygonatum cyrtoneura*, *Rhizoma paridis*, *Rhodiola crenulata*, *Notopterygium incisum*, and *Astragalus membranaceus*, together with their rhizosphere soils were collected in the Sichuan province, southwest of China. The method and media used for isolating endophytic actinobacteria of plant samples were as described by Zhao et al. (2011). Isolation from the rhizosphere soils was done by serial dilution method, and  $10^{-3}$ – $10^{-5}$  dilutions were used.

In order to confirm the success of the sterilization process, aliquots of the sterile distilled water from the final rinse were inoculated on International *Streptomyces* Project-2 (ISP2) medium plates, incubated at 28 °C, and observed for microbial growth. If no microbial growth occurred on the surface of the medium, the sterilization was considered complete (Qin et al. 2009).

Incubation of endophytic strains was performed at 28 °C until single colonies could be detected. The aerial mycelium of colonies were isolated and purified on ISP2 medium. The stocks were prepared on ISP2 and kept at –70 °C (under 30 % of glycerol) for long-term storage and at 4 °C as source cultures.

Preliminary screening of the antimicrobial activities of the isolates

To select antagonistic strains, the antimicrobial activity of each isolate was tested against the pathogenic fungi (*F. oxysporum* f. sp. *cucumerinum*, this strain has been preserved at the Microbial Natural Product Library, Institute of Microbiology, Chinese Academy of Sciences, with the accession number SCPF151). The spores of actinomycetes were spread on ISP-2 and cultivated for 7 days (three repetitions). The pathogenic fungus (*F. oxysporum* f. sp. *cucumerinum*) was inoculated on potato dextrose agar (PDA) medium for 1 week (Shomura et al. 1979). A mycelia disk (6-mm diameter) of each pathogen was transferred to the

center of freshly prepared PDA medium on different plates, and three disks of actinomycetes (6-mm diameter) were inoculated onto the margin areas 3 cm from the central pathogen colony, which was followed by incubation at 28 °C for 5–7 days.

Antimicrobial spectrum of CNS-42

In order to investigate the antimicrobial spectrum of the strain CNS-42, which demonstrated the strongest inhibition of *F. oxysporum* on agar medium, six pathogenic fungi (including *F. oxysporum*, *Colletotrichum orbiculare*, *Rhizoctonia solani*, *Alternaria solani*, *F. graminearum*, and *Curvularia lunata*), *Saccharomyces cerevisiae*, and three bacteria (including *Escherichia coli* ATCC35218, *Bacillus subtilis* ATCC31785, and *Staphylococcus aureus* ATCC25923) were used in this study. The pathogenic fungi and yeast have been preserved at the Microbial Natural Product Library, Institute of Microbiology, Chinese Academy of Sciences, with the accession number SCPF151, SCPF152, SCPF153, SCPF154, SCPF155, SCPF156, and SCYF151, respectively. Mycelia disk of strain CNS-42 (6 mm diameter) was cut from the plates and inoculated in a 250 mL conical flask containing 50 mL liquid medium. The flask was then placed in a rotary shaker at 200 rpm and 28 °C for 7 days. After incubation, the culture medium was separated from the mycelium by centrifugation at 5,000 rpm for 10 min. The supernatant was used for the antimicrobial activity test using the method described previously (Thakur et al. 2007). The zones of inhibition were determined after 24 h of incubation at 37 °C for bacteria. Antifungal activity of CNS-42 was tested by the methods described in the previous section. Each experiment was replicated three times and the average zone of inhibition was calculated.

In vivo biocontrol assay using CNS-42

Four treatments (A1, B1, C1, and D1) were carried out to evaluate the ability of strain CNS-42 to control the severity of disease and promote growth of cucumber. The soil used in the experiment was steam-sterilized for 2 h at 121 °C. In the treatment A1, the spore suspension of *F. oxysporum* f. sp. *cucumerinum* (2 mL,  $3 \times 10^4$ – $4 \times 10^4$  cfu/mL) was spread in the soil when cucumber was planted. In the treatment B1, both the mycelia suspension of strain CNS-42 (2 mL,

$3 \times 10^6$ – $4 \times 10^6$  cfu/mL) and the spore suspension of *F. oxysporum* f. sp. *cucumerinum* (2 mL,  $3 \times 10^4$ – $4 \times 10^4$  cfu/mL) were spread in the soil at the same time when cucumber was planted. In the treatment C1, no microorganism solution was added to the soil. For the treatment D1, the mycelia suspension of strain CNS-42 (2 mL,  $3 \times 10^6$ – $4 \times 10^6$  cfu/mL) was spread in the soil as cucumber was planted. For each treatment, five replicates of cucumber plantlets (2-week old) were cultivated in each 15 cm diameter plastic pots. The experiment was carried out in a greenhouse at an average temperature of 25 °C, with relative humidity at about 60 % and 12 h of illumination per day at 11.8 W/m<sup>2</sup>. The plantlets were watered once every 2 days from the second day onwards and no fertilizers were added to the soil. After 6 weeks of cultivation, five plants of each treatment were removed from the pot randomly and their shoot fresh weights and heights were recorded. Disease evaluation was graded according to the wilt index (WI) scale and internal examination for the extent of rhizome discoloration graded according to the vascular discoloration index (VDI), as reported by Saravanan et al. (2003).

#### Preliminary identification of CNS-42

The morphological and cultural characteristics of the CNS-42 were determined by Gram staining and light microscopy of 14-day-old cultures grown on various International *Streptomyces* Project (ISP) media (Shirling and Gottlieb 1966). Colors of aerial and substrate mycelia were determined with the ISCC-NBS centroid color charts. The main physiological characteristics of CNS-42 were performed as described by Nishimura et al. (2002). Morphological features of spores and mycelia were observed by scanning electron microscopy (Quanta 200, USA) after desiccation and coating the cells with gold (Brunk et al. 1981). This strain has been preserved at the China General Microbiological Culture Collection Center (accession no. 4.7127).

#### DNA extraction, sequencing and analysis

Genomic DNA extraction of strain CNS-42 was performed as described by the TINAamp Bacteria DNA Kit (TIANGENG BIOTECH, Beijing, China), and PCR amplification of the 16S rRNA gene was carried out with universal primers (27f: 5'-

AGAGTTTGATCCTGGCTCAG-3'; 1492r: 5'-TACG GCTACCTTGTTACGACTT-3'). The conditions used for thermal cycler (TaKaRa) was according to the following amplification profile: 95 °C for 5 min followed by 30 cycles consisting of denaturation at 95 °C for 30 s, primer annealing at 55 °C for 30 s, and primer extension at 72 °C for 1 min 45 s. At the end of the cycles, the reaction mixture was kept at 72 °C for 7 min and then cooled to 4 °C. The PCR products were purified by electrophoresis in 1.0 % agarose gels with ethidium bromide staining, and were isolated from the agarose gel by using a E.Z.N.A.<sup>TM</sup> Gel Extraction Kit (OMEGA bio-tek, China). The extracted PCR product was connected with pMD 18-T Vector (TaKaRa) and cloned. The recombinant plasmids were sequenced by TSINGKE Biological Technology Ltd. (Beijing, China).

The 16S rRNA gene sequence determined was compared with the GenBank databases by using the BLAST search program. Phylogenetic and molecular evolutionary analyses were conducted using the software included in MEGA 4.0 package. The 16S rRNA gene sequence of the strain CNS-42 and 13 type strains of actinobacteria were aligned using the Clustal X 1.83 program (Thompson et al. 1994). The evolutionary tree was inferred by using the neighbor-joining method (Saitou and Nei 1987) from the evolutionary distance data corrected by Kimura's two-parameter model (Kimura 1980) and the topology of the phylogenetic tree was evaluated by bootstrap resampling method with 1,000 replicates (Felsenstein 1985). The 16S rRNA gene sequence reported in this article was assigned by the GenBank with the accession number KJ596488.

#### Fermentation, isolation, and purification of staurosporine

Strain CNS-42 was cultivated on an ISP2 agar (Nishimura et al. 2002) plate at 28 °C for 5 days. Ten 250 mL Erlenmeyer flasks containing 40 mL of AM2 medium (starch 0.5 %, glucose 2 %, soybean meal 1 %, peptone 0.2 %, yeast extract 0.2 %, NaCl 0.4 %, K<sub>2</sub>HPO<sub>4</sub> 0.05 %, MgSO<sub>4</sub>·7H<sub>2</sub>O 0.05 %, CaCO<sub>3</sub> 0.2 %, sea salt 3.4 %, pH 7.8) was inoculated with pieces of well-grown agar cultures of the strain CNS-42 and incubated at 28 °C, 220 rpm for 4 days, which were used as seed cultures for the large-scale fermentation. Aliquots (15 mL) of the seed cultures

were aseptically transferred to 135 × 1 L Erlenmeyer flasks, each containing 300 mL of AM2 medium, and the flasks were incubated at 28 °C, 220 rpm for 7 days. The broth was combined and centrifuged (8,000 rpm, 15 min, 4 °C) to yield a supernatant and a mycelial cake. The supernatant was extracted with EtOAc (v/v: 1:1) for three times, while the mycelial cake was extracted with acetone (3 × 1 L). The organic layers were combined and dried to afford a crude extract (Figure S1a). The crude extract (14 g) was fractionated by ODS-MPLC using gradient elution from 20 to 80 % aqueous MeOH for 80 min to provide 11 fractions (Frs. 1–11). The active fraction (Fr. 5, 810 mg) was fractionated by ODS-MPLC (35 × 2.2 cm) using an elution of 32 % acetonitrile (ACN) in water to give four sub-fractions (Frs. 5–1 to 5–4). The active sub-fraction (Fr. 5–2, 450 mg) was further purified by ODS-MPLC (35 × 2.2 cm) using a gradient elution from 35 to 70 % ACN/H<sub>2</sub>O in 120 min to provide an antifungal compound **3** (90 mg) with an MIC value of 6.25 µg/mL (Figure S1b). Fr. 6 (65 mg) showed antifungal activity and were subjected to ODS-MPLC (35 × 2.2 cm, 20 % ACN/H<sub>2</sub>O), followed by reversed phase HPLC preparation (Zorbax SB-C18 250 × 9.4, 5 µm column, 2 mL/min, gradient elution from 35 to 70 % in 60 min) to afford **2** (21 mg) showed antifungal activity with an MIC value of 3.125 µg/mL (Figure S1b). Fraction 9 (2.4 g) was purified first by recrystallization to obtain staurosporine (**1**, 200 mg), and this compound showed the best anti-fungal activity (MIC = 0.1 µg/mL).

#### Evaluation of the growth promoting potential of staurosporine on cucumber

The *in vivo* growth promoting activity of staurosporine on cucumber was evaluated. Four treatments (A2, B2, C2 and D2) were carried out with four concentrations of 0, 50, 150 and 300 µg/L of staurosporine, respectively. The soil used in the experiment was steam-sterilized for 2 h at 121 °C. These experiments were carried out in a greenhouse at an average temperature of 25 °C, with relative humidity of about 60 % and 12 h of illumination per day at 11.8 W/m<sup>2</sup>. The plantlets were watered once every 2 days from the second day onwards and no fertilizers were added to the soil. After 6 weeks of cultivation, all plants of each treatment were removed from the pot randomly and

the weights of fresh and dry biomass were recorded. The dry biomass was measured after the deactivation of enzymes at 105 °C for 40–60 min and drying to constant weight at 75 °C.

#### General antimicrobial assays

Antimicrobial assays were performed according to the Antimicrobial Susceptibility Testing Standards outlined by the Clinical and Laboratory Standards Institute (CLSI) using the bacteria *S. aureus* (ATCC 6538), *P. aeruginosa* (PAO1) and *B. subtilis* (ATCC 6633). For each organism, a loopful of glycerol stock was streaked on an LB-agar plate, which was incubated overnight at 37 °C. A single bacterial colony was picked and suspended in Mueller–Hinton Broth to approximately 1 × 10<sup>4</sup> cfu/mL. A two-fold serial dilution of each compound to be tested (4,000–31.3 µg/mL in DMSO) was prepared and an aliquot of each dilution (2 µL) was added to a 96-well flat-bottom microtiter plate (Greiner). Vancomycin and ciprofloxacin were used as positive controls and DMSO as the negative control. An aliquot (78 µL) of bacterial suspension was then added to each well (to give final compound concentrations of 100–0.78 µg/mL in 2.5 % DMSO) and the plate was incubated at 37 °C aerobically for 16 h. Finally, the optical density of each well at 600 nm was measured with an EnVision 2103 Multi-label Plate Reader (Perkin-Elmer Life Sciences). MIC values were defined as the minimum concentration of compound that inhibited visible bacterial growth. All the experiments were performed in triplicates.

#### Antifungal assays

Antifungal and bioassays were performed according to a previously described (Zhang et al. 2007) protocol modified from the Clinical and Laboratory Standards Institute M-27A methods (National Committee for Clinical Laboratory Standards 2002) using the fungus *Candida albicans* SC5314. A colony of *C. albicans* was picked from a YPD agar plate and suspended in RPMI 1640 to a concentration of 1 × 10<sup>4</sup> cfu/mL. A two-fold serial dilution of each compound to be tested (4,000–31.3 µg/mL in DMSO) was prepared and an aliquot of each dilution (2 µL) was added to a flat bottom, 96-well microtiter plate (Greiner). Ketoconazole was used as the positive control and DMSO as the negative control. An aliquot (78 µL) of the fungal suspension was added to each well (to give final



compound concentrations of 100–0.78  $\mu\text{g/mL}$  in 2.5 % DMSO) and the plate was incubated at 35 °C for 16 h. Finally, the optical density of each well at 600 nm was measured with an EnVision 2103 Multilabel Plate Reader (Perkin-Elmer Life Sciences) and the antifungal MICs were defined as the minimum concentration of compound that inhibited visible fungal growth.

#### Statistical analysis

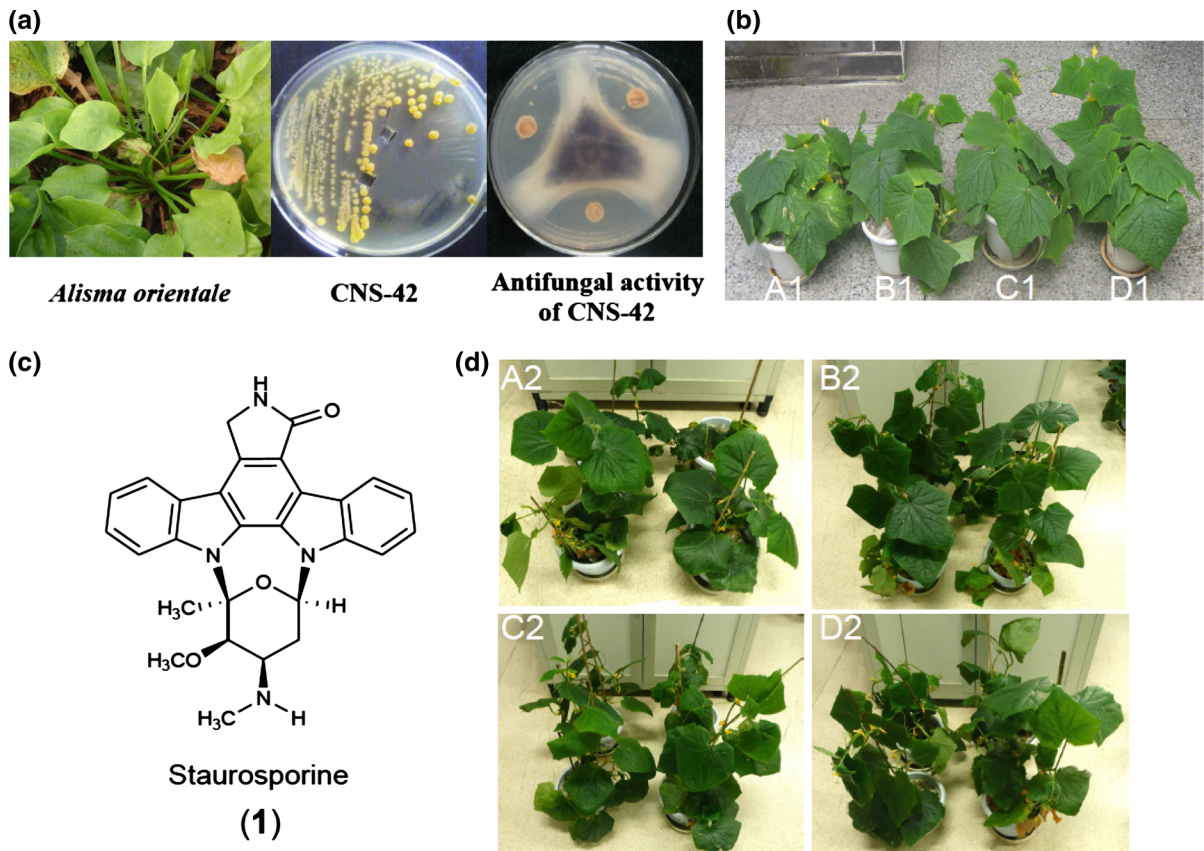
Analysis of variance (ANOVA) was conducted on collected data, and the mean values of plant shoot fresh weight, root fresh weight, height and disease index were statistically analyzed using the LSD test.

Differences were considered to be significant when the probability was  $<0.05$ .

## Results

#### In vitro antagonistic activities

Among the 180 isolates, only 21 strains showed antimicrobial activity to the pathogenic fungi (Table S1). The endophytic strain CNS-42 showed the strongest inhibitory activity against *F. oxysporum* f. sp. *cucumerinum* with inhibition zone larger than 20 mm (Fig. 1a). In addition, CNS-42 showed antimicrobial activity to *F. oxysporum*, *C. lunata*,



**Fig. 1** Overview of assessment of strain CNS-42 as a potential biocontrol agent of *Fusarium oxysporum* f. sp. *cucumerinum* and a plant growth promoter. **a** Screening antifungal activity of microbes from TCMs and their rhizosphere soil (the host medicinal plant *A. orientale*; strain CNS-42; inhibition zone of strain CNS-42 against *F. oxysporum* f. sp. *cucumerinum*, grown on PDA medium for 7 days). **b** Growth promotion of cucumber inoculated with strain CNS-42 and/or *F. oxysporum* f. sp.

*cucumerinum*. Cucumber plants grown in soil inoculated with (**A1**) *F. oxysporum* f. sp. *Cucumerinum* alone, (**B1**) *F. oxysporum* f. sp. *Cucumerinum* and CNS-42, (**C1**) neither *F. oxysporum* f. sp. *cucumerinum* nor CNS-42 as a control, (**D1**) CNS-42. **c** Structure of active compound staurosporine (**1**) isolated from CNS-42. **d** Growth promotion of cucumber treated with staurosporine (**1**). Cucumber plants grown in soil containing 0 (**A2**), 50 (**B2**), 150 (**C2**), and 300 (**D2**)  $\mu\text{g/L}$ .

*C. orbiculare*, *S. cerevisiae*, *R. solani*, *E. coli*, *A. solani*, *B. subtilis* and *S. aureus* indicating a good antimicrobial spectrum.

#### In vivo biocontrol assay of strain CNS-42

In the greenhouse experiment, the average shoot fresh weight and height of cucumber plants were measured a month after planting and the results are shown in Table 1 and Fig. 1b. With the application of the endophytic actinomycete strain CNS-42, the shoot fresh weight and height of cucumber were increased ( $P < 0.05$ ). The average shoot fresh weight and height of cucumber (49.40 g, 78.88 cm) with treatment D1 were higher than those of treatment C1 (45.22 g, 74.94 cm), which clearly demonstrated the promotion effect of CNS-42 on cucumber growth.

The application of endophytic actinomycete CNS-42 could also decrease the impact of *F. oxysporum* f. sp. *cucumerinum* on cucumber shoot fresh weight and height significantly ( $P < 0.05$ ). In treatment A1, the average shoot fresh weight and height were 31.10 g and 52.76 cm, respectively, with the introduction of the spore suspension of *F. oxysporum* f. sp. *cucumerinum*. In treatment B1, when CNS-42 was introduced to inhibit the spore suspension of *F. oxysporum* f. sp. *cucumerinum*, the shoot fresh weight and height were 41.66 g and 71.90 cm, respectively.

Even though some visual external wilt symptoms (yellowing of leaves) of plantlets in both A1 and B1 were observed and leaf chlorosis emerged firstly in older leaves and then spread to younger leaves, after 3 weeks of inoculation, treatment B1 which had endophytic actinomycete CNS-42 applied to the plant could effectively decrease the impact of *F. oxysporum*

f. sp. *cucumerinum* on cucumber growth as seen from the observation of the severity of the disease a month after their planting. For A1 experiment, a total of 16 (80 %) plantlets showed wilt symptoms and the disease index reached 50. Otherwise, for B1 experiment only 9 plantlets (45 %) showed wilt symptoms and the disease index was reduced to 25 due to the introduction of CNS-42. In the case of the controls, the plantlets for D1 and C1 experiments remained healthy, without any signs of leaf chlorosis.

#### Characterization and identification of the isolate CNS-42

Colonies of CNS-42 grown on GT (soluble starch 2.0 %, L-asparagine 0.05 %,  $\text{KNO}_3$  0.1 %,  $\text{K}_2\text{HPO}_4 \cdot \text{H}_2\text{O}$  0.05 %, NaCl 0.05 %,  $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$  0.05 %, pH 7.5) media at 35 °C for 4 days were observed to be 2–6 mm in diameter. The colonial and morphological properties (Fig. 2) and physiological characteristics (Table S3) suggest that CNS-42 is a member of the genus *Streptomyces*. Substrate mycelium is light yellow in color. Aerial mycelium produced on GT medium was gray. No diffusible pigments are obviously produced. Aerial hyphae differentiate into short, curly spores chains with smooth surface.

Through 16S rRNA gene sequence analysis, an amplified fragment of about 1,500 bp was obtained and compared with sequences of the reference species of actinobacteria contained in GenBank. The results indicated that strain CNS-42 clustered within the genus *Streptomyces* and exhibited the closest phylogenetic affinity (Figure S2) and highest sequence similarity to *Streptomyces fradiae* NBRC 12773.

**Table 1** Shoot fresh weight,height, number of infected plantlets and disease severity indices (DSI) with each treatment in greenhouse experiment

Treatments	Shoot fresh weight (g)	Height (cm)	Disease index	Infected plantlets
A1	31.10 ± 1.92 d	52.76 ± 1.74 c	50	16 (80 %)
B1	41.66 ± 2.37 c	71.90 ± 2.22 b	25	9 (45 %)
C1	45.22 ± 3.07 b	74.94 ± 2.89 b	0	0
D1	49.40 ± 2.69 a	78.88 ± 2.39 a	0	0

Average shoot fresh weight and height of five plantlets for each treatment (mean ± SD). Different letters in the same column indicate significant differences ( $P < 0.05$ ). Cucumber plants grown in soil containing A1 the spore suspension of *F. oxysporum* f. sp. *cucumerinum* (2 mL of  $3\text{--}4 \times 10^4$  cfu/mL); B1 the mycelia suspension of strain CNS-42 (2 mL of  $3\text{--}4 \times 10^6$  cfu/mL) and the spore suspension of *F. oxysporum* f. sp. *cucumerinum* (2 mL of  $3\text{--}4 \times 10^4$  cfu/mL); C1 no microorganism (2 mL of sterile tap water); D1 the mycelia suspension of CNS-42 (2 mL of  $3\text{--}4 \times 10^6$  cfu/mL)



**Fig. 2** Scanning electron micrograph of CNS-42 grown on GT agar at 35 °C for 4 days (the arrow indicates the spores chains)

#### Structure elucidation and antibacterial activity of staurosporine

The structure of staurosporine was elucidated based on UV, MS analysis,  $^1\text{H-NMR}$ ,  $^{13}\text{C-NMR}$ , and comparison with those of reported literature (Li et al. 2011). Staurosporine showed typical UV–Vis spectra (Figure S3) similar to those of known indolecarbazoles. A molecular formula  $\text{C}_{28}\text{H}_{26}\text{N}_4\text{O}_3$  of staurosporine was deduced from its HRESIMS ( $[\text{M} + \text{H}]^+$   $m/z$  467.2099, cald. 467.2078) (Figure S4). The structure of staurosporine for the identification of specific staurosporine type could be determined by further comparing its  $^1\text{H-NMR}$  (Figure S5) and  $^{13}\text{C-NMR}$  (Figure S6) with those reported previously (Li et al. 2011) (Table S2).

The pure compound staurosporine showed activities with MIC values of 0.1  $\mu\text{g/mL}$  against *C. albicans* and 12.5  $\mu\text{g/mL}$  against *B. subtilis* respectively.

#### Growth promoting activity of staurosporine

Three doses of staurosporine, 50 (B2), 150 (C2), and 300 (D2)  $\mu\text{g/L}$ , together with a negative control (A2), were applied to the growth promotion experiments. The results demonstrated that the height, size, and number of leaves were increased in B2, C2, and D2 treatments (Fig. 1d). The height, fresh and dry biomass in the treatments of B2 and D2, i.e. 50 and 300  $\mu\text{g/L}$  of staurosporine, respectively, were significantly higher than those of the control treatment ( $P < 0.05$ , Table 2). However, the results indicated that there was no growth promoting effect of staurosporine on the root biomass of plants.

#### Discussion

As compared to actinomycetes in the rhizosphere, endophytic actinomycetes are not subjected to competition with soil bacteria and can colonize plants well. Thus the endophytic actinomycetes can be considered as potential biocontrol agents (Cao et al. 2004). The endophytic *Streptomyces* plays important roles in the development and health of plants by enhancing nutrient assimilation or producing secondary metabolites (Kizuka et al. 2002; Matsukuma et al. 1994). Endophytic *Streptomyces* have been investigated as potential biological control agents of soil-borne fungal plant pathogens in previous studies and been revealed as a promising resource for the development of agricultural supplies (Cao et al. 2004; Nishimura

**Table 2** The plant promotion effects of staurosporine (1) isolated from CNS-42

Treatments	Height (cm)	Fresh biomass (g)		Dry biomass (g)	
		Plant shoot	Root	Plant shoot	Root
A2	36.55 ± 7.61 b	27.45 ± 5.99 b	1.48 ± 0.46 a	3.54 ± 1.11 b	0.21 ± 0.09 a
B2	67.61 ± 22.12 a	48.31 ± 17.39 a	2.45 ± 0.93 a	6.90 ± 4.34 a	0.27 ± 0.10 a
C2	54.64 ± 9.56 ab	31.30 ± 6.84 b	1.13 ± 0.53 a	3.44 ± 0.53 b	0.14 ± 0.07 a
D2	61.13 ± 12.19 a	45.16 ± 9.06 a	1.33 ± 0.50 a	6.67 ± 2.42 a	0.19 ± 0.04 a

Average height, shoot fresh weight and root fresh weight of four plantlets for each treatment (mean ± SD). Different letters in the same column indicate significant differences ( $P < 0.05$ )



et al. 2002; Shimizu et al. 2006; Taechowisan et al. 2005). Previous studies on endophytic actinobacteria discovered several *Streptomyces* sp. with a variety of activities against *Micrococcus luteus* or *F. oxysporum* (Sardi et al. 1992), bacteria, yeast, and filamentous fungi (Cidaria et al. 1993; Shimizu et al. 2001). Actinomycin X2 and fungichromin produced by endophytic *S. galbus* R-5 conferred some level of protection from diseases on seedlings of rhododendron (Shimizu et al. 2001).

However, our study revealed that the substances produced within tissues might not be the primary cause of acquired resistance of seedlings treated with strain CNS-42. Resistance in cucumber seedlings induced by the spread of CNS-42 in the soil may not be as a result of the production of antibiotics within tissues. As Yuan and Crawford (1995) indicated, there is no general relationship between in vitro antagonism of biocontrol candidate and in vivo disease suppression. Further field tests on the biocontrol effects of CNS-42 are underway to confirm the usefulness of the isolate under actual biocontrol situations.

Strain CNS-42 had a broad spectrum of activities against bacteria, yeasts and pathogenic fungi, and could protect cucumber seedlings from *F. oxysporum* f. sp. *cucumerinum* disease to a desirable degree when spread on the soil surface in pots. Based on cultural, physiological, morphological characteristics and the 16S rRNA gene analysis, CNS-42 was determined as one strain of *Streptomyces* sp. Further bioassay-guided isolation and purification revealed that staurosporine was responsible for its antifungal and plant growth promoting activities, which was undiscovered in previous reports. The in vivo assay was further evaluated and indicated that staurosporine showed good growth promoting effect on the aboveground biomass of cucumber growth. In the previous studies, staurosporine had been confirmed as a potent inhibitor of protein kinase C (McGlynn et al. 1992) and can induce cell apoptosis (Couldwell et al. 1994).

Our study is the first to demonstrate and report antifungal and plant growth promoting activities of staurosporine from endophytic actinobacteria. In addition, this is the first report of screening and identification of endophytic actinobacteria associated with the Chinese medicinal plant *A. orientale*. The present study indicates the great potentials of strain CNS-42 as a biological control agent against *F. oxysporum* f. sp. *cucumerinum*, which causes the root and

stem rot disease of cucumber. The mechanism involved in plant growth promotion and the range of the plant species that respond to staurosporine are still under investigation in our lab.

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