

Full Length Research Paper

# Isolation and characterization of a novel *Streptomyces* strain Eri11 exhibiting antioxidant activity from the rhizosphere of *Rhizoma Curcumae Longae*

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In the present study, the phylogenetic analysis of the *Streptomyces* strain Eri11 isolated from the rhizosphere of *Rhizoma Curcumae Longae* and the antioxidant activity of the broth cultured with Eri11 were investigated. Analysis of 16S rDNA gene sequences demonstrated that the strains Eri11 was most closely related to representatives of the genera *Streptomyces*. The total phenols and flavonoids contents in cultured broth were detected to be  $13.59 \pm 0.17$  mg gallic acid equivalent/g and  $9.93 \pm 0.83$  mg rutin equivalent/g, respectively. The cultured broth showed the antioxidant activity against the ABTS (2, 2'-Azinobis-3-ethyl benzthiazoline-6-sulfonic acid) free radicals and hydroxyl free radicals with IC<sub>50</sub> (The half-inhibitory concentration) of  $223.81 \pm 24.50$  µg/ml and  $582.42 \pm 83.10$  µg/ml respectively. So, it was suggested that the isolated *Streptomyces* strain Eri11 could be a candidate for the nature resource of the antioxidants.

**Key words:** *Streptomyces*, phylogeny, cultured broth, antioxidant activity.

## INTRODUCTION

Actinomycetes play a significant role in the pharmaceutical industry for their capacity to produce secondary metabolites with diverse chemical structures and biological activities. Tens of thousands of such compounds have been isolated and characterized, many of which have been developed into drugs for treatment of wide range of diseases in human, veterinary and agriculture sectors (Castillo et al., 2002; Singh et al., 2003; El-Shatoury et al., 2009). Searching for novel actinomycete that metabolized an essential component in natural product-based drug is becoming more and more interesting and meaningful.

*Streptomyces*, the Gram positive filamentous bacteria, belongs to actinomycetales, are widely

distributed in a variety of natural and manmade environments, constituting a significant component of the microbial population in most soils (Hwang et al., 1994; Watve et al., 2001). These bacteria produce about 75% of commercially and medically useful antibiotics and approximately 60% of antibiotics, which have been developed for agriculture use (Miyadoh et al., 1993; Takana et al., 1993). Meanwhile, previous studies showed that the *Streptomyces* could metabolize the compounds with antioxidant activity such as isoflavonoids (Komiyama et al., 1989), diphenazithionin (Hosoya et al., 1996), dihydroherbimycin A (Chang et al., 2007), poly-saccharide (He et al., 2008) and protocatechualdehyde (Kim et al., 2008).

Actinomycetes are known to form intimate associations with plants and colonize their internal tissue. *Streptomyces* spp., *Microbispora* spp., and *Streptosporangium* spp., were isolated from roots of different plant species in Italy and from maize in Brazil,

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which showed antagonistic activities against Gram positive bacteria and fungi (Sardi et al., 1992; Araujo et al., 2000). Previous studies showed that a variety of actinomycetes inhabit a wide range of plants as either symbionts or parasites (Sardi et al., 1992; Okazaki et al., 1995; Yuan and Crawford, 1995). They may have important role in plant development and health because they can affect plant growth either by nutrient assimilation or through secondary metabolite productions. The actinomycetes mainly inhabit the soil, and a large number of actinomycetes have already been isolated and described. Recently, the rate of discovery of new actinomycete has decreased, whereas the rate of re-isolation of known material has increased. Therefore, many efforts have been made to select and isolate from other biotopes, such as sea water, plant surface and plant tissues.

*Rhizoma Curcumae Longae* is the one kind of typical Chinese medical plants, which has many pharmacological functions, such as hypolipemic, antitumor, anti-inflammatory, antibacterial, antioxidant, chologagic, etc (Yang et al., 2010). There are rare reports about isolated streptomycetes associated with the rhizosphere of *R. Curcumae Longae* based on literature survey. In the present study, we isolated one streptomycetes strain Eri11 from the rhizosphere of *R. Curcumae Longae* in Ya'an city, Sichuan province, Southwest of China, and investigated the phylogeny analysis and antioxidant activity of Eri11 strain.

## MATERIALS AND METHODS

### Chemicals

2,2'-Azinobis-3-ethyl benzthiazoline-6-sulfonic acid (ABTS), gallic acid, rutin and potassium persulfate were purchased from Sigma-Aldrich (St. Louis, MO). 1,10-Phenanthroline monohydrate, hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>), Folin-Ciocalteu reagent, sodium carbonate (Na<sub>2</sub>CO<sub>3</sub>), sodium nitrite (NaNO<sub>2</sub>), aluminium trichloride (AlCl<sub>3</sub>), and ferric sulfate (FeSO<sub>4</sub>) were purchased from Zheng Chang Glass and Reagents Co., Ltd., Sichuan.

### Isolation of microorganism

Isolation of *Streptomyces* associated with rhizosphere of *R. Curcumae Longae*. In the present study, the rhizosphere of the Chinese medicinal plant *R. Curcumae Longae* was collected in Ya'an city of Sichuan province, southwest of China.

S medium (Baker, 1990), ISP-2 medium (Shirling and Gottlieb, 1966), trehalose-proline media (Jiang et al., 2006) and modified Gause II media (Jiang et al., 2006) were chosen for the isolation of actinomycetes. Nalidixic acid (15 µg/ml, w/v) and potassium dichromate (25 µg/ml, w/v) were added to suppress the growth of non-actinobacteria, and then incubated at 28°C (Cao et al., 2005).

An amount of 5 g of rhizosphere of *R. Curcumae Longae* was soaked in the 250 ml flask with 50 ml sterile 6% peptone broth. The sample was incubated at 28°C, 180 rpm for 30 min, and then 1 ml suspension of sample was diluted with sterile distilled water. An aliquot of 0.1 ml with 10<sup>-4</sup> and 10<sup>-5</sup> dilutions of the suspension were

inoculated on the isolation media and incubated at 28°C for 14 days, and then checked for microbial growth (Schulz et al., 1993). The colonies were inoculated on S medium for purification. Stocks were prepared on S medium and kept at -70°C (under 30% of glycerol) for long-term storage and at 4°C as source culture.

### Morphological identification

Morphological characteristics are the basis for identifying actinobacteria. Isolated strains were identified according to the traditional morphological criteria including characteristics of colonies on the plate, the presence of aerial mycelium, spore mass colour, distinctive reverse colony colour, diffusible pigment, and sporophore and spore chain morphology (Goodfellow and Cross, 1984).

### Physiological and biochemical characteristics

The utilization pattern of carbon sources by the strains was carried out according to the methods of Gottlieb (1961) since it can be used as an aid for species determination (Pridham and Gottlieb, 1948). Tolerance of the strains to NaCl concentration was also evaluated (Tresner et al., 1968). The actinomycete strain was tested for its ability to produce H<sub>2</sub>S and melanin pigments (Holding and Collee, 1971).

### Phylogenetic analysis (16S rDNA)

Genomic DNA extraction was done using the procedure described by Xu et al. (2003) and PCR amplification of the 16S rDNA gene was carried out with primers A and B (Xu et al., 2003). The amplified products were sent to Invitrogen Company for purification and sequencing. The quality of the sequences was verified by sequencing both strands. These sequences were submitted to the GenBank database.

Phylogenetic trees were constructed using the Neighbor-Joining method in MEGA program version 4.1 (Kumar et al., 2004). The topology of the phylogenetic tree was evaluated by bootstrap resampling method of Felsenstein with 1000 replicates (Felsenstein, 1985). The 16S rDNA gene sequences of the type strains of the various genera used in this study were retrieved from the nonredundant GenBank database and used for cladistic analysis. These sequences have been chosen as reference sequences, in which unidentified and unpublished sequences were not included.

### Extraction of the cultured broth

The spore suspensions of the culture were inoculated on S liquid media and incubated at 28°C, 180 rpm for 14 days. S broth with cultures was collected and centrifuged (3000 rpm, 30 min) to remove the *streptomyces* spore, and the cultured broth without the streptomycetes spore was collected to antioxidant test.

### Determination of total phenols and flavonoids contents

Total phenols were determined by using Folin-Ciocalteu method (Velioglu et al., 1998). Analyses were performed by visible spectrophotometry at 750 nm after reaction with Folin-Ciocalteu's reagent. In brief, an amount of 0.1 ml extract samples with different dilution were mixed with 2 ml of 20 mg/ml Na<sub>2</sub>CO<sub>3</sub> for 2 min, and then 0.9 ml of Folin-Ciocalteu's reagent (previously diluted 2-fold with distilled water) was added. The absorbance of reaction was

**Table 1.** Cultrual characterization of the *streptomyces* strain Eri11 cultured on the S solid medium.

Mycelial colouration		Melanoid pigmentation	Spore surface	Spore morphology
Aerial	Substrate			
Grey	Blue	+	Smooth	Rectiflexibilis

+ = Pigments produced; - = Not produced.

**Table 2.** Biochemical characteristics of isolated strain Eri11.

Catalase	Glucose	Xylose	Arabinose	Rhamnose	Fructose	Galactose
+	+	+	+	-	+	+
Raffinose	Mannitol	Inositol	Sucrose	1.5% NaCl	3% NaCl	5% NaCl
-	+	+	+	+	+	+

+ = Positive utilization; - = utilization negative.

measured at 750 nm by using the MAPADA V-1100D spectrophotometer (Xinke Instruments Co., Ltd., Sichuan, China) after 30 min of incubation at room temperature. Total phenol contents were calculated as gallic acid from a calibration curve:  $Y = 1.373 X - 0.044$ ,  $R^2 = 0.997$ , where Y was the absorbance and X was the gallic acid equivalent (mg gallic acid/g extract).

Total flavonoids were determined by using spectrophotometrical methods (Ordonez et al., 2006). To 0.1 ml of extract samples with different dilution, 2 ml of distilled water was mixed with 0.1 ml of 5%  $\text{NaNO}_2$  for 6 min, and then 0.2 ml of 10%  $\text{AlCl}_3$  were added and mixed for 5 min. The total volume was made up to 3 ml with distilled water. The absorbance of reaction was measured at 420 nm against a prepared blank by using the MAPADA V-1100D spectrophotometer. Total flavonoids contents were calculated as rutin from a calibration curve:  $Y = 0.525 X$ ,  $R^2 = 0.999$ , where X was the absorbance and Y was the rutin equivalent (mg rutin/g extract).

#### ABTS radical scavenging activity

ABTS radical scavenging activity was determined according to a modified method (Re et al., 1999). In brief, 19 mg of ABTS was reacted with 3.3 mg of potassium persulfate overnight in the dark at room temperature. The working solution was prepared by diluting it with water to get absorbance around 0.70 at 734 nm. An amount of 30  $\mu\text{l}$  of test sample was reacted with 2.97 ml of diluted ABTS and absorbance was recorded within 30 min at 734 nm. Trolox was used as a positive. The activity was expressed as the concentration of sample necessary to give a 50% reduction in the original absorbance ( $\text{IC}_{50}$ ).

#### Hydroxyl radical ( $\cdot\text{OH}$ ) scavenging activity

Hydroxyl radical scavenging activity was determined according to a modified method (Li et al., 2010). To 1.0 ml of 0.75 mM 1,10-phenanthroline monohydrate, 2 ml of pH 7.4 PBS buffer were mixed with 1 ml distilled water and 1 ml of 0.75 mM  $\text{FeSO}_4$  was then added. An amount of 1 ml of 0.01%  $\text{H}_2\text{O}_2$  was added after mixed. The absorbance of reaction was measured at 536 nm after 1 h of incubation at 37°C. The percentage (%) of radical scavenging activity (RSA) was calculated using the following equation:

$$\text{RSA}(\%) = [(A_S - A_P) / (A_B - A_P)] \times 100$$

where  $A_S$  is the absorbance of the sample at 536 nm, and  $A_P$  is the absorbance of the positive control which 1 ml distilled water instead of 1 ml sample in the reaction system at 536 nm, and  $A_B$  is the absorbance of the control which 1 ml distilled water was added to the reaction system instead of 1 ml of 0.01%  $\text{H}_2\text{O}_2$  at 536 nm.

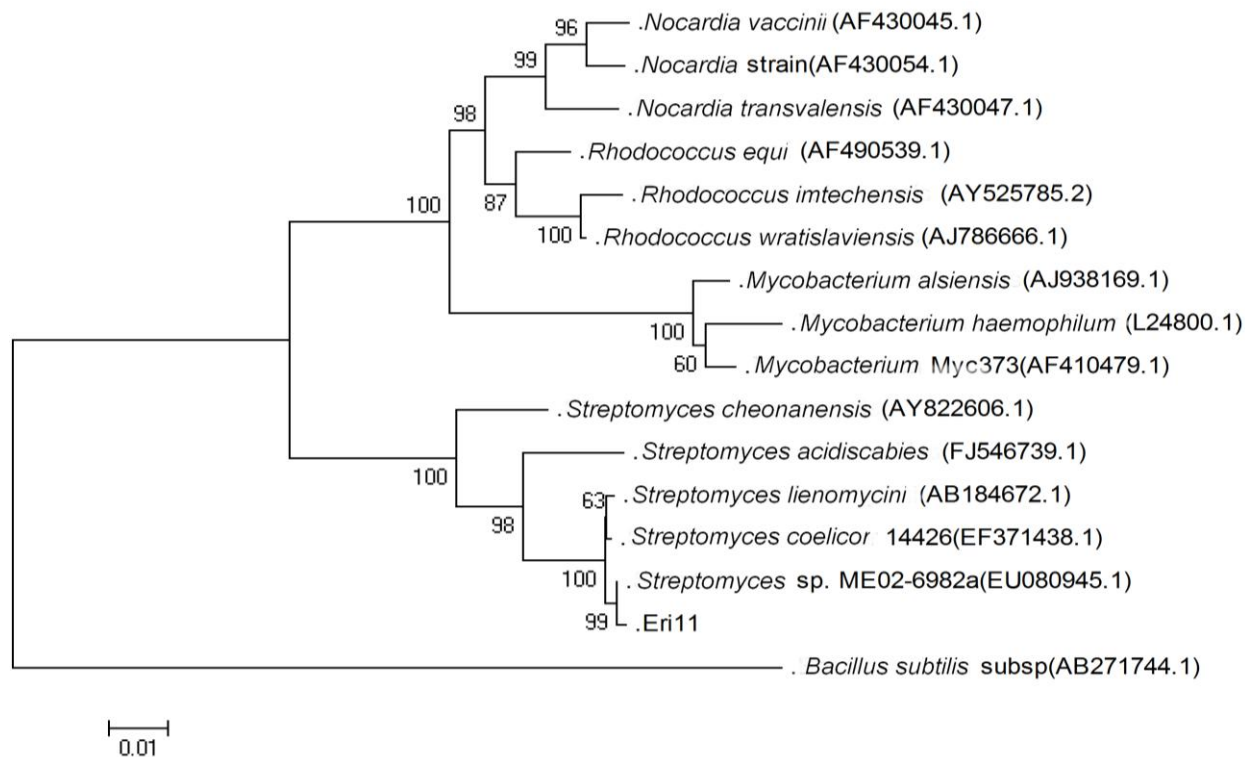
#### Statistical analysis

The data of all experiments were recorded as means  $\pm$  standard deviations and were analyzed with SPSS (version 17.0 for Windows, SPSS Inc.). Differences were considered significant at  $P < 0.05$ .

## RESULTS AND DISCUSSION

In the present study, the *Streptomyces* strain Eri11 was isolated from the rhizosphere of *R. Curcumae Longae*, one of the Chinese medical plants. The morphology of the strain Eri11 sporophore after cultured on S medium for 7 days showed rectiflexibilis pattern. The colonies were aerobic and grew slowly. The colour pattern of the strain Eri11 mycelia grown on S medium was totally shown in Table 1. The colour of arial mycelia was grey, and the colour of substrate mycelia was blue. The strain Eri11 was Gram-positive according to the result of the Gram's staining. The strain produced melanoid pigments (Table 1). A great variety of biochemical features of the strain Eri11 were also studied. As shown in Table 2, the strain Eri11 could utilize the carbon sources efficiently including glucose, xylose, arabinose, fructose, galactose, mannitol, inositol and sucrose. Tolerance of the strains to NaCl concentration also served as an important character for species identification. The maximum tolerance of the strain Eri11 to NaCl was 5%.

In order to classify the isolated strain based on its biochemical traits, the phylogeny analysis of the strain Eri11 was investigated. Phylogeny of the strain based on



**Figure 1.** Phylogenetic tree based on the 16S rDNA sequences of the showing affiliation of Eri11 strain with closely related members in GenBank. *Bacillus subtilis* (AB271744.1) was used as an outgroup. Phylogenetic trees were generated using MEGA version 4.1 with default parameters, K2P distance model and the Neighbor-Joining algorithm. The numbers at the branching prints are the percentages of occurrence in 1000 bootstrapped tree.

**Table 3.** Total phenols and flavonoids in the cultured broth extract and its antioxidant activity.

Total phenols contents	13.59 ± 0.17 mg gallic acid equivalent/g cultured broth
Total flavonoids contents	9.93 ± 0.83 mg rutin equivalent/g cultured broth
ABTS radical scavenging activity	IC <sub>50</sub> = 223.81 ± 24.50 µg/ml
Hydroxyl radical scavenging activity	IC <sub>50</sub> = 582.42 ± 83.10 µg/ml

the sequencing of the 16S rDNA, had become the method of choice for tracing bacterial phylogenies and defining taxonomy (Yin et al., 2008). To elucidate the taxonomic positions of the isolated strain, we sequenced the full length of 16S rDNA gene of the strain Eri11. A continuous stretch of 1561 bp (including gaps) was used for building the phylogenetic Neighbor-Joining tree by using MEGA version 4.1 with default parameters, K2P distance model and the Neighbor-Joining algorithm. The confidence of the grouping was verified by bootstrap analysis (1000 replications). *Bacillus subtilis* (AB271744.1) was used as an outgroup. From the phylogenetic tree (Figure 1), the actinomycete isolated from *R. Curcumae Longae* was classified into *Streptomyces*. The 16S rDNA gene sequence of Eri11 showed high similarity of 99.4% to that of *Streptomyces*

*coelicor* 14426 (EF371438.1). Phylogenetic tree revealed that Eri11 is strongly related to *Streptomyces* forming a distinct cluster (Figure 1).

Cultured with the isolated strain Eri11 in the S medium broth for 14 days, there were some kinds of antioxidants produced. It has been reported that many flavonoids and flavonols exhibited prominent antioxidant behavior as a result of their ability to scavenge radicals (Torres et al., 2002). Therefore, we investigated the total contents of phenols and flavonoids of cultured broth by using the Folin–Ciocalteu method and AlCl<sub>3</sub> colorimetric method, respectively. By using the Folin–Ciocalteu method, the total phenols contents in the cultured broth were calculated to be 13.59 ± 0.17 mg gallic acid equivalent/g cultured broth (Table 3). Meanwhile, by using AlCl<sub>3</sub> colorimetric method, the total flavonoids contents in the

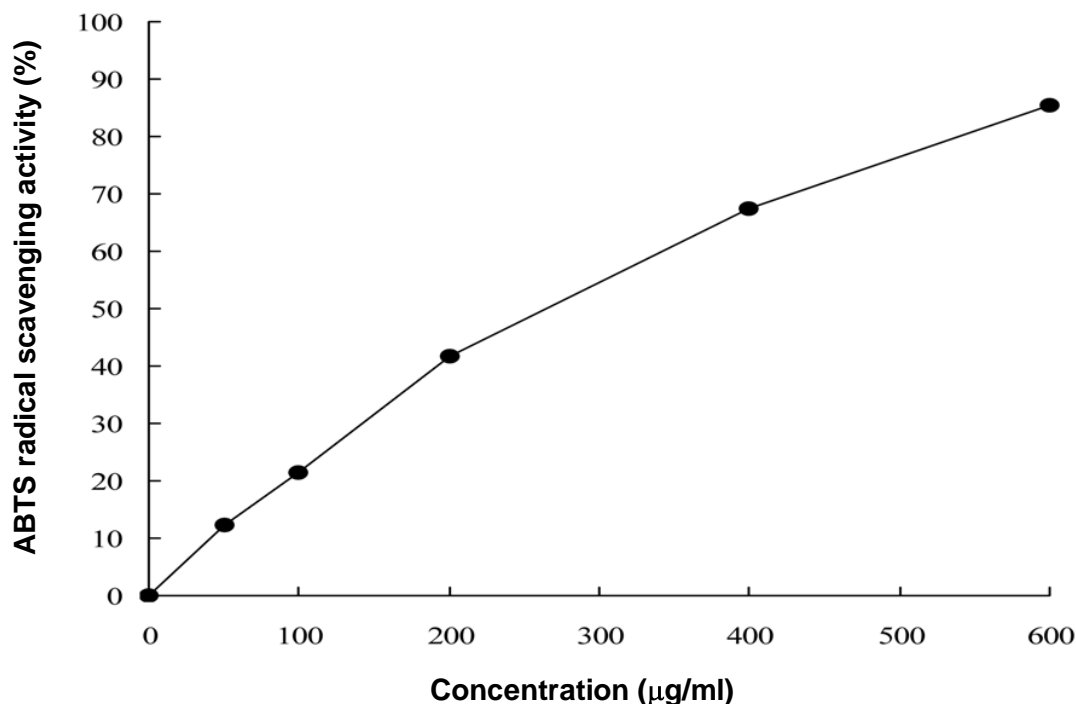


Figure 2. ABTS radical scavenging activity of the cultured broth extract from *Streptomyces* strain Eri11.

cultured broth were calculated to be  $9.93 \pm 0.83$  mg rutin equivalent/g cultured broth.

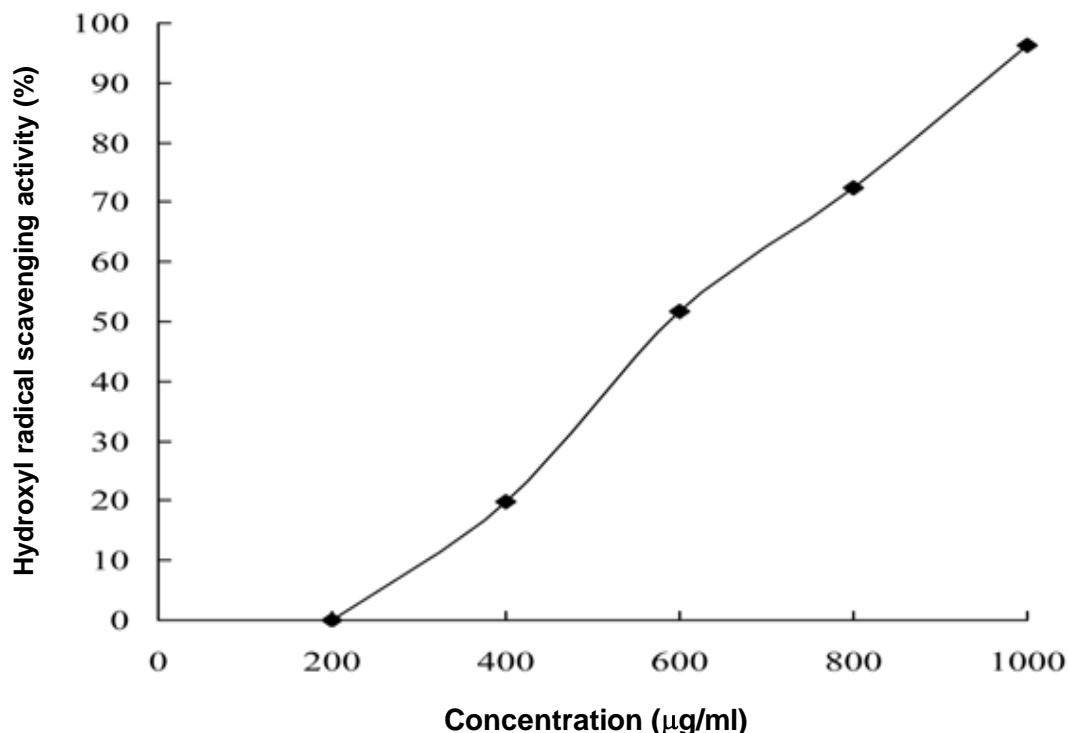
The ABTS radical cation decolorization assay is one of methods for the screening of the antioxidant activity (Re et al., 1999). The pre-formed radical monocation of 2,2'-azinobis-(3-ethylbenzothiazoline-6-sulfonic acid) (ABTS<sup>•+</sup>) is generated by oxidation of ABTS with potassium persulfate and is reduced in the presence of such hydrogen-donating antioxidants. Therefore, the ABTS radical scavenging activity of the cultured broth was determined.

After reacting for 30 min, the cultured broth showed ABTS radical scavenging activities of 12.41% at 50 µg/ml, 21.42% at 100 µg/ml, 41.61% at 200 µg/ml, 67.53% at 400 µg/ml, and 85.54% at 600 µg/ml, respectively. The results indicated that the cultured broth showed a lesser tendency to decay ABTS radicals at low concentrations of reaction than at high concentrations. As shown in Figure 2, the cultured broth scavenged ABTS radicals in a concentration-dependent manner. The half-inhibitory concentration (IC<sub>50</sub>) was estimated at  $223.81 \pm 24.50$  µg/ml with a linear regression. The antioxidant activity of the cultured broth was about 300 times lower than that of the positive control (Trolox, IC<sub>50</sub> = 0.76 µg/ml) (Table 3). This different half-inhibitory concentration between the cultured broth and the standard antioxidant can be explained in terms of the fact that the active components in the cultured broth comprise only a fraction of the cultured broth. When considered total phenols or flavonoids contents, the cultured broth showed ABTS

radical scavenging activity with IC<sub>50</sub> values of 3.04 µg/ml (calibrated with total phenols presented in the cultured broth) and 2.22 µg/ml (calibrated with total flavonoids presented in the cultured broth), respectively.

To determine the antioxidant activity of the cultured broth, hydroxyl radicals scavenging activity test, another typical method, was adopted. Phenanthroline monohydrate-Fe<sup>2+</sup> is a common indicator of redox reaction. The change of Phenanthroline monohydrate-Fe<sup>2+</sup> color could indicate the redox reaction of the reagent. Hydroxyl radical is one kind of strong oxidizers, which could be produced after the chemical reaction between hydrogen peroxide and Fe<sup>2+</sup>. The content of hydroxyl radical of the reaction system is determined by the change of the indicator (Phenanthroline monohydrate-Fe<sup>2+</sup>) color. Therefore, the hydroxyl radical scavenging activity of the cultured broth was determined according to the change of the indicator (Phenanthroline monohydrate-Fe<sup>2+</sup>) colour.

After reacting for 1 h, the cultured broth showed hydroxyl radical scavenging activities of 0.11% at 200 µg/ml, 19.71% at 400 µg/ml, 51.80% at 600 µg/ml, 72.30% at 800 µg/ml, and 96.28% at 1000 µg/ml, respectively. The results indicated that the cultured broth at low concentrations showed a lesser tendency to decay hydroxyl radicals than at high concentrations. As shown in Figure 3, the cultured broth scavenged hydroxyl radicals in a concentration-dependent manner. The half-inhibitory concentration (IC<sub>50</sub>) was estimated at  $582.42 \pm 83.10$  µg/ml with a logarithmic regression. The



**Figure 3.** Hydroxyl radical scavenging activity of the cultured broth extract from *Streptomyces* strain Eri11.

antioxidant activity of the cultured broth was about 400 times lower than that of the positive control (Vitamin C,  $IC_{50} = 1.37 \mu\text{g/ml}$ ) (Table 3). This different half-inhibitory concentration between the cultured broth and the standard antioxidant can be explained in terms of the fact that the active components in the cultured broth comprise only a fraction of the cultured broth. When considered total phenols or flavonoids contents, the cultured broth showed hydroxyl radical scavenging activity with  $IC_{50}$  values of  $7.92 \mu\text{g/ml}$  (calibrated with total phenols presented in the cultured broth) and  $5.78 \mu\text{g/ml}$  (calibrated with total flavonoids presented in the cultured broth), respectively.

## Conclusion

The present study indicated that the strain Eri11 isolated from the rhizosphere of *R. Curcumae Longae* was a novel *streptomyces* strain based on the morphological characteristics, biochemical traits and phylogeny analysis. According to the results of the antioxidant activity tests, it suggested that the strain Eri11 could metabolize antioxidants in S liquid medium. Thus, it will prove to be a good candidate as the antioxidant products for industry. Further work is in progress in our laboratory to elucidate the identity of compounds responsible for the antioxidant activity.

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